

Advances in Delivery Science and Technology

Kenneth A. Howard
Thomas Vorup-Jensen
Dan Peer *Editors*

Nanomedicine



Advances in Delivery Science and Technology

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Nanomedicine

 Springer

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*Editor Kenneth A. Howard would like to
dedicate this book to his sister Jacqueline
Teer.*

Preface

Nanomedicine is set to meet medical challenges such as early and rapid diagnosis, targeted and effective drug delivery, and novel ways of providing organ and tissue replacement that work towards new personalized medicines. Utilizing properties occurring at the nanoscale for unique medical effects as well as understanding and controlling cellular molecular processes and clinical translation are key elements in nanomedicine. These skills are strongly bound by the threads that connect nanoscience, physics, material science, immunology, molecular biology, pharmacology, pharmaceuticals, and medicine. The book *Nanomedicine* sets to capture this broad spectrum of disciplines whilst still maintaining focus on the key elements that establishes it as a *bona fide* unique field. The inclusion of adequate background information for understanding the basic principles makes this text an excellent learning tool for students as well as a useful reference for more senior scientists and other practitioners in the field.

The opening chapter defines the current field of nanomedicine by addressing the constituent key elements and describing major nanomedicine initiatives and programs, as well as product landscape to understand the development process and the impact of nanomedicine. The following three chapters focus on immune recognition with emphasis on the response by the innate immune system, notably complement activation of relevance in removal of nanomedicines as well as in disease pathogenesis. Chapter 2 focuses on complement factor inhibitors developed by structural biology determination. Chapter 3 addresses processes leading to nanoparticle-induced innate immune activation and discusses consequences for nanomedicine-based therapeutics. Chapter 4 focuses on protein polyvalent interactions with molecular recognition patterns at the nanoscale which is relevant in design criteria for the reduction of immune recognition of particulate nanomedicines. The next section focuses on nanomedicine-based diagnostics. Chapter 5 describes microfluidics-based tools for single cancer cell analysis and the use of nanotechnology for improved signal responses. Nanotheranostics and in vivo imaging modalities are addressed in Chapter 6. Chapters 7–11 describe a range of nanoscale nanopharmaceuticals. Chapter 7 gives an overview of a range of nanocarriers used for cancer treatment with a focus on targeting approaches. The ability to influence interaction

with the extracellular and intracellular biological environment by particle geometry is the focus of Chapter 8. Nano-drug delivery systems for delivery of protein/peptide drugs to the brain are described in Chapter 9, and polymer-based DNA delivery to dendritic cells for cancer immunotherapy in Chapter 10. The section concludes with Chapter 11 on the use of the biopolymer silk in nanomedicine. The clinical translation of nanomedicines is dependent on safety and regulatory approval, addressed in Chapter 12. The final section describes nanomedicine-based tissue regeneration and cell therapy with the application of nanotechnology for medical implant drug release in Chapter 13, and Chapter 14 describes guided cellular responses by surface cues for nanomedicine applications.

The book *Nanomedicine* is relevant both as a textbook of an emerging field and as an essential read that gives insight and clarity to this interdisciplinary subject.

Acknowledgements

The editors would like to express their sincere gratitude to all the authors for the time and effort they have given in the preparation of the excellent chapters contained in the book. We are grateful to the CRS and especially Series Editor Michael Rathbone for giving us the opportunity to put together a book we feel is needed in the field. Our special thanks also go to our publisher Springer and their staff members Carolyn Honour, Renata Hutter, Sarah McCabe, Jacob Rosati, and Michelle Feng He, all who worked with both timely attention and patience during the preparation of this book.

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Editor's Biography

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Dan Peer is a translational scientist and Professor that leads an NIH-funded lab in the Faculty of Life Science and the Faculty of Engineering at Tel Aviv University (TAU). He is also the Director of the Focal Technology Area (FTA) on Nanomedicines for Personalized Theranostics, a National Nanotechnology Initiative, and the Director of the Leona M. and Harry B. Helmsley Nanotechnology Research Fund. Prof. Peer's work was among the first to demonstrate systemic delivery of RNAi molecules using targeted nanocarriers to the immune system, and he pioneered the use of RNA interference (RNAi) for in vivo validation of new drug targets within the immune system. He is an editor of several books in the field of nanomedicine, Editor of *Molecular and Cellular Therapies* (Springer), Editor of *Biology and Medicine in Nanotechnology* (IOP), and an Associate Editor of the *Journal of Controlled Release* (Elsevier), *Journal of Biomedical Nanotechnology*, and *Biochemistry*; he is on the Editorial Boards of the *Biomedical Microdevices* (Springer), *Cancer Letters* (Elsevier), *Nanomedicine: Nanotechnology, Biology and Medicine* (Elsevier), and *Bioconjugate Chemistry* (ACS). Prof. Peer is currently the President of the Israeli Chapter of the Controlled Release Society and a Member of the Israel Young Academy of Sciences and Humanities.

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Chapter 1

Nanomedicine: Working Towards Defining the Field

Kenneth A. Howard

Abstract An aging population and poor clinical solutions for many disease types has fuelled the rapid emergence of nanomedicine. Defining what constitutes nanomedicine is crucial to its recognition as a distinct field and directing future research focus towards clinical translation. This chapter sets out to define the field by addressing the important constituent key elements and describing major nanomedicine initiatives and programmes, and product landscape to understand the development process and current status.

Keywords Nanomedicine • Definition • Nanoscale • Molecular processes • Clinical translation • Nanomedicine products

1.1 Introduction

Poor clinical solutions for many disease types and rising healthcare cost due to an ageing population, has fuelled the rapid emergence of nanomedicine to meet challenges such as early and rapid diagnosis, effective targeted drug delivery and discovery, organ and tissue replacement and personalized medicines. Immense expectation is the motivation behind the establishment of global roadmaps in nanomedicine [1, 2] with high industrial and governmental investment. Dedicated nanomedicine journals and global societies [3–6] have been established over recent years that perpetuate the wave of activity and interest in the field with a rapid increase in published articles. The field is extremely broad and multidisciplinary exemplified by the commonly used definition “*The medical application of nanotechnology for diagnosis, treatment and management of human health*” that spans nanoscience, physics, material science, molecular biology, pharmaceuticals, medicine and clinical translation. Whilst utilization of properties occurring at the

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nanoscale having unique medical effects, is central to nanomedicine, understanding and controlling cellular molecular processes and clinical translation are other key elements that need to be considered for inclusion. Defining the field is crucial for the establishment of nanomedicine as a *bona fide* field of its own rather than rebranding of traditional disciplines, a measure of its societal impact, and identifying current status and steps needed in shaping its future development. This chapter sets out to clearly define the field by describing major nanomedicine initiatives and programmes and addressing the important constituent key elements, and marketed examples to give a global perspective and the impact of nanomedicine.

1.2 Nanomedicine Initiatives and Approaches

This section will describe some major initiatives in Europe and the United States of America and their respective views on what constitutes “nanomedicine”, in order to give an historical perspective and understand the process and current status, and importantly, work towards consensus and a uniform definition required for focused efforts and the future development of the field. Whilst these serve to exemplify the process and status, there are numerous global initiatives in Asia and elsewhere in the nanomedicine field.

1.2.1 European Nanomedicine Initiatives

The European Medical Research Council (EMRC) within the European Science Foundation (ESF), initiated a process in 2003 towards preparation of a Scientific Forward Look report on Nanomedicine. Academic and industrial experts met to define the field, discuss future societal impact, identify Europe’s strengths and recommend funding priorities and national and pan-European infrastructures needed for a coordinated scientific activity. This process was facilitated through a series of specialized workshops held during 2004 culminating in a final consensus meeting in November 2004 to prepare the ESF-EMRC forward look report published in 2005 [7]. In this report the field of nanomedicine was defined as: “*the science and technology of diagnosing, treating and preventing disease and traumatic injury, of relieving pain, and of preserving and improving human health, using molecular tools and molecular knowledge of the human body*”.

Nanostructures from one nanometre to hundreds of nanometres, included in a device or a biological environment, were highlighted as an important component with key sub-disciplines; analytical tools, nanoimaging, nanomaterials and nanodevices, novel therapeutics and drug delivery systems, and clinical, regulatory and toxicological issues highlighted.

In September 2005, a European Commission Nanotechnology for Health Vision Paper was published that recommended the establishment of a European Union

European Technology Platform on NanoMedicine and a Strategic Research Agenda to be focused on nanotechnology-based diagnostics imaging, targeted drug delivery and release, and regenerative medicine [8]. It was an industrial-led consortium including academic participants, with a focus on strengthening competitive scientific and industrial positioning in the area of nanomedicine and to improve EU citizen's quality of life and healthcare. In this paper nanomedicine was defined as: *“The application of Nanotechnology to Health. It exploits the improved and often novel physical, chemical, and biological properties of materials at the nanometric scale. NanoMedicine has potential impact on the prevention, early and reliable diagnosis and treatment of diseases.”*

In contrast to the aforementioned ESF report, unique properties at the nanoscale were included within the definition. Following on from the Vision Paper, *Nanomedicine nanotechnology for health European technology platform strategic research agenda (SRA) for nanomedicine* [9] was published in November 2006, with defined objectives directed to its member states and the commission with the main aim to provide information “for decision making processes for policy makers and funding agencies”. Although the document stressed technology development was driven by healthcare needs, it stated “nanomedicine is a strategic issue for the sustainable competitiveness of Europe”. Cardiovascular diseases, cancer, musculoskeletal disorders, neurodegenerative diseases and psychiatric conditions, diabetes and bacterial and viral infectious diseases were identified as target diseases based on mortality rates, prevalence, and potential impact of nanotechnology in their diagnosis and treatment. Specific key research areas were diagnostics and imaging, targeted delivery-multi-tasking medicines and regenerative medicine that may overlap e.g. theranostics that include both diagnostic and therapeutic components.

The aforementioned initiatives, in combination with academic and industrial expert workshops in 2009, led to a joint European Commission/ETP nanomedicine expert report published in October 2009 *Roadmaps in nanomedicine towards 2020* [1]. This report focused on nano-diagnostics, nanopharmaceuticals and regenerative medicine and technological timelines towards the treatment of diseases highlighted in the SRA 2006, and to provide a future research funding focus in EU programmes. The industry-led and application driven approach was preferred with emphasis for detailed specific recommendations. Focus was placed on translational research and commercial potential and the necessity for industrial/academic interaction for turning academic results into products directed towards better healthcare for patients. The report highlighted a need for improved knowledge and communication between academics, small medium enterprises and industry for successful implementation of the roadmap and a requirement to exploit industries strengths in product development pathway and regulatory issues. The European nanomedicine roadmap was commissioned by the European Commission with economic considerations a factor. The involvement of industry has led to an application and product-driven approach that is focused and defined.

The ETP in nanomedicine is the main player in providing advice and suggestions including identification of priority areas to the European Commission [10]. It provides strategic documents and recommendations from the nanomedicine community

including a nanomedicine white paper, the new EU Framework Programme for Research and Innovation Horizon 2020 [11], and dissemination of knowledge, regulatory and IPR issues, standardisation, ethical, safety, environmental and toxicity concerns. The ETP have prepared an interactive European Nanomedicine Map of the private and academic nanomedicine community to facilitate coordination and collaboration in Europe [12]. A key component of both the ESF and ETP is the focus on European strengths, a strategy that may neglect other potential important areas.

Most European member states have designated nanomedicine calls in its research funding programmes independent from the EU funded projects and allows diversity in the focus and approach. For example, in the UK, the Engineering and Physical Sciences Research Council (EPSRC) had a priority area in therapeutics and nanomedicine in the 2013 Centres for Doctoral Training (CDTs) doctoral training call. In 2008, “nanotechnology to medicine and healthcare grand challenge programme” was awarded by the EPSRC as part of the cross council programme Nanoscience through Engineering to Application. The £15 million call invited proposals for large-scale, integrated projects focused on Nanotechnologies for the targeted delivery of therapeutic agents and Nanotechnologies for healthcare diagnostics. The call was structured to support 8–10, 3 year projects in a first phase 2009–2012, followed by extension to selected projects prioritized on potential to attract alternative external funding in reaching a clinical aim. Interestingly, as part of the same scheme public engagement and dialogue workshop sessions discussing key issues such as research priorities and public perception were organized. The public particularly supported early diagnostic devices that would allow lifestyle changes managed by the patient, and drug delivery systems for difficult to reach organs and cure serious conditions. The public, however, were opposed to treatment controlled by the device and not the healthcare worker or patient, for example, theranostics based on insertion of devices that control drug release based on chemical levels detected by that device.

A model comprising of a specific cluster of dedicated nanomedicine hubs has been established in Denmark in which three new Nanomedicine Research Centers were established in 2010 through \$17.7 million funding by the Lundbeck Foundation for an initial period of 5 years [13]. The Lundbeck Foundation Nanomedicine Centre for Individualized Management of Tissue Damage and Regeneration (LUNA) (Aarhus University), Lundbeck Foundation Center for Biomembranes in Nanomedicine (Copenhagen University) (CBN) and, the Lundbeck Foundation Nanomedicine Research Center for Cancer Stem Cell Targeting Therapeutics (NanoCAN) (University of Southern Denmark) were the three centres. The three centres of excellence investigate novel strategies for exploiting nanoscaled biostructures in combating major human diseases, including cancer as well as neurological, infectious, cardiovascular and musculoskeletal disorders. The centres are highly interdisciplinary including nanotechnology, biotechnology, molecular biology, biomedicine, chemistry, and physics with a key element to combine basic and clinical research. Independent funding from the Danish Research Council facilitates a sustained nanomedicine drive in the country.

1.2.2 USA Nanomedicine Initiatives

The National Institutes of Health have a common fund programme in nanomedicine also referred to as the NIH roadmap [2]. It is a 10 year programme, initiated in 2005, based on “High Risk, High Reward” with two main goals (1) study nanoscale cellular structures and processes using tools to probe and manipulate (Phase 1) and, (2) utilize this information to understand and treat specific disease (Phase 2). The programme started with a national network of eight nanomedicine centers, reduced to four in 2010 based on the most translational potential. The programme first phase focused on basic science and the second phase to clinical translation. The programmes definition of nanomedicine “*refers to highly specific medical intervention at the molecular scale for curing disease or repairing damaged tissues, such as bone, muscle, or nerve. A nanometer is one-billionth of a meter; too small to be seen with a conventional lab microscope. It is at this size scale – about 100 nanometres or less – that biological molecules and structures operate in living cells.*” Emphasis is placed on size and relationship with nanoscale biological structures, using the strengths of NIH-funded researchers to investigate and understand cellular mechanisms towards research directed at characterisation of molecular cellular components and re-engineering those processes for medical treatments. Funded centers included, the nanomedicine center for nucleoprotein machines, center for protein folding machinery, nanomedicine center for mechanobiology directing the immune response, cellular control: synthetic signalling and motility systems. Funding will be discontinued from the common fund in 2015 and funded from other sources as more focus is directed towards specific disease types.

Other initiative include the National Cancer Institute \$144.3 million (2005–2010) *alliance for nanotechnology in cancer* that started in 2004, comprising of nine centers of excellence focused more towards clinical applications than basic science to use nanotechnology to diagnose, treat and prevent cancer [14]. Research includes tools to identify new biological targets, agents for early disease diagnosis using in vitro assays and in vivo imaging tools, multi-functional targeted devices to deliver multiple therapeutics to cancer, tools for real-time assessment of therapeutic and surgical efficacy. Emphasis is on multidisciplinary research between diverse institutions and organizations, with an alliance framework that includes nine centers of cancer nanotechnology excellence focused on discovery and tool development of nanotechnology in clinical oncology; 12 cancer nanotechnology platform partnerships that support individual projects and use nanotechnology to address fundamental questions on cancer; and six cancer nanotechnology training centers educating and training researchers from different fields in the use of nanotechnology-based approaches in cancer. This is a two phase programme, with the first phase 2005–2010, to establish multidisciplinary teams in focused areas and established a Nanotechnology Characterisation Laboratory as a nanomaterials characterisation hub for standardization of materials to better enable technology transfer from academia to companies to the clinic [15]. The second phase, 2010–2015 aims to translate nanotechnology findings into the clinic.

1.3 Nanomedicine Key Elements

Utilization of properties at the nanoscale having unique medical effects is central to nanomedicine programmes and easy to appreciate its inclusion. Understanding and controlling cellular molecular processes and clinical translation are, however, other components that constitute the field. This section will address each in turn towards a definition encompassing all these key elements.

1.3.1 Nanoscale Properties That Have Unique Medical Effects

Harnessing novel physicochemical and biological properties exhibited by nanoscale materials to improve human health is perhaps the fundamental element of nanomedicine. As far back as 1959 the future societal impact of nanotechnology was predicted by Nobel Laureate Richard Feynman “*Many of the cells are very tiny.....; they do all kinds of marvelous things – all on a very small scale.....Consider the possibility that we too can make a thing very small which does what we want – that we can manufacture an object that maneuvers at that level*” [16]. The rapid emergence of nanoscience in the new millennium has provided a wealth of knowledge into the physicochemical properties and biological performance of nanoscale materials that can be used to utilise nanotechnology for medical applications (Fig. 1.1). Examples of unique physical properties occurring at the nanoscale include distinct light-scattering and optical properties exhibited by metal nanoparticles of different size, shape and compositions that can be utilised in diagnostic biosensors [17]. The magnetic to superparamagnetic shift found at the nanoscale can be used for molecular imaging applications [18]. Furthermore, localized exposure of superparamagnetic iron oxide nanoparticles (SPION) to an oscillating magnetic field induce extreme polarity fluctuations that results in an increase in core temperature that has been utilised for thermal-induced destruction of tumours [19]. The increased solubility occurring at the nanoscale has been utilised in Nanocrystals for dissolution of poorly soluble drugs widely used within the Pharmaceutical Industry [20]. The saturation solubility becomes a function of particle size below a critical size ~1000 nm where solubility increases with decrease in size.

Nanoparticles used as drug delivery nanocarriers exhibit unique biological properties that can be utilised to create therapeutics with higher efficacy and reduced toxicity in patients. The ability for nanocarriers to circulate and extravasate across endothelial barriers into tissue and enter cells by endocytosis is a size dependent process [21] (Fig. 1.1). Furthermore, the large surface area of nanoscale particles potentiates surface functionalisation characteristics such as targeting and stealth. Nanoscale particles consequently have been shown to increase drug payload to diseased sites [22].

The ability to control cellular interactions, proliferation and stem cell differentiation by nanotopography and biomolecule patterning has great potential in tissue

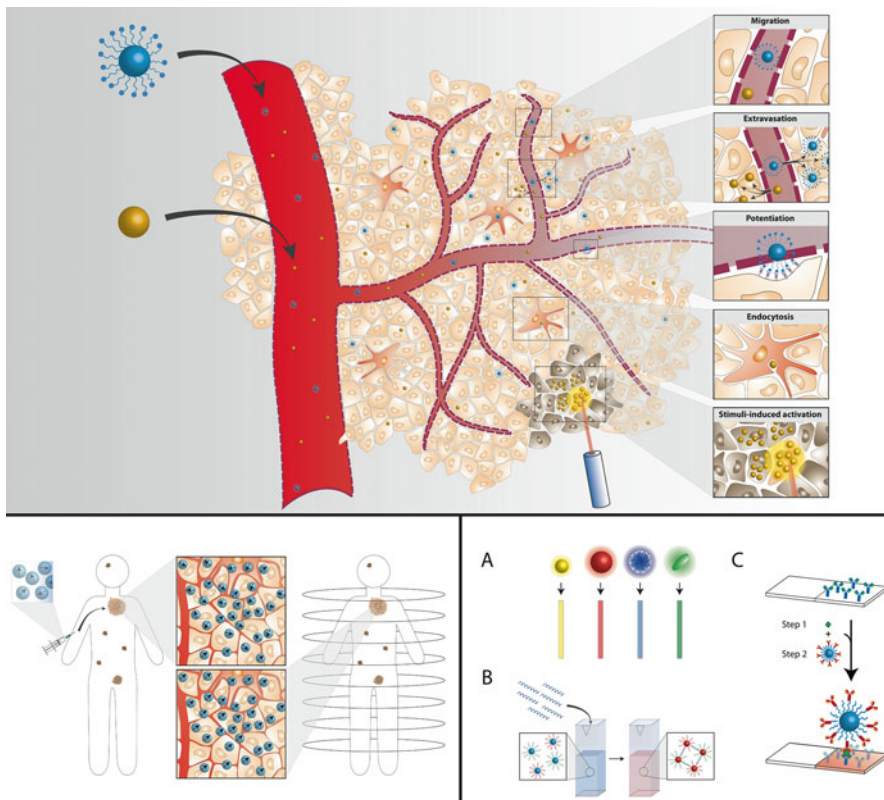


Fig. 1.1 Properties occurring at the nanoscale that exhibit unique medical effects. *Upper panel:* Nanoparticle-based drug delivery. Nanoscale particles can migrate within the bloodstream facilitated by stealth coatings and extravasate across endothelium into normal and diseased tissue. Surface engineered nanoparticles functionalised with targeting agents engage with cellular surface receptors potentiated by the nanoparticle high surface to volume ratio. Size-mediated endocytosis allows cellular entry and delivery of the drug cargo. Inorganic nanoparticles composed of gold or iron oxide can be used for thermal-induced disease death activated by thermal or magnetic stimuli, respectively. *Lower right panel:* In vitro diagnostics. The optical properties dependent on nanoparticle size, composition and shape (a) can be utilised for diagnostic purposes. Complementarity of specific oligonucleotide to oligonucleotide coated on particles induces aggregation and a detectable spectral shift (b). An antibody on a chip and a nanoparticle that target different epitopes on the same antigen can be used detect pathogenic antigens (c). *Lower left panel:* Molecular imaging. The shift to paramagnetism at the nanoscale where all the magnetic domains are aligned in the same direction can be used for improved disease contrast in magnetic resonance imaging. Graphic prepared by Morten Tobias Jarlsted Olesen, Aarhus University

engineering. Cells respond to nanoscale chemical patterns and topography cues provided by different protein formats such as protein folding or collagen banding found in the tissue. This can be mimicked by nanostructures functionalized onto implant surfaces with different biomolecule patterns [23]. The importance of interactions at the nano-bio interface is exemplified in recognition of materials by components of the

immune system. Induction of the immune cascade is dependent on the structural conformation of pattern recognition molecules (PRM) for engagement with pathogenic associated recognition molecules (PAMPS). Geometry and curvature of surfaces determines the capability of PRM to adopt these conformations [24] that can potentially be used to fabricate nanomedicines with increased biocompatibility.

1.3.2 The Use of Molecular Tools and Molecular Knowledge of the Human Body for Medical Applications

The definition of nanomedicine in the ESF-EMRC forward look report [7] places focus on the use of molecular tools and molecular knowledge of the human body as a key component, whilst, the USA Roadmap focuses on understanding and characterisation of nanoscale cellular structures and processes and re-engineering those processes for medical treatments. Traditional disciplines such as cell biology, structural biology, proteomics, genomics and molecular biology to investigate the key molecular processes in disease, therefore, are relevant in nanomedicine. Furthermore, nano-characterisation and cellular investigatory tools such as scanning tunnelling microscopy (SPM) and real-time microscopic cellular imaging at the single molecule level allow a greater understanding of the materials and cellular processes. Examples within the NIH roadmap [2], is the investigation of “nucleoprotein machines” involved in DNA replication and repair, RNA synthesis and protein translation. At the Nanomedicine Center for Nucleoprotein Machines, GA Tech Research Corp. tools are being developed to better investigate and understand these fundamental processes, structure-function relationship such as assembly and disassembly, signaling and control mechanisms. The study of chaperone proteins, or “nanomachines” involved in folding proteins crucial to the cell homeostasis, and removal of damaging misfolded polypeptides towards identification of therapeutic or diagnostic targets is the focus at Center for Protein Folding Machinery at the Baylor college of Medicine.

The Nanomedicine Centre for Individualized Management of Tissue Damage and Regeneration (LUNA), Aarhus University, Denmark [25] combines basic with applied science using traditional and nano-approaches including X-ray diffraction, liquid- and solid-state NMR, magnetic resonance imaging (MRI), small-angle X-ray scattering, and cryo-electron microscopy and scanning tunnelling microscopy. The aim is to identify the central molecular processes involved in the inflammation and de/regenerative imbalance associated with certain cardiovascular and musculoskeletal diseases with focus on the role and regulation of pattern recognition molecules such as collectins and toll-like receptors. This information is to be used for new approaches in drug design, drug delivery, bioimaging, tissue regeneration towards therapeutics and early diagnostic tools. An example is the structure-based drug design for identification of complement blocking agents. X-ray diffraction has been used to study the atomic structure of C5 and its cleavage fragments C5a and C5b involved in innate immune-mediated autoimmune response. The aim is to identify novel blocking drugs, as an alternative to C5 antibody, with the possibility

for specific inhibition of downstream cleavage fragments in autoimmunity without detriment to overall immune protection. The application of nanoparticle-based enabling technologies developed within a drug delivery component of the center can then be used for in vivo delivery, emphasising the link between the key elements of molecular knowledge and nanoscale properties. The functional importance to immunity of nm-scaled conformational changes in the large proteins of the immune system is a focus [26]. The capability to regulate ligand nanoscale spacing of carbohydrates such as mannose and N-acetyl glucosamine can be used to investigate by real-time monitoring using atomic force microscopy and small-angle X-ray scattering of the spacing of mannan-binding lectin (MBL) required for complement activation. This will allow insight into the conformational nature of the patterns recognized by MBL and allow design of antagonists to block MBL-mediated complement.

1.3.3 Nanoscience Merged with the Biomedical Sciences in a Clinical Setting

The application of nanotechnology is fundamental to nanomedicine and, thus, requires a nanoscience environment. The diverse disciplines ranging from nanoscience, physics, material science, molecular biology, pharmaceuticals, medicine and clinical translation, however, necessitates interdisciplinary/translational research that are key to successful nanomedicine programmes. The environment is unlikely to be found in a single institution, and so calls for interdisciplinary programmes outlined in earlier section such as pan-European initiatives and NIH networks. The involvement of clinicians early in the programmes is needed to provide input and focus towards unmet clinical needs.

1.3.4 Nanomedicine: Redefined

This section has identified components of nanomedicine. The general definitions fall short of inclusion of all these key elements. A more precise, all inclusive definition could be “New scientific concepts through a greater understanding of human processes utilised for development of nano-based strategies in a clinical setting for prevention, diagnosis and treatment of diseases”. Whilst this may not be appropriate for the public or press sound bites, it is important that scientists consider all elements and includes relevant participants in order to fulfil the potential of nanomedicine.

1.4 Nanomedicine Products

The necessity for a uniform definition of nanomedicine is crucial for an accurate assessment of the current status of nanomedicines on the market. A lack of consensus that includes differing size range criteria used, however, complicates clear

identification of what constitutes a nanomedicine, exemplified by the different number and type described, and a lack of a definitive list. Furthermore, the inclusion of products approved before the “nanomedicine era”, for examples, the liposomal Doxil[®] (FDA approval 1995) [27], and the PEGylated proteins Adagen[®] (FDA approval 1990) and Oncaspar[®] (FDA approval 1994) interferes with evaluating the impact of nanomedicine. An early review of the nanomedicine commercialization landscape was reported by Wagner et al. [28] based on science citation, medical and patent databases, and the business literature. Physical effects occurring at the nanoscale was the basis for inclusion such as iron oxide MRI contrast agents that exhibit paramagnetism, however, structures up to 1000 nm were also considered due to unique physiological behaviour. Products were categorized into drug delivery, biomaterials, in vivo imaging, in vitro diagnostics, active implants and, drugs and therapy. The predominant drug delivery category was highly represented by PEGylated proteins and liposomal-based systems. PEGylation that confers extended plasma half-life of proteins by Mw increase-mediated reduction in renal clearance can be viewed as an enabling technology, rather than a unique physical effect that occurs at the nanoscale. Similarly, the proposed ability for liposomes such as the liposomal doxorubicin formulation Doxil to extravasate across tumour endothelium is not dependent on transitions in physicochemical properties at the nanoscale. Both, however, are examples of changes in physiological interactions of particles of a certain size range. This has been highlighted by Etheridge et al. [29], and considerably expands the range of nanomedicines solely from a size range where quantum effects occur at ~100 nm occur to materials up to 500–100 nm e.g. cellular transcytosis or endothelial penetration which can occur, with particles of this size range. Nanocrystals formed by the micronization of drug to increase solubility has a size range from a few nm to microns and are widely used in the Pharmaceutical industry. This, however, reduces the distinction between micro and nano for categorization of a nanomedicine. Nanostructures to improve the mechanical strength of medical implants such as nanoparticle-based dental fillers and nanohydroxyapatite-based bone repair products are predominant. The therapeutic nanomedicine landscape continues to be dominated by liposomes and PEGylated proteins with the addition of Abraxane[®] by Celgene that is nanoparticle albumin bound paclitaxel (nab-paclitaxel) approved for cancer treatment [30]. It is proposed the nanoparticles disassemble into individual albumin drug-bound molecules after injection that results in high drug plasma levels and tumour accumulation attributed to the high metabolic uptake of albumin with a suggestion that SPARC expression in tumours is involved in an active albumin targeting process. There is, however, still a significant absence of targeted nanoparticles. Weissig et al. [31] focused on therapeutic nanomedicines commonly referred to as nanopharmaceutics in a more recent review. In this review nanopharmaceutical categories included liposomes, PEGylated proteins, polypeptides and aptamers, nanocrystals, protein-drug conjugates, polymer-based nanoformulations, metal-based nanoformulations such as imaging agents. A lack of a clear definition and designated data base is an issue, exemplified by the necessity to assign nanomedicines into categories of relevance ranging from “confirmed” to “questionable” in the Etheridge et al. review. There are no specific

regulatory requirements for products containing nanoparticles that may reflect the current uncertainty of what constitutes a nanomedicine. The ability of nanoscale materials to penetrate tissue is of particular toxicology concerns for non-biodegradable metal nanoparticles rather than organic-based systems such as liposomes. The aforementioned initiative by the National Cancer Institute with the Nanotechnology Characterisation laboratory to standardize the characterisation of nanomaterials is needed for the clinical approval process as novel materials arise.

1.5 Conclusion

A clear definition of nanomedicine is required to identify it as a distinct field. Physical properties occurring at the nanoscale having unique medical effects would certainly serve this purpose but neglect other key elements. Inclusion of traditional disciplines such as cell biology, structural biology, proteomics and genomics, may first appear to limit its distinction, however, are relevant for understanding, controlling and re-engineering key molecular processes for disease treatment. Nanomedicine is a fledgling field trying to “find its feet”; knowledge of unique physical and biological properties of an expanding panel of new materials along with a greater understanding of molecular processes is part of its transformation into a distinct established field. The current product landscape reflects its early status, with inclusion of some materials above nanoscale. Whilst, a limited repertoire of products is expected in a field still emerging, it is important not to include materials that reduce the distinction between nano and micro and obstruct its unique field status and measuring the true impact of nanomedicine. Furthermore, it is important to resist “scientific rebranding” that can be damaging to its perception and acceptance as a unique field; “nanomedicine is more than a name”. The recruitment of scientists, from various backgrounds and fields that “repurpose” respective technologies, towards development of nanomedicine, however, is a unique opportunity to take the field of nanomedicine forward. An example is the application of material scientist technologies for use in medical science. Defining the field is important to direct and focus research and education programmes, and standardisation of products. Key elements outlined in this chapter work towards consensus in the field required for focused efforts needed for realising the tremendous potential of nanomedicine.

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Chapter 2

Complement Regulators and Inhibitors in Health and Disease: A Structural Perspective

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Abstract The complement system is an important effector within the innate immune system as a defence against pathogens and maintaining homeostasis. Detection of pathogen- and damage-associated molecular patterns triggers the proteolytic cascade in complement. In healthy self-tissues effector proteins are tightly controlled by proteins acting as regulators of complement activation, and absence or malfunction of these regulators contribute to pathogenesis in a number of disease conditions in humans. Complement is highly relevant to nanomedicine due its role in adverse reactions on polymers and nanoparticle drug carriers, but also since complement hyperactivation contributes to pathogenesis in many disease conditions that are frequently addressed within nanomedicine. We review here the regulatory mechanisms that modulate complement activation and some of the most prominent cases linking complement dysregulation/deficiencies to pathogenesis as well as the strategies that have been considered for the development of therapeutic complement inhibitors and modulators to alleviate complement-mediated detrimental effects. In addition, this chapter summarizes the wealth of strategies adopted by pathogens to evade complement, such as inhibition of the proteolytic cascade, degradation of complement effector molecules and interference with transmembrane signaling by effectors, and highlights how structural and functional insight into their mode of function now provides leads for the development of novel complement therapeutics.

Keywords Innate immunity • Complement • Structural biology • Therapeutics

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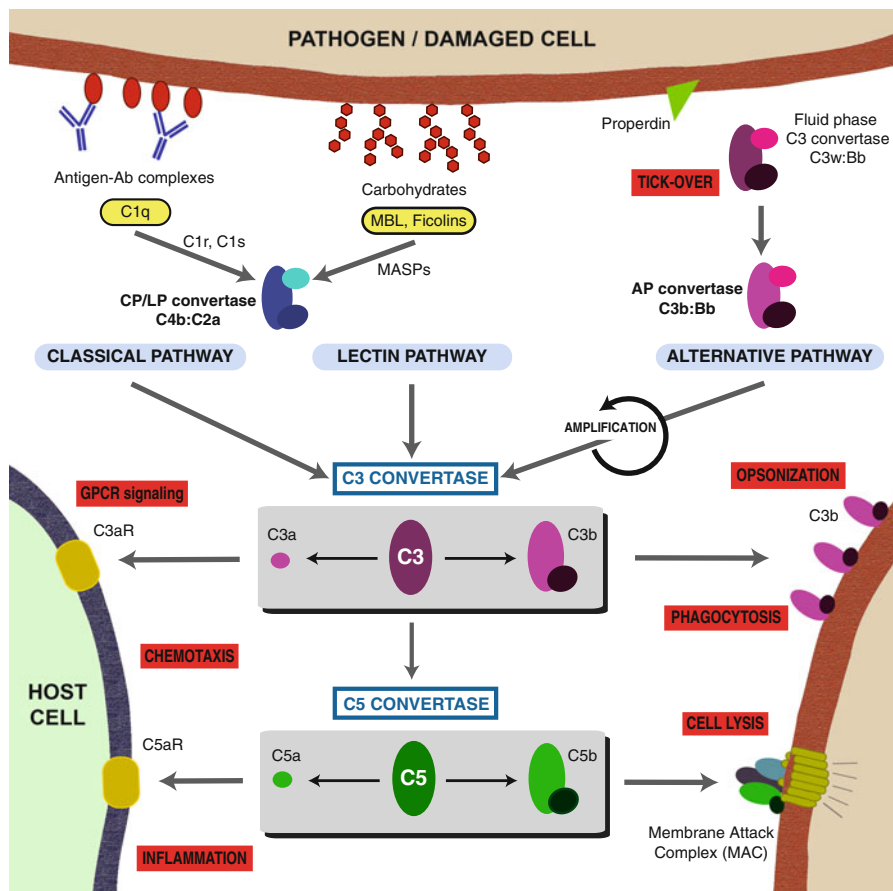


Fig. 2.1 General overview of the complement cascade emphasizing the initiation of the proteolytic cascade upon pattern recognition and the effector molecules acting on both host cells and pathogen/damaged cells

2.1 Introduction to Complement and Associated Diseases

2.1.1 Overview of the Complement Cascade

Complement is one of the major effectors of the innate immune system and is in the first line of defence against invading pathogens. Complement not only protects against infectious organisms, but also disposes of immune complexes, products of inflammatory injury and bridges the innate and the adaptive immunity [1–3]. Complement is a complex network of more than 50 circulating and membrane-bound proteins that can be activated through three different pathways: the classical (CP), lectin (LP) and alternative (AP) pathways (Fig. 2.1). The classical pathway is

initiated upon recognition by the C1 complex of antibody (IgG or IgM):antigen complexes or pentraxins (CRP, PTX3 and SAP) binding directly to activators [4, 5]. The C1 complex consists of the pattern-recognition molecule C1q and the associated serine proteases C1r and C1s that are activated upon pattern recognition [6]. C1s then initiates the cascade by proteolytic cleavage of C4 into C4a and C4b. Through an internal reactive thioester C4b is covalently linked to the activator [7], and C2 subsequently joins to form the proconvertase C4bC2, which is cleaved, also by C1s, resulting in the appearance of the CP C3 convertase C4bC2a [8]. The lectin pathway is similar to the CP but differs in the molecular patterns activating it. The recognition is achieved by mannan-binding lectin (MBL), ficolins, and collectin-11 binding to glycan moieties of a variety of glycoproteins and glycolipids specific to pathogens (bacteria, viruses and fungi) and damaged self [9, 10]. The pattern recognition molecules are associated to serine proteases MASP-1 and MASP-2, and recent work has established that MASP-1 may autoactivate and then cleaves MASP-2 resulting in an activated MASP-2 which can cleave C4. Further processing of the proconvertase may then be carried out by either MASP-1 or MASP-2 [11, 12].

C3 convertases are proteolytic complexes able to cleave the central complement component C3 into C3a and C3b [13]. The anaphylatoxin C3a recruits immune cells to the site of infection and initiates an acute inflammatory response [14] whereas C3b is the major opsonin of the complement system. Like C4b, it covalently attaches to the activator through its thioester [15] thereby “tagging” foreign and altered-self objects leading to opsonization. This also leads to initiation of the alternative pathway as activator-bound C3b recruits and binds factor B which is then cleaved by factor D, yielding the AP C3 convertase C3bBb [16]. The C3 convertase in turn generates more C3b from C3 and in this way creates a powerful amplification loop that accounts for 80–90 % of the outcome in the terminal pathway (see below) when complement is activated through the CP [17] or the LP [18]. However, AP activation may also occur spontaneously upon hydrolysis (tick-over) of the thioester bond within C3 generating a water-reacted molecule, C3(H₂O) capable of forming the fluid-phase C3 convertase [19], which can then be stabilized on microbial surfaces and apoptotic/necrotic host cells by properdin [20].

The C3 convertases may recruit a second molecule of C3b yielding the AP C5 convertase C3bBbC3b or the CP C5 convertase C4bC2aC3b [21]. These cleave C5 into C5a and C5b thereby initiating the terminal pathway (TP) [22]. The C5a anaphylatoxin signals through the G-protein coupled receptor, C5aR, and initiates inflammation (see below). C5b is devoid of the thioester and does not attach to activating surfaces but instead binds to C6, C7, C8 and multiple copies of C9, forming the (C5bC6C7C8)C9_n complex known as the membrane attack complex (MAC) [23]. Active MAC is able to insert into membranes and form pores resulting in cell lysis. Although MAC is potentially a powerful weapon against invading pathogens, deficiencies in the TP proteins primarily leads to *meningococci* infections [24].

2.1.2 Complement Regulation

Obviously, uncontrolled C3b deposition causes inflammation and cytolysis. For this reason complement has to be tightly regulated on healthy tissues. Host cells express cell-surface and soluble regulators (Fig. 2.2). Low concentration/absence or mutations in these proteins are at the heart of pathogenesis when complement is involved. The serpin C1 esterase inhibitor (C1-INH) blocks the C1 complex and also the LP proteases MASP-1 and MASP-2 [25], but it is not specific to complement as it also targets plasmin, thrombin, factor Xa and kallikrein. In the AP, most regulators function at the level of the C3 convertases by stimulating their dissociation (decay acceleration activity) or by promoting the proteolytic degradation of C3b into iC3b by the serine protease factor I [26] (co-factor activity). iC3b cannot associate with factor B and is thus irreversibly unable to form the C3 convertase. iC3b may be further degraded into C3dg and finally C3d by factor I and plasmin [27].

Factor H (fH) is the major AP regulator and exhibits both decay and co-factor activity for both C3bBb and C3(H₂O)Bb C3 convertases [28]. A pathogen cell opsonized with C3b but not capable of stabilizing fH binding will bind fH weakly preventing inactivation of C3b, whereas a non-activating host cell presenting the appropriate glycosaminoglycans will associate efficiently with fH yielding protection through C3b degradation [29]. fH consists of 20 complement control protein (CCP) domains (Fig. 2.2a). X-ray crystallography and NMR have deciphered the three-dimensional structures of all fH CCP domains but CCPs 9, 14 and 17, while full length fH has been studied by solution scattering [30–34]. Most of the structure-function research has focused on CCPs 1–4 and 19–20 binding C3b, and CCPs 6–8 and 19–20 associating with self-surfaces [34–36].

There are six other proteins related to fH that bind to C3b or C3d, the fH-like protein 1 (a splice variant of fH) and five fH-related proteins (CFHR1–5). All these are also composed entirely of CCP domains with different degrees of sequence identity with fH (Fig. 2.2a). CFHR1 can associate into a homodimer as well as into a heterodimer with CFHR2 or CFHR5 [37]. CFHR1 inhibits the C5 convertase as well as MAC formation [38] by binding to C3b through its N-terminal homodimer-forming moiety, but CFHR1 lacks both co-factor and decay activity [39]. CFHR2 inhibits the formation of the C3 convertase [40] but does not compete with fH for the binding to C3b [40]. CFHR3 competes with fH for the binding to C3b but its role in complement control is not settled [41]. The activity of CFHR4 is also unclear, but it binds C3b and has cofactor activity. CFHR5 possesses both cofactor and decay activities and is also recruited to damaged self-surfaces [42].

C4b-binding protein (C4BP) is a large plasma-circulating glycoprotein. The major form of C4BP consists of seven identical α chains and one β chain, both consisting of CCP domains [43]. The chains are connected by disulfide bridges and the structure is spider-like with seven elongated subunits attached to a relatively small central body [44]. Each α chain contains a binding site for C4b. Once bound to C4BP, C4b serves as a substrate for factor I. C4BP can also act as cofactor in factor I-mediated proteolysis of C3b, although fH and CR1 (see below) are more efficient

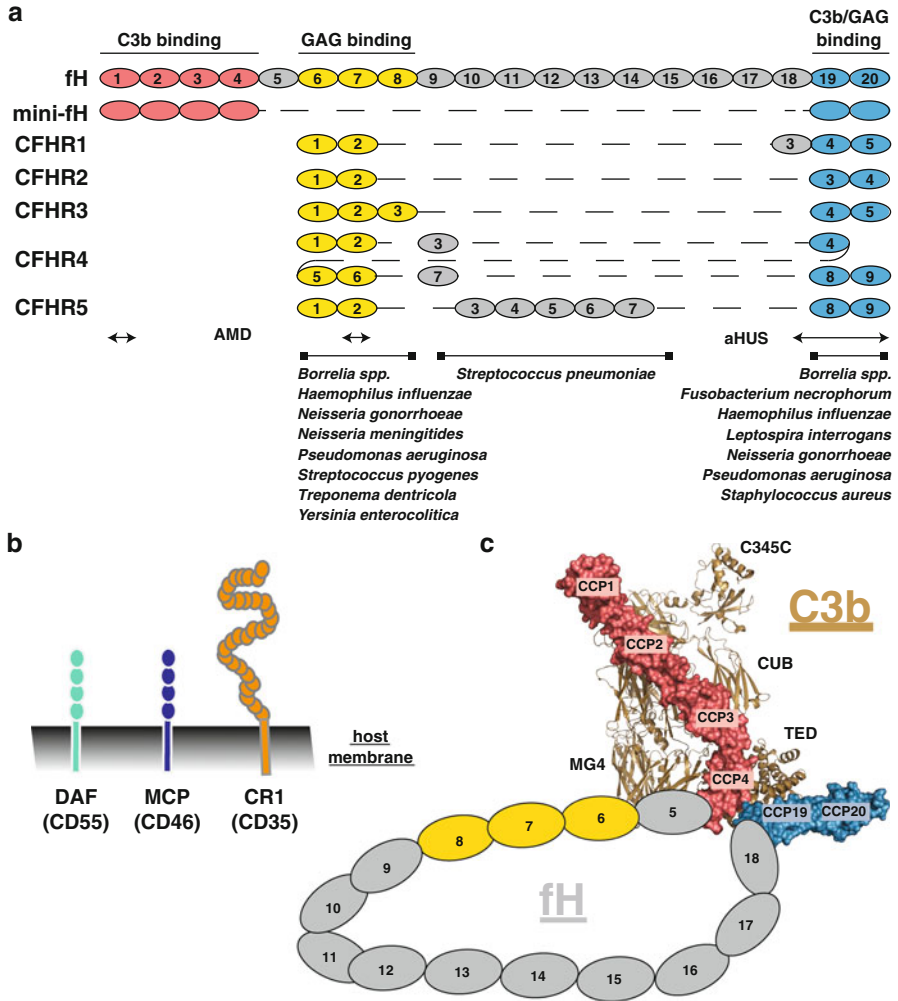


Fig. 2.2 Complement regulators built on complement control protein (CCP) modules. **(a)** Domain organization of human factor H. CCP domains involved in C3b binding are colored *red*, glycosaminoglycan (GAG) binding moieties *yellow*, CCPs binding C3d/GAG *blue*. Therapeutic molecule mini-fH is presented below. Factor H CCPs are conserved in CFHRs and are presented directly below. Regions mutated in AMD and aHUS are delimited with *arrows*. Bacterial species evading complement by binding fH are listed below with the respective fH binding regions. **(b)** Domain organization of the most prominent membrane-bound host regulators. **(c)** Structural model of fH binding to C3b [34, 36] (Color online)

cofactors for C3b. Humans have three CCP-based membrane-bound complement regulators as well (Fig. 2.2b). The first is CD46, also known as the membrane cofactor protein (MCP). It is a cofactor for factor I-mediated cleavage of C3b and C4b into iC3b and iC4b. CD46 is ubiquitously expressed on all nucleated cells, thus only

erythrocytes lack CD46 [45]. The extracellular part of CD46 contains four CCP domains that harbor the C3b and C4b binding sites. The second membrane-bound complement regulator is CD55 or decay accelerating factor (DAF). This is a 70 kDa GPI-anchored glycoprotein expressed on a variety of cells and tissues [46]. Membrane-bound DAF exerts its complement-inhibitory properties by disrupting both AP and CP C3 and C5 convertases, and this function resides within its four CCP domains [47].

Another important complement regulator is complement receptor 1 (CR1). It is a type 1 transmembrane protein composed of numerous CCP domains (44 for the longest allelic variant). CR1 is expressed on almost all peripheral blood cells except platelets, NK- and T-cells [48, 49]. Apart from peripheral blood cells, CR1 is found in some tissues. It plays an important role in the germinal centers of the lymph nodes where it is found on follicular dendritic cells capturing complement-opsonized antigens that serve to stimulate B-cells [50, 51]. CR1 can bind to both C3b and C4b with high affinity and to iC3b and C3d(g) with somewhat lower affinity [52]. Both CP and AP C3 and C5 convertases are inhibited by CR1 via its decay-accelerating activity. It also serves as cofactor for factor I-mediated degradation of C3b and C4b.

Three proteins not based on CCP domains function as MAC assembly inhibitors. Vitronectin (S-protein) binds to C5b-7 through the C5b-7 membrane-binding site and the resulting SC5b-7 complex associates with C8 and three molecules of C9 to form the 1 MDa soluble SC5b-9 complex [53, 54]. Another regulator is clusterin (also called SP-40,40 or apolipoprotein J) which impedes C5b-7 membrane association and the addition of C9 to C5b-8 and C5b-9 [55]. Whereas vitronectin and clusterin are soluble proteins, the third inhibitor of MAC assembly, CD59, is a 20 kDa, GPI-anchored and heavily glycosylated protein widely expressed on almost all tissues and circulating cells [56].

2.1.3 Complement-related Diseases

Uncontrolled and excessive complement activation leads to tissue damage and pathogenesis. The molecular details of how complement proteins contribute to a wide variety of disease conditions and how control of complement may be re-established has recently been extensively reviewed [57–59]. Here we will present some prominent conditions in which animal models, studies of individuals with mutations in complement proteins, and clinical usage of a C5 antibody suggest that therapeutic control of complement is clinically relevant.

The most frequent cause of blindness (50 %) affecting elderly persons is age-related macular degeneration (AMD) with more than 30 million people affected worldwide. AMD causes significant changes in the retinal anatomy and deposition of drusen. This can develop into either neovascular/wet AMD with invasion of blood vessels into the retina or dry AMD with constriction of blood vessels, photoreceptor degeneration, and geographic atrophy [60]. The dry form is currently untreatable whereas wet AMD is treated with intravitreal injections of VEGF antibody.

AMD is strongly correlated with insufficient control of the AP caused by mutations in fH, C3, and fI but also in other complement proteins [61–64]. The best characterized mutant, fH Tyr402His, accounts for 50 % of the heritability of AMD, and causes the altered fH to bind more weakly to host cell glycans as the tyrosine is located in the ligand binding site of fH [32]. Since fH has AP convertase decay and co-factor activity for fI in C3b degradation, host cells are less protected than in the presence of normal fH. The same effect is caused by two C3 mutations for which fH co-factor activity is reduced [65], and fI variants which are less expressed and secreted [66]. Further supporting the link between AMD and high convertase activity, mutations in fB causing weaker association with C3b provide protection against AMD [67]. Despite these connections between AMD and complement, clinical trials with complement inhibitors targeting either the AP (C3 and factor D) or the TP (C5) have so far been disappointing, and so far no complement based therapeutic for AMD has reached phase III clinical trials [60].

Sepsis is a systemic inflammatory condition established by infectious agents, which leads to an excessive immune response resulting in host damage [68]. A 'cytokine storm' occurs and together with intravascular coagulation triggers multi-organ failure. The two C5a receptors C5aR and C5L2 have been implicated in the pathogenesis of sepsis by contributing to the cytokine storm and suppression of the oxidative burst in neutrophils, thereby, hampering the elimination of the infectious agent [69]. In a rat cecal ligation and puncture (CLP) sepsis model, C5a antibodies effectively decreased bacteremia, increased survival, restored H₂O₂ release by blood neutrophils and reduced coagulation [70, 71]. Neutrophil function was also restored in a mouse CLP model upon treatment with a cyclic peptide C5aR antagonist [72]. Furthermore, in a CLP model of sepsis simultaneous blockade of both C5aR and C5L2 with antibodies or a C5aR/C5L2 antagonist (C5a-based antagonist (see below)) increased survival compared to inhibition of the receptors one at a time [73].

Ischemia–reperfusion (I/R) injuries are caused by a reduction of blood flow to tissues and organs followed by re-establishment of the blood flow during which there is an accumulation of leukocytes in the vascular epithelium, upregulation of vascular pro-inflammatory molecules and reactive oxygen species [74]. Complement activation has been observed in many different organs undergoing I/R, including the gastrointestinal system, brain, lung, and kidneys [75]. I/R damage occurs during kidney transplantation, and locally-produced complement proteins are the important mediators of damage in this case [76, 77]. The molecular mechanism of complement-mediated damage during I/R appears to be intricate and tissue-dependent. In renal I/R, the TP plays a central role since both inhibition of C5b-9 assembly in C6 deficient mice and inhibition of C5aR with a small molecule antagonist reduce I/R damage [78, 79]. Whereas MBL-deficient mice are protected [80], absence of C4 does not provide protection [78], suggesting that C3 cleavage bypasses the C4b-based CP C3 convertase. In mouse models of myocardial and gastro-intestinal I/R injury both knockout of MASP-2 and MASP-2 inhibition with antibodies conferred protection and again C3 deposition was found to be independent of C4 [81]. Once C3b has been generated, further amplification takes place through the AP explaining the benefits in a mouse ischemic stroke model of fB absence,

but with no protection afforded by C6 knockout [82]. In a mouse model of myocardial I/R injury absence of MBL and MASP-2 [81, 83] attenuates injury, and interestingly, when given at pharmacological doses, the naturally occurring MAP-1 (Map44) protein competing with MASP-1/2 for binding to MBL and ficolins also inhibits I/R injury in addition to inhibiting thrombosis [84].

Rheumatoid arthritis (RA) is a chronic autoimmune inflammatory condition conferring synovial joint damage [85] to which complement TP has been acknowledged as an important contributor [86]. Aggregation of self-antigen-antibody complexes leads to the formation of complement-activating immune complexes in the synovial tissue [87]. Several complement proteins are involved in RA pathogenesis, but C5 is likely to play a pivotal role in complement-mediated tissue damage within RA [86]. Studies on C5-deficient murine models with type II collagen-induced arthritis (CIA), suggest a role of C5 in the pathogenesis [88], and administration of anti-C5 antibodies in C5-sufficient murine models prevents the onset of CIA and significantly reduces the severity of the disease during active CIA [89]. Signaling of C5a through C5aR and C5L2 is believed to play a significant role in the pathogenesis of RA [90–92] and vaccination with a C5a-fusion protein reduced arthritis severity and incidence in a mouse model [93]. However, somewhat disappointing considering these results from animal models, administration of the C5aR antagonist PMX53 failed to reduce synovial inflammation in RA patients [94].

In humans the terminal pathway and MAC formation is an important contributor to pathophysiology in paroxysmal nocturnal hemoglobinuria (PNH), paroxysmal cold hemoglobinuria [95], and atypical hemolytic urelytic syndrome (aHUS) [96]. PNH is caused by a *PIGA* gene mutation in haemopoietic stem cells resulting in deficiency of all GPI-anchored proteins on progeny cells [97–99] including CD59 and DAF. In PNH patients up to 90 % of erythrocytes are lysed by MAC assembly [100] resulting in anemia, intravascular hemolysis, and thrombosis. PNH is routinely treated by administration of a humanized monoclonal antibody (Eculizumab), preventing C5 cleavage and, thus, the assembly of MAC and the formation of C5a [101]. Eculizumab-treated patients are much more susceptible to neisserial infections and need to be immunized prior to treatment with the C5 antibody.

Asthma is a chronic airway disease characterized by inflammation of the upper respiratory tract, reversible airway obstruction, mucus cell hyperplasia and airway hyperresponsiveness (AHR) [102]. These outcomes are mediated by a T helper type 2 (Th2) polarized immune response, and the anaphylatoxins C3a and C5a have been implicated in both the sensitizing and the effector phases of the disease by regulating the adaptive immune response to allergens [103]. Through studies targeting C5aR by monoclonal antibodies or a C5a-based antagonist, it was suggested that C5a signaling mediates tolerance to aeroallergens by altering the ratio of immunogenic myeloid dendritic cells (mDCs) to tolerogenic plasmacytoid dendritic cells (pDCs), suppressing the Th2 immune response [104]. In contrast to C5a, C3aR knock-out mice develop less pronounced AHR when treated with ovalbumin (OVA) [105], and blocking of complement activation with the recombinant soluble form of the rodent-specific complement regulator Crry decreases airway inflammation in already sensitized mice, by a decrease in both pulmonary eosinophils and immunogenic

Th2 cytokine levels in bronchoalveolar lavage fluid (BALF) [106]. Increased levels of both C3a and C5a were found in BALF from asthmatic patients when compared to normal individuals [107], suggesting that both anaphylatoxins function during the effector phase [106, 108].

2.2 Inhibitors Targeting Complement

2.2.1 Inhibitors Targeting the Convertases

Convertases are obvious targets for complement inhibition, although therapeutics targeting these will in most complement-related disease conditions have to be given systemically, thus, requiring high doses. Convertases targeting has been exploited by pathogens to evade complement and in research and drug development for controlling complement activation at either C3 or C5 cleavage stage [109]. To understand in detail the mechanism of known convertase inhibitors and how to develop new ones, it is helpful to investigate the structure of convertases, their substrates and the complexes these molecules form with their inhibitors (Fig. 2.3). Considerable structural information has been generated concerning the AP C3 convertase. Structures of the proconvertase C3bB by EM and crystallography revealed how fB, in a MIDAS-Mg²⁺ dependent manner, associates with the C-terminal C345c domain of C3b, that is flexibly attached to the remaining relatively rigid part of C3b [110, 111]. The proconvertase can exist in two states, open and closed, differing by a rotation of the fB SP domain, and only in the open conformation is the scissile bond region exposed and accessible to fD, which binds primarily through a fB exosite located 25 Å from the scissile bond [110]. The relevance of this exosite is evidenced by the ability of the anti-factor D mAb, developed for localized complement inhibition in the eye by Genentech, to prevent fD binding to this exosite by steric hindrance [112].

Once the activation of the convertase has taken place it dissociates in minutes, but the *S. aureus* protein SCIN (see below) binds tightly to the AP C3 convertase and prevents it from binding C3. A dimeric form of the C3bBb-SCIN complex in which two SCIN molecules bridge two C3bBb complexes were crystallized and revealed the basic architecture of C3bBb. The only contact here is between the C3b C345c domain and the Bb von Willebrand factor type A (vWA) domain, whereas, the SP domain is extending away from C3b [113]. Deeper insight into substrate recognition by the convertases was obtained with the structure of C5 in complex with the C3b homolog cobra venom factor (CVF). C5 and CVF interact in a head-to-head manner with the long axis of the two molecules aligned and with two separated points of contact. The largest of these is formed between the MG4 and MG5 domains from both proteins. The remaining intermolecular contacts are formed between the C5 MG7 domain and the CVF MG6 and MG7 domains [114]. Compared to the structure of both unbound C5 [115] and human C3 [116], CVF-bound C5 undergoes a significant conformational change that is necessary to establish the two-points interaction, and this conformation has also been captured by crystal

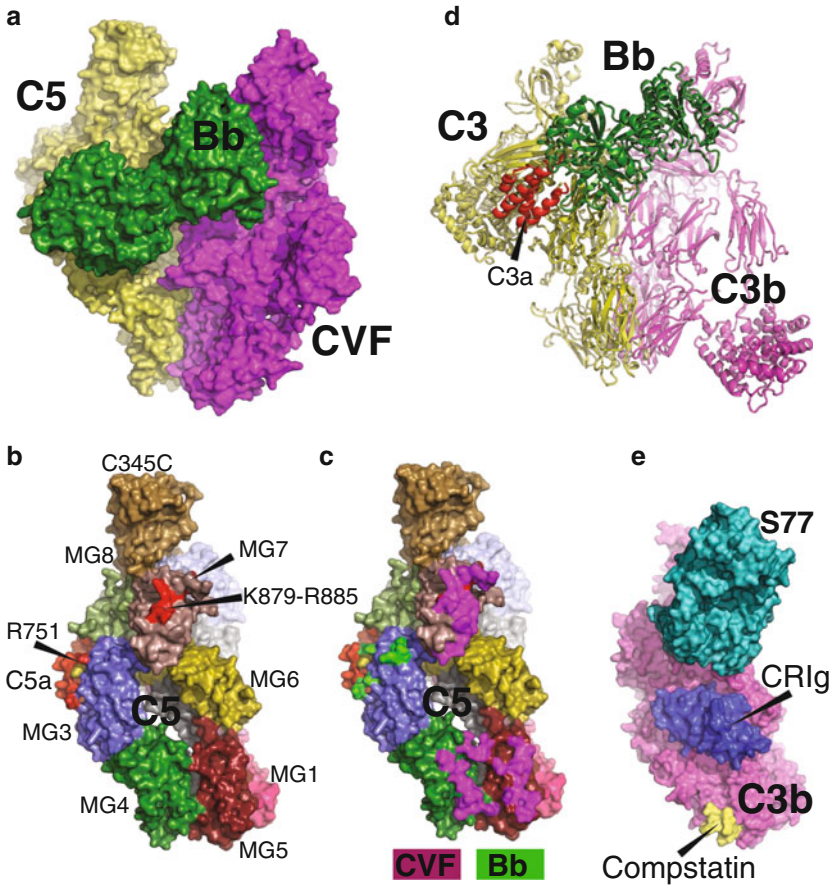


Fig. 2.3 A general model for convertase-substrate complexes explains the inhibition mechanism for convertase inhibitors. **(a)** The model of the CVFBb convertase bound to C5 [114]. **(b)** Complement C5 colored and labelled according to its domains as seen from CVF, the Eculizumab epitope K879-R885 is indicated. **(c)** As in *panel b*, but surface areas in contact with CVF and Bb in *panel a* are colored *magenta* and *green*, respectively. Notice the overlap between the Eculizumab epitope and the CVF-interacting area of the C5 MG7 domain explaining how the antibody prevents C5 cleavage. **(d)** A model of the C3 substrate bound to the AP convertase C3bBb. The C3a domain released is marked in *red*. **(e)** C3b within the AP C3 convertase as seen from the C3 substrate. The three convertase inhibitors S77, CRIg and Compstatin effectively prevent substrate recognition by steric hindrance (Color online)

packing for bovine C3 [117]. The combination of the SCIN-stabilized C3bBb structure and the C5-CVF structure led to the formulation of a general model for convertase-substrate interactions (Fig. 2.3). Since the catalytic subunit C2a/Bb is common for the CP/AP C3 and C5 convertases, it was suggested that the orientation of the substrates C3 and C5 with respect to the catalytic subunit is similar in the two types of convertases [114], although covalent or non-covalent association of either C3

convertase with C3b switches the specificity from C3 to C5 [118–120]. The additional C3b molecule lowers the K_m value for the C5 substrate by a factor of 100–1000 [121, 122], but whether this is through a direct interaction with C5 or induced conformational change in the C3 convertase remains open. The validity of this convertase-substrate model is emphasized by its ability to rationalize how some of the best characterized man-made and microbial complement inhibitors exert their effect. Prominent examples are the antibodies S77 and Eculizumab, the CR1g ectodomain, and the cyclic compstatin peptide, as described in the following.

The S77 mAb interferes with both AP convertases [123], which can be explained by its binding to C3b MG6 and MG7 domains (Fig. 2.3e), that are both predicted to recognize the substrate C3/C5 MG7 domain [114]. A similar inhibitor is the ectodomain of the CR1g complement receptor. This receptor is found on tissue resident macrophages and plays an important function in clearance of pathogens from circulation through interaction with C3b and iC3b on complement-opsonized activators [124]. The binding site on C3b has been mapped to the MG3, MG4, MG5, MG6 domains and the LNK region (Fig. 2.3e), and the ectodomain inhibits AP C3 and C5 convertase activity [125]. The Compstatin peptide is frequently used as a general complement inhibitor and through its binding to C3 it blocks both the CP and AP C3 convertases by interfering with binding of C3 to the convertases. Compstatin also binds to C3b, iC3b and C3c [126]. One disadvantage of compstatin is the high C3 plasma concentration (7 μM) and the rapid clearance of peptides, but new compstatin analogues with sub-nanomolar K_d have substantially increased half-lives [127]. The peptide binds in a groove between the MG4 and MG5 domains of C3 or its fragments [128] (Fig. 2.3e). As these domains in the substrate C3 are predicted to be recognized by the convertase, this explains the ability of compstatins to suppress C3 cleavage by either AP or CP convertases. Like compstatin, the C5 antibody Eculizumab hinders the binding of the substrate C5 to the convertase and, thereby, prevents its cleavage. However, Eculizumab binds to an epitope far from the convertase cleavage site in the MG7 domain centered on residues 879–885 [129, 130], which overlaps substantially with the area of C5 in contact with CVF in their complex [114] (Fig. 2.3b, c).

2.2.2 Regulator-derived C3 and C5 Inhibitors

Over the past years, the development of complement-targeted therapeutics has been inspired by natural complement regulators. The idea behind the rational design of such therapeutics is the use of naturally-occurring host proteins. Strategies involving complement targeting with systemically-administered inhibitors of such type have, however, a few drawbacks. Firstly, inhibiting complement systemically could have detrimental consequences in regard to the host immune defence. Secondly, due to the relatively low affinity of these molecules (μM range) the administered dose would need to be very high. Thus, a more focused strategy is needed. Indeed, direct targeting of the surfaces where complement activation occurs is a more intelligent

and less costly approach. fH would be very attractive in therapeutics as it is a complement modulator that is a self-molecule, acts in the fluid-phase as well as on surfaces and does not interfere with complement defence against pathogens. Purified fH was shown to be able to control the C3 convertase in a mouse model of fH deficiency [131]. In vitro assays in the presence of aHUS-associated anti-fH autoantibodies showed that fH protected self-cells from complement. However, due to its size (155 kDa), presence of glycans and 40 disulfide bridges, recombinant production of full-length fH is challenging for routine administration to patients.

By the use of protein engineering it has been possible to develop mini-fH, a compact version of fH harbouring fH CCP 1–4 linked to CCP 19–20 (Fig. 2.2a). This molecule contains fH binding sites to C3b/iC3b/C3d, retains its decay acceleration and cofactor activity and, very importantly, has the ability of fH to recognize host surface-specific glycans. Interestingly, this molecule surpasses fH in both affinity towards C3 activation products and the ability to control AP activation [132, 133]. Mini-fH has potential as a therapeutic molecule in complement-mediated diseases such as aHUS, PNH or C3 glomerulopathies although its clinical properties remain to be investigated. Another inhibitor based on the fH ability to bind C3b is TT30 [134]. It is a chimeric protein developed by Alexion containing fH regulatory domains CCP 1–5 and CR2 CCP 1–4 responsible for binding to the C3b thioester domain. TT30 was extensively tested in the case of complement-mediated hemolysis and found to inhibit MAC formation resulting from AP but not CP or LP activation.

Part of the recent research in complement therapeutics has been focused on preventing complement activation on blood-exposed materials, such as implants. A great deal of research has been evolving around heparin as it has been expected to recruit fH and inhibit complement activation on the surface of biomaterials. As expected, coating the biomaterials with high densities of heparin resulted in inhibition of complement activation [135–137] but it was later found that this inhibition was fH-independent [138]. An alternative approach, employed recently, consists of coating the bioimplant surfaces with fH binding peptides but not interfering with its activity as complement regulator [139] but this approach remains to be tested in a clinical setting.

Since CR1 possesses cofactor activity for factor I-mediated proteolysis of C3b and C4b, its potential in therapeutics has been extensively studied. Soluble recombinant CR1 (sCR1) was studied in animal models of autoimmune and inflammatory disorders such as glomerulonephritis [140], myocardial infarction [141] and autoimmune thyroiditis [142]. In mice, administration of sCR1 resulted in resolution of inflammation and these encouraging results drove Avant Immunotherapeutics to develop sCR1 (TP10) for managing complement activation following coronary artery bypass graft surgery. TP10 was shown to be safe and well-tolerated in patients. However, its efficacy was far greater in male patients, and this formulation has been withdrawn from the market but its potential as a therapeutic agent has not been disputed. In fact, it has been successfully used in I/R injury when administered intravenously [143]. Nevertheless, since it is produced in Chinese hamster ovary cells as a 240 kDa glycoprotein [141, 144] and has to be injected systemically, it is

an expensive therapeutic agent. An alternative approach has been used for the development of Microcept [145], a molecule readily produced in *E. coli* containing only the first 3 CCP domains of CR1, sufficient for inhibition of C3 and C5 activation by the AP and CP. Microcept contains a synthetic thiol-reactive myristoylated basic peptide tail attached through a C-terminal cysteine residue allowing its insertion into biological membranes and, thus, targeting and protecting the cells directly.

CD59 is an important complement regulator as evidenced by PNH patients, and the protein has been explored as a therapeutic agent since it only blocks MAC formation. Although CD59 has little effect as MAC inhibitor when administered systemically in a soluble form, if targeted to the sites of MAC formation, recombinant CD59 could be an effective complement modulator. Several strategies have explored generating CD59 chimeric proteins. One of them consisted of fusing CD59 to DAF and a GPI anchor, thus allowing its insertion into the membranes and its localization where C3 and C5 convertases are present [146]. Another strategy involved fusing CD59 to CR2 and DAF to CR2 [147]. These CR2 fusion proteins were efficient complement modulators in mouse models of lupus nephritis. In conclusion, fusing CD59 to a cell-surface targeting moiety may be an efficient therapeutic strategy in disease conditions where only MAC formation needs to be inhibited.

2.2.3 *Inhibitors Targeting the Anaphylatoxin/Receptor Axis*

The anaphylatoxins C3a and C5a produced by the complement proteolytic cascade and their associated receptors are important targets for the therapeutical treatment of inflammatory disorders. C3a and C5a mediate their pro-inflammatory effects by signaling through the G-coupled protein receptors C3aR and C5aR, respectively. This leads to chemotaxis, oxidative burst, production of pro-inflammatory molecules and activation of the adaptive immune system [148]. C3a is generally considered as a weaker pro-inflammatory inducer than C5a. To control the signaling exerted by both anaphylatoxins, carboxypeptidases cleave their C-terminal arginine resulting in C5a-desArg and C3a-desArg. C5a-desArg partially maintains C5aR binding and signaling activity whereas C3a-desArg is devoid of any signaling through C3aR [149, 150]. The 7TM receptor C5L2 binding C5a, C5a-desArg, C3a, and C3a-desArg, has long been considered as a decoy receptor, but there are now reports suggesting that it actively participates in orchestrating pro-inflammatory events [151]. Plasma levels of anaphylatoxins are elevated in various disease settings, and they can be used as biomarkers in a number of inflammatory disorders [148].

Targeting of the anaphylatoxin-receptor axis provides a selective way of down-regulating complement-mediated inflammation and is, therefore, an attractive therapeutic strategy since opsonization and MAC formation are preserved. Focus has so far been given to the C5a-C5aR axis, although C3aR antagonists are also in development as extensively reviewed recently [152]. In humans, C5a is a glycosylated protein that adopts either a compact four-helix bundle core with a flexible C-terminal extension [153] or a three-helix bundle for C5a-desArg [154]. The proposed binding

interface for C5aR on C5a is hidden within the C5 molecule [115]. Thus targeting molecules can mostly be directed either towards the exposed C5a surface present on C5 or towards the C5aR binding surface only available on the released anaphylatoxin. The first strategy presents more risks since it can generate global C5 inhibitors, if large inhibitory molecules are employed impairing C5 cleavage and, thereby, the entire TP. This effect is in some clinical contexts undesirable, e.g. in the treatment of sepsis. Inhibitors directly interfering with C5a-C5aR interaction while preserving other C5 functions are, therefore, more appealing.

Several groups and companies have developed potent C5a monoclonal antibodies [155]. The use of C5a antibodies for preventing multiple organ failure and improving survival rate in sepsis has been documented since the 1980s [156]. Of interest among these, the anti-C5a mAb 137-26 directly binds to the C5a moiety on C5 without inhibiting C5 cleavage and subsequent MAC formation [157]. It is commercialized as TNX558 by Tanox/Genentech and has been in preclinical development for inflammatory diseases [158]. Another humanized C5a monoclonal antibody, CaCP29 (IFX-1), developed by InflaRx GmbH, has passed Phase I clinical trials in Germany for human sepsis [159]. Aptamer approaches have also been considered for the blockade of C5a function. NOXXON Pharma has developed a class of aptamers called Spiegelmers® [160] built on nucleotides containing L-ribose making them the mirror images of D-ribose containing RNA. As therapeutics, Spiegelmers are much more resistant to nuclease degradation compared to conventional aptamers giving high stability in the blood. C5a has been efficiently targeted by Spiegelmers® NOX-D19 and NOX-D20, which have shown promising results in reducing vascular injuries after transplantation and in attenuating organ damage during sepsis [151, 161]. NOX-D20 binds to both human and murine C5a with picomolar affinities but also human C5 with similar affinity although NOX-D20 does not prevent C5 cleavage. These mirror-image L-RNA aptamers, therefore, appear to be promising anti-C5a therapeutics. An alternative strategy involving immunization with a recombinant MBP-C5a fusion protein resulted in the production of neutralizing C5a antibodies [93]. A new study instead used a recombinant C5a molecule modified with unnatural amino acids. A single replacement was sufficient to induce the production of anti-C5a antibodies capable of blocking the C5a-C5aR interaction, leading to significant relief of the clinical symptoms in a mouse model of rheumatoid arthritis [162].

In relation to C5a-targeting molecules the anaphylatoxin receptors have also been a focus for inhibitor development. A large number of C5aR inhibitors have been developed. A well described cyclic peptide antagonist known as PMX53 or 3D53 targets C5aR and competes with C5a binding to human polymorphonuclear neutrophils (PMNs) with an IC_{50} value of 300 nM, but does not bind human C5L2 [148, 163]. The antagonistic activity of PMX53 was measured, showing an inhibition of myeloperoxidase release from PMNs with an IC_{50} of 20 nM [163]. NMR studies of PMX53 suggests a β -turn motif in the molecule [164]. Mitsubishi Pharmaceuticals Company developed the orally active small molecule C5aR antagonist W54011 with a K_i of 2.2 nM. It inhibits calcium mobilization in human neutrophils and C5a-induced neutropenia in gerbils, and shows no inhibition of

C5a-binding to C5L2 [165, 166]. Another small molecule, NDT9513727, specifically targeting C5aR, reduces the constitutive GTP γ [³⁵S] binding of human C5aR-coupled G-proteins, and, therefore, functions as an inverse agonist [167].

Proteins have also been selected to target C5aR. Developing human C5aR knock-in mice responding to endogenously produced C5a facilitated the selection of anti-C5aR antibodies showing promising results in preventing and reversing serum-induced inflammation in the knock-in mice. The most potent anti-C5aR antibodies bind to the second extracellular loop of the receptor [90], which was previously identified as important for balancing the activity of C5aR, since mutations in this region resulted in constitutively active receptors [168]. The C5a molecule has also been exploited as a scaffold to generate potent competitors of C5a receptors, and from a phage library an antagonist called Δ pIII-A8 was selected which reduced intestinal injury and lung vascular permeability and increased survival of (I/R) injured mice [169]. A shortened version (A8 ^{Δ 71-73}) targeting both C5aR and C5L2 was developed later and, interestingly, it was found that one particular residue determines agonism versus antagonism of A8 related proteins. In C5a, this amino acid is an aspartate and in A8 ^{Δ 71-73} it is mutated to an arginine [170].

For C3aR inhibitor developments have mainly focused on peptide analogs of the C-terminal region of C3a. Hexapeptides mimicking the C-terminus of C3a were synthesized resulting in both agonists and antagonists. The most potent agonists shared an N-terminal phenylalanine, a tryptophan or leucine at the second position and the highly conserved C-terminal sequence Leu-Ala-Arg. Substituting the fourth leucine to the bulky cyclohexylalanine resulted in antagonists [171]. By NMR, one of the most potent agonistic peptides was found to adopt a β -turn motif similar to C5aR ligands [172]. As an alternative to peptides, small-molecule compounds targeting C3aR have also been discovered, with an example being the functional antagonist SB290157 with an IC₅₀ of 200 nM [173]. Although this drug showed anti-inflammatory activity in a guinea pig model [174], a more recent paper found this molecule to have partial agonistic activity, a conflict that might be explained by differences in receptor density in the systems used [175]. Numerous other drugs including proteins, peptides and small molecules have been developed targeting the anaphylatoxin receptors and have been extensively reviewed elsewhere [148, 152, 176].

2.3 Bacterial Strategies for Immune Evasion: What Can Be Learnt from Them

Complement primary function resides in the host defence against pathogens, but many pathogens successfully evade complement [177]. One can learn from these evasion strategies on how to inhibit complement at various stages of the cascade and apply this knowledge to design inhibitors for therapeutic applications. Many pathogen inhibitors have also proven useful to gain structural insights into complement mechanistic by freezing complement proteins and their complexes in specific functional states with the ability of the *S. aureus* protein SCIN to stabilize the

otherwise rapidly dissociating C3 convertase as the best example. We will review here some of the best characterized pathogen inhibitors and try to comprehend how their interaction with complement may inspire attempts to make potent, selective inhibitors of complement.

2.3.1 Pathogen Inhibitors Targeting C3

Staphylococci are quite versatile organisms with respect to complement evasion. Targets of choice for staphylococcal proteins are the C3 and C5 convertases [178] as already described for SCIN above. Three other potent complement inhibitors from *Staphylococcus aureus* - Efb, the closely related Ehp, and Sbi - are binders of C3 and its cleavage products, C3b and iC3b, through preferential interaction with the C3 thioester domain, with the C3b degradation product C3d as the minimal binding partner. Crystal structures of the Ehc:C3d, Efb-C:C3d and Sbi:C3d complexes revealed that their primary binding site on C3d is partially inaccessible within the intact C3b molecule [179–181] (Fig. 2.4a). As a consequence of their binding to this hindered site, Efb and Ehp induce an altered overall conformation of C3 and C3b. Displacement of the C3d thioester domain is relayed to the rest of the molecule, therefore preventing its further participation in the downstream events of the complement cascade, including formation of the AP convertases and covalent deposition on the pathogen surface. In the case of Sbi, a secondary binding site on C3d allows the inhibitory protein to form a covalent adduct with activated C3 forms (C3b and/or C3(H₂O)), thereby directly interfering with AP activation [181] (Fig. 2.4a). In addition, all three inhibitors compete with complement receptor 2 for the binding of C3d and, therefore, impede the stimulation of B cells mediated by this receptor [182].

2.3.2 Pathogen Inhibitors Targeting C5

Several pathogen inhibitors block the C5 convertase by interacting with C5. OmCI, a small protein from the soft tick *Ornithodoros moubata*, binds directly to C5 and inhibits its cleavage into C5a and C5b [183]. Structural studies of the OmCI:C5 complex suggested that OmCI binding to C5 fixes the C5 C345c domain and inhibits convertase binding to C5 by impairing the conformational flexibility of this domain [115]. A recombinant version of OmCI, rev576, has shown promising results in models of auto-immune neuromuscular diseases and sepsis [184–186]. The SSL7 protein from *Staphylococcus aureus* likewise binds C5 and prevents its proteolytic processing by the C5 convertase, thereby impairing MAC-mediated bacteriolysis and C5a release. Its mechanism of action was elucidated through structural analysis of the C5:SSL7 complex [187]. SSL7 binds to a surface patch quite distant from the C5 cleavage site suggesting a more complex mechanism than

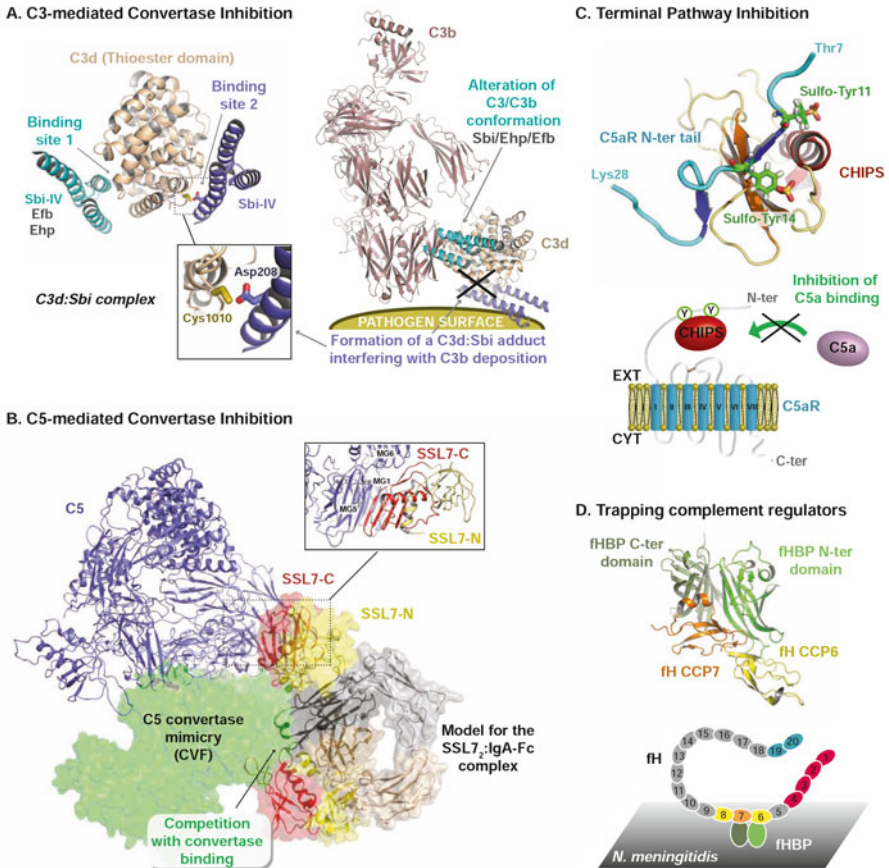


Fig. 2.4 Complement evasion by pathogens. (a) The two binding modes of staphylococcal protein Sbi onto C3d [181] and their docking onto C3b [210], revealing the structural model for their inhibition of the C3 convertase function. (b) Structural model for SSL7 inhibition of the C5 convertase by dual recruitment of C5 and IgA, based on the C5:SSL7 [187], C5:CVF [114] and SSL7:IgA-Fc [211] complex structures. (c) Structure of the inhibitor CHIPS in complex with a peptidic fragment of C5aR [193] and model for its inhibition of C5a binding by blocking the C5aR N-terminal docking site around the two sulfo-tyrosines. (d) Structure of the complex between fH CCPs 6–7 and fH-binding protein (fHBP) from *N. meningitidis* [199] and scheme for fH trapping on the pathogen surface

simple steric hindrance of the C5 cleavage site. This was later confirmed by the structure of the ternary complex between C5, SSL7 and CVF [114]. Together these studies explained how simultaneous recruitment of C5 by SSL7 C-terminal domain and IgA by SSL7 N-terminal domain prevented convertase recognition of C5 (Fig. 2.4b). Interestingly, the isolated C-terminal domain of SSL7 devoid of affinity for IgA, did not prevent C5 cleavage and permitted bacteriolysis while preserving a very low hemolytic activity on erythrocytes [187] suggesting that this domain may

indicate a direction for development of a therapeutic agent targeting complement-dependent hemolytic diseases.

Bacterial proteins that directly target MAC formation have also been reported. Streptococcal Inhibitor of Complement (SIC) and variants from other *Streptococcus* strains bind to the C5b67 complex hindering membrane insertion [188]. An outer membrane protein from *Escherichia coli* K12 strain, TraT, also impairs MAC formation by direct binding to C5b6 preventing C7 recruitment [189]. *Borrelia burgdorferi* produces a CD59-like protein that blocks C9 polymerization [190]. Pathogen proteases are also potent effectors in complement evasion and C5a peptidases are found in various organisms. In particular, group A streptococci produce a cell-envelope proteinase (ScpA or ScpB) cleaving off the seven last residues from the C5a C-terminus, thereby shifting C5a from agonistic to antagonistic activity [190]. Proteinases from *Porphyromonas gingivalis* are also capable of shedding off the N-terminal region of C5aR [191], thus inactivating the receptor. Direct inhibition of C5aR is also achieved by pathogenic virulence factors. A well-studied case is the secreted chemotaxis inhibitory protein of *Staphylococcus aureus* (CHIPS) binding C5aR with a K_d of 1.1 nM [192]. An NMR structure of CHIPS in complex with a peptide mimicking the N-terminus of C5aR containing *O*-sulfated tyrosines revealed how CHIPS competitively antagonizes the interaction of the core structure of C5a with the receptor [193] (Fig. 2.4c). CHIPS or derivatives of this was suggested to serve as an anti-inflammatory therapeutic agent but this was questioned due to the high immunogenicity of CHIPS [194]. However, mutated or shortened versions of CHIPS with lower immunogenicity maintain the C5aR antagonizing effect [195, 196].

2.3.3 Inhibition/Trapping of the Complement Regulators

Bacteria and viruses have also developed approaches to hijack complement components to their own advantage with complement regulators fH, fHL-1 and C4BP being the preferred victims. The plethora of bacterial fH binders includes complement regulator-acquiring surface proteins (CRASP) and outer surface protein E (OspE) from *Borrelia burgdorferi*, M proteins, Fba, Scl1.6 and Hic from *Streptococci*, and fH binding protein (fHBP) from *Neisseria meningitidis*. CRASP-1 binds to CCP5–7 of fH as a dimer and clamps the bound CCP domains, thereby enhancing scavenging of the complement regulator on the pathogen cell surface [197]. OspE and neisserial FHBP, on the other hand, use a protein mimicry of the host carbohydrates to sequester fH by targeting either CCP19–20 [198] or CCP6–7 [199] (Fig. 2.4d). In both cases, the binding epitopes on fH are similar to the ones proposed for glycosaminoglycans binding on endothelial cells. Finally, certain viruses also possess complement regulators, such as the vaccinia virus complement-control protein (VCP) and the smallpox inhibitor of complement enzyme (SPICE) which both show cofactor activity for degradation of C3b [200]. <http://www.nature.com/nrmicro/journal/v6/n2/full/nrmicro1824.html-B27>

2.4 Discussion and Concluding Remarks

Although the concept of complement as a cornerstone in defence against microbial invasion and in homeostasis still holds true, there is also clearly a dark side to this highly complex machinery. The capacity of complement to recognize danger-associated molecular patterns from both pathogens and host cells and its interconnectivity with other branches of the innate and adaptive immune system and even the coagulation system [2], adds on to difficulty of maintaining tight control of the system. Evidently, any breach in its regulation may disturb the fine balance between protection and damage. The list of diseases that implicate complement as one of the causative elements keeps growing and encompasses a broad panel of pathologies ranging from inflammatory diseases to neurodegenerative disorders and cancers [59, 201]. Our knowledge of the intricate mechanisms at work during complement activation and complement crosstalk with parallel defence systems has gained considerable depth over the last decade, not least due to the increasing amount of available structural and biochemical data allowing a comprehension of the complement cascade in atomic details.

The success of Eculizumab [101] has encouraged many new initiatives in the field. A constantly growing number of molecules arising from both academic and industrial research efforts are now under development, with already promising preliminary results in clinical studies for several of them, as reviewed here. Many classes of molecules have been considered for these drug candidates, including antibodies, aptamers, small chemical compounds and recombinant versions of naturally occurring proteic inhibitors, e.g. based on bacterial inhibitors or host regulators. The constantly improved understanding of the mode of action of complement regulators and receptors has led to the conception of new generations of multimodular inhibitors incorporating functionalities of one or several complement regulators for efficient convertase decay and/or opsonin degradation [57]. Although many regulatory concepts have already been exploited, new ideas are emerging thanks to the availability of both functional and structural data allowing comprehending the mechanistic of complement activation at the molecular level. As one example, recent structures of C4 and its complex with MASP-2 revealed an important exosite interaction of MASP-2 with the C4 C345c domain and it could be demonstrated that a recombinant version of this domain functions as a CP/LP pathway inhibitor [202].

Evidently, targeting of complement at specific stages of the cascade is highly desirable to allow selective containment of the dysregulated pathway while preserving protective functions of the overall immune system. While targeting of central complement components such as the C3 convertase will lead to complete shutdown of the system, more refined approaches can be directed towards the initiation or the terminal steps. Interfering with the initial steps of complement activation may, for example, offer a way to specifically target one of the activation pathways while retaining normal complement defence functions through the other

untargeted pathways—granted that the identified pathway is the major contributor to the pathological condition.

Targeting of the terminal steps of complement cascade has already been extensively exploited through the inhibition of C5a-mediated signaling [70, 152] and the blockade of C5 by Eculizumab [95, 101]. Nevertheless, improvement of the known strategies may still be needed as exemplified by Eculizumab. Although the antibody effectively prevents hemolytic activity by impairing MAC formation, it does not interfere with AP complement activation and subsequent opsonization of PNH cells, which thus are still preferentially marked for extravascular lysis [203]. Furthermore, the increased susceptibility to neisserial infections of the patients treated with Eculizumab still constitutes a disadvantage [204]. Another major concern for life-long Eculizumab treatment for PNH and for some aHUS patients is the annual cost of €460,000 for an adult [130]. Despite these drawbacks, no efficient substitute has been produced yet and clinical trials for the use of Eculizumab in acute inflammatory disorders, such as antibody-mediated transplant graft rejection, are currently conducted [205]. In such settings where C5aR-mediated signaling is seen as a major contributor to the underlying inflammation, a C5a or C5aR antagonist might simply be a more promising drug candidate [203].

An emerging idea is that the inhibitor design has to be rethought for each particular disease [57]. During acute inflammation (e.g. in sepsis), large amounts of complement effectors (C3b, C4b, C5b, anaphylatoxins) will be produced. Thus, efficient inhibitors should have fast, high affinity binding capacities towards their target and a slow dissociation rate, to allow rapid and complete blocking of the complement cascade. In chronic inflammation, on the other hand, a milder modulation of complement may be sufficient to re-establish a proper balance between protective function and injury—complete shutdown of the system being avoidable in that case. Another trend within complement inhibitors is the specific cell/tissue targeting of these [206]. Local delivery at the site of injury is highly desirable if one wants to preserve systemic complement function. Efforts in that sense have already been made for example by fusing regulator-mimicking molecules to targeting modules in order to deliver engineered versions of these inhibitors to sites of complement activation, with the factor H-CR2 fusion protein TT30 as an excellent example [134, 207, 208]. Such targeting approaches may in the future be combined with nanoparticle-based delivery systems for delivery of anti-inflammatory drugs. More thoughts are to be put in this design strategy and organ- or tissue-specific delivering strategies used for other systems should be addressed as well in the complement therapeutics field. Finally, the control of complement activation may become an important issue for the future success of nanomedicine as surfaces considered as “foreign” to the complement system are introduced into the human body and often elicit activation of the system [209] and deeper structural knowledge of the mechanisms at play during complement activation will undoubtedly provide new tools to successfully overcome these challenges.

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Chapter 3

The Art of Complement: Complement Sensing of Nanoparticles and Consequences

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Abstract The complement system is a complex network of plasma and membrane-associated proteins and represents one of the major effector mechanisms of the innate immune system. The function of complement in innate host defence is accomplished through highly efficient and tightly orchestrated opsonisation, lytic and inflammatory processes. Nanoparticle-based medicines may trigger complement and a number of consequences ensue from complement activation. These comprise both beneficial and adverse reactions, depending on the extent and severity of complement activation as well as microenvironmental factors. These concepts are briefly discussed in relation to therapeutic applications of nanoparticles and anti-cancer nanomedicines.

Keywords Adjuvanticity • Adverse reactions • Complement system • Immune responses • Nanomedicine • Nanoparticles • Opsonization • Phagocytes

3.1 Introduction

Synthetic nanoparticles and particulate drug carriers (e.g., liposomes, polymeric nanoparticles and nanocapsules) by virtue of their size, shape and surface characteristics (e.g., display of architecture with repetitive epitopes such as those arising from surface projected polymers or polyanionic/polycationic clusters) may resemble viruses or microorganisms. These ‘pathogen-mimicking’ properties make nanoparticles prone to interception by different components of the host defences following entry into the body [1–4]. Accordingly, the type and the extent of immune responses will depend on physicochemical characteristics of nanoparticles,

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administered dose, frequency and route of administration [2]. A key component of immunity that plays a central role in the sensing of particulate matters, their processing and elimination is the complement system [5].

3.2 The Complement System

The complement system, which consists of at least 30 soluble and membrane-bound proteins, is a key effector of both innate and cognate immunity [6]. Complement acts in a wide variety of host defence, inflammatory, homeostatic and immune reactions [6]. Complement recognizes danger signals primarily through pattern recognition. Indeed, many particulate drug carriers and nanomedicines are composed of polymeric components and other patterned nanostructures that makes them prone to complement sensing [5]. Complement activation proceeds through a series of enzymatic reactions that lead to the assembly of the lytic membrane attack complex (MAC) for target destruction [6]. There are three established pathways of complement activation: the classical, the lectin and the alternative pathways. Each pathway is triggered on binding of complement initiating molecules such as C1q, mannan-binding lectin (MBL), ficolins and properdin (either individually or in combination) to the target, but they converge to generate the same set of effector molecules [6]. For details of complement activation pathways and enzymatic cascades the reader is referred to excellent reviews elsewhere [5, 6]. Here, we limit our discussion to sensing of nanoparticles by complement initiation molecules and consequences of nanoparticle-mediated complement activation.

3.2.1 *Complement Initiating Molecules in Nanoparticle-Mediated Complement Activation*

The initiating molecule of the classical pathway is C1q, a pattern-recognition molecule composed of six identical subunits with cationic globular heads and long collagen-like tails [6, 7]. The classical pathway activation occurs when the globular head regions of C1q binds either directly to polyanionic surfaces (e.g., certain classes of negatively charged liposomes and polymeric nanoparticles such as polystyrene nanospheres) [5, 8, 9] or particulate materials that have become coated with immunoglobulins (e.g., IgM, IgG1, IgG2 and IgG3 classes) [10, 11] or C-reactive protein [12]. For instance, liposomes on binding to natural anti-phospholipid and anti-cholesterol antibodies, can trigger C1q-dependent classical pathway of the complement system [5, 8, 9], whereas lysophosphatidylcholine vesicles do so through initial binding of C-reactive protein [13].

MBL (also known as mannan-binding lectin, mannan- or mannose-binding protein) and ficolins are the initiating molecules of the lectin pathway and recognize nanometre scale oligosaccharide-based or acetyl-based molecular patterns on surfaces [5, 6, 14].

The role of lectin pathway in complement activation by nanoparticles displaying surface projected poly(ethylene oxide) has been demonstrated [15–17]. Indeed, surface projected poly(ethylene oxide) chains in close proximity may form dynamic ‘pathogen-mimicking’ clusters transiently resembling structural motifs of the D-mannose, which serves as a platform for MBL/ficolin docking. Also, in mouse sera MBL was found to bind to dextran-coated iron oxide nanoparticles [18], although the significance of this finding to complement activation and pharmacokinetic of diagnostic iron oxide nanocrystals in humans remains to be evaluated.

Initiation of the alternative pathway activation requires the presence of preformed C3b (arises from enzymatic cleavage of the third complement protein, C3), or C3(H₂O) (a meta-stable form of C3 arising from spontaneous hydrolysis of the thioester in C3, which has C3b-like properties) [5, 6]. It is the internal thioester bond in the α -chain of C3b, which undergoes a nucleophilic attack in the presence of a foreign surface structure rich in nucleophilic groups [6]. There are many examples of liposomes, oil-in-water emulsions, polymeric nanoparticles and metallic nanoparticles (e.g., gold nanoparticles and iron oxide nanoparticles, etc.) that activate complement through the alternative pathway [5]. For instance, a recent study has shown that nanoparticles grafted with a glucose-containing polymer trigger the alternative pathway of complement at a lower grafting density compared with nanoparticles grafted with a galactose-containing polymer [19]. However, no explanation was provided on the mechanisms of the initiation steps, but galactose polymer modified nanoparticles adsorbed more factor H (a negative regulator of complement) [6], than the glucose surface, providing a reason for its lower level of complement activation. In contrast to these observations, others have demonstrated a unique set of conformational changes related to a target adsorbed form of intact C3, for instance through transformation into C3(H₂O), leading to initiation of complement activation via the alternative pathway [19]. For example, zwitterionic dimyristoylphosphatidylcholine (DMPC) liposomes trigger the alternative pathway through contact activation of C3(H₂O) and not C3b binding [20]. However, these studies were performed with liposomes that were frozen inappropriately without the use of cryo-preserves. Accordingly, on thawing these vesicles may contain many membrane defects and altered morphologies that may explain the binding of C3(H₂O). In our hand, complement activation in human serum/plasma by freshly prepared DMPC unilamellar vesicles of 100 nm in size only proceeds through the classical pathway (Moghimi, unpublished data). Another mode by which the alternative pathway turnover can be enhanced is through sufficient surface deposition of properdin. Properdin is a highly cationic pattern-recognition protein that binds to certain polymers, nucleic acids, anionic clusters and apoptotic cells [15, 21, 22]. For instance, polymeric nanoparticles displaying surface projected poly(ethylene oxide) chains accelerate alternative pathway turnover through binding of both nascent C3b and properdin, but the latter is controlled by poly(ethylene oxide) surface density and conformation [15].

The literature surrounding the role of nanoparticle size in complement activation pathways and processes is somewhat confusing [2, 5, 23–27]. The reported discrepancies may be a reflection of misunderstanding of the effect of surface

curvature in modulating the affinity and topological strain of complement initiating molecules such as IgM and C1q [25, 26, 28, 29].

3.3 Consequences of Nanoparticle-Mediated Complement Activation and Fixation

3.3.1 Opsonisation and Phagocytic Elimination

Complement activation and fixation remains a central component for efficient clearance and destruction of particulate invaders at the hand of phagocytic cells. The C3 molecule is central to opsonisation [6, 30]. Its first cleavage product, C3b, acts as an opsonin and becomes covalently bound to the activating nanoparticle surface and facilitates nanoparticle binding to phagocytes via complement receptor (CR) 1 [6, 30, 31]. C3b is further degraded to iC3b, C3c and C3dg, products that serve as ligands for other complement receptors on leukocytes [31]. For instance, iC3b is the primary ligand of CR3 (Mac-1, CD11b/CD18) and CR4 (CD11c/CD18, p150,95), but CR3 is the predominant receptor for phagocytic recognition and safer elimination of complement-opsonised particles and complexes [31].

Complement opsonisation and subsequent macrophage recognition of particulate drug carriers offers an unprecedented opportunity for intravenous delivery of therapeutic agents and immunomodulators to phagocytic cells in the liver (Kupffer cells) and the spleen (marginal zone and the red-pulp macrophages) [1–3]. In murine, Kupffer cells express a CR of the immunoglobulin superfamily (CRIg) that binds C3b- and iC3b-opsonized particles [32]. Depending on the species, complement opsonized particles may further interact with CRs of erythrocytes (CR1 in primates) [33] and platelets (CR1 in rats) [34, 35]. These modes of interaction may further modulate nanoparticle pharmacokinetics and biological fate.

Some studies have now confirmed that stealth nanoparticles such as PEGylated liposomes and polymeric nanospheres although fixing complement, still show considerable resistance to macrophage phagocytosis [36, 37]. This is due to the strong steric barrier of PEG on the nanoparticle surface, which interferes with the binding of surface localized C3b and iC3b to their corresponding phagocyte CRs.

3.3.2 Nanoparticle Integrity in the Blood

Complement activation can significantly affect the integrity of certain classes of drug carriers in the blood. For example, insertion of MAC into the liposomal bilayer may lead to substantial leakage of entrapped (aqueous) cargo [38], but this may be minimized or prevented through surface PEGylation [37]. Similarly, MAC may also affect the integrity of colloidal dispersions of cubosomes and hexosomes

(liquid crystal phases of cubic and hexagonal structures in water [39]), which also act as drug carriers.

3.3.3 *Adjuvanticity*

Nanoparticles have been used as immune potentiators or adjuvants triggering elements of innate immunity that subsequently assist the generation of potent and persistent adaptive immune responses [40]. Most of these efforts are being directed to enhance the immunogenicity of subunit vaccines through both antigen protection and targeting to antigen-presenting cells as well as immunostimulation. The role of the complement system in some of these processes is rather intriguing. For instance, some complement activation products (e.g., C3d) can induce B lymphocyte activation [41]. Likewise, simultaneous binding of CR3 and CR2 to C3 fragments may modulate trafficking of complement-opsonised immune complexes and nanoparticles from macrophages to B lymphocytes and follicular dendritic cells in lymph nodes [42].

3.3.4 *Nanoparticle-Mediated Infusion-Related Adverse Reactions*

Acute allergic-like reactions with haemodynamic, respiratory, cardiovascular, cutaneous and gastrointestinal manifestations, which are not initiated or mediated by pre-existing IgE antibodies, have been reported to occur in approximately 45 % of individuals within a few minutes of infusion of nanomedicines (e.g., liposomal drugs such as Abelcet[®], Ambisome[®], DaunoXome[®], Doxil[®], Myocet[®] and Visudyne[®]; micelle-solubilized drugs such as Taxol[®], Taxotere[®] and Vumon[®]) and diagnostic nanoparticles (e.g., dextran-coated iron oxide nanocrystals such as Feridex[®] and Combidex[®]) [43, 44]. In some isolated cases this has been fatal (e.g., Taxol[®]) [44]. Compelling evidence suggests that inadvertent activation of the complement system is an important factor in eliciting these reactions [43–46]. This is partly due to the liberation of potent complement bioactive products (e.g., C3a, C5a and C5b-9) with the ability to modulate the function of a variety of immune cells (e.g., monocytes, polymorphonuclear cells, platelets, mast cells) and vascular endothelial cells [43–46], and partly to cross-talk with Toll-like receptors (TLRs) [47]. The latter may include TLR2, 4 and 9. For instance, complement synergistically enhances TLR-induced production of proinflammatory cytokines such as TNF- α , IL-1 β , and IL-6 through anaphylatoxin (C3a and C5a) signaling [48]. The pathways of the aforementioned TLRs converge with anaphylatoxin signaling at the level of mitogen-activated protein kinases ERK1/2 and JNK63 [48]. Some nanoparticles may directly interact with TLRs and TLR activation can also modulate expression of complement components or receptors [49].

3.3.5 Nanoparticle-Mediated Tumour Growth

A recent study in immunocompetent mice, as well as in C5 and C5a receptor knockout animals bearing a syngeneic tumour, has strongly indicated that intratumoral accumulation of complement activating long-circulating nanoparticles can accelerate tumour growth through C5a liberation [50]. Tumour growth was more significant with nanoparticles capable of directly enhancing the alternative pathway turnover of the complement system compared with nanoparticles that triggered complement predominantly through the lectin pathway. These observations are consistent with the findings that concentration of local C5a within the tumour microenvironment is a critical factor in determining tumour progression [51]. Indeed, C5a has been suggested to enhance tumour growth by promoting the recruitment of regulatory T cells (resulting in deregulation or suppression of CD8⁺ cytotoxic T cell activity), immunosuppressive monocytes and alternatively activated macrophages into malignant tumours [51, 52]. Complement activation, therefore, is of serious concern for successful development of intravenous anti-cancer nanomedicine and other anti-cancer nanotechnology initiatives [52]. Possible therapeutic function of complement inhibitors (e.g., small molecule C5a receptor antagonists) in cancer treatment is an unexplored avenue, and, perhaps, a worthy strategy for exploration and better engineering of anti-cancer nanomedicines.

3.4 Conclusions

The interaction between medically relevant nanomaterials and the complement system is complex and regulated by inter-related factors comprising morphology, dimension, chemical makeup and surface characteristics. Understanding of nanomaterial properties that incite complement is a prerequisite for design and engineering of immunologically safer nanomedicine and biomedical devices. Many pathogenic microorganisms have deployed intriguing strategies that bypass complement activation [52, 53]. Translation of microbial strategies could provide effective means for design and engineering of ‘complement-safe’ as well as ‘phagocyte-resistant’ particulate drug carriers and biomaterials [52, 54]. However, it is imperative that such engineered nanoparticles must be structurally simple with attributes that will allow for production of affordable, viable (e.g., considering scaling up and Good Manufacturing Practice) and clinically acceptable pharmaceutical products.

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Chapter 4

The Nanoscience of Polyvalent Binding by Proteins in the Immune Response

Thomas Vorup-Jensen

Abstract Recent research has demonstrated that the successful use of nanometer-scaled material, such as nanoparticles, as medicines is often challenged by the host immune system. Mechanisms of the innate immunity seem to provide a swift response to administration of particulate nanomedicines, which may clear or in other way incapacitate the function of these drugs. To rationalize, why and how, the innate immune system especially interacts with nanomedicines, this chapter points to the prominent role of polyvalent interactions by large, immunoactive proteins with the surfaces of nanoparticles. From addressing the thermodynamics and ultrastructural properties of these interactions, it is proposed that the nm-scaled ligand presentation and symmetry on such surfaces is a determinant in the binding of these proteins. Better control over nanomedicine ultrastructure is consequently likely to provide important ways of regulating the interactions, wanted or unwanted, with the innate immune system.

Keywords Polyvalent interactions • Avidity entropy • Innate immunity • Pattern recognition • Complement • Mannan-binding lectin

4.1 Introduction

Any new field such as nanomedicine suffers a challenge to establish reasonably rigid borders to permit a definition what is inside and outside of its realm. In this process, it is easy to embody parts of other scientific areas by essentially a change of nomenclature, which will not, however, add to the scientific insight previously established. Following more than a century of investigations, one might suspect that the classic topic of protein structure in consequence cannot rightfully be claimed to

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be part of nanoscience or nanomedicine. Nevertheless, it is one of the purposes of the present chapter to argue that, in particular, the connection between structure and function of proteins in many ways can be treated as a *bona fide* nanoscience. This serves both as a better understanding of how proteins interact with their ligands and what methodologies or theoretical approaches are suitable for analyzing protein function. With the prominent role of proteins in human health, development of diseases, and, more recently, as therapeutics, it seems wholly appropriate to consider the nanoscience of proteins as an important part of nanomedicine. In choosing the immune system as a particular focus, the effort is greatly supported by the vast insight now available on the structure of the proteins associated with functions of the immune system.

4.1.1 *The Nanoscience of Protein Chemistry and Structure*

Proteins in the human body are polymers made from a selection of 20 of zwitterionic amino acids, all composed of sulfur, carbon, oxygen, nitrogen, or hydrogen atoms. The chemical structure of the amino acids involves a central carbon atom (C_{α}) bonded to a hydrogen atom, a carboxyl group, and an amine group as well as in most cases a side chain. The cellular synthesis catalyzes the formation of the peptidic bond between a carboxyl group and an amine group, this way creating the peptidic “back bone” of proteins. In a classic report by Pauling [1], the length of the peptide bond was determined to 0.272 nm. This simple structural information is a determinant to why proteins are part of the nanoscience since many proteins contain hundreds of residues. As just one example, the abundant plasma protein human serum albumin contains 585 residues and, hence, a contour length reaching $585 \times 0.272 \text{ nm} \sim 160 \text{ nm}$. In its normally folded state albumin takes approximately a diameter of 8 nm whereas the molecule under certain circumstances can be found in an extended, but still folded, form with a length of 22 nm [2]. Taken together, in quantitative terms these numbers certainly classify albumin as a type of nanocarrier and, consequently, the engineering of medicines to enable such carriage as an important part of nanomedicine [3].

To rationalize the three-dimensional properties of proteins, it is useful to introduce the hierarchical ordering of their structure [4]. The primary structure represents the sequence of amino acid residues. Through hydrogen bonds formed between $-\text{NH}$ and $-\text{CO}$ groups of in the peptide back bone, proteins may form (alpha) helical or (beta) sheet structure, these elements being referred to as the secondary structure. The spatial organization of alpha helices and beta sheets within a globular domain make up the tertiary structure. Finally, the quaternary structure describes the spatial organization of the multiple subunits. In some cases, the protein structure is formed by contacts between multiple protein chains. A striking example is the collagens. Here, the primary structure contains a repeated motif of glycine followed by any selection of two residues (typically referred to as a Gly-X-Y motif) enables three

chains to form a helical structure. Covalent bonds between cysteine residues are another important part of the architecture of large proteins, which permit formation of stable complexes through intermolecular bonds.

The proteinous molecules encountered in the body are, of course, highly heterogeneous with regard to size. For that reason, it is helpful to consider a more narrow group of proteins, which will be discussed further below in terms of polyvalency and interactions with nm-scaled materials. A detailed enquiry into the size of plasma proteins forming a part of the complement system has been made earlier [5]. The complement system is part of innate immunity largely acting to defend the body by catalyzing the deposition of complement proteins on, for instance, viral or bacterial surface. Thereby, these agents are targeted for uptake by phagocytic leukocytes, e.g., macrophages, or directly inactivated through insertion of pores in their membranes in addition to multiple other antimicrobial functions of the complement system. The hydrodynamic radii (R_G) of these proteins range from 4 to 12 nm with some of the smaller proteins being the part of the pore-forming agents while, strikingly, proteins initiating the deposition of complement are larger with R_G at ~ 12 nm and cross sectional diameter of ~ 30 – 35 nm.

These latter molecular species are easily comparable with engineered nanoparticles (NPs) as can be outlined in at least three ways.

First, both large molecules of the complement system and NPs up to about a diameter of 100 nm will not sediment without the aid of high-speed centrifugation, typically exceeding $100,000\times g$. Indeed, the major forces affecting their colloidal stability are the physicochemical properties of their environment, e.g., ionic strength, viscosity, and availability of electric charge interactions. This also implies that hydrophobic or hydrophilic properties of their surfaces are determinants in both colloidal stability and ability to interaction with host tissue *in vivo*. Second, their sizes are a critical component in their tissue distribution inside the body of experimental animals or humans. This includes the penetration of physiological barriers such the endothelium of blood vessels, the blood-brain barrier, mucosal epithelial surfaces, or retention in certain tissues, e.g., zones of inflammation or malignant tissue. Third, the dimensions of NP and proteins both enable and limit the types of interaction these can form with other molecules or tissue presenting ligands. In the case of polyvalent interactions, an appropriate distance between ligands for a polyvalent molecule is a critical determinant in the binding as discussed further in Sect. 4.2.2.

4.1.2 Affinity, Avidity, and Polyvalent Interactions

The function of proteins can be quantified in a number of ways depending on what precise functional properties are supported by that protein. A classic example is the rate of catalysis of certain chemical reaction, e.g., the cleavage of nucleotide phosphates or protein chains. Another example is the binding strength in interactions

between proteins and their ligands. This section will explore, in more detail, this topic as some recent reports have linked binding strength to nm-scaled properties of proteins. Furthermore, the binding strength is also an important parameter when using proteins as drugs or when considering their interactions with drugs.

For quantification of the binding strength between molecules, it is necessary to address what type of reaction is under consideration. A simple scheme is the 1:1 binding reaction, where



The constant K describes the concentrations at chemical equilibrium of reactants R and L as well as the formed complex $R \cdot L$. In this case, the association constant is defined by

$$K_A = \frac{[R \cdot L]}{[R] \cdot [L]}. \quad (4.2)$$

However, usually in molecular biology the dissociation constant, $K_D = 1 / K_A$, is the preferred measure of binding strength. This preference is elegantly explained from considering the Langmuir Equation [6]

$$\theta = \frac{c}{c + K_D}, \quad (4.3)$$

where θ is the fractional saturation of binding sites at the concentration c of species binding these binding sites with a dissociation constant K_D at chemical equilibrium. At this condition, the formula implies at $c = K_D$, 50 % of all binding sites would be occupied. In other words, the K_D value is a practical measure of what concentration would be required to reach a significant fractional saturation in chemical binding experiments. Equation (4.3) becomes especially useful in situations when one of the reactants is in vast stoichiometric excess over the other reactant, i.e., $[L] \gg [R]$. For the reaction scheme in Eq. (4.1), it follows that $[L]$ can then be approximated by the total concentration of L , $[L]_0$. Obviously, both technical and theoretical aspects limit the number of analyses where it is practicable to “flood” the chemical reaction with one of the reactants. A very useful alternative are methodologies generating a steady-state equilibrium by fixing the concentration by continuous supply of one of the reactants, for instance as performed with the popular surface plasmon resonance (SPR) sensors supplied with the BIAcore instruments [7]. The “steady-state scenario” could be thought of as mimicking the encounter between a microbe and blood proteins of the human immune system, which are subject to continuous synthesis and, hence, often remains at a essentially constant concentration. In this case Eq. (4.3) also applies. With the fixation of one reactant concentration it is also

simple to describe the time course of the formation of the complex $R \cdot L$ by the differential equation

$$\frac{d[R \cdot L]}{dt} = k_a [R]_0 \cdot ([L]_0 - [R \cdot L]) - k_d \cdot [R \cdot L], \quad (4.4)$$

where k_a and k_d are the association and dissociation rates, respectively. As discussed further below, the integration of Eq. (4.4) permits determination of these rates from time-resolved data on the binding between R and L .

Many molecules in the immune system permit a polyvalent binding, typically due to an oligomeric structure with identical repeated binding domains. Important examples present in blood are the immunoglobulins (Ig), certain soluble lectins, and other lectins expressed in the cell membrane of leukocytes. This repeated presentation of binding domains enables a surprisingly stable and strong interaction with ligand presenting surfaces, where the binding strength grows *quasi* exponentially with the number of interactions. The resulting “functional affinity” [8], also referred to as avidity, has been studied for almost 100 years [9–11]. Specifically, several attempts were made to relate the equilibrium constant for the monovalent interaction between R and L , as defined in Eq. (4.1), to the equilibrium constant, K_{Avidity} , for the polyvalent interaction

$$R_m + L_n \xleftrightarrow{K_{\text{Avidity}}} R_m \cdot L_n (i). \quad (4.5)$$

The indices m and n are here meant to indicate the number binding sites in the receptor and ligand, respectively, while i denotes the actual number of bonds formed between the two species.

A satisfying thermodynamical description of polyvalent interaction was provided by Kitov and Bundle [12]. A starting point is made by dividing the polyvalent binding into a sequentially ordered series of binding events [12, 13]. The initial step is an intermolecular interaction, with a Gibbs free energy of $\Delta G_{\text{inter}}^0$, followed by multiple intramolecular binding step for simplicity assumed to contribute same Gibbs free energies, $\Delta G_{\text{intra}}^0$ (Fig. 4.1). To further characterize these interactions, the chemical binding is deconstructed into macroscopically and microscopically different interaction. Macroscopically different interaction differ in the number of bonds formed, i.e., i , while microscopically different interactions share the number of bonds formed, but differ with regard to exact interactions sites joined in R and L (Fig. 4.2). In the context of the present writings, it should perhaps be noted that the distinction between macro and microscopically different interactions are purely a nomenclature to enable distinction. The differences between these types of complex formation either within the class or between the classes are all on a nm-scale when dealing with biological macromolecules. This permits writing of the chemical binding reaction for polyvalent interaction as:

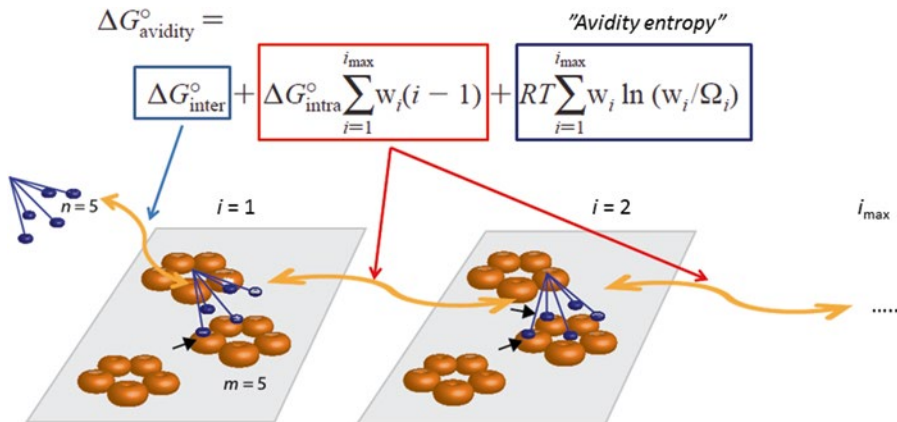


Fig. 4.1 Sequential binding model for polyvalent interactions according to Kitov and Bundle [12]. A polyvalent molecule with five ligand binding domains ($n=5$) and a polyvalent ligand with five binding sites ($m=5$) forms an initial, intermolecular contact ($i=1$), characterized by a Gibbs free energy of $\Delta G^{\circ}_{\text{inter}}$. All subsequent, intramolecular contacts are characterized by the second term of the equation, which includes the Gibbs free energy of $\Delta G^{\circ}_{\text{intra}}$ and summed to the number of maximal bond formation, i_{max} . Each bound state (i) is weighed by its occupancy, w_i . A third term constitutes the contribution by the avidity entropy. This involves the degeneracy factor, Ω_i , which may be calculated as shown in Fig. 4.3 under assumption of certain simple geometries of the polyvalent binding molecule and ligand

$$R + L \leftrightarrow R \cdot L(1) + R \cdot L(2) \dots R \cdot L(i), \quad (4.6)$$

i.e., where the product of the interaction between R and L is an ensemble of macroscopically different interactions. It is straight forward to suggest that the number of bonds formed is the critical component in regulating the binding strength of the interaction and hence the K_{Avidity} can also be written as a sum of equilibrium constants for the macroscopic identical interactions, i.e., those complexes with same i (Fig. 4.2).

$$K_{\text{Avidity}} = \frac{\sum [RL(i)]}{[R][L]} = \frac{[R_{\text{bound}}]}{[R][L]} = \frac{[R]_0 - [R]}{[R][L]} = \sum_{i_{\text{max}}} K_{RL(i)} \quad (4.7)$$

Obviously, the number of bonds formed between these species, i , cannot exceed $\min(m, n)$.

Even though the microscopically different interactions have identical equilibrium constants, and hence are energetically identical, they nevertheless form a crucial contribution to the binding strength of polyvalent interactions through a phenomenon known as avidity entropy. This entropic contribution to the binding

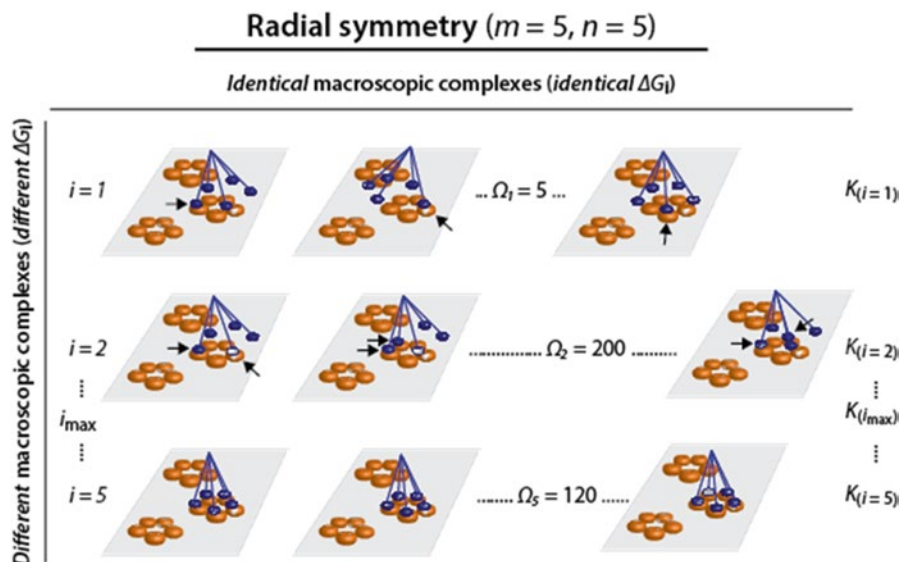


Fig. 4.2 Macroscopic and microscopic differences in complex formation. A molecule with radial symmetry, i.e., free rotation of the “arms” around the connecting point, with five binding sites binds a ligand with five binding sites. Complexes with identical macroscopic bonding (identical i) but with microscopic differences (the exact position of the molecule-ligand bonds) are compared with complexes with differences in the macroscopic bonding (different i). For each number of bond formation ($i = 1-5$), the number of possible macroscopically identical complexes (Ω_i) are calculated according to the formula shown in Fig. 4.3. According to equation for the Gibbs free energy in polyvalent interactions derived by Kitov and Bundle [12] shown in Fig. 4.1, the Gibbs free energy is identical for complexes with the same number of bond formation. This enables the assignment of a single equilibrium constant, K_i , for each bound state, i

increases with the number of ways a molecule can bind a ligand. The exact number of such possible bonding depends on the valency of R and L (i.e., m and n), the number of bonds formed, i , and geometry of the interaction (Fig. 4.3). This makes polyvalent interactions uniquely sensitive to the geometry of the binding molecules and the presentation of ligands, for instance as found on a microbial surface. The implications of this observation are explored further below.

4.1.3 Quantification of Protein-Ligand Binding Kinetics of Polyvalent Interactions

The possibilities for determining equilibrium constants for the binding of large bio-macromolecule, such as proteins, to their ligands are supported by a large and increasing number of methodologies. Among the routinely used techniques are

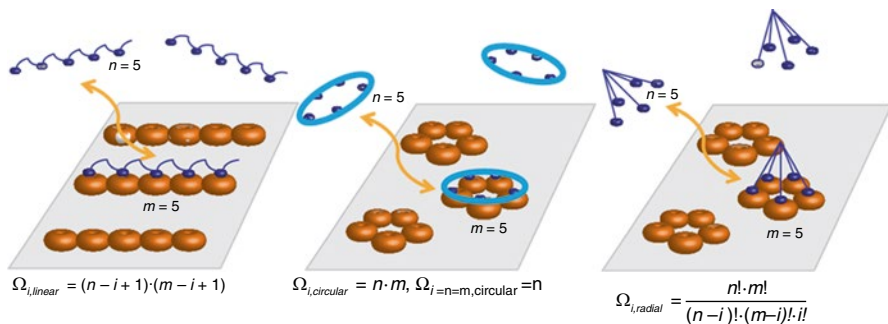


Fig. 4.3 Degeneracy in the binding by polyvalent molecules to polyvalent ligands. For simplicity all interactions show a pentavalent molecule binding a pentavalent ligand. The number of degenerate states (macroscopically identical complexes) depends on the geometry of the interaction, here shown as either a mutually linear interaction between molecule and ligand, a mutually circular interaction with binding domains within the molecule in a locked position, or with a molecule with radial symmetry where binding domains may free rotate around the central axis of the molecule. For each type of interaction, the formula for calculating Ω_i is shown. Drawings are modified from Kitov and Bundle [12] as shown earlier in Ref. [10]

various isothermal calorimetry, analytical ultracentrifugation and, as a more recent addition, microscale thermophoresis. It is beyond the scope of the present chapter to go into detail on how these methods are capable of affinity measurements. However, none of these techniques enable measurements of the underlying binding kinetics. Indeed, only two techniques routinely used are capable of measuring binding kinetics of large biomacromolecules, namely biosensors based on the principle of SPR or the quartz crystal microbalance (QCM). Excellent reviews describe the precise functioning of these sensors [7, 14, 15]. In both cases, a ligand is immobilized on the sensor surface followed by injection of a binding molecule, typically as provided in flow stream keeping the concentration of the binding molecule constant. The sensors will provide real-time data on the binding of the injected molecules to the sensor surface. In the case of SPR instruments, the binding is measured as a change in response level, S . S is proportional to the formation of the complex $R \cdot L$ in Eq. (4.1).

Binding kinetics are estimated from both the association phase, when the binding molecule is supplied to the surface, and the dissociation phase, when injection of the binding molecule is ceased and the release of the binding molecule from the surface can be followed. For the simple 1:1 interaction in Eq. (4.1), integration of Eq. (4.4) describes the expected deposition of binding molecule on the ligand-coated sensor surface

$$S(k_d, K_D, c, t) = \frac{c \cdot S_{Max}}{c + K_D} \cdot \left(1 - e^{-\frac{k_d}{K_D} \cdot (c + K_D) \cdot (t - t_0)} \right), \quad (4.8)$$

where S_{Max} is the signal at binding saturation, t is the time since start of the experiment, and t_0 the time point for injection of the binding molecule. During the dissociation phase, S is described by a simple exponential function:

$$S(k_d, K_D, c, t) = \left(\frac{c \cdot S_{\text{Max}}}{c + K_D} \cdot \left(1 - e^{-\frac{k_d}{K_D} \cdot (c + K_D) \cdot (t_c - t_0)} \right) \right) \cdot \left(e^{-k_d \cdot (t - t_c)} \right), \quad (4.9)$$

where t_c is time point for the end of the association phase with a duration of $t_c - t_0$. Fitting of these equations to experimental data will return K_D and k_d . k_a may then be calculated from $K_D = k_d/k_a$.

While often used, the simple 1:1 binding scheme can be compromised by experimental aspects not always appreciated in the studies using these methodologies. One such aspect is the valency of the binding molecule. Experimental studies have, in particular been made on immunoglobulins such as IgG. The IgG molecule is bivalent. Nevertheless, the binding of IgG to surfaces with multiple ligands (formally referred to as epitopes) has in multiple cases been studied by applying the simple 1:1 binding scheme. Karlsson et al. solved the equations to better describe bivalent binding [16]. However, until recently, no description was available for the more general problem of how to quantify the binding kinetics of polyvalent molecules. A solution to this problem has been proposed [10] by combining the experimental observation of Gjelstrup et al. [17] with the theoretical insights offered by Kitov and Bundle [12].

Gjelstrup et al. [17] investigated experimentally the binding of the human plasma protein mannan-binding lectin (MBL) to carbohydrate ligands using SPR. While the 1:1 binding scheme failed to fit the binding isotherms from the SPR biosensor, this was achieved when applying a more recently described algorithm for resolving heterogeneous binding. In the context of SPR instruments or similar devices such as the QCM, heterogeneous interactions are observed when the injected binding molecules interact with the immobilized ligands with multiple association and dissociation rates [18]. This can happen when, for instance, the ligand is randomly coupled to the surface, thereby, at least in principle, partially damming the binding site within the ligand. Other examples involve binding molecules capable of forming multiple interactions with one ligand molecule. Here, a recently studied case is certain types of adhesion molecules in the cell membrane of leukocytes, namely integrins [19–21]. Some of these receptors appear to be able to bind the $-\text{COOH}$ groups in the side chains of aspartate and glutamate [20]. Since such side chains are abundant in most proteins, each protein will provide multiple binding sites. Even though there is probably little other contact with other parts of the ligand protein, when binding involves especially the short side chain of aspartate there could be a contribution by the nearby residues, at least in a way which would limit the accessibility of the aspartate $-\text{COOH}$ [21]. In this way, an ensemble of largely similar, yet, with regard to the detailed binding kinetics, heterogeneous, interactions are provided by single ligand proteins with multiple acidic side chains. A solution to the analysis of such heterogeneous interactions has been contributed by Schuck and

his colleagues [18, 22]. In the algorithm by Svitel et al. [18], the distribution of interactions, $P(k_d, K_D)$, is formulated such that the integral $P(k_d^*, K_D^*)dk_d dK_D$ is the fractional abundance of interactions with a dissociation rate between k_d^* and $k_d^* + dk_d$ and a dissociation constant between K_D^* and $K_D^* + dK_D$. The time-course of analyte binding in the association and dissociation phase, now a sum of multiple interactions, is then

$$S_{Tot}(c, t) = \int_{K_{D,max}}^{K_{D,min}} \int_{k_{d,max}}^{k_{d,min}} S(k_d, K_D, c, t) P(k_d, K_D) dk_d dK_D, \quad (4.10)$$

where the function S is defined by Eqs. (4.8) and (4.9) during the injection and dissociation phases, respectively. A robust and physically meaningful determination of the function P is a critical part of this analysis.

In considering the application of this strategy for resolving the binding kinetics of polyvalent interactions, a starting point is the observation that $K_{Avidity}$ may be written as a simple sum of equilibrium constants for each macroscopically distinct complex (Eq. 4.7) [12]. This suggests that polyvalent interactions share an intrinsic property with the heterogeneous interactions analyzed by Svitel et al. [18], namely that they are an ensemble of energetically, and hence binding kinetically, distinct interactions. In general, this algorithm has been used to resolve heterogeneity in the immobilized ligand. Nevertheless, as pointed out by Svitel et al. [18], with some assumptions mentioned below the algorithm can also handle binding heterogeneity as a consequence of a heterogeneous composition of the injected analyte. It is helpful to combine the nomenclature of Eqs. (4.7) and (4.10) by noting at binding equilibrium that

$$\sum_i [RL(i)] = \sum_i [R_{bound}] \cdot P(k_{d,i}, K_{D,i}) \propto S_{Tot}, \quad (4.11)$$

where $k_{d,i}$ and $K_{D,i}$ are dissociation rate and constant for the $RL(i)$ complex. P is again the fractional abundance of the i th class of interaction such that $\sum P = 1$. Since $[R_{bound}] \propto S(k_{d,i}, K_{D,i}, c, t)$ Eq. (4.11) is similar to Eq. (4.10) with regard to the calculation of S_{Tot} except that the classes of interactions are assumed to be discrete while Eq. (4.10) represents a continuous distribution of binding kinetics. Consequently, it would be expected that Eq. (4.10) is an acceptable approximation of the binding kinetics of polyvalent molecules. Importantly, this approximation rests on the assumption of non-competing interactions between the components of the analyte. Considering the strong binding by highly polyvalent molecules compared to molecules with lesser capacity for polyvalent interactions it is actually difficult to rule out some level of competition. Also, although reports are now emerging on binding data for polyvalent interactions resolved by the use of Svitel et al.'s algorithm [17, 23], a more theoretical approach to the topic is needed to clarify the results from the analysis. Specifically, since a precise estimation of the P_i s depends on a precise estimation of S_{Tot} while non-competition between binding species is required, it seems unavoidable that S_{Tot} must remain determined from extrapolation.

4.2 Examples of Polyvalent Binding by Proteins in the Immune System

The above section briefly outlined how large proteins of the immune system may be envisaged as part of nanoscience and how the binding may be detected and quantified. In this section, two aspects of polyvalent interactions are explored. One is the link to conformational change in the surface-bound protein, which is critically linked to the polyvalent interaction as well as the surface topology. The other involves how the binding strength is regulated by the polyvalent interaction, here especially taking into account the geometry of ligand presentation.

4.2.1 *Conformational Change and Polyvalent Interactions*

Conformational changes in protein structure is probably an over-used explanation of functional regulation [24]. In reality, conformational changes are far from trivial to record or just, in any sense, to document. One problem is the time scale, which typically does not permit the use of biosensors such as those based on SPR for detecting conformational changes when protein are, or become, surface bound. The past several decades of research clearly seems to suggest that conformational changes in large proteins are best investigated through means of characterizing protein structure, most notably through the use of electron microscopy (EM), X-ray crystallography, and small-angle X-ray scattering (SAXS). Below, conformational changes studied in the immunoglobulin IgM and mannan-binding lectin (MBL) are provided to highlight the link between these changes and polyvalency.

IgM is part of the immunoglobulin family of plasma proteins, which also include IgG and IgA and IgE. The Igs serve an important role in the immune system. Their cellular sources are selected through complex processes of maturation and genomic recombination to form Ig protein products that bind certain structures (epitopes) present on, e.g., microbial threats to the body while avoiding binding to the body's own molecules. The cellular maturation process eventually provide Igs with high affinity and specificity for the target structure as well as a compartment of memory cells, which can easily be enrolled if the body is subsequently challenged with a previously met microbial challenge. The “immunological memory” is one hall mark of so-called adaptive immunity, since past exposures of the immune system to non-self antigens will shape the abundance and reactivities of both immune cells and antibodies. However, while the formation of antibodies in this sense is classic part of adaptive immunity, recent research has identified especially IgM antibodies in mice, which seems to be generated and provide immunity in a ways, which are closer to the innate immune system [25].

Structural evidence suggests that IgM is capable of undergoing a significant conformational change when binding to epitope-presenting surfaces. In a classic study using EM, Feinstein and Munn reported a staple-like conformation of the molecule

as imaged when bound to a bacterial flagellum from *Salmonella* that likely constituted an antigen for the IgM molecule [26]. By contrast, analysis with SAXS suggested a planar conformation of the molecule in solution, i.e., in the unbound state [27]. The dimensions of planar IgM compared to the staple-like conformation indicated that the C1 complex would only form efficient contact with the stable-like conformation. This elegantly rationalized how complement activation through the classic pathway is regulated by conformational changes in surface bound IgM while avoiding activation in the unbound, soluble state. More recently, investigations by Czajkowsky and Shao using AFM prompted the suggestion that such complement activation does not involve conformational change in the IgM molecule [28]. In their model, IgM-mediated complement activation is driven by surface binding in a planar or at least stabilized conformation as a consequence of the surface attachment. This will act to restrict structural fluctuations of the molecule compared to the soluble molecule such that strong binding by the C1 complex is enabled. Unfortunately, unlike the study by Feinstein and Munn [26], no direct observation of epitope-bound IgM was reported by Czajkowsky and Shao. Furthermore, the ability of Perkins et al. to obtain a full-size envelope from the SAXS studies [27] suggests a level of rigidity in the soluble molecule, which would be inconsistent with a highly fluctuating structure. Finally, functional data supported by structural calculations presented by Pedersen et al. [29] also support the idea that appropriate conformational change in IgM is critical for complement activation. In these studies, human sera were incubated with iron oxide nanoparticles (NPs) coated with dextran. Antibodies to dextran are commonly found in human sera [30] although the levels (usually expressed as a titer) differ. It was possible to show that complement activation by these particles correlated with the IgM titer to dextran suggesting that this protein was particularly important in the activation [29]. A striking finding was that the complement activation differed between particles depending on the size of these particles. Particles presenting the dextran antigen and with a diameter of 100–250 nm activated complement while smaller (50 nm) or larger (600 nm) particles failed in this activation [29]. From the work by Perkins et al. [31], it was possible to relate these experimental findings to the expected conformation of IgM on these particle surfaces. To take the staple-like conformation in a way permitting C1 binding, the bound IgM molecule must bind the Fab fragments in an angle of approximately 68° relative to the plane of the Fc regions. The resulting angle of the surface-bound molecule is probably influenced by several factors, for instance the precise location of the contact between the Fab and its epitope as well as the orientation of the epitopes on the surface. From geometrical considerations it is possible to show, however, that a major factor in regulating the structure of the surface-bound molecule is the curvature of the particle surface, most notably when the curvature are comparable to the dimensions of the molecule (Fig. 4.4) [29]. Flat or moderately curved surfaces will not be able to deflect the planar conformation of IgM in solution into the staple-like conformation. By contrast, more curved surface, e.g., as found on particles with diameters in the range of the cross-sectional diameter of the

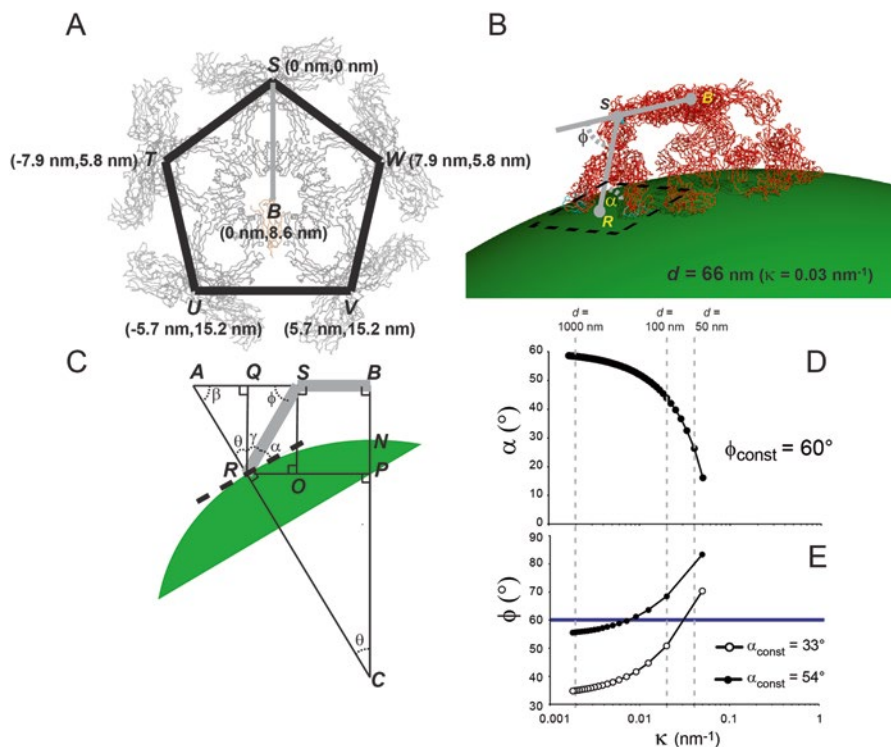


Fig. 4.4 Model of the structural requirements for the binding of IgM in the staple-like conformation [26, 62] to epitopes exposed on the surfaces of nanoparticles with differing curvatures. (**a, b**) A pseudo atomic model of the IgM pentamer in the staple-like conformation was made from the planar solution structure of IgM reported by Perkins et al. [27]. The regions corresponding to the five Fab₂' were rotated 60° downwards with respect to Fc₅/J chain disc. As shown in Panel A the centroid (**B**) of the Fc₅ disc, including the J chain indicated in green, was determined from considering the distances in two dimensions between equivalent residues (indicated with **S, T, U, V, and W**, respectively) located in the C μ 2 domains, i.e., the hinge between the C μ 1 domain and the Fc₅/J chain disc. From these calculations the distance **SB** was 8.6 nm. In *panel b* the IgM molecule in the staple-like conformation is shown on the surface of particle with a diameter of 66 nm, i.e., with a curvature ($\kappa=0.03 \text{ nm}^{-1}$); the variable loops are indicated in blue. The point **R** on the particle surface was placed equidistant between the variable loops of a Fab₂' on a line through the C μ 1 domains as indicated in the panel. The distance **SR** equals 10.6 nm. (**c**) Two-dimensional representation of geometric relationship between angles and intramolecular distances of the IgM molecule (sketched in grey). The angle ϕ measures bending of the C μ 1 and variable domains relative to the Fc₅ disk while α measures the angle between the line **SR** and the tangent to the circle with a radius **CR** (i.e. the particle surface) at the point **R**. (**d, e**) Relationship between the angles α , ϕ , and the curvature of the particle κ . With ϕ at a fixed angle (60°) there is an analytical solution of Eq. (4.6) for α as a function of κ : $\alpha = \arcsin\left(\frac{SR \cdot \cos \phi_{const} + SB}{CR}\right) - \phi_{const}$. Based on this function α was plotted as a function of κ in *panel d*. In *panel e* values of ϕ was plotted as a function of κ with α at a fixed value of either 33° or 54°. For each plot in *panels d* and *e* the shortest distance from centroid in the Fc₅/J chain disc to the particle surface (the distance **BN** in *panel c*) is indicated for the minimum and maximum values of κ . Figure and legend were first published in Pedersen et al. [29] Copyright 2010. The American Association of Immunologists, Inc.

IgM molecule itself, would significantly deflect the planar structure of IgM when this molecule binds such as surface. Finally, some surfaces may be so steeply curved that they may not accommodate IgM binding, at least not in a way accommodating full engagement of all Fabs on the surface. Taken together, these three settings may well explain why certain sizes of particles will activate the complement system through IgM polyvalent binding while other sizes will not or prove considerably less efficient. It is evident from the proposed geometry of the activation that the sensitivity toward surface curvature is wholly a result of the polyvalency of the IgM binding. These findings encourage the expectation that other types of polyvalent protein interactions with surface-bound ligands may show a similar dependence on surface curvature.

A second example of conformational changes as part of complement protein binding involves the MBL molecule. MBL is a well-established part of the innate immune system and deficiency is associated with susceptibility to infections although rarely in a way that is life threatening [32]. Effector mechanism of innate immunity are not subject to immunological memory although the cross-talk between innate and adaptive immunity probably facilitates development of memory in the adaptive immune system through activation of the innate immune system [33]. The proteins of the innate immunity are encoded by germ-line genes, which remain unaltered in somatic cells. The gene products have been selected through the eons of biological evolution to provide protection against infectious agents. In many cases the protein architecture permits recognition of certain structures, or, as famously coined by Charles Janeway, “patterns”, present on these microbial agents but not on host cells [34]. MBL is an excellent example of such a protein. Through the lectin domains MBL binds carbohydrates, notably *N*-acetyl glucosamine or mannose, but also other types. The MBL proteins chains consist of a cystein-rich N-terminal segment, enabling covalent inter chain binding, a collagen-like region which form homo trimers of MBL polypeptide chains, and finally a Ca^{2+} -dependent carbohydrate recognition domain (CRD) binding ligands [32]. The collagenous trimerization of polypeptide chains creates a structural unit (here designated MBL_3). These units are able to form larger oligomers, typically with 3–8 units ($3\text{--}8 \times \text{MBL}_3$) as judged from biochemical characterization and AFM imaging [17, 35]. The MBL oligomers are associated with proteases (MBL-associated serine proteases, or MASP), which permit activation of the complement system once the MBL oligomers are bound to target surfaces. The biochemical mechanism of distinguishing “self” from “non-self” is discussed further below. At this point, the studies on the MBL ultrastructure are helpful to outline the challenges in documenting conformational changes in large proteins. The MBL ultrastructure is actually similar to both IgM and C1q in the sense that these proteins are approximately 40 nm in cross sectional diameter, contains a pseudo rotational symmetry, and possesses multiple binding domains [35–37]. For this reason, it is tempting to suggest that complement activation by MBL involves a conformational change when the molecule attaches to a target surface as also proposed for IgM. To make a study on this topic Dong et al. characterized the structure of MBL in solution by SAXS [37]. The ultrastructure of $3 \times \text{MBL}_3$ oligomer is moderately bend and highly ramified with an internal maximum distance not exceeding 32 nm and a cross sectional diameter of 37 nm.

The CRD₃ domains are flexibly attached to the collagenous region [37, 38]. To provide ultrastructural data on the ligand-bound state, AFM imaging was made on the same preparation of MBL applied to a surface with carbohydrate ligand [37]. In this case, the cross sectional diameter was found to be roughly 60 nm on average when liquid-state imaging was performed. Higher resolution was obtained by dry-state imaging, but under these conditions the molecular structure deteriorated. Taken together, these measurements would nevertheless suggest that some kind of unbending or stretching (or both) of the molecule occurs in the surface-bound state.

Are conformational changes in MBL then critical for complement activation? Degn et al. noted recently [39] that a simple accumulation of MBL/MASP complexes would probably permit autoactivation of the MASPs, and, hence, activation of the complement cascade, as judged from what is known about enzymology of these proteins. In this model, the ligand binding activity of MBL simply acts as a conveyor of MASPs to the target surfaces eventually raising the enzyme concentration beyond a level critical for enzymatic activation. In principle, no conformational change in MBL is required per se for MASP activation. On the other hand, few, if any, of the surfaces, which were routinely used for testing complement activation by the MBL/MASP complexes, are flat compared to the size of the MBL oligomers. *Staphylococcus aureus* was used as complement-activating agent in studies suggesting catalytic activation of MASP by juxtaposition of complexes carrying these proteases [40]. *S. aureus* synthesizes its cell wall in such a way that the surface presents ridges organized in concentric circles [41]. The ridge height seems to vary considerably, with amplitudes from ~1 to 5 nm depending on the age of the material. Other structural features of the cell wall involve central depressions, where novel cell wall is synthesized. In a radius of ~30 nm from such depressions, the ridge height reaches 30 nm. When MBL binds through multiple points of contact to a surface wrinkled in these ways, it seems likely to involve some sort of structural change in the bound MBL compared to the soluble state. This suggestion lends support from the consideration made above on IgM, namely that polyvalent interactions—unlike their monovalent equivalents—are sensitive to surface topography of the bound surface. The study by Dong et al. [37] seems at least to suggest that such changes in MBL can be accommodated by the molecule. To settle the specific role of conformational changes in MBL as part of complement activation will consequently remain a challenge in a foreseeable future.

4.2.2 Avidity, Polyvalency, and Immune Target Recognition by Soluble Molecules

One of the most striking consequences of polyvalency is the increase in binding strength of a polyvalent molecule compared to its monovalent source. The polyvalent nature of the interaction is doing more to target recognition, however, than merely increasing binding strength or serving as anchor points to convey conformational changes as can be appreciated from as few cases already introduced.

In the case of MBL, it was suggested that the CRD trimer permit a critical polyvalent interactions with polymeric carbohydrates [42, 43]. The trimerization creates a distance between the CRD ligand binding sites of roughly 5 nm. Binding of these multiple CRDs to carbohydrate chains consequently requires branching of the glycan or a glycan of some length, longer than those found on mammalian cells but typical of fungal cell wall carbohydrates. This forms a biochemical principle for pattern recognition by this molecule although the exact target organisms recognized by such principles can be disputed [10, 44]. Binding data provided by Gjelstrup et al. [17] seems to suggest that the major source of avidity is the oligomerization of structural MBL₃ units while the CRD₃ unit only binds weakly to glycosylated targets. In any case, it is possible to rationalize this type of pattern recognition as “topological patterns”, where the actual structural presentation of ligands will enable or discharge polyvalent interactions and hence MBL binding [10]. From the evolutionary selection of MBL to form an efficient tool to bind such patterns, it may be speculated that the ultrastructure of this, and other, similar pattern recognition molecules is imprinted by the size and topology of their binding targets, i.e., the patterns found on microbes. In this sense the topological patterns are of a fundamentally different structure than most of those recognized by other pattern recognition molecules, namely Toll-like receptors (TLR). The critical part for recognition by these molecules is not thought to involve any major ultrastructural presentation of ligands from microbial organisms such as parts of the cell wall (peptidoglycan) or flagella (flagellin) [10, 45, 46]. For instance, the surface recognition involved in TLR5 binding of flagellin, a protein component of bacterial flagella, would cover ~13 nm² in the flagellin [46]. By contrast, MBL pattern recognition may cover ligands presented over an area of as much as ~3000 nm² [37]. As judged from the work by Feinstein on Munn on IgM [26] something similar would probably be the case for this large molecule as well. In this context avidity entropy, mentioned in Sect. 4.1.2, is a particularly interesting phenomenon. Microbial organisms are made “bottom-up” implying that repeated biomacromolecular elements are found in microbial surfaces with certain symmetries in their presentation. In particular, work on nanomicrobiology by Yves Dufrière and colleagues [47] as well as the work cited above on the cell wall structure of *S. aureus* [41] have unraveled a previously unappreciated ultrastructure of microbial cell surfaces. It seems certainly possible that an important part of topological pattern recognition would involve symmetries of ligand presentation on these cell surfaces and hence link the binding strength to aspects of avidity entropy.

4.3 Nanomedicines and Polyvalent Proteins of the Immune System

Almost three decades of immunological research has highlighted the importance of the innate immune system in acting as a first-line of defense against invasive microorganisms. As briefly discussed above, innate immunity distinguishes “friend

from foe” by the ability of certain germ-line encoded proteins to bind patterns exposed by non-self surfaces or molecular constituents. Whilst there is plenty of evidence that such a system is required to protect the body from infections with pathogenic microbes, it is also now clear that the innate immune system also constitutes a formidable barrier to nearly any kind of drug delivery involving particulate materials. Below, the connection is made between the nanoscience of polyvalent interactions and the challenges in nanotechnology to develop efficient drug delivery vehicles.

4.3.1 How Nanomedicines Resemble Targets for the Innate Immune System

It is a truism that nanoscience is a science of the importance of size. In nanomedicine, this is extended to include biological materials, where nearly any phenomenon involving biomacromolecules, in principle, is a variant nanoscience. In this line, with the molecular basis of the pathologies of diseases these could be argued both to be rationalized in terms of nanoscience as well as be treated by agents capable of acting on a nm scale. This notion was famously captured by the physicist Richard Feynman’s vivid foresight in 1959 of a future, where the patient could “swallow the doctor” to enable this nano-sized medical practitioner to directly interfere with undue molecular mechanisms at a similar size scale [48]. The best examples of medical therapies related to this vision are probably the remotely similar application of NPs in medicine. Below, a few of examples are identified and put in a context of why they challenge, in particular, the innate immune system.

If one follows the European Union’s definition of NPs as “with one or more dimensions of the order of 100 nm or less”, several infectious agents, notably viruses are also on this list. Unsurprisingly, this has prompted the idea that the sheer size of viruses is their strongest resemblance to NPs and that this property somehow is in a major way responsible for NP recognition by the immune system [5, 10, 49]. However, arguably the mechanistic insight to support this end has been lacking. Indeed, at least two other properties are likely to be important, particularly in the context of the recognition of NP by polyvalent proteins of the innate immune system.

First, although it is possible to make certain types of NPs top-down by grinding and subsequent appropriate liquid chromatography (e.g., [50]), in more general such material is manufactured by precipitation or aggregation reactions (e.g., [51]). These “bottom-up” strategies basically assemble the particles from smaller metallic or organic polymer compounds, in some cases producing a semi-crystalline material. While this material is not necessarily highly ordered, the limited types of constituents are likely to create some degree of repeated structures in and on the surface of the resulting particles. An important example of this type is ultra small

paramagnetic iron oxide particles (USPIO), which amongst other applications are used as contrast reagent in magnetic resonance imaging. These NPs are usually modified with carbohydrates (dextran) [52]. This is not unlike viral capsids, which also are rigid, semi crystalline particles, in many cases with diameters of ~100 nm [53]. Obviously, viral particles are also assembled in host cells “bottom up” from their macromolecular constituents. In this case, a high degree of ordered presentation of these constituents is the results [53]. Also, the viral capsids may be covered with a lipid envelope as well as integrate glycosylated proteins, as for instance known from human immunodeficiency virus, further strengthening the similarity to USPIO formulations. Second, the attempt to create biocompatible NPs has prompted a choice of materials for making the particles or their surface coating from biological sources. While this in many cases would ensure degradability of the material and avoidance of toxic responses from hepatic, pulmonary or renal accumulation, it does not prevent immune responses. It has become increasingly clear biologically-derived materials create a significant problem from immune recognition as “biomolecules” of apparently innocuous origin [54], e.g. chitosan derived from the shells of shrimp, triggers an instantaneous immune response from the innate immune system [55, 56]. Chitosan is made from deacetylating chitin, thereby, creating highly positively charged carbohydrate polymers of repeating glucosamine residues, however, with varying degrees of deacetylation leaving behind unmodified *N*-acetyl glucosamine residues as well. Since this latter constituent is an efficient ligand for, e.g., MBL and induces chitosan triggers antibody production [56], it not surprising that chitosan NPs trigger complement activation, nor that complement activation is likely to inflict serious challenges in keeping the particles functional or even in circulation. One of the reasons that the immune system to this extent attacks chitosan is that a similar material is generated by the enzymatic breakdown of fungal cell walls, which may contain chitin [57]. In brief, the choice of chitosan will make the particles effectively mimic surfaces of a known treat to the human body, i.e., fungi. Presumably, the innate immune system has been subject to selection for enabling a response to these organisms and the biological origin of the material is, thus, of no avail in avoiding immune responses. A similar point can be raised for dextran. Dextran-like carbohydrates are made by gram-negative organisms [58], which, in turn, may cause formation of host antibodies to this material. Human blood abounds with high level of dextran antibodies [30], which would also effectuate the clearance of the dextran-coated particles. Some success of engineering so-called stealth particles was achieved with synthetic polyethylene glycol layers [51]. Nevertheless, also in this case certain MBL-like molecules of the immune system, namely the ficolins, may be capable of triggering a response through complement activation [59]. These examples clearly advocate a more refined strategy, if these materials, particularly biomaterials, are to be used as drug delivery vehicles.

4.3.2 Surface Ultrastructure, Polyvalent Interactions and the Functionality of NPs

The point has been made above that many nanoparticulate systems by mode of synthesis and choice of materials are targets for molecules of the innate immune system as extensively reviewed elsewhere [54, 60, 61]. The view in the present text is that a considerable part of the challenges induced by the immunoreactivity reflects the polyvalent interactions often supported by the immune system. Of course, this begs the question if our current insight on how polyvalent interactions are made can be used to improve NP drug functionality.

The work by Pedersen et al. [29] highlight the point that size, or specifically curvature, of NPs is important when conformationally labile molecules such as IgM bind their surfaces. Whilst it is not surprising that certain particles are too small to accommodate this binding and will escape triggering IgM immune functions, the contrary finding that large particles with a lower curvature can do so, seems to offer new avenues of drug design. At this point it is, of course, difficult to determine if appropriate particle size will enable concomitant escape from multiple types of immune recognition such as phagocytic capture. At least Pedersen et al.'s work offers a rationale to explain why the size of NPs, i.e., particles with a size close to the dimensions of the large biomolecules of the immune system, is a critical parameter. Indeed, when wrongly chosen, this will impede the function of the NPs with respect to clearance or other incapacitation of function by the immune system. Polyvalent interactions seems here particularly important consider as they are highly sensitive to the topology of the surface.

A more surprising aspect relates to the thermodynamic aspects of polyvalent interactions, especially the sensitivity to the symmetry of ligand presentation via avidity entropy. From a simple experimental enquiry, Gjelstrup et al. [17] reported that ligand density may dramatically regulate polyvalent binding of MBL to surfaces coated with variable amounts of coated ligand [10]. It seems likely that avidity entropy plays a role in the binding of these surfaces, but arguably a more carefully engineered surface would be required to better analyze this point. In this context, it is important to remember that the schematic representation and associated calculations of degeneracy of ligand binding (Fig. 4.3) is a theoretical approach, which may not capture the full complexity of the actual ultrastructural properties of, for instance, microbial surfaces as discussed above [10, 12]. Nevertheless, it is clear that any complete or distorted symmetries of ligands on such surfaces, or the engineered surfaces of NPs, are likely to be an important factor in polyvalent interaction. While the current state-of-art in designing NPs mostly involves processes that only permits a partial control over the surface ultrastructure, it would be envisaged, from the considerations made above, that such control would enable better control over what molecules would bind the NP surfaces. Not only would this address the important aspect of avoiding particle clearance of the particles, it could also contribute novel means drug targeting, e.g., in drugs targeting immune cells or

functions of the immune system by facilitating the design of antagonists. This potential was actually experimentally demonstrated by Kitov and Bundle by use of dendrimers targeting IgM [12].

4.4 Concluding Remarks

This chapter wants to enable the Reader to understand basic principles of polyvalent binding by large, immuno active proteins to ligands, mainly presented on surfaces of microbes or NP look-alikes. The few examples of such mechanisms are far from exhaustive, nor scientifically well understood. In consequence, the Reader is encouraged to more broadly consider in what sense and where polyvalent interaction are part of nanoscience and nanotechnology.

At this stage, it seems safe to conclude that the topic of large proteins and their interactions with NPs and other nanostructured materials is treatable as part of nanoscience with significant implications for nanomedicine. It is an interesting question if our insight on the structure, or simply the size, of proteins of the immune system can benefit the design of, for instance, NPs, which are better tolerated and with better performance in the body. At least two aspects come to mind from the current presentation. First, the chemical nature of polyvalent interactions, specifically with regard to the entropy-driven part of the binding, seems to advocate against highly symmetric systems. Second, the conformational influence of molecules bound or otherwise adsorbed onto surface can be critical for the outcome of such adsorption, in particular when the molecules in question are immunoactive. While adsorption affects conformational change in the finer protein structure, this chapter points to changes at the ultrastructural level—at the nm scale—as also potentially important. This is at least technically significant as nanoscience now offers multiple approaches to studying the ultrastructural changes in molecular architecture. In this sense, an outlook on future studies on the interaction between the immune system and nanostructured materials would include a vast increase in our knowledge about protein structure influenced by nanostructured materials. It appears as a fair expectation that such insights will benefit the design of efficient nanomedicines that are capable of better escaping unwanted activation of the immune system than current methodologies.

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Chapter 5

Microfluidics-based Single Cell Analytical Platforms for Characterization of Cancer

Emil Laust Kristoffersen, Morten Leth Jepsen, Birgitta R. Knudsen, and Yi-Ping Ho

Abstract Individual cancer cells in a tumor are very diverse both genetically and functionally. Moreover, in many cancers, whether a tumor flourishes or dies after a given treatment depends on a small fraction of cells in the tumor, for instance, the cancer stem cells, rather than the bulk population. Traditionally, scientists only obtain averaged information from the entire population of cells in a bulk tumor, but it is critical to develop an effective and user-friendly platform capable of interrogating single cells in order to understand and treat the disease better. This chapter reviews recent progress of microfluidics-based tools for single cell analysis that are relevant for cancer characterization, as well as how nanotechnology may advance the analysis with improved signal responses. We hope that this general introduction may catalyze the adoption of these advanced single-cell analysis approaches for cancer studies.

Keywords Nanosensors • Circulating tumor cells • Genomic analysis • Enzymatic activities

5.1 The Importance of Single Cell Analyses for Cancer Characterization

Cells are defined as the basic functional unit of living organisms. They sense environmental stimuli, respond and adapt to the environmental changes chemically or physically in order to form and maintain the distinct tissues in complex biological organisms. Therefore, even cells of identical genetic identities can display a wide disparity of physical responses (such as cell morphologies) or chemical responses (such as RNA or protein expression level regulation). These asynchronous responses

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to the microenvironments, often due to random fluctuations, or noise in gene expression [1–3], make precise characterization of cells in a population difficult, especially when techniques with single cell sensitivity are not available. Hence, currently our understanding of cells are in many aspects limited by the inherent ensemble average obtained by analyzing bulk populations of cells, which does not necessarily precisely represent the individual behavior of each cell within a population.

While many population-level studies using bulk assays are presented, it remains ambiguous whether the population data may faithfully reflect the dynamic response of individual cells. In fact, cell-to-cell heterogeneity has been evidenced in many examples. For instance, the transcription events in mammalian cells are subject to random fluctuations and lead to large variation in messenger RNA (mRNA) copy numbers [4, 5]. Moreover, cell heterogeneity is observed to impact cell-fate decisions in mouse multipotent progenitor cells [6]. In particular, cancers are typically characterized by a very high level of cell-to-cell variation in regard to both morphological, physiological and genetic features [1, 7, 8]. The obvious reason for such heterogeneity is that cancers arise as a consequence of multiple and successive mutations in the genome of the cell leading to the generation of many different subpopulations of cells constituting the tumor. According to the cancer stem cell theory, which was presented a few decades ago, cancer initiates by the accumulation of mutations in stem cells [9]. This leads to the transformation of the stem cells into cancer stem cells, with self-renewal potential and the capability to differentiate into other types of cancer cells. Due to these characteristics, isolation and characterization of individual cells within a population of cancer cells (e.g. a tumor) are envisioned of particular value for understanding the origin and progression of this disease.

Especially as a predictive tool for individualized cancer treatment, single cell characterization may prove of tremendous value. Recent studies suggest that the drugs developed by measuring therapeutic responses from traditional bulk analysis, tend to suppress the growth of bulk tumor cells but not as effective in eliminating cancer stem cells (CSCs), where the relapse of tumor may occur from [10]. The different drug responses between bulk tumor and CSCs further demonstrate that the fidelity of bulk analysis is insufficient for the evaluation of antitumor treatments. Furthermore, a potential innate consequence of the heterogenic nature of tumor cells is great fluctuations in the amount/activity of cellular drug targets or important determinants of drug efficiency, such as growth rate or metabolic activity.

For example, the heterogeneity of prostate cancer poses a significant challenge to effective targeted therapy [11, 12], such as those involved introduction of a homologous double-stranded RNA (dsRNA), or RNA interference (RNAi) [13], simply because the individual cancerous cells respond very differently to the same treatment. Survival of only a few cancer cells after initial systemic treatment may result in recurrent tumors, which may be even more aggressive or chemoresistant than the initial tumor. Therefore, the identification of these rare cells may be imperial for tailoring treatment to the individual patient in the best possible way.

Finally, when it comes to detection of cancers for diagnostic purposes, the capture and registration of single tumor cells from the bloodstream may prove a powerful tool. Indeed, in combination with protocols for fast and reliable single-cell characteristics of captured cells, such techniques may pave the road for future diagnosis integrated with individually tailored treatment.

Previous development of microfluidics has shown great success in biomedical applications, but mainly centered on the analysis of targeting biomolecules from minute amount of samples, typically nanoliters to hundreds of picoliters. More recently, the exploitation of microfluidics for cell manipulation and analysis, taking advantages of the matching length scale of microfluidics and the size of individual cells (tens of micrometers) has increased. Of particular importance is the detection and analysis of diseased cells, which are typically rare among the whole cell population, difficult to detect and retrieve, and yet critical for accurate analysis of the biological processes. In this chapter, we intend to review the existing development of microfluidics-based tools for single cell analysis that are relevant for cancer characterization, as well as how nanotechnology may advance the analysis with improved signal responses. We hope that this chapter will give the audience a general introduction and catalyze the adoption of these advanced single-cell analysis approaches [2, 4, 14] for cancer studies.

5.2 Enrichment of Circulating Tumor Cells

5.2.1 *What Are Circulating Tumor Cells?*

Metastasis, the spread of cancer cells from the primary site to other organs in the body, represents the major cause of cancer-related patient death. Previous evidence suggests that the tumor cells are shed from the primary tumor at an early stage of metastasis development, break through the vascular wall, travel via the peripheral blood to sites distant from the primary tumor, and form a secondary tumor [15]. Cells escaping from the primary tumor, are called circulating tumor cells (CTCs). The significant role of CTCs in the metastatic spread of tumor has rendered them valuable biomarkers for both detection of the onset of cancer metastasis and clinical evaluation of treatment outcome. Moreover, detecting CTCs as a cancer marker is advantageous in the clinics because it makes noninvasive detection possible through capturing CTCs in a liquid biopsy, such as a blood sample. Recent evidence on how the CTCs may reflect the molecular features of the primary tumor cells further displays the importance of CTCs in cancer biology [15, 16]. For instance, the presence of mesenchymal markers on CTCs envisages more accurate prognosis than the expression of cytokeratins alone, implying that the currently used assays based on epithelial antigens may overlook the most aggressive subpopulation. However, CTCs are extremely rare and appear in very low concentrations down to one per millions of normal blood cells. Therefore, their detection remains a great challenge in cancer characterization [15, 17].

5.2.2 *State-of-Art*

CTC isolation is typically evaluated by many factors, including the capture efficiency (i.e. 100 % capture efficiency being isolation of all of the CTCs in the liquid biopsy, therefore, allowing identification of the cancer occurrence), isolation purity (i.e. 100 % isolation purity being isolation of only the CTCs, with no other cell types), isolation speed and the required sample volume. A large panel of macro-scale sorting techniques have been previously reported for CTC enrichment, such as immunomagnetic beads separation [18], laser scanning cytometry [19], fiber-optic array scanning technology (FAST) [20], as detailed in previously published reviews [18, 21]. In general, these approaches utilize the differences between CTCs and normal hematologic blood cells in physical (size, density, electric charges, deformability) or biochemical (surface protein expression, invasion capacity) properties, as illustrated in Fig. 5.1. For example, separation without labeling through the physical properties of the cells is adopted in the isolation by size of epithelial tumor cells (ISET) [22–24]. Dielectrophoretic field-flow fractionation (DEP-FFF) utilizes membrane resistance in combination with size to sort different responses to dielectrophoresis. Biochemical separation relies on immunological procedures using antibodies against tumor-associated antigens and common leukocytes antigens. For instance, the CellSearch® and the Ariol® select CTCs by utilizing magnetic beads coated with antibodies against genes that are highly expressed in CTCs [18, 21, 25]. Subsequently, the antigen-antibody complex is separated from the liquid phase *via* exposure to a magnetic field. However, many of the currently available approaches remain relatively ineffective in isolation efficiency with CTC identification in 50–90 % of patients, while 5–10 mL of sample volume is typically required [21]. Therefore, the search for sensitive, specific, and economical analytical techniques continues.

5.2.3 *Microfluidics Based CTC Enrichment*

Microfluidics based CTC enrichment has garnered considerable attention, due to the matching length scale of microfluidic channels to the cell sizes. Secondly, the micron-sized geometric features used in microfluidics greatly reduce the sample consumption. To date, microfluidics-based CTC enrichment has shown great promise by identification of CTCs in close to 99 % of patients [26], while requiring very minute sample size of 10 μ l [27].

The very early demonstration of microfluidics based CTC enrichment is the so-called “CTC-chip” developed by Toner’s group [26]. The CTC-chip consists of an array of microposts coated with anti-EpCAM antibodies, as shown in Fig. 5.2, where the positive selection is implemented by the antibodies against the epithelial cellular adhesion molecule (EpCAM), relevant to epithelial growth and differentiation. Over expression of EpCAM has been observed in many human carcinomas including prostate, colon and rectum, breast, lung, esophagus, and pancreas.

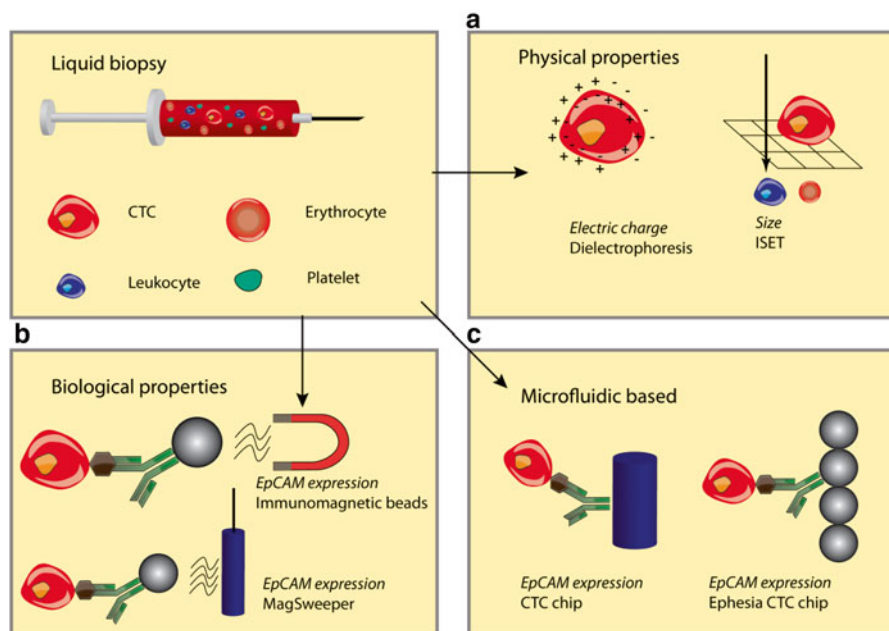
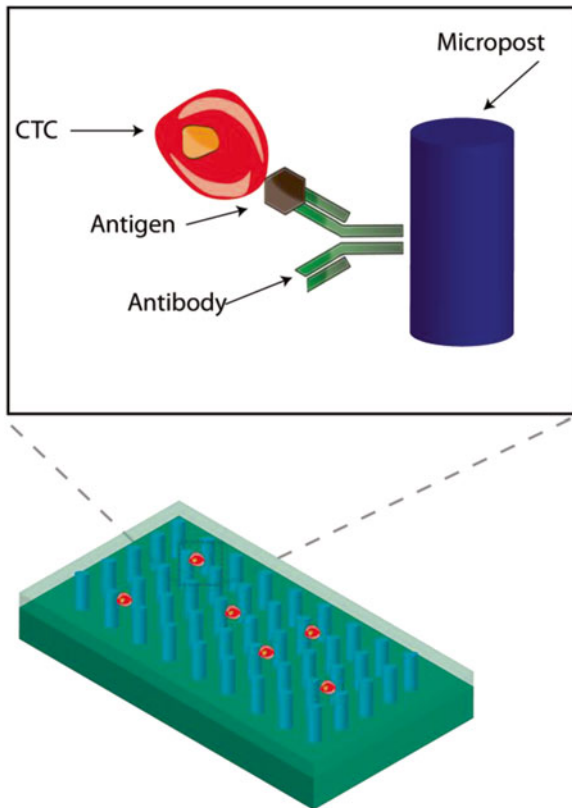


Fig. 5.1 Strategies to isolate circulating tumor cells (CTCs): Isolation of CTCs from normal hematologic blood cells, such as leukocyte, lymphocyte and erythrocyte, typically relies on dissimilar physical or biochemical properties of CTCs. (a) Separation by physical properties normally does not involve additional labeling, such as isolation by size of epithelial tumor cells (ISET), or the dielectrophoretic field-flow fractionation (DEP-FFF) that isolates cells by their responses to dielectrophoresis, which is determined by the size and membrane resistance. (b) Separation by biological properties: This category of separation usually involves immunological procedures which involve antibodies against tumor-associated antigens and common leukocytes antigens. The CTCs-specific antibodies are typically bound to micron-sized magnetic beads, which allow a separation when applying a magnetic field. (c) Microfluidics-based separation: Typical form of microfluidics-based CTC enrichment consists of microposts functionalized with antibodies specific to the surface-markers of tumor cells, such as epithelial cellular adhesion molecule (EpCAM). The microposts may be constructed by a solid structure, or by a pile of magnetic beads as recently demonstrated in the Ephesia CTC chip

In contrast, hematologic cells do not express EpCAM [28, 29]. Hence, EpCAM appears as an effective cancer biomarker and an appropriate target molecules for CTCs enrichment in a liquid biopsy. Combining immunolabeling with controlled laminar flow conditions in the microfluidic chip, the CTC-chip has shown to successfully identify CTCs in peripheral blood from 99 % of patients carrying metastatic lung, prostate, pancreatic, breast, and colon cancer, or more precisely, 115 identifications out of 116 investigated samples [26]. The capture efficiency, as defined previously, is improved by two essential parameters operated in microfluidics: (a) Low flow speed: permitting cells to interact with the microposts for extended duration and, thus, increasing the likelihood of cells sticking to the posts and (b) Low shear stress: enabling the cells to flow through the channel with minimum physical

Fig. 5.2 Typical design of the CTC-Chip: The microfluidic-based separation takes advantages of the comparable size of microfluidic channel and the cell sizes, that allows effective capture of CTCs. This chip usually consists of an array of micron-sized pillar posts functionalized by antibodies specific to CTCs. The microposts are used to disturb the flow-streamline and enhance cell-microposts interactions. Current development has centered on the design and production of microposts to scale up for large-scale clinical applications



distress. The CTC chip appears very gentle to the cells with a shear force of less than 0.4 dyn/cm^2 . The low shear supplied in the CTC-chip even allows capture of T-24 cells, which have a relatively low expression of EpCAMs. Other than EpCAM, current characterization of CTCs is typically through immunostaining of the cells with markers such as KLK3 (prostate specific antigen) for prostate cancer or TTF-1 (thyroid transcription factor-1) for lung adenocarcinoma. Furthermore, to accelerate the isolation speed, Stoot et al. later demonstrated an automated system for prostate cancers with enrichment and quantitative analysis of positive CTCs for prostate specific antigen (PSA) [30].

As an alternative to the microposts based CTC-chip, an innovative geometric improvement termed the Herringbone-chip (HB-chip) [31], has been demonstrated to prime the enrichment of CTC. The chevrons, or the herringbones, as depicted in Fig. 5.3, function by disrupting the laminar flow, which would then enhance the mixing between streamlines and encourage the collisions of cells and the antibody-coated herringbone structures. The HB-chip has shown a 26.3 % improvement in capture efficiency compared to the previously discussed CTC-chip, along with significantly higher purity of the captured CTCs. Moreover, the use of transparent and chemically stable materials allows imaging of the captured cells with standard staining or assays such as Fluorescent In Situ Hybridization (FISH).

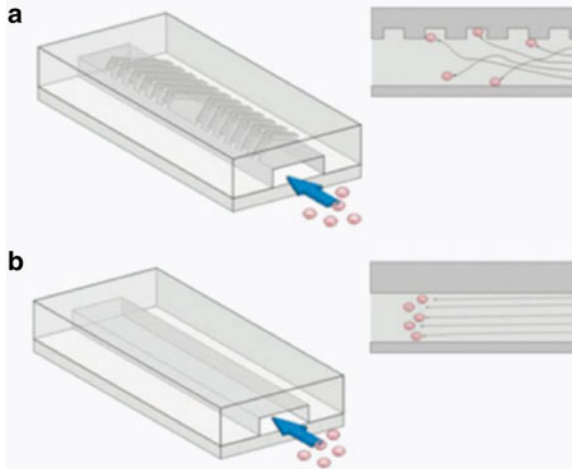


Fig. 5.3 The Herringbone (HB)-Chip: Conventional design of CTC-Chip relies on laminar flow, which limits the interactions of target cells with surfaces, and the complex micropost structure is also challenging to scale up for high-throughput production. The HB-chip represents an alternative strategy, which takes advantages of the design of herringbones, hence the name. (a) The surface ridges help to break up streamlines, maximizing collisions between target cells and the antibody-coated walls. (b) As a comparison, there is a lack of mixing under low Reynolds number regime in traditional flat-walled design. Figure adapted from Ref. 31 with permission

Saliba et al. have recently developed a system, named Ephesia, in which they combine superparamagnetic beads with microfluidic technology [27]. In this platform, superparamagnetic beads are pre-functionalized with antibodies. When introduced into the chip, upon application of an external magnetic field, the beads would stack up due to dipole-dipole interactions, forming microposts out of stacks of beads. To reduce production cost and technical complexity, the magnetic pattern is generated through microcontact printing with water-based ferrofluid, or “magnetic ink”. Major benefits of this method, compared to the conventional design of CTC-chip, are the greatly reduced production cost and possible batch-functionalization of the magnetic beads. It is also of particular note that the proposed self-assembly process offers an aspect ratio beyond the most sophisticated nanofabrication techniques. In summary, the microfluidic technologies have emerged as an attractive micro-scale CTC isolation system. The unique features of fluidic mechanics at the micro- and nano-scale, such as channel dimensions, flow operated at the laminar flow regime, and surface area to volume ratio, have enabled improved capture efficiency and isolation purity of CTC. However, perhaps similar to the challenges for other microfluidics-based applications, it is necessary to validate the reproducibility and robustness of technology with extensive clinical relevant testing. Further, it is expected that the reduced technical complexity, together with low cost of production and testing, would encourage the widespread adoption of microfluidics-based CTC enrichment in the clinics. Audiences interested in the latest development of microfluidics-based CTC detection are referred to recently published reviews [32–34].

5.3 Nanoscaled Molecular Techniques for Analysis of Single Cells

Nanotechnology based molecular techniques, including nanoscale molecular manufacturing, nanosensors, and single molecule detection, represent a significant evolution towards investigation of cell population heterogeneity. As described above, single cell analysis and cell population heterogeneity investigation is necessary in order to elucidate how the rare cells may contribute to tumor development and treatment outcome. In the following section, we overview and comment on how nanoscaled molecular techniques may join the effort of microfluidics in single-cell analysis for the characterization of cancers from three perspectives: The genomic-, transcriptomic- and proteomic-level (as illustrated in Fig. 5.4).

5.3.1 Single Cell Genome Analysis for Cancer Characterization

Cancers are typically caused by errors, or mutations, induced in the genome, which subsequently prompt the cells to malfunction, or grow uncontrollably. Such mutations can either be subtle genomic alterations (such as single base mutations) or gross

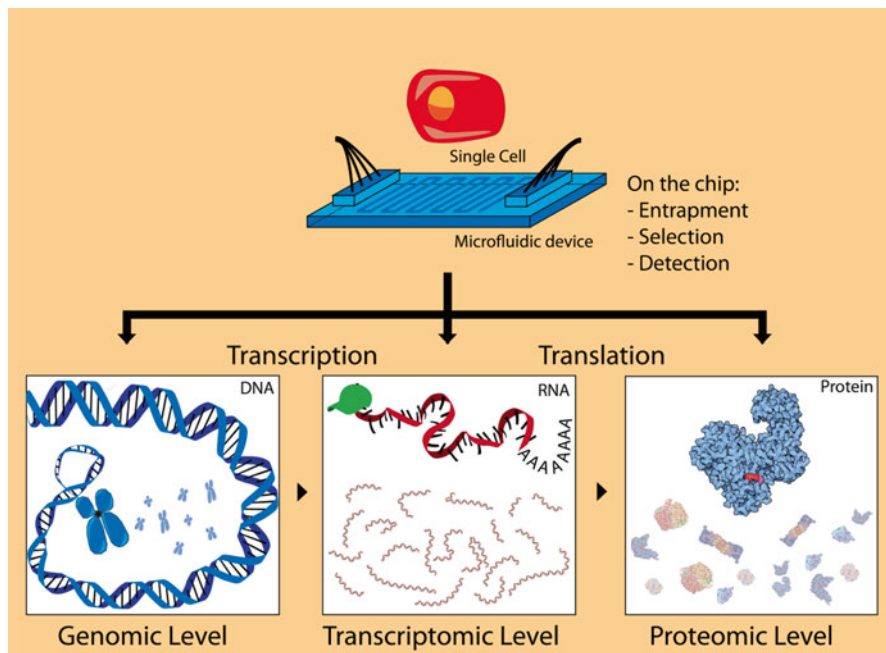


Fig. 5.4 Overview of how microfluidic systems may assist in single cell analysis: High-throughput cell-based screens can benefit considerably from the unique liquid-handling capabilities offered by microfluidic systems

genomic alterations (such as deletions, translocations, insertions, rearrangement or even loss or gain of entire chromosomes) [35]. As the disease progresses, the evolutionary accumulation of cancer causing mutations result in a high degree of genetic diversity between or among cancer cells [36–38]. Furthermore, it is becoming established that CSCs, which account for only a minor part of the bulk tumor, exist in a variety of cancers, such as colon, ovarian, and small cell lung cancers [39]. CSCs are known to have a distinct gene expression phenotype, rendering them highly resistant to chemotherapeutic treatments [39–42].

Identification of genomic alterations at the single cell level are routinely accomplished today by standard cancer diagnostic techniques such as chromosome staining and FISH [43–45], or FISH based on nanoparticles-labeled probes [46]. These refined techniques are based on microscopic readouts, and can be used to identify large genomic alterations in single cells. In some cases FISH can even be used to detect Single Nucleotide Polymorphisms (SNPs) if the site and polymorphism is known [47]. In recent years, genomic sequencing has been adapted for single human cell genome analysis [48, 49]. This late arrival of DNA-sequencing as a tool for single cell analysis reflects the preceding barriers: (1) sufficiently sensitive sequencing techniques for single cell analysis have emerged not until the last decade, (2) the data obtained by sequencing is often very comprehensive and the analysis of which requires well-established bioinformatics tools and/or statistics as well as appropriate references. With this said, a combination with microfluidics might provide a unique means for isolation and/or enrichment of specific populations of cells, for example the CSCs.

As a pioneer in the full genome sequencing of single human cells using single cell based approaches, Wigler's group combined flow-sorting, Whole Genome Amplification (WGA) and next-generation sequencing for the investigation of single breast cancer cells [49]. More than 400 cells were sequenced with around 6 % genome coverage of each cell. Due to the suboptimal genome coverage (~6 %), sequencing data was clustered in 54 kilo base (kb) sized "bins", to obtain proper statistical significance. These bins were then used to determine the copy numbers of genomic areas mapped to the healthy genome, which enabled the analysis of the evolutionary history of the cancer. Another example of genome sequencing of single cells is provided by Frumkin and colleagues [48] however, this was done without the use of any microfluidics or flow based techniques. In this study, cancer lineage relations were investigated by cutting out single tumor cells from tissue sections of mouse lymphoma by microdissection. Around 50 single cells were genotyped using Sanger sequencing and data was used to produce a "lineage tree" for the analyzed cells populating the tumor.

These studies provide a demonstration of the need of single cell sequencing in characterization of cancer, along with pros and cons of the utilized approaches. For example, microdissection on tissue samples allows the selection of specific cells. However, microdissection holds a general limitation in relation to number of analyzed cells: ~50 in Frumkin's study compared to >400 in the study by Navin et al. using a flow based single cell analysis. Moving forward, microfluidics based approaches are expected to ease the handling of sample and enhance the enrichment efficiency prior to the genome analysis. Moreover, the automation offered by

microfluidics also makes the experiments less prone towards human errors, such as the biased selection of cells by manually picking, and contamination of sample by human handling.

5.3.2 Single Cell Transcription Analysis for Cancer Characterization

The genetic information held by the genome is transcribed into RNA either as coding RNA such as messenger RNA (mRNA), or as non-coding RNAs such as transfer RNA (tRNA), ribosomal RNA (rRNA) and the regulatory RNAs (miRNA, siRNA and other types) [50]. The set of all the RNA molecules is henceforth termed transcriptome. The noncoding RNAs play direct roles in the cellular functions in e.g. constituting the core elements of protein synthesis, regulating gene expression or protection against viral infection. In contrast, the coding RNA, such as mRNAs in most cases are categorized as mediator molecules without direct cellular effect of their own. Rather, mRNAs convey the genetic information from DNA to the ribosome, where they are then translated into a polymer of amino acids, or a protein, as stated in the central dogma of molecular biology. The transcriptome in a cell, regardless of their functions, is extensively regulated by the cellular microenvironments and the request of the cells. Therefore, probing the transcriptome would provide important information about the cell. But why is “single cell” transcription analysis necessary? It has been reported that the cells encode a subtle set of analogue parameters to modulate the responses to environmental stimuli, e.g. Tay et al. have observed a heterogeneous activation of mouse fibroblast (3T3) cells in response to the signaling molecule tumor-necrosis factor (TNF)- α [51]. In this scenario, bulk measurements provide relatively limited biologically relevant information in the development of cancers, especially how individual cells respond to the changes of environment differently. Moreover, considerable cell-to-cell variation with regard to transcription is an inherent feature of cancers [52, 53]. Therefore, methods for analyzing the transcriptomic level of single cells is of critical need to bring forth new and insightful information about cancer development and the potential treatment against the diseases.

In the clinical settings, RNA based methods, such as quantitative polymerase chain reaction (qPCR), are widely used in cancer diagnosis and prognosis [54–56], where the expression level of specific genes is analyzed mainly in bulk set-ups. Hitherto, transcription analyses in the single cell manner have not yet found its way to the clinic. Nevertheless, microfluidics based single cell studies at the transcriptomic level, made possible by the commercialized system supplied by Fluidigm, have been widely adopted for scientific purposes by combining quantitative Reverse Transcriptase Polynucleotide Chain Reaction (qRT-PCR) and microfluidics in cancer characterization [57–63]. The commercialized single cell qRT-PCR platform is based on the prototype reported by Quake’s group (Fig. 5.5) [64]. The basic principle is to trap single cells sorted by Fluorescence-Activated Cell Sorting (FACS) in

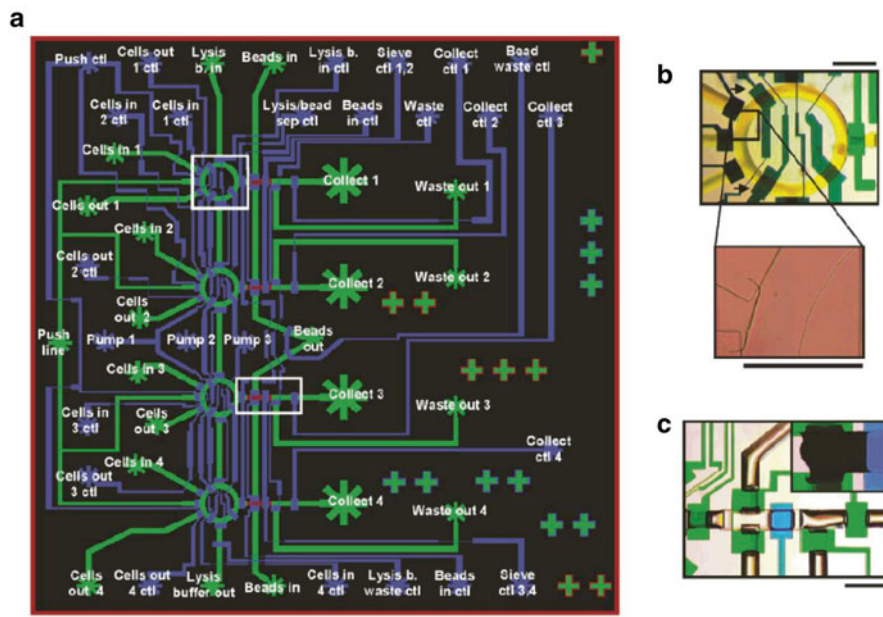


Fig. 5.5 Single-cell mRNA isolation and cDNA synthesis: (a) Overview of the device, which implements five steps, including cell capture, cell lysis, mRNA purification, cDNA synthesis, cDNA purification, in one integrated device. Flow channels and control channels are depicted in green and blue, respectively. The red unrounded (rectangular profile) flow channels are where designed for affinity column construction. White boxed regions are zoomed in (b) and (c), individually. (b) The lysis ring and an NIH/3T3 cell captured in the ring. (c) The affinity column construction area and a stacked column. Scale bars are 400 μm . Figure adapted from Ref. 62 with permission (Color online)

confined wells, and perform qRT-PCR reactions in the individual wells. The tiny nanoliter wells are generated by a fluidic circuit composed of a fluidic layer and a valve layer. These are separated by a thin elastomeric rubber member, as shown in Fig. 5.6. When pressurized, gas is applied to the valve layer, the membrane deflects and interrupts the flow in the fluidic layer [65]. As a result, trapped single cells may be analyzed within the individualized well without cross-contamination. Using this platform, Dalerba and colleagues have shown that human colon cancer tissues contain distinct cell populations of which the transcriptional identities mirror those of the different cellular lineages of the normal colon tissue [57]. In their work, more than 230 genes were evaluated in 336 single cells of three different colon epithelial cells and cancer cell lineages. Other reports have used a similar approach to investigate transcription in breast cancers [58, 59] and leukemia [60]. As an alternative to the commercial qRT-PCR platform, White and colleagues have developed a fully integrated microfluidic qRT-PCR device that implements all steps, including cell entrapment, lysis, reverse transcription and qPCR analysis in one single device, which produces a throughput of 300 cells per run [66]. The device is capable of

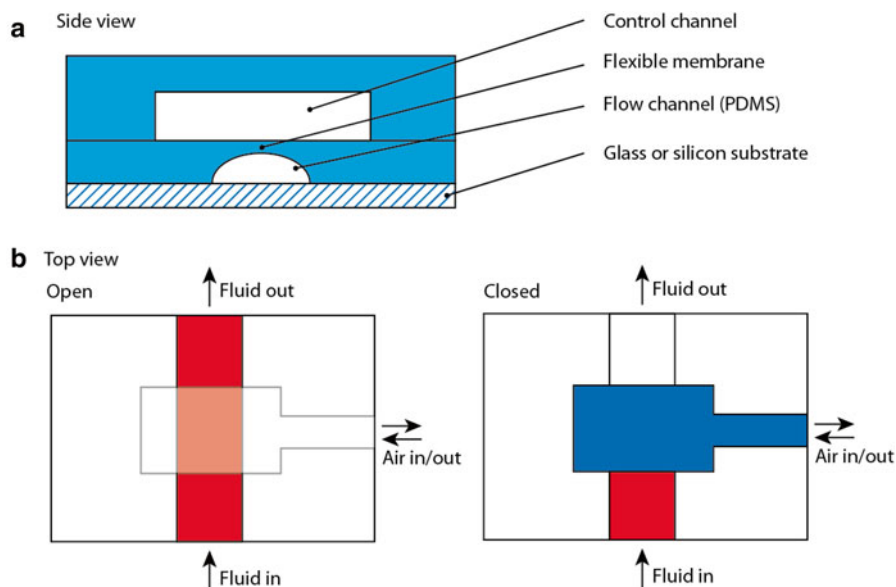


Fig. 5.6 Valve control in microfluidics: Monolithic valves in microfluidic devices are typically produced by soft-lithography techniques using polydimethylsiloxane (PDMS). Shown here is a typical two-layer PDMS “push-down” microfluidic valve. **(a)** Side view: A thin elastomeric membrane is placed in between the flow and control channels. **(b)** Top view: When the control channel is pressurized, the thin membrane would be “pushed” downward and close the flow in the flow channel. The flow channel is typically positioned orthogonal to the control channel

measuring RNA levels, as well as performing single nucleotide variant detection of single cells in a high throughput manner.

In contrast to the techniques mentioned above, capable of handling many cells simultaneously but only allowing a limited number of RNAs per cell (up to 96 RNAs) to be analyzed, Ramsköld et al. have recently presented a non-microfluidic based whole transcriptomic analysis of 12 single human CTCs using mRNA-sequencing [67]. The mRNA-sequencing study was performed on fewer cells compared to the microfluidic based qRT-PCR studies (12 cells in Ramsköld et al. [67] compared to >500 by qRT-PCR in Dalerba et al. [57]). In future studies, incorporation of microfluidics with mRNA-sequencing is expected to improve the throughput of transcriptional analyses of single cells.

5.3.3 Single Cell Protein Analysis for Cancer Characterization

The human genome codes for ~25,000 genes that are transcribed and translated into proteins [68]. The process of mRNA splicing makes the number of different proteins expressed by the cell many fold higher than the number of genes [69].

These proteins, which include active enzymes, are the main determinants for the cellular processes. Many currently used drugs in cancer therapy function by targeting specific enzymatic reactions in the cell [70–73]. Consistently, changes in the activity of the target enzyme may result in elevated chemo-resistance. Since the activity of enzymes are often modified on the post-translational level, such changes may not always be recognized at the genomic or transcriptional level [74], posing a need for analysis at the post-translational protein level.

Various methods have been applied for analysis of proteins at the single cell level, including mass spectrometry, enzyme-linked immunosorbent assay (ELISA), enzymatic detection or optical approaches [75–80]. Such analyses can involve measurement of the “expressed protein amounts” or “protein activity”. For example, Qihui et al. have proposed a microfluidic device that can isolate 0–5 cells in a 2 nL volume chamber and assay up to 11 different proteins per chamber on the chip using immunostaining procedures [78]. Regarding the throughput, the microfluidic device is able to isolate and assay 100 single cells per chip.

The strategy of detecting protein amount by immunohistochemistry is well established and antibodies can be designed to recognize almost all proteins and even specific features of a protein such as posttranslational modifications (e.g. phosphorylation and other modifications) [81–84]. However, in most cases it is the function of the proteins, such as the catalytic activity for enzymes, and not the amount per se, that determines the effect of the given protein on cellular conditions, such as health, drug response etc. From that perspective, our group has developed an array of DNA nanosensors that measure cancer relevant enzymatic activities, such as human topoisomerase I (hTopI) [85–87], topoisomerase II (hTopII) [88], tyrosyl-DNA phosphodiesterase 1 (Tdp1) [89, 90], which are emerging targets for anticancer therapy. The major working principle, termed rolling circle enhanced enzyme activity detection (REEAD), utilizes the catalytic activity of the target enzyme to generate an intrinsic amplification. Take the detection of hTopI activity for example (Fig. 5.7a), the DNA nanosensors are designed as a linear DNA substrate which fold upon itself, forming a dumbbell shape. hTopI is able to recognize the substrate, cleave from the 3' end and religate the 5' hydroxyl group, turning the linear DNA into a circularized one. Subsequent process is designed to differentiate the circles from the linear pieces by isothermal rolling circle amplification (RCA). As a result, the measured fluorescence represents the individual cleavage-religation event generated by active hTopI. Perhaps the best example demonstrating the potential of combining the microfluidics with nanotechnology for single cell analysis, water-in-oil droplets generated by a flow-focusing type of droplet generator is introduced to encapsulate individual cells along with the above-mentioned DNA nanosensors. The droplets provide a confined environment for the serial of biochemical reactions, enabling the enzymatic activities to be observed at the single cell level.

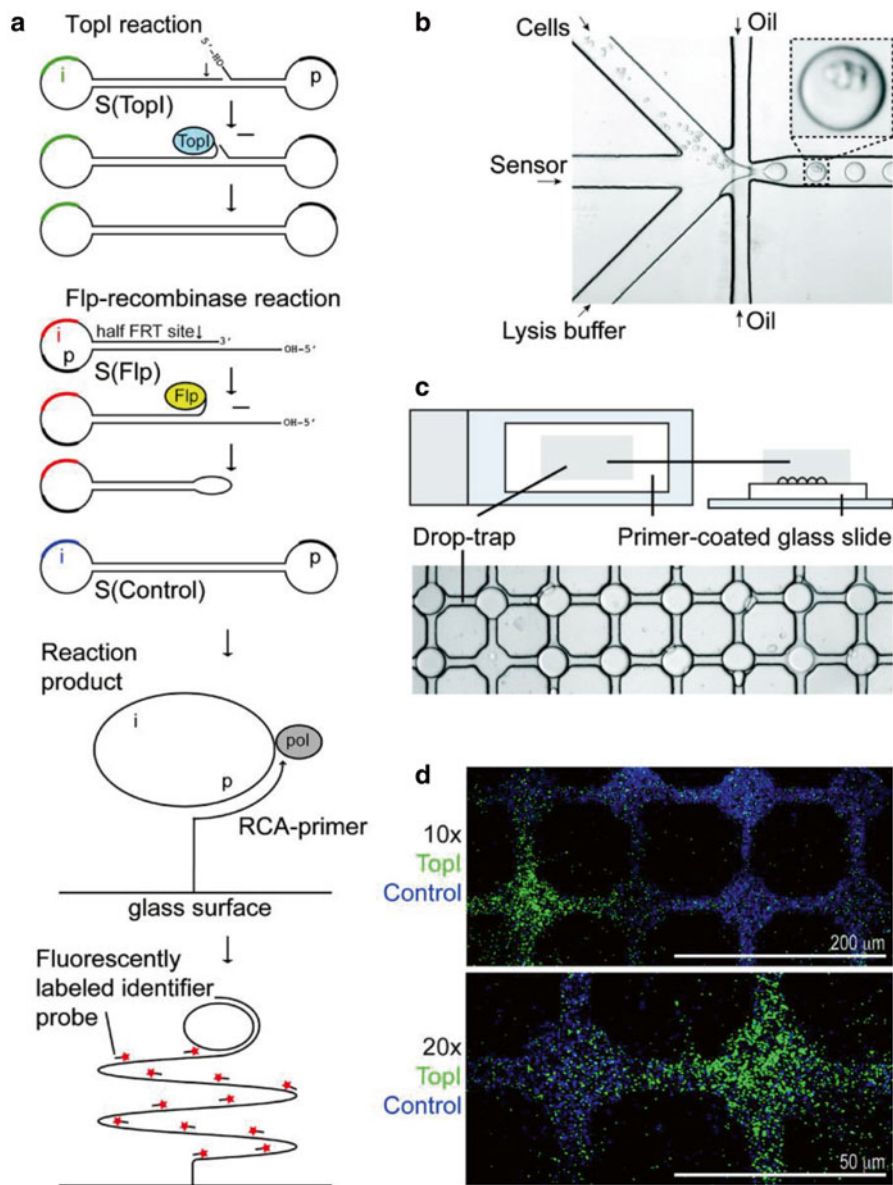


Fig. 5.7 Combining DNA nanosensors and microfluidics for single cell based analysis of enzymatic activities: **(a)** DNA substrates S(TopI) and S(Flp) are oligonucleotides that target cleavage-ligation by human topoisomerase I (TopI) and Flp, respectively. The detection, termed rolling circle enhanced enzyme activity detection (REEAD), initiates by recognition of enzymes to the substrates, which results in a circularized product. Subsequently, the circles allow a solid-support rolling circle amplification (RCA), which generate $\sim 10^3$ tandem repeat of amplified products. The results are visualized by fluorescence microscopy at the single-molecule level by hybridization of fluorescently labeled probes. **(b)** The droplet microfluidics is introduced to encapsulate individual cells along with DNA substrates and lysis buffer, in picoliters of water-in-oil droplets. **(c)** The droplets containing circularized DNA are then confined in a drop-trap on a primer-coated glass slide, where the RCA takes place. **(d)** The observed enzymatic activities from single cells are visualized as fluorescence signals: *green* (TopI) and *blue* (control). Figure adapted from Ref. 78 with permission (Color online)

5.4 Conclusion

Microfluidics-based approaches have been promoted for many biochemical applications, such as drug screening [91], nucleic acid amplification [92], and analysis of chemical reactions [93]. Therefore, it becomes a natural extension to take advantage of microfluidics for single cells interrogation. The device can be tailored to exploit physical and/or biological differences for isolation of particular cell types from a population of cells, such as enrichment of CTCs, and separation of CSCs from non-stem cells. The small dimensions of microfluidic devices have also enabled many unique features, such as gentle capture of live or rare cells [34], which avoids possible interferences for the subsequent analysis. Microfluidics also presents an opportunity to integrate many functions, such as isolation, biochemical reaction and detection, in a single device. For example, recent evidence has shown a high degree of heterogeneity even within a population of CTCs [94]. Therefore, combining CTC enrichment and molecular analysis at the single cell level is clinically important for the characterization of rare cell phenotypes, including CTCs in various stages during the cancer progression or perhaps study of how CSCs behave differently compared to non-stem cells. Furthermore, the isolated cells can be directed to next-stage analysis on-chip (e.g., genomic, transcriptomic or proteomic analysis), or on-chip cell culture as part of the analysis. Such integration will speed up the cancer characterization process while eliminating the intermediate sample transfer procedures typically required in macro-scale approaches.

Furthermore, advancement in cellular, microscopic, or nanoscaled molecular techniques is pivotal to the single cell analysis. The development of reliable biomarkers shall closely follow, if not precede, the emergence of microfluidics. For this purpose, immunostaining remains the prevailing technique for cellular or protein recognition. Recent progress in nanotechnology has joined the league by providing immunomodulatory agents engineered with nanostructure materials, including metallic nanoparticles, quantum dots, or nanotubes, to either improve the efficiency immunorecognition or enhance the detection sensitivity [95]. On the other hand, nanoparticles possessing unique photophysical properties, such as semiconductor nanocrystals and noble metal nanoclusters, also serve as unique fluorescence analogs in illuminating various forms of biological analytes through different signal transduction pathways. Taken together, the latest development of both nanotechnology and microfluidics is expected to encourage new excitement in the field of single cell analysis.

Despite the large variety of available approaches targeting and analyzing single cells, we are still in the infancy to uncover the implications of cellular heterogeneity. Many challenges have yet to come. One of the foremost, existing available approaches for single cell analysis is still expensive and technically complex, in particular those require high precision pumps or delicate valving control, which prevent the adaptation in the clinical studies. Future development of microfluidics-based single cell analysis is expected to reduce the production cost, sample consumption and to avoid human operational error by integration of sample preparation and subsequent data acquisition onto one single device. However, tradeoff of microfluidics remains on the

analysis speed, especially when it comes to large sample volume. The capability of parallel processing, when made possible, will hopefully accelerate the adoption of microfluidics for large-scale single cell analysis. Accessible single cell analysis, however, requires collective efforts including further engineering optimization of microfluidic systems and suitable nanoscaled molecular analysis approaches on the chip. The new possibility to reveal the characteristics of rare cells is expected to ultimately lead to clinical implications in the combat of cancers.

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Chapter 6

Nanotheranostics and In-Vivo Imaging

Brandon Buckway and Hamidreza Ghandehari

Abstract Advances in imaging and nanotechnology have provided the opportunity for simultaneous delivery and diagnosis. Modalities such as positron emission tomography (PET), single photon emission computerized tomography (SPECT), magnetic resonance imaging (MRI) and optical imaging have allowed researchers to visualize nano-sized drug delivery vehicles which carry payloads in order to coordinate disease treatment. This important tool can be termed “Nanotheranostics.” This chapter describes the potential utility of the combined approach. The importance of selecting the correct components for a particular disease will also be discussed allowing for researchers to design effective delivery systems in order to accelerate the development in this field.

Keywords Theranostics • Polymeric drug delivery • Image-guided delivery • Personalized medicine • Nanomedicine

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6.1 Introduction

One of the main challenges to successful treatment of disease is the difficulty in site specific delivery. Historically, patients have been administered treatments in a “one-size-fits-all” approach [1]. For example, many cancer patients are given a cocktail of chemotherapeutic drugs and are dosed based on the maximum tolerance levels of these drugs and the fact that these levels have worked previously in other cases. This approach often leads to toxic effects which can severely harm the patient. Although there have been many strategies developed both pre-clinically and clinically for improved delivery to diseased tissue, there has been relatively few successful or approved technologies by the regulatory agencies [2].

Targeted therapies, such as monoclonal antibodies have been developed which take advantage of the high expression of related proteins to selectively treat the disease and avoid off target toxicity. However, response to these therapies has been limited. This can primarily be attributed to a lack of improvement in efficacy in the cancer population based on the wide variety of tumor characteristics present in, not only an individual patient, but in the population as a whole. Targeted therapies today rely on a specific phenotype of a tumor and that phenotype may be different from tumor-to-tumor, thus, leaving some anti-tumor response in one tissue while lack of treatment in another. This leads to non-efficient patient care and disease recurrence in many patients. There remains a need to develop diagnostics which can predict targeted therapeutic efficacy in a patient by assessing whether the patient’s disease is expressing the specified target.

Theranostics, combine both diagnostic and therapeutic modalities and can be used for personalized therapy [3]. This includes the use of diagnostic tests including genetic testing, histology and/or imaging which can estimate potential response, predict safety and monitor progress of a specific therapy. One of the earliest examples of this type of approach involves the use of an excised biopsy tissue being evaluated for HER2 expression using the HercepTest[®] and other HER2 assays from the tumor of a breast cancer patient which gives prognostic information when treating with Herceptin[®] [4]. The patient can then be qualified or disqualified (stratifying patients) for the therapy based on the outcome of this diagnostic test, thus assuring that the patient is receiving a treatment that has a much greater chance of efficacy. However, diagnostics such as HER2 assays for breast cancer have their limitations. They rely on invasive tissue biopsies, sometimes leading to false negative results simply because of the limited sampling of the tissue. The use of modern imaging technologies can more accurately assess the tumors for prognostic factors because the whole diseased tissue is visualized in its natural environment.

Advances in modern imaging technologies have provided clinicians with the ability to actively assess in real-time the status of disease. High-resolution techniques such as X-ray computerized tomography (CT) and magnetic resonance imaging (MRI) can enable anatomical placement of tumors along with size and shape that can be progressively measured during treatment to monitor response. Through the advanced molecular imaging techniques used in nuclear medicine modalities such as

single photon emission computerized tomography (SPECT) and positron emission tomography (PET) imaging, clinicians can also understand the underlying processes effecting tumor treatment response, including metabolic state, hypoxia or proliferation that are occurring in each tumor within an individual [5, 6]. Each of these modalities can be combined to give a more accurate and precise treatment plan for patients, leading to better overall treatment. Personalized medicine can be further accomplished by combining these imaging technologies with advanced drug delivery approaches, thus, providing a more accurate and precise method for a theranostic system.

Among the many potential drug delivery vehicles for use in theranostics, one promising area is the use of nano-sized biocompatible nanomaterials. Nanomaterials can be functionalized with drugs, targeting moieties and imaging agents to provide multifunctional carriers for assessing both real-time localization of delivery to the tumor via imaging modalities. One example of a nano-sized material that has been demonstrated in the clinic is monoclonal antibodies. Zevalin[®], a radioimmunotherapy based on a radiolabelled monoclonal antibody, ibritumomab tiuxetan, recognizes CD20 receptors highly expressed on B-cells. This approved therapy for treatment of lymphoma has an imaging version and a therapeutic version. The patients are administered a low dose of gamma emitting ¹¹¹In-ibritumomab tiuxetan to confirm normal biodistribution observed by SPECT imaging. If no abnormal biodistribution is observed, the patient is administered a high dose of beta emitting ⁹⁰Y-ibritumomab tiuxetan for an effective radiotherapy. This example demonstrates the potential selection and qualification of a patient for a given therapy. Many other synthetic biocompatible nanomaterials can also be designed in a similar approach. Because of their macromolecular nature they can be tuned for long-term circulation and site specific delivery by targeting to various diseased tissues.

In this chapter, the historical aspects of imaging and theranostics, including several approaches in which nanomaterials can be utilized, will be discussed. The various imaging modalities will be presented along with each modality's potential for use with nanomaterial based theranostics. The basis for selecting appropriate nanomaterial carriers for delivery of both an imaging agent and a therapeutic also will be described along with their specific advantages and disadvantages. Therapeutic choices are also of particular importance when considering a theranostic system and, therefore, will be presented with technologies for site specific activation. Finally, the future directions and summary of the current status of theranostic nanomaterials for imaging will be discussed.

6.2 Imaging and Theranostics

6.2.1 *Historical Advances in Imaging Technology*

The beginning of imaging in the modern sense started in nuclear medicine with researchers using hand-held Geiger Counters to measure biodistribution of radioactivity without obtaining an actual image [7]. The first device for full body imaging was known as the rectilinear scanner which was fitted with a photographic

component for recording [8]. This led to the development of the gamma-camera for scintigraphic imaging of gamma-emitting radionuclides that were administered to living subjects. Various advances in gamma-cameras improved the quality and utility of the images through the development of 3D reconstructions of images using SPECT imaging. Further advances led to the development of images obtained from positron emitting isotopes also known as PET imaging beginning in the early 60s which provide even more sensitivity and higher resolution images of these radioisotopes. MRI was initially developed in the 70s and the use of paramagnetic contrast agents to visualize molecules distributing throughout the body has become one of the most useful tools in imaging [9]. Researchers have continued to develop other methods of detecting materials that are administered to subjects non-invasively through the use of fluorescence emission in optical imaging [10–12] and also sound transmission using ultrasound techniques [13–16]. Every one of these advances has opened a large field of research for the development of materials which can be safely administered to a patient and provide accurate real-time assessment of disease status or diagnostics.

Diagnostics have been utilized for treatment planning for many years. Diagnostic testing provides genomic, proteomic and anatomical information related to a disease state. These tests come from a wide range of sources, including tissue biopsies and blood tests for in vitro diagnostics or even more sophisticated minimally invasive methods such as imaging. Diagnostic testing utilizes these results to assay the disease status in order to take advantage of weak points in the disease that can be treated by a particular drug or therapeutic intervention. In vitro diagnostics for receptor expression have been found to be useful in qualifying patients for various targeted therapies. However, these assays are invasive and suspect to false positive or false negative results due to sampling error. A sample may or may not be reflective of the entire diseased tissue's expression profile. Also some may be too small and not detected within the patient and, therefore, tissue sampling for a diagnostic test cannot be performed in this case. Therefore, diagnostic methods involving imaging are extremely advantageous because they accurately assess the whole tissue for target expression.

6.2.2 *Theranostics*

Combining diagnostics with therapeutics in order to increase safety and/or efficacy was first described by Funkhouser and coworkers in 2002 [17] as theranostics. Successful development of theranostics has the potential of improving the treatment of diseases such as cancer, and can potentially lower healthcare costs and streamline regulatory approval of targeted therapies [18]. This concept of combining treatment with a diagnostic is gaining the attention of pharmaceutical companies. Many companies focused on development of therapeutics are able to partner with diagnostic companies to accelerate the drug development process by improving clinical trial outcomes. This is done by selecting patients based on potential therapeutic success on a “companion diagnostic.” Early examples of

this were Herceptin® and the HercepTest® [4]. The FDA simultaneously approved both Genentech's Herceptin® and Dako's HercepTest® for treatment and diagnosis of Stage IV breast cancer.

6.2.3 Modern Diagnostics and Use in Theranostics

Modern advances in medical imaging have brought theranostics to a whole new level of potential possibilities. Molecular imaging methods such as MRI, X-ray computed tomography (CT), ultrasound, SPECT and PET imaging are prevalent in many medical centers today and can accurately monitor disease status [7, 19, 20]. Theranostics are being accelerated in development with these imaging modalities. Imaging provides noninvasive and real-time information for diagnosis of a patient. MRI and CT are used for anatomical information and can be combined with other nuclear medicine modalities (SPECT and PET) to provide quantification of radiolabeled targeted probes and determine their localization within the patient. Another advantage to image guided therapies is the potential to use platforms which include both diagnostic and therapeutic on one material. Real-time assessment of both target and therapeutic can be measured simultaneously. As discussed earlier Zevalin® [21] is a clear clinical example of how theranostics can improve the treatment of not only cancer but other diseases as well. It improves the safety of radioimmunotherapy because the imaging reduces the risk of more radiation exposure to patients who would not see any benefit from the therapeutic version due to abnormal biodistribution. This strategy was also helpful for regulatory approval because it qualifies a patient for therapy and increases the overall chance for efficacious treatment of lymphoma without risking the patient to unnecessary dangerous ionizing radiation treatment due to lack of targeting efficiency.

6.2.4 Theranostic Methods

The methods that can be applied to nano-based materials for theranostics can be summarized in three main approaches (Fig. 6.1). Although many more approaches could potentially be imagined [22], these three main approaches basically encompass the methods utilized with nanomaterials with imaging and therapeutics. The first method utilizes the basic concept of monitoring deposition of a single nanomaterial containing both the imaging agent and the therapeutic within the entire subject. Pharmacokinetic and biodistribution information then provides a basis on which one can determine if enough of the therapeutic component reaches the intended site. If the therapeutic dose required at the site is known, the efficacy of the administered nanotherapeutic may be predicted. If the amount of carrier is insufficient, the subject can receive an increased dose on subsequent administrations. The second method utilizes imaging of the known diseased site for carrier accumulation and non-invasive triggering using

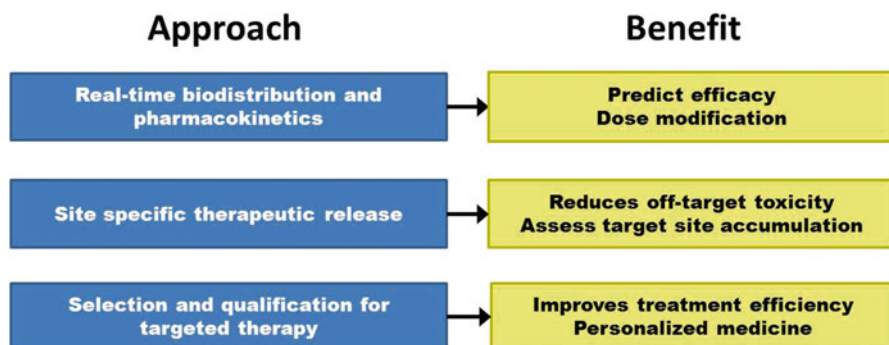


Fig. 6.1 Theranostic approaches for image based nanomaterials and their clinical benefit [22]

light, sound or heat to activate or release the therapeutic component. As long as the trigger is administered only in the required site, off-target toxicity is greatly diminished. If designed appropriately, therapeutic release can also be visualized by an imaging modality. The third approach for theranostics using nanomaterials with imaging is selection and qualification for targeted therapy. Targeted therapy relies on disease status which can change not only from patient-to-patient but from tumor-to-tumor and at different stages or times of the disease. An imaging version of the nano-carrier without the therapeutic can be administered before therapeutic versions are considered, to observe if the targeted receptor is adequately expressed. Thus reducing cost and optimizing time by allowing patients to be treated only with materials of which the desired therapeutic will be optimally effective. If the imaging results also demonstrate off-target accumulation that may compromise the subject's health, then the subject can also be eliminated from the current treatment and receive alternative treatments with better safety. These three methods describe the basic ideas and advantages behind the use of theranostics nanomaterials in imaging.

Imaging theranostics have great potential for improving the treatment of diseases. Many strategies can be utilized with these systems. However, there are several challenges for the development and successful use of these constructs when treating a disease. A wide range of imaging modalities, carriers, targeting strategies and therapeutics have been studied and currently only a select few have overcome the hurdles required for clinical use. The following sections will focus on the design criteria and components of theranostic nanomaterials for imaging.

6.3 Selecting the Appropriate Imaging Modality

Imaging modalities are becoming more sensitive, accurate and precise at detecting tracers in the body. Availability of sophisticated imaging devices is increasing; therefore, there is a growing need to rapidly develop probes for use in these different imaging modalities, especially in the area of theranostics. Personalized medicine is

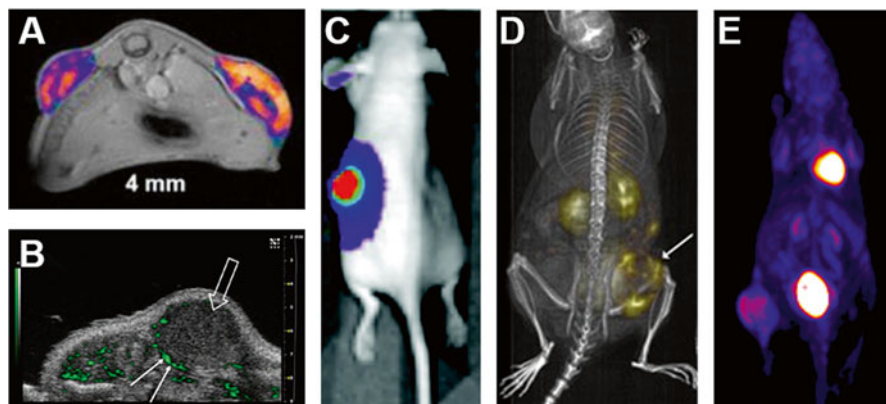


Fig. 6.2 Representative preclinical cancer images from common imaging modalities. *Arrow* denotes tumor. (a) Contrast enhanced MRI detailing the enhanced localization and tumor penetration of a gadolinium polymer conjugate treated with hyperthermia [23]. (b) Microbubble contrast imaging in a tumor model representing angiogenesis in the periphery of the tumor (*green*) [24]. (c) Representative luminescence imaging (optical imaging) of gene transfection events using silk-elastinlike polymer gene carrier [25]. (d) SPECT/CT image of ^{111}In -labeled HPMA copolymer localization in a tumor model [26]. (e) Representative PET image of tumor bearing mouse model injected with a proliferation biomarker (Courtesy of Jeffrey T. Yapp, Ph.D. Center for Quantitative Cancer Imaging at the Huntsman Cancer Institute, University of Utah) (Color online)

the ultimate goal of imaging in the clinic and its full potential is yet to be realized. This section will present the different strengths and weaknesses associated with various imaging modalities. Many of the modalities can be combined in such a way as to compensate for lack of sensitivity or accuracy in detection of different probes. The development of nanoparticles which have multiple methods of detection for multimodality imaging have been investigated providing information that is not possible with the use of one alone. Figure 6.2 is an overview of some of the available modalities for imaging. Table 6.1 shows a brief summary comparison of the different modalities. A more in-depth discussion is provided for each modality in the following subsections.

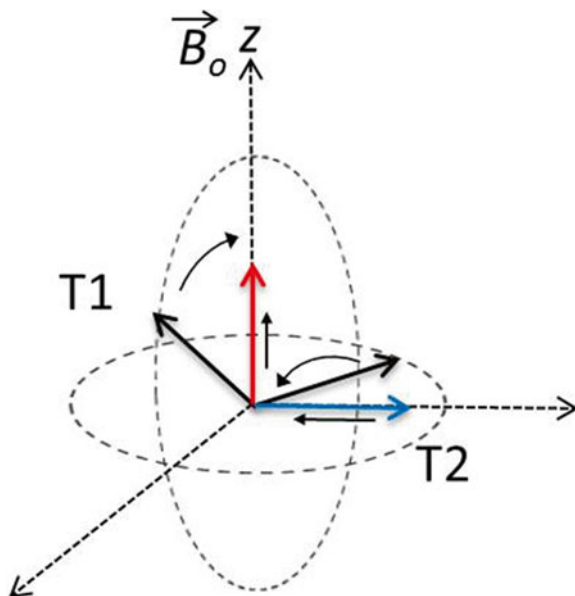
6.3.1 Magnetic Resonance Imaging

MRI is primarily used in the clinic for anatomical imaging. MRI signals are produced from changes in magnetic orientation from radiofrequency pulses of aligned protons in a strong magnetic field [28]. Signals from MRI are measured based on two typical responses, T1 and T2 relaxation demonstrated in Fig. 6.3. Typical use of MRI measures proton T1 or T2 relaxation signals in the body for high resolution images containing soft-tissue anatomical information. Differences in each proton's environment influence the T1 or T2 relaxation rate which in turn causes the variations that can be

Table 6.1 Summary comparison of the main imaging modalities [7, 27]

Modality	Temporal resolution	Spatial resolution	Quantitation	Depth of penetration	Probe sensitivity	Imaging agents	Clinical use	Relative cost
X-ray computed tomography (CT)	Minutes	0.5–1 mm	No	Limitless	Not determined	Iodine, barium suspensions	YES	\$\$
Magnetic resonance imaging (MRI)	Minutes-hours	~1 mm	Semi-quantitative	Limitless	10^{-3} – 10^{-5} M	Gadolinium, iron oxide nanoparticles, ^{19}F -fluorine	YES	\$\$\$
Fluorescence molecular tomography (FMT)	Seconds-minutes	1–3 mm	Quantitative	<2 cm	10^{-9} – 10^{-12} M	NIR fluorochromes	NO	\$
Ultrasound (US)	Seconds-minutes	0.01–2 mm	No	mm-cm	$\sim 10^{-12}$ M	Microbubbles	YES	\$
Single photon emission computed tomography (SPECT)	Minutes	8–10 mm	Semi-quantitative	Limitless	10^{-10} – 10^{-11} M	Gamma emitting isotopes	YES	\$\$
Positron emission tomography (PET)	Seconds-minutes	5–7 mm	Quantitative	Limitless	10^{-11} – 10^{-12} M	Positron emitting isotopes	YES	\$\$\$

Fig. 6.3 Diagram of T1 and T2 relaxation in MRI. T1 relaxation is the rate at which the magnetic vector realigns or net alignment (*red arrow*) increases on z-axis which aligns with the external magnetic field after a 90° radiofrequency pulse. T2 relaxation is the rate at which the magnetic vector disappears or net x–y vector (*blue*) decreases in the x–y plane after a 90° radiofrequency pulse [29] (Color online)



converted into an image. Different materials used as MRI contrast agents also have various effects on protons in the body with these two relaxation signals [7, 28]. Gadolinium has been used as a contrast agent for T1 relaxation in image based theranostic systems [30]. An additional method for MRI imaging is the detection of other paramagnetic elements such as ^{19}F called magnetic resonance spectroscopy [31]. ^{19}F has been incorporated into many carriers [32–35]. One example is the system developed by Porsch et al. containing ^{19}F conjugated to amphiphilic polymers [36]. These polymers forming a micelle structure were subsequently loaded with doxorubicin for a theranostic drug delivery approach that exhibited MRI suitable signal-to-noise ratio (SNR) in phantoms [36]. However, one of the main challenges for MRI contrast agents is the lack of sensitivity of the MRI scanner. Large amounts of contrast are needed in order to produce a signal distinguishable from background. This makes MRI very difficult to use quantitatively. Although some techniques are in development to improve quantitative capacity of MRI, the physical nature of sensitivity is a large hurdle to overcome. Nonetheless, MRI is a valuable tool for anatomical imaging that can be used in conjunction with other imaging modalities which are more quantitative.

6.3.2 X-Ray Computed Tomography

X-ray computed tomography or CT is another method in which anatomical information is provided. CT uses X-ray projections that interact with high electron dense materials through a subject in multiple planes [7]. These planar images can then be reconstructed based on computer algorithms which produce a high-resolution image of the body.

Resolution depends on the electron density of the material and, therefore, primarily returns high-resolution images of hard-tissues. Contrast agents are based on heavier atoms such as iodine or barium. Iodine contrast agents are commonly used for angiography studies but require large frequent doses for a sustained signal output. The need for highly dense materials for contrast is the main limitation for use of CT with image-guided delivery. Therefore, CT is utilized in a multimodality approach which provides 3D anatomical information in tandem with other imaging information modalities. Risk due to ionizing radiation from prolonged CT exposure is of some concern. However, CT has made a substantial impact in detection and measurement of tumor sizes [37].

6.3.3 *Optical Imaging*

Optical imaging is based on light emitting probes which can be detected by a camera [10, 12]. Optical imaging has been successfully used for many years for the detection of molecular processes in *in vitro* assays. However, *in vivo* the challenge becomes increasingly difficult when trying to penetrate tissues which rapidly attenuate light signals. The attenuation of light prevents absolute quantification and limits resolution [38]. A modern technique called fluorescence molecular tomography (FMT) can measure signals from the visible to near-infrared spectrum (500–900 nm) of fluorescent probes in multiple orientations and use mathematical models which predict attenuation in the subject to produce three-dimensional images of probe localization [19, 39]. FMT is currently available for small animal imaging research, but clinical translation is yet to be viable. The tissue penetration in a human subject is more difficult. Therefore, FMT and other optical imaging techniques remain at the preclinical level. However, one of the advantages to using fluorescent probes in the development of theranostics is the potential to visualize drug release from a nano-construct. Recent developments in dyes which are conjugated to nanoparticles can be activated once released by targeted mechanisms from the nanoparticle surface [40]. Fluorescent dyes when in proximity (i.e., conjugated to the surface of a nanoparticle) lead to a strong fluorescence quenching due to fluorescence resonance energy transfer (FRET). Once the dyes are released and quenching is stopped, the signal will be visible and the resulting signal can be related to a mechanistic process [40]. Other imaging modalities are not capable of having signals that are activated in relation to cellular responses *in vivo* such as SPECT or PET imaging. Radionuclide signals are constitutively active and cannot be suppressed unlike fluorophores. One example of such a system provides information of cathepsin B protease activity [41]. This is performed by a FRET designed peptide sequence conjugated with two terminal fluorophores. When the peptide sequence is cleaved, FRET interactions cease and the signal representing protease activity can be visualized via FMT imaging. A similar type of system could potentially be used to visualize drug release by containing enzymatically degradable linkers conjugated with drug and FRET capable fluorophores. The released fluorophore would produce the optical signal that could be related to the release of drug from the linker.

6.3.4 *Ultrasound*

Ultrasound is probably the most cost effective and safe imaging modality available in the clinic today [7, 27]. Ultrasound utilizes high-intensity ultrasonic waves mechanically produced from a transducer. The sound waves then reflect or scatter from different tissues which can be detected by the transducer and converted into images. One advantage that ultrasound has is actual real-time imaging. Images are returned within seconds and, therefore, provide the highest temporal resolution available among the different modalities. Temporal resolution is a term describing the ability to distinguish between individual events [7]. Contrast agents for ultrasound are limited to gas bubbles which have specific properties that resonate in the 1–20 MHz frequency range, producing highly specific signals that can be recognized by the ultrasound device [19]. Ultrasound is limited to only microbubbles for visualizing probes and is, therefore, relatively limited in application to theranostics. It also suffers from a lack of penetration (Table 6.1) and requires contact of the device to the subject. Ultrasound resolution is highly dependent on the type of tissue and its depth and is, therefore, difficult to directly compare against other modalities. In the best case scenario, resolution is on the order of 10–100 μm [7]. Microbubbles are used as carriers for theranostic delivery. Further refinement in manufacturing is needed for clinical translation of this approach for theranostics.

6.3.5 *Single Photon Emission Computerized Tomography*

Nuclear medicine techniques such as single photon emission computerized tomography (SPECT) detect gamma emission from radioisotopes which are administered to a patient for purposes of diagnosis and treatment [42]. SPECT is a descendent of older gamma scintigraphic methods which were only capable of producing planar images that were not quantifiable and had very poor resolution. SPECT imaging takes modern advances in scanner and computer technology to obtain single γ -ray emissions using two to three gamma cameras that rotate around a patient who has been administered a gamma-emitting isotope tracer. Gamma emission is detected through thick collimators, plates of lead or tungsten, with small holes between the subject and gamma detector, that only allow photons emitted in the 90° direction of the gamma camera to interact with the detector. SPECT relies on reconstructing these multiple projections into a 3D image that gives precise localization of radioisotope biodistribution. Because only a select few gamma emissions are detected by the camera, the sensitivity is compromised to some degree due to lack of sufficient detector events [42]. Images are also dependent on the energy of the γ -rays emitted from the radioisotope. Lower energy radioisotopes have more attenuation than higher energy radioisotopes. Attenuation increases the scatter and noise detected from the camera, thus, compromising the resolution of the image. The quality and usefulness of images for image-guided approaches with SPECT are highly dependent on the type of radioisotope being utilized.

Table 6.2 Radioisotopes for SPECT imaging [19, 46, 47]

Radionuclides	T _{1/2}	γ-Ray energy
^{99m} Tc	6.01 h	140 keV
¹¹¹ In	2.83 days	173, 247 keV
⁶⁷ Ga	3.26 days	93, 184, 300, 393 keV
¹²³ I	13.3 h	159 keV
¹³¹ I	8 days	365 keV
⁶⁷ Cu	2.58 days	184.6 keV
²⁰¹ Tl	3 days	69–81, 167 keV
¹³³ Xe	5.2 days	81 keV

Radioisotopes for SPECT imaging must be selected based on the length of time needed to acquire information; otherwise, the half-life of the radioisotope must match the biological process that is being monitored. A list of clinically utilized radioisotopes for SPECT imaging with their emission properties are described in Table 6.2. Many isotopes can be utilized in SPECT for detection and the majority of them can be attached to nanoparticles via metal chelation, ionic interaction or covalent linkage. Transition metal based radioisotopes are easily conjugated to nanoparticles using stable bifunctional chelators of metals [43]. Some radioisotopes can be associated with a nanoparticle by ionic charge interactions [44]. Others, including halides, form stable covalent bonds in order to radiolabel a nanoparticle [45]. Many strategies exist to radiolabel a nanoparticle. Consideration for the method of labeling must not interfere with other functions (i.e., drug and targeting) for theranostics. One interesting strategy for SPECT imaging is the ability to image two different radionuclides with different energies at the same time. This may allow for imaging of distinct processes within a given construct. For example, a study was conducted using an ¹¹¹In labeled targeted peptide and ¹⁷⁷Lu labeled control nontargeting peptide injected into the same mouse in order to visualize differences in accumulation without inter animal variations [48]. The dual isotope approach could be applied in mixed micelles where the hydrophobic and hydrophilic components are labeled with different isotopes and could potentially give information as to the breakdown and ultimate fate of the two components.

SPECT is a promising modality for theranostics because it can provide accurate information on the relative location of carriers. However, there is the risk of ionizing radiation exposure that limits the amount of radioactive exposure a patient can receive. It also does not provide anatomical reference. This has been overcome by multimodal approaches such as combining images with both CT and MRI. The combined modalities can more accurately pin-point where the carriers are accumulating and give accurate information predicting both potential efficacy and safety of image guided constructs. The major challenge for SPECT imaging is obtaining accurate quantitative results. Detection events required for accurate quantitation are limited due to the use of collimators and a large range of gamma emission energies that are scattered and attenuated differently in tissue. Longer imaging times are also required for SPECT which can exaggerate motion effects leading to increased signal noise. There are methods for both attenuation and scatter correction which are being developed to make SPECT more

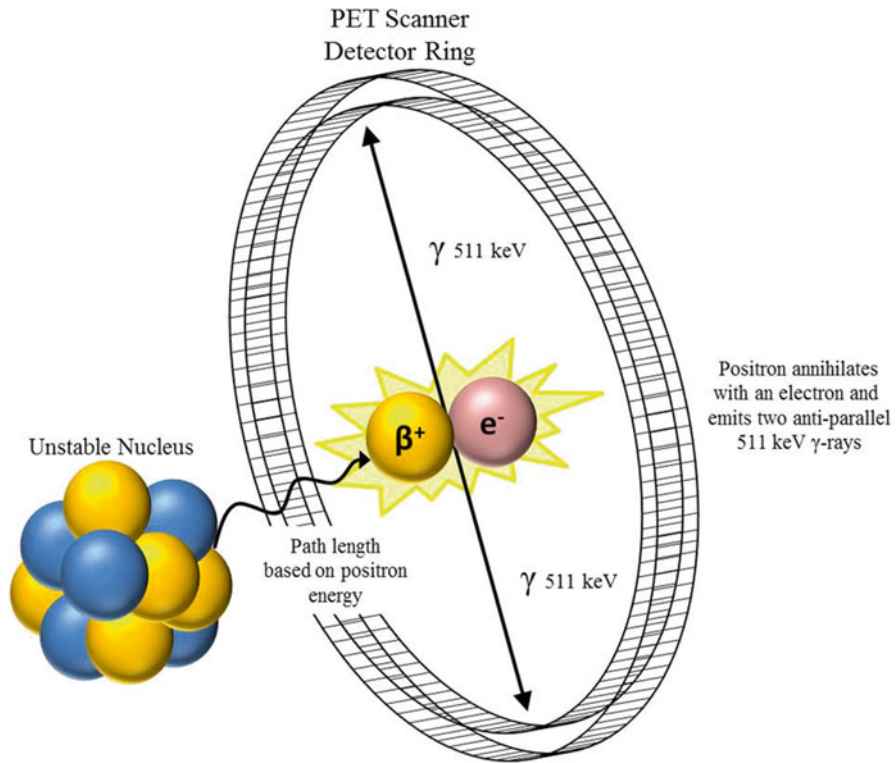


Fig. 6.4 Positron emission tomography [5]

quantitative. However, these methods and capacities vary greatly among scanners and groups. Ideally, further development and common adoption of correction techniques may eventually lead to more routine quantitative capability of SPECT.

6.3.6 Positron Emission Tomography

Positron emission tomography (PET) is another nuclear medicine technique which has made a major impact in the field of cancer treatment [5, 49]. PET detection is based on radionuclides emitting positrons which interact with nearby electrons and annihilate into two antiparallel gamma photons each with signature 511 keV energy. This allows for coincidence detection by a circular array of detectors that can trace back along the line of response to the origin of emission and produce high-resolution images of radioisotope probes. A basic diagram of PET is shown in Fig. 6.4. PET can detect radioisotopes down to the pico-molar (10^{-12}) range and has limitless depth penetration due to the high energy of its 511 keV gamma-rays [7].

Table 6.3 Physical characteristics of common positron emitting radioisotopes [5, 19, 50]

Radionuclides	T _{1/2}	Max β ⁺ energy (MeV)	Availability ^a
¹⁸ F	110 min	0.69	+++
¹¹ C	20.4 min	0.96	+++
¹⁵ O	122.2 s	1.7	+++
¹³ N	9.97 min	1.20	+++
¹²⁴ I	4.2 days	2.14	+
⁶⁴ Cu	12.7 h	0.65	+
⁸⁹ Zr	3.3 days	0.897	+
⁶⁸ Ga	67.7 min	1.90	++
⁸² Rb	75 s	3.18	+

^aAvailability based on estimated cost and production facilities

In comparison with SPECT, PET has a much higher count rate and better resolution, thus, providing the ability for accurate quantitation of imaging agents. PET scanners are also combined with CT and other techniques in order to provide attenuation and scatter correction, thus, increasing its quantitative ability in comparison to SPECT. Due to the fact that only one gamma energy window is needed for detection around 511 keV, these correction techniques can be simplified when compared to SPECT. Radioisotopes used in the clinic for PET are generally short-lived and for the most part must be produced locally using a cyclotron. This increases the costs and availability of PET radionuclides. Like SPECT, ionizing radiation also limits the ability of patients to be continually administered radioisotopes for research studies. Table 6.3 lists some of the isotopes used in PET imaging with details of the characteristics and properties. PET also does not provide anatomical reference; thus, almost all PET imaging devices are coupled with CT detectors for precise localization of signals with the patient. The CT images can also be used to correct for attenuation in the imaged subject for improved resolution and quantification.

Selection of radionuclides for PET imaging is important in that it can affect the resolution and relevance of the research study. The initial positron energy determines the path-length from the parent nuclide to the annihilation event. The higher the energy, the longer the path-length, thus, increasing image noise and reducing resolution [19]. Like SPECT, isotope half-lives must match the relevant biological process intended to be measured. This represents a significant challenge in imaging with PET, since most of the clinically available radionuclides have ½-lives on the order of minutes (¹⁵O ~2 min ½-life) to a couple of hours (¹⁸F ~110 min ½-life) which does not match the biological half-life of many macromolecular systems which circulate in the bloodstream for hours and days before being taken up by the target or eliminated. However, the majority of these short-lived isotopes are basic elements found in biology and, therefore, can be incorporated into drugs, sugars and other biological entities without influencing their structure.

A recent study using a positron emitting ⁶⁴Cu radionuclide conjugated to HPMA copolymers containing angiogenesis targeting peptides demonstrated measurable

increased tumor localization by PET imaging in prostate tumor bearing mice [51]. Due to ^{64}Cu also having some beta-emission, the radionuclide can serve a dual purpose for imaging and therapy. This is an example of how radioisotopes can be utilized in theranostics. Clearly, more research and effort in this area need to be conducted to improve the availability of radionuclides. PET represents the future of nuclear medicine imaging and has already influenced the treatment of cancer profoundly [52]. However, costs and lack of radionuclide availability are detrimental for rapid development of theranostics based on PET imaging.

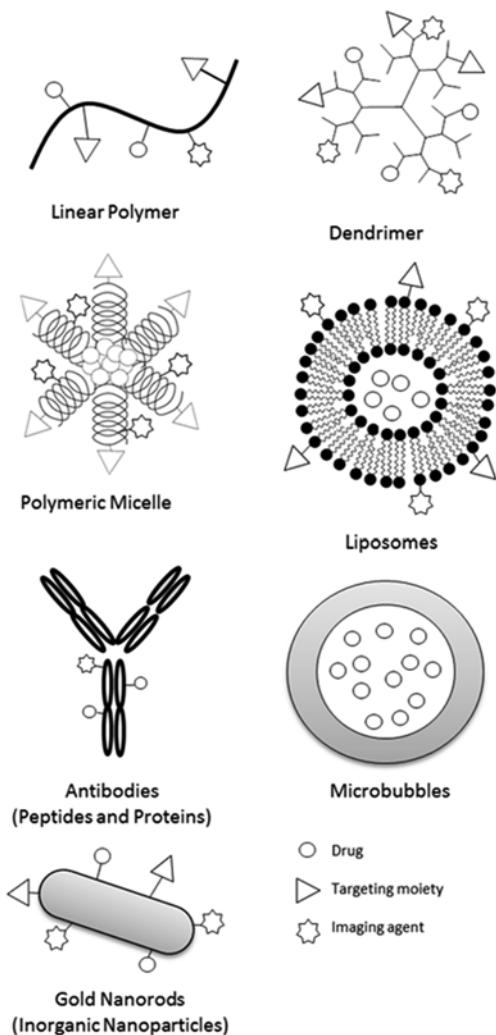
6.4 Choosing the Appropriate Nanomaterial Carrier

A large investment in research has been focused on the design and development of materials for targeted delivery. Each has their advantages and disadvantages for delivery of imaging and therapeutics. Some have had extensive clinical experience or testing. Recently, constructs based on nanomaterials have become a promising area of research [53–55]. The use of nano-scale constructs of 1–100 nm in size as therapeutic delivery systems has generated a promising venue for image-guided theranostics [56–58]. This property allows for multiple components to be incorporated on the surface or within these materials for targeted delivery of both imaging agents and therapeutics. They are also able to be tailored in ways as to interact with fenestrations, channels and surfaces in unique ways because of their small size [54]. Many types of carriers are synthesized in the nano-size range. Figure 6.5 displays some of the most investigated systems that have been utilized for image-guided delivery. The following are examples of nanoparticle based carriers that can be used as theranostics.

6.4.1 Water Soluble Polymers

Some examples of water soluble polymers that have been extensively studied are poly(ethylene glycol) (PEG), *N*-(2-hydroxypropyl)methacrylamide (HPMA) and poly-L-glutamic acid (PGA) [59–61]. A wide body of research has been conducted using these polymers. These polymers can have a linear or branched structure. PEG and HPMA copolymers are inherently non-degradable unless degradable sequences are included within the backbone of these copolymers [62–64]. However, PGA is naturally degradable in the body. The main advantage of polymer carriers is the ease to control size and afford multifunctionality [61]. Water soluble polymers, such as those described above, are biocompatible and have limited recognition by the immune system [65, 66]. This provides an advantage because they can circulate in the body for an extensive period of time, thus, increasing the likelihood of their ability to interact with the targeted tumor sites. PEG and HPMA copolymers also have properties which impart steric hindrance to degradation of the attached payload of drug and targeting moieties [67]. One advantage that HPMA copolymers have is the ability to

Fig. 6.5 Examples of carriers for nanotheranostics



incorporate multiple therapeutics and targeting agents into the side chains. Linear PEG has limitations in the amount of payload because it is limited to end group functionalization [60]. Some of the main disadvantages to traditional HPMA copolymers are nonbiodegradability and potential long-term exposure in the body leading to possible toxicity effects [68, 69]. PGA is biodegradable; however, this may not be ideal for a combined image-guided therapeutic because breakdown of the PGA backbone will eventually lead to imaging agent being separated from the carrier. HPMA copolymer doxorubicin drug conjugates were evaluated in clinical trials [70]. A matching HPMA copolymer was available with attached radioisotopes for imaging the biodistribution of the copolymers in patients [71]. The drug conjugate failed due to the lack

of efficacy in some patients [72]. However, the trial may have had more success if they had used the imaging version of the HPMA copolymer for preselection of patients that would have been more susceptible to the HPMA copolymer-drug conjugate [22]. Recently, Yuan et al. designed integrin targeted HPMA copolymer-DOTA-⁶⁴Cu conjugates for PET imaging of angiogenesis [51]. These conjugates demonstrated increased uptake in tumors in-vivo. This approach could be applied to many HPMA copolymer drug conjugates for a combined imaging and drug delivery approach.

6.4.2 Polymeric Micelles

Micelles are constructed from a combination of hydrophobic and hydrophilic components or segments [73–75]. The structure is formed in aqueous solutions by self-assembly of a hydrophobic core and hydrophilic shell. Many chemotherapeutics are hydrophobic leading to solubility issues [73]. Hydrophobic drugs can be associated with the hydrophobic core and improve their solubility and protect them from metabolic enzymes in the bloodstream while associated with the nanoparticle [76]. Imaging agents and targeting groups can be associated with the hydrophilic components and allow for imaging of micelle biodistribution within the body [76, 77]. Another advantage that micelles have is their ability to have triggered therapeutic release based on pH change [76, 78]. Micelles however, have limited stability in the body and progressively breakdown to the initial components, especially when encountering biological milieu [54]. Therefore, imaging of the nanoparticle has limitations due to the eventual breakdown of the micelle. One example of a block copolymer micelle image-guided drug delivery system incorporated folate targeting, pH sensitive drug release of doxorubicin and Cy5.5 for optical imaging [79]. The polymeric conjugate was able to successfully image targeted conjugate localization and treatment in HeLa tumor cell bearing nude mice [79]. Another example of a promising micelle carrier was developed by Decato et al. [80]. This construct incorporates several perfluoro-*tert*-butyl groups as a fluorophilic tail in combination with hydrophilic PEG groups which self-assemble into stable spherical micelles. They demonstrated that these conjugates can potentially be detected by ¹⁹F-MRI and in the future could be used for delivery of hydrophobic drugs and serve as potential imaging and drug delivery vehicles.

6.4.3 Dendrimers

Dendrimers are multibranched polymeric systems with a central initiator core [81, 82]. Dendrimer size can be precisely controlled by successive additions of layers of branched units. Dendrimers have exceptionally low polydispersity and in some cases can be unimolecular [83]. Because of their unimolecular structure, regulatory hurdles to clinical approval can be significantly less challenging compared to other polydisperse polymers [84]. The branched structure allows for specific control of

the amount and density of functional groups on the surface of the dendrimer. This can be utilized for surface decoration with various imaging agents, targeting moieties and drugs. With larger generations of dendrimers, a hollow core can be formed which can also be loaded with hydrophobic drugs. Several limitations, however, exist with dendrimers. Large dendrimers become difficult to synthesize, because the larger the dendrimers become, the more steric hindrance prevents chemical ligation of the branching units and surface modifications [85]. The dendrimers that have been extensively investigated for clinical development are poly(amido amine) or PAMAM dendrimers [82, 86]. Depending on surface charge and generation, PAMAM dendrimers can have toxicity based on the charge density of the surface when reacting with cells [87, 88]. One of the key aspects of PAMAM dendrimers is the alternating generations terminating with either primary amines or carboxyl groups. One limitation with PAMAM dendrimers is that their typical branched layered structure contains only one type of surface group for conjugation of targeting moieties, imaging agents and drugs. This limits the degree of control over how much of each component is incorporated unless different dendrons are constructed during conjugation. Several groups have begun strategically synthesizing “Janus” dendrimers which are able to have several different types of surface functional groups, thereby providing better surface conjugation control [89]. One particular example shows great promise as a platform for a theranostic [90]. Although the study only demonstrated the dendrimer with a near-infrared agent for optical imaging, the conjugate did not show any toxicity in presence of T98G human cells and the unique trifunctional surface groups showed potential for development as a theranostics treatment.

6.4.4 Liposomes

Liposomes are spherical lipid bilayer constructs with an aqueous core which can contain drugs or other therapeutic agents [91, 92]. Their lipid bilayer structure mimics the biological environment of a cell but can be made of many different materials with the majority using phospholipids. Liposomes are among the most studied drug delivery systems with many clinically approved formulations [93, 94]. They are primarily formulated to entrap hydrophilic drugs within their aqueous core or associate hydrophobic drugs into their lipid bilayer [91]. Due to their macromolecular nature, they can passively target tumors via the EPR effect [95]. Of particular mention is Doxil[®], a liposome-based formulation of doxorubicin for treatment of advanced ovarian cancer [93]. Liposomes protect their encapsulated drugs from metabolism in the bloodstream and fuse with their biologically similar membranes of targeted cells. Once fused with cells, they release the payload into the cell for increased therapeutic efficacy. Phospholipids can be modified with different imaging agents and targeting moieties and be inserted into the lipid bilayer in order to make theranostic nanoparticle systems [92]. One particular challenge recognized by the FDA with liposomes is reproducibly controlling the size and polydispersity [96]. They are also subject to

rapid degradation or reticuloendothelial system (RES) clearance from the body [97]. Methods have been developed to overcome some of the issues such as incorporation of PEG for “stealth” liposomes [98]. However, the instability of these systems in-vivo remains a particular challenge when considered for theranostics. Regardless, complex systems have been attempted to overcome these issues. For example, magnetic resonance high-intensity focused ultrasound (MR-HIFU) has been used with temperature sensitive liposomes loaded with doxorubicin [99]. MR imaging was used to guide the placement of localized hyperthermia within the tumor for triggered release of the liposomes in Vx2 rabbit models. This study shows the potential application of a liposomal formulation for image guided therapy.

6.4.5 Antibodies and Other Proteins

Nature has developed its own methods of recognizing diseased tissue which can be utilized for delivery of both drugs and imaging agents in a theranostic approach. Antibodies are of particular interest in delivery because of their biocompatibility and intrinsic ability to target antigens within the body, including those highly expressed in tumors. Many monoclonal antibodies (MoAbs) can be therapeutically active and have been approved, or are in, clinical trials for the treatment of cancer [100]. Therapeutics such as drugs and radionuclides conjugated to MoAbs have been shown to be effective for targeted delivery and image guided therapy [101]. One of the major concerns in using antibodies as image-guided therapies is their method of manufacture. MoAbs are produced from nonhuman sources (typically mice) and this can induce immune recognition within a patient by human anti-mouse antibodies (HAMA) that can either reduce the effectiveness of the therapeutic conjugate or cause severe life-threatening immune reactions [102]. However, advances in this technology have minimized mouse component of the antibodies, thus minimizing the potential for HAMA. Image guidance helps to select patients which may suffer from this effect as demonstrated in radioimmunotherapy [21]. Antibodies are not the only proteins that are used in therapeutic delivery. Albumin, a native protein in the blood-stream, serves as a macromolecular delivery system that is biocompatible and capable of being imaged. For example, a photosensitizer, chlorin e6, covalently linked to albumin formed nanoparticles that were used for image guided drug delivery [103]. Tumor localization was observed by optical imaging of the photosensitizer and then subsequently irradiated with light in the tumor for anticancer treatment in a mouse model.

6.4.6 Inorganic Nanoparticles

Nanotheranostics can also encompass an array of inorganic materials such as quantum dots, gold and iron oxide nanoparticles. Metal elements exhibit interesting properties when synthesized in the nano-size ranges. For example, gold nanoparticles can absorb

light in the near infrared spectrum and produce heat for localized hyperthermia [104]. Quantum dots are semiconductor nanocrystals made of transition metals that have tunable emission spectra that are much stronger than organic dyes [105]. Iron-oxide nanoparticles can be detected using MRI. All of these materials can be decorated with targeting agents, imaging agents and therapeutics for image-guided therapeutic delivery in cancer [106]. One of the main concerns with these constructs is their toxicity. Typically, these constructs do not breakdown in the body and, therefore, are deposited in tissues for an extended period of time [106]. The nature of their toxicity is still being investigated, but the long-term effects of some of these materials raise many questions as to whether they can be used safely in the clinic.

6.5 Therapeutic Considerations for Nanotheranostics

The appropriate therapeutic type for a delivery system is an important part of designing an effective theranostic. In this section we will describe three major classifications of therapeutics that have been used in theranostics with considerations for their advantages or disadvantages that need to be considered before selecting one for any given system.

6.5.1 Chemotherapeutics

Chemotherapeutics are the main-stay of treatments for advanced stages of cancer. While surgical methods to remove known tumor sites are preferred, small invasive tumors are difficult to detect and remove. Therefore, they require small molecule agents to penetrate and destroy tumor sites which are not accessible. However, many small molecules are nonspecific in their cytotoxic action. Therefore, they cause a variety of side effects that reduce quality of life and endanger the patient. Most chemotherapeutics can be broken down into a few classes: alkaloids, antibiotics, platinates, antimetabolites, topoisomerase inhibitors, mitosis inhibitors and others [107]. In discussing chemotherapeutics for image-guided drug delivery it becomes necessary to discuss what has been performed in the past. Chemotherapeutic selection for drug delivery systems has been based primarily on clinically approved drugs which have limitations due to solubility, dose limiting toxicity, instability in the bloodstream or poor efficacy. For FDA approval, selecting drugs that are already approved for use is one less hurdle to cross and is, therefore, beneficial in image-guided therapeutics.

One of the many challenges for chemotherapeutic nanoparticle delivery is the incorporation methods and subsequent release. Some carriers entrap the chemotherapeutic cargo like liposomes, microbubbles and potentially micelles. The important part of image-guided drug delivery is that the nanoparticle drug complex must stay intact until it arrives at the targeted tumor site. Otherwise, the imaging of

the carrier does not represent the actual drug localization. This is a challenge for nanoparticle formulations which have limited stability over time and “leak” drug. Covalent linkages to nanoparticles can overcome this but methods must be developed in order to realize site-specific drug release. Linear carriers such as HPMA copolymers have utilized lysosomally degradable GFLG peptide sequences for drug release once the nanoparticle has been endocytosed into the cell [108]. Success of site-specific drug release has been variable using this approach. In the case of HPMA copolymers conjugated with docetaxel, rapid release was observed in cell culture media and, therefore, expected to observe similar behavior in vivo [109]. In another study, release kinetics was variable when HPMA copolymers were conjugated with both gemcitabine and doxorubicin via GFLG linkers [110]. Gemcitabine was rapidly released in the presence of cathepsin-B, a lysosomal enzyme, but docetaxel showed a very slow release even with the enzyme. These examples demonstrate the balance that is needed with site-specific drug release. Drugs must be stable enough before getting to the targeted site but have the ability to empty its payload rapidly for a maximally effective image-guided delivery.

6.5.2 Radiotherapeutics

Radioisotopes have been utilized for cancer treatment for a considerable amount of time. Some of the first clinically used radioisotopes were produced in the 1930s such as ^{131}I iodine [111]. Radiotherapy is primarily performed using isotopes which emit alpha and beta radiation. Both forms of radiation are efficient at forming radicals (usually with water) which exert damage to the DNA of cells and cause irreversible damage leading to cell death. However, the range of tissue in which these effects occur is different depending on the type and energy of particle [47]. Many of the radioisotopes in the clinic for radiotherapy have beta emission with a range of 50–5000 μm . More rare isotopes with alpha emission have a much shorter range of 40–90 μm . Radioisotopes with auger electrons have the shortest range of 0.01–1.0 μm . The greater the range the more risk to nontargeted tissues near tumors. However, for larger tumors, a larger range may be more advantageous. Selecting the proper isotope is a matter of $1/2$ -life and the range of emission that is ideal for the targeted tumor. Table 6.4 is a list of radioisotopes for potential use in radiotherapy. The key aspect required for successful radioisotopes is site-specific localization and ample radiation dose to cause sufficient irreversible DNA damage leading to tumor cell death. This is a challenge that requires tumor-targeted approaches which do not cause off-site accumulation. One of the first clinically available theranostic delivery systems is based on a radiotherapeutic approach. Iodine uptake is especially high in the thyroid and for more than 50 years, radiation oncologists have exploited this feature to treat thyroid cancer with ^{131}I -iodine [112]. Imaging can be performed using SPECT or simple gamma scintigraphy of ^{123}I , a gamma emitter, which gives information on the areas of accumulating iodine in the patient. Safety and efficacy concerns can be utilized to select the patient for the subsequent ^{131}I therapy. Other examples

Table 6.4 Isotopes for theranostic radiotherapy [47, 113]

Radionuclides	T _{1/2}	Type of emission	Average energy (keV)
¹³¹ I	8.0 day	β ⁻	181
⁹⁰ Y	2.7 days	β ⁻	935
¹⁵³ Sm	1.94 days	β ⁻	280
⁶⁴ Cu	12.7 h	β ⁻	1670 (max)
⁶⁷ Cu	2.58 days	β ⁻	141
²¹¹ At	7.2 h	α	5867
²¹³ Bi	46 min	α	6000 (max)
⁶⁷ Ga	3.26 days	Auger	0.04–9.5
¹²⁵ I	60.5 days	Auger	27 (max)

were discussed early on in this chapter, specifically the radioimmunotherapies, Bexxar[®] and Zevalin[®] for lymphoma patients. Theranostic approaches have been extremely useful in the approval of these therapeutics in the clinic. Selecting the proper isotope is highly dependent on the ability to stably link the radionuclide to the nanoparticle platform. For efficacious and safe radiotherapy, loss of the radioisotope must not occur. Using nanoparticles to carry isotopes prevents them from accumulating in radiation sensitive areas such as bone marrow as long as the chelated radioisotope is stable in the bloodstream. The main advantage of using radioisotopes in nanoparticles is that their effect on cancer is independent of release, unlike chemotherapeutics. Nanoparticles in this case do not need to penetrate deep within the tumor to exert their anticancer effects. This advantage is profound because typically nanomedicines cannot penetrate deep within tumor tissues.

Several successful image guided radiotherapeutics have been developed and have shown promising results in treating cancer [113]. Wang et al. synthesized a multifunctional lipid-polymer hybrid nanoparticle that contained docetaxel and chelator for both ¹¹¹In and ⁹⁰Y radioisotopes [114]. The prostate tumor-targeting nanoparticle demonstrated superior antitumor efficacy of the combination chemotherapy and radiotherapy particle. Although the approach has promise, control over all components especially between loaded drug and ⁹⁰Y may prevent reproducible synthesis required for eventual translation.

6.5.3 Nucleic Acid Delivery

A promising area still in development for therapeutics is antisense oligonucleotides. This can include short strands of either DNA or RNA which are complementary to a chosen sequence. When introduced into the cell, they can knock-down expression of a disease-related gene. One such technology involves small interference RNA or siRNA which has the higher therapeutic index when compared to other gene modifying therapies [115]. Due to the siRNA charge, it cannot cross

biological membranes and, therefore, polymers have been utilized to shield the charge, protect the nucleic acid in the bloodstream and transport them into the cytosol of tumor cells. Optical imaging offers an interesting opportunity with FRET-like systems based on siRNA delivery. Cationic quantum dots were conjugated with poly(ethylene imine) (PEI) in order to complex with siRNA that had been labeled with a fluorochrome [116]. Once complexed, the fluorescence signal was quenched and upon siRNA release, an increase in fluorescence was observed. This system could potentially be utilized to observe release and essentially be correlated with efficacy within tumor cells. However, systems such as these are in early-development. For example, Medarova et al. demonstrated the utility of using MRI and optical imaging to observe the nanoparticle uptake and subsequent knockdown of GFP expressing tumors by complexed siRNA [117].

6.6 Methods of Release or Activation

The therapeutic component of a theranostic system may or may not require the need to be released from the carrier in order to exhibit its function in the target tissue. For example, doxorubicin requires uptake into the cell and subcellular localization to the nucleus in order to intercalate into the DNA to inhibit biosynthesis of the cell [118]. In this case the carrier requires some type of release mechanism that promotes delivery to the nucleus of the cell to be most effective. On the other hand, therapeutic radioisotopes do not necessarily require detachment from the carrier because the ionizing radiation that causes cell death can penetrate cellular and subcellular walls depending on the amount of energy in the emitted particle [119]. Therefore, careful consideration must be given in choosing a proper mechanism of release for a therapeutic in order to exert its potential action against the diseased tissue of interest depending on where the therapeutic must exert its action. This section will describe some of the technologies that have been developed for site specific release or activation of a therapeutic component on a nano-carrier.

6.6.1 *Photo-activation*

One exogenous source of energy that can be used to activate or release drug within a diseased tissue is the use of light. Some of the first examples of light triggered therapy used photosensitizers mainly for photodynamic therapy (PDT). A photosensitizer is a light sensitive molecule that when activated at the specific wavelength of light creates reactive oxygen species which can interact in a destructive manner within cells [120–122]. PDT has been used for treatment of skin disorders, cancer and other various diseases. One benefit of photosensitizers is that their fluorescence emission can be observed by optical imaging, therefore, they can both provide detection and therapy. The potential for PDT is attractive, however, limitations in tissue penetration of light

limit this therapy to superficial tumors or through endoscopically accessible organs such as the lung [123]. There is potential to use small light emitting optical fiber probes through guided needles into deeper tissues such as the pancreas but this remains challenging and invasive [124]. Multimodality imaging using PET or SPECT is also a possibility using nanoparticles conjugated with photosensitizers, thus, identifying accurately the location and depth of required light penetration for PDT [125]. However, PDT is generally limited to superficial and local treatment, thus, limited against metastatic cancer [126]. Despite this, PDT can be used for cancers which are not restricted by the limitations of this approach. Several examples of photosensitizers have been conjugated to nano-carriers for use in a theranostic approach [103, 127–132]. In a recent example, a multi-block based polymeric micelle containing folate for tumor targeting and a mitochondrial targeted porphyrin based photosensitizer was examined in-vitro and demonstrated enhanced uptake in HeLa cells and subcellular localization to the mitochondria [131]. Site specific toxicity relies on the lack of cytotoxicity of the construct until the appropriate wavelength of light interacts with the photosensitizer. The polymeric micelle mentioned above accomplished this goal of having cytotoxicity only after treatment with light.

6.6.2 *Thermal-activation*

Another potential exogenous source that can be used to release or activate a therapeutic drug delivery is the use of heat. Hyperthermia has been used for treatment of various diseases [133–135] and could potentially be used in conjunction with a nanotheranostic for targeted release or activation. However, it must be able to be administered only in the diseased tissue in order for it to be utilized for site specific delivery. One such method of localized delivery is the use of gold nanorods for plasmonic photothermal therapy (PPT) in cancer. PPT utilizes the surface plasmon resonance of gold nanorods to produce heat when activated at the appropriate laser light wavelength [136, 137]. Gold nanorods, depending on the size, can passively accumulate into tumors and, therefore, when treated with light in a controlled manner efficiently heat the surrounding tissue. However, because light is utilized, penetration for deep tissue administration remains a challenge. Another method that may be closer to clinical translation is high-intensity focused ultrasound (HIFU). HIFU utilizes ultrasonic waves to generate heat within biological tissues [138]. It can also be combined with MRI imaging with gadolinium thermal sensitive liposome conjugates for precise imaging of the nanoparticle localization within the tumor and subsequent drug release [139–141]. One example of thermal activation for a drug release involved the use of a temperature sensitive HEMA mono/dilactate grafted liposome which was tuned to the desired temperatures for release [142]. This was accomplished by varying the content of grafted copolymer and demonstrated efficient drug release in-vitro using HIFU. Hyperthermic delivery based on HIFU represents a promising future for targeted delivery of nanoparticles for therapeutics and imaging.

6.6.3 *Acoustic-activation*

Another method for site specific drug release that can be controlled locally at the disease site is the use of ultrasound with microbubbles. Microbubbles for therapeutic delivery are a special case of vesicular structure which encompasses gas and can be designed for image-guided therapeutic delivery [14]. Small gas bubbles, typically perfluorocarbons, are imaged via ultrasound based on their difference in echogenicity than liquid media [15]. In most cases, microbubbles are stabilized by a lipid bilayer which surrounds the gas bubble. The surrounding liposome can be loaded with drugs and targeting agents for tumor delivery. With increased sonic waves these liposomal bubbles can burst, thus, releasing their contents. This can be focused in the areas where the microbubbles are accumulating, including tumors. Imaging and drug release is, therefore, controlled by ultrasound, and, thus, is a promising theranostic. One such example is the stabilization of perfluorocarbon nanodroplets using block copolymers containing paclitaxel [143]. Due to the high ^{19}F -fluorine content, ^{19}F -MR spectroscopy in conjunction with ultrasound was utilized to determine precise anatomical location of the nanoconstructs. Complete tumor regression was observed in a pilot study with tumor focusing ultrasound in tumor bearing mice. The promise of ultrasound-mediated delivery using microbubbles is being investigated in several clinical trials [13]. However, challenges remain in reproducibly manufacturing these constructs, especially when complicated by associating targeting agents and drugs into their structure. Their size distribution also remains a challenge. Microbubbles span a large range in sizes affecting their distribution in the body. Strategies are needed to refine microbubble manufacturing process for more reproducible formulations, smaller size and less polydispersity. Stabilization of the liposomal structure is also important to ensure that drug leakage does not occur. Polymersomes made of block copolymers can be designed to encapsulate drug in the hydrophobic shell and form a stable liposomal structure for gas for detection with ultrasound. Specific delivery to the site can be accomplished by rupturing the shell structure with cavitation forces using the ultrasound device in a similar fashion as lipid counterpart. Polymersomes have been developed for this purpose, and the surface tuned to have functional groups which can be utilized to attach targeting moieties [144], thus, enhancing site specific delivery. The combined approach of targeting and site specific delivery is a promising direction in tumor treatment.

6.7 Challenges for Development of Image Based Theranostics

The previous sections have presented the different aspects, advantages, disadvantages and requirements for image-based theranostics. In order to overcome some of the challenges and limitations of certain imaging modalities, researchers have designed materials that can be visualized by more than one way. For example, Chakravarty et al. developed ^{69}Ge -superparamagnetic iron oxide nanoparticles for both MRI and

PET imaging [145]. The advantage of this system is increased sensitivity provided by the PET isotope component but high resolution of the MRI iron oxide contrast component for accurate localization within the targeted area. One must also consider for clinical translation the regulatory hurdles that need to be overcome for manufacturing of nanomaterial delivery systems with imaging agent, therapeutic and targeting.

Perhaps one of the more challenging hurdles to overcome is the fact that no true nanotheranostic system has been successfully tested in the clinic with the exception of monoclonal antibodies. This may be due to the fact that many questions about safety still need to be answered regarding the long-term effects of nanoparticles within the body. Many of the nanoparticles with the intent of being used as a theranostics agent have been developed. The literature is full of nanoparticle systems designed with both therapeutic and diagnostic components which have been well characterized and evaluated in-vitro. However, what is missing is well-designed studies using appropriate animal models that truly evaluate how advantageous these systems could be in the clinic. More studies are needed that demonstrate how nanotheranostics actually can predict efficacy or correlate imaging results with actual efficacy. Several researchers have designed and synthesized nanoparticle conjugates with evidence of efficacy but study designs often lack the main goal which is to determine whether imaging of the nanoparticle localization to the target does or does not correlate with efficacy. Steps toward translation will accelerate, if researchers utilize appropriate animal models that can mimic various stages of disease and utilize the imaging information from the nanotheranostic localization in the subject to correlate with the corresponding therapeutic.

Generally, the simpler the design is, the better the approach. Good manufacturing practices (GMP) required for FDA approval require that each component of the formulation have complete characterization at the full range of possibilities within the construct or formulation. With multiple components having multiple cross-interactions possible, the regulatory hurdle for just one image-based theranostic is significant. Much is still needed to address how each component of these nanotheranostics will be deposited in the human body and what long-term effects might occur. Thus, a simpler approach is more likely going to be logistically and economically feasible. Selecting components especially drugs with ample clinical experience will also lessen the burden because previous experience can yield much information in regards to potential problems. The FDA has recently begun to address these challenges by characterizing these materials using criteria provided by the National Nanotechnology Initiative [146]. The Center for Drug Evaluation and Research within the FDA has recognized that this technology will need specific guidance in regards to their eventual approval but have also acknowledged that they will be treated as any other diagnostic or therapeutic.

6.8 Conclusion

In summary nanotheranostics are in their earliest stages of development. With the exception of MoAbs, no nanotheranostic system has been approved for use in the treatment of diseases. As the industry evolves it is clear that imaging combined with

therapeutic development with nano-sized carriers will be promising in personalized medicine. Questions, however, will still need to be answered regarding their overall safety. Many advances with nanomaterial design and imaging technologies have taken the field closer to realization. However, much is still to be done to make these systems widely applicable in the field of medicine.

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Chapter 7

Targeting Cancer Using Nanocarriers

Dalit Landesman-Milo, Shahd Qassem, and Dan Peer

Abstract Nanotechnology is an emerging multidisciplinary field that offers unprecedented access to living cells of target (i.e. cancer cells) and promises the state of the art in cancer detection and treatment. Development of nanocarriers that target cancer for diagnostics and therapy draws upon principles in the field of chemistry, medicine, physics, biology, and engineering. Given the zealous activity in the field as demonstrated by over 7000 published journal articles on the topic and given the promise of recent clinical results, nanocarrier-based approaches are anticipated to soon have a profound impact on cancer medicine and as a consequence on human health. The versatility in size, material, and targeting agents of nanocarriers permits potential targeting for individual cancer cells. This chapter addresses nanocarriers spanning liposomes to polymeric nanoparticles, inorganic nanoparticles, polymers conjugates and dendrimers. The targeting approaches include conjugation of molecules such as receptor-specific ligands, antibodies and aptamers to the surface of the carrier. Targeting cancer with nanocarriers represent the next important milestone that is already impacts the lives of millions around the world. One such example, is of DaunoXome (Liposomal Daunorubicin) for Acute myeloid Leukemia (AML) treatment that shows an increased intratumor and intracellular levels of the drug, while normal tissue toxicity, including cardiotoxicity, may be reduced.

Keywords RNAi • Nanoparticles • Targeting agents • mAbs

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7.1 Introduction

Current therapy for cancer involves the cyclical administration of one or more chemotherapeutic drugs, with or without, surgical intervention and radiation. The effectiveness of the treatment is directly related to the ability to selectively kill the cancer cells without damaging healthy cells. By administering bolus doses of cytotoxic drugs, adverse effects are commonly observed and patients may only experience a marginal improvement on long-term survival. Therefore, more effective cancer therapies are needed both in terms of new classes of drugs and in terms of developing “smart delivery vehicles” to minimize the adverse effects and to enhance the survival of patients with metastatic cancer.

One approach to target the delivery of drugs specifically to cancer cells is through directing cellular events at the nanometer scale [1–6]. Where current technologies, mainly based on anatomical and physiological alteration require hundreds of thousands of cells to detect the presence of a tumor, while nanotechnology approaches utilize metabolic and molecular unique features of cancer cells (e.g. Glucose fluorescent conjugates) that could radically lower this requirement, enabling much earlier diagnosis/treatment regimens. Through working on the nano scale it becomes possible to differentiate between healthy and cancerous cells, thus, significantly offering a wide therapeutic index with reduced side effects. One major clinical advantage of nanocarrier-based strategies over drugs that are directly linked to a targeting moiety is the specific delivery of large amounts of chemotherapeutic agents per recognition event.

Typically, nano-carrier based approaches include a carrier, a targeting moiety that is bound to the carrier via specific conjugation chemistry, and a drug (Fig. 7.1). Carriers may be composed of liposomes, polymeric nanoparticles, inorganic nanoparticles, polymers or dendrimers. Targeting moieties may include high affinity ligands, peptides, antibodies, and nucleic acids, and they may be conjugated to the carriers utilizing a variety of chemistries. Given the wide array of potential nanocarrier-based strategies for targeting cancer, this review will focus on drug delivery strategies for supplying a lethal chemotherapeutic dose to target cells and nucleic acid drug delivery system with a focus on siRNA delivery platforms.

7.2 Targeting

There are generally two approaches for nanocarrier-mediated systemic drug targeting from the site of administration (i.e. injection):

1. Passive targeting to tumor tissues—via enhanced permeability of tumor vasculature.
2. Active targeting at the cellular level—to cancer cells without affecting normal cells.

Passive targeting to the tumor tissues relies on ‘leaky’ microvascular or the enhanced permeation and retention (EPR) effect [7, 8], which permits selective

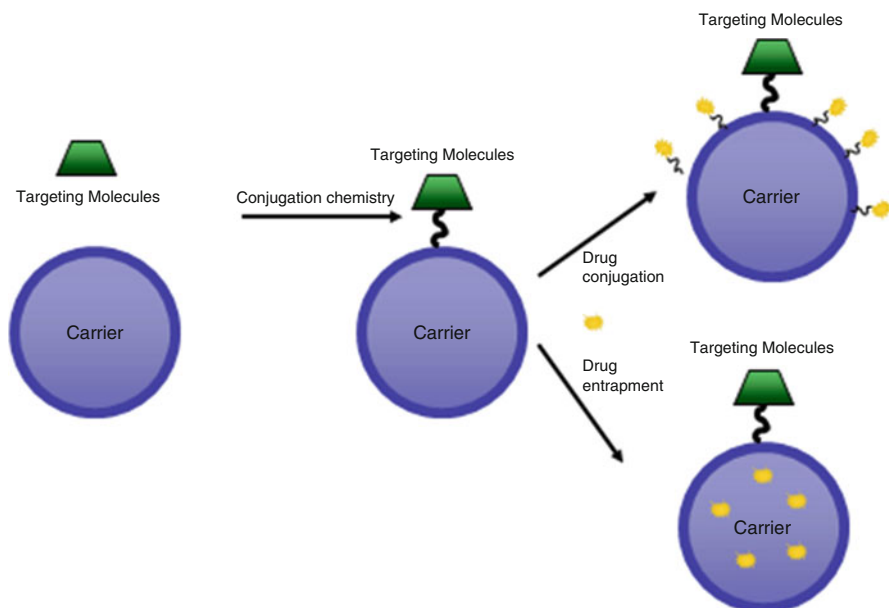


Fig. 7.1 Nanocarrier components for targeting cancer. Construction of the vehicle consists of a nano-scale carrier, a targeting moiety conjugate to the nanocarrier and the cargo (the desired drug)

permeation of nanoparticles into the desired tumor tissue. Active targeting to cancer cells is possible through promoting specific interactions with targeting ligands and surface receptors. An engineered nanoparticle should be able to overcome the physiological challenges of systemic circulation including tumor vascular biology and organs of mononuclear phagocytic system (MPS) whose function is to remove foreign material from the circulation [9].

Once at the target site, nanoparticles may release the drug in close proximity to cancer cells (though passive targeting), attached to the cancer cells, or within the cell following internalization (Fig. 7.2). Cell surface targets (often receptors), may recognize a variety of targeting agents such as ligands, antibodies, and nucleic acids.

7.3 Ligands

Targeting agents that may be broadly classified as ligands include proteins, peptides, vitamins and carbohydrates. One of the commonly used ligands for cancer targeting is the high affinity vitamin folic acid (folate) since folate receptors (FRs) are frequently overexpressed in a range of tumor cells [10, 11]. Folate is a small molecule (MW 441.40 Da) that strongly binds to FRs. Folate is water soluble and can be easily conjugated using its gamma-carboxyl group and, thus, it has been used as a targeting moiety in a wide array of drug delivery vehicles including liposomes, polymeric nanoparticles, linear polymers, and dendrimers to deliver drugs into cancer cells

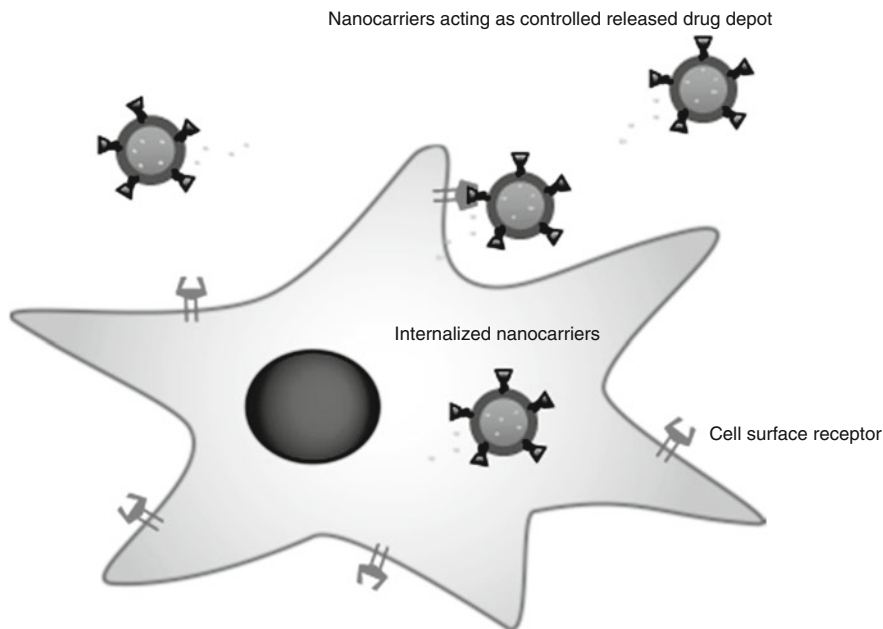


Fig. 7.2 Targeted nanocarriers-cell interactions. The Nanocarrier can release its content in close proximity of the target cells (passive targeting); attach to the membrane of the cell and act as an extracellular sustained release drug depot; or internalize into the cells

using FR-mediated endocytosis [11–16]. Another ligand with selectivity for cancer cells is transferrin (Tf) protein [17]. Tf receptors (TfR) are overexpressed on many tumor cells and coupling Tf directly to the liposomes, and other nanoparticles entrapping chemotherapies has resulted in improved intracellular delivery and therapeutic outcomes in cancer models [18–22]. Recently, Tf has also been used to facilitate siRNA delivery through transferrin receptors allowing a specific killing of tumor cells [23] and for effective brain tumors targeting [24]. In addition to cell surface antigens, extracellular matrices (ECM) such as heparin sulfate, chondroitin sulfate, and hyaluronan (also called Hyaluronic Acid, or HA), are usually responsible for the tumor's cells communication and loss of homeostatic control, and, thus, play an important role in the targeted drug delivery niche. HA is a naturally-occurring glycosaminoglycan that binds to CD44 and CD168 receptors. Both receptors are highly expressed on the surface of cancer cells. The potential of using ECM as targeting agents was demonstrated through intravenous injections of Paclitaxel loaded HA nanoparticles into tumor bearing mice, which reduced tumor size [25–28].

This effect is likely a result of competition of HA with its receptors on tumor cells. Although HA is ubiquitously expressed in normal tissues, normal cells bind it with low affinity, and seem to require activation to bind HA more efficiently [28–32]. This may help limit non-specific interactions thereby increasing the effectiveness of

approaches that aim to utilize ECM components as targeting agents. Over the last 10 years, Chitin and Chitosan, amino sugar N-acetyl-glucosamines and other sugar residue (e.g. mannose), biocompatible, biodegradable, low cost polysaccharides and their derivatives have drawn attention as vehicles and targeting moieties for drug delivery [33]. Kawakami et al. have previously demonstrated an efficient mannose receptor-mediated gene transfer into macrophages [34]. In addition to proteins, peptides such as Arginine–Glycine–Aspartic acid (RGD) that target integrin $\alpha_v\beta_3$ on tumor cells' surface show an increased efficacy against different murine tumor models [35, 36]. Although RGD based peptide approaches are commonly employed, this peptide also binds to other integrins such as $\alpha_5\beta_1$ and $\alpha_4\beta_1$ and, thus, is not highly specific to cancer cells only. Alternatively, vasoactive intestinal peptide (VIP), which only binds to VIP receptors has been used to deliver paclitaxel to VIP receptors on the surface of breast cancer cells that resulted in inhibition of tumors in vivo [37–39].

Another example for targeting agent is TH10 peptide, which was isolated from a phage display library that targets neural/glial antigen 2 (NG2), a proteoglycan highly expressed in tumor-derived vascular pericytes lining the inside of blood vessels [40].

Interfering with pro-cancer and metastasis cellular triggering activities such as cell proliferation, chemotaxis and gene transcription by blocking a critical chemotactic axis interaction in tumor/metastasis progression is another targeting option. Blocking the chemokine receptor 4 (CXCR4) and its ligand (CXCL12) CXCR4/CXCL12 is an example for this attitude [41].

7.4 Antibodies

One of the earliest cancer targeting is using a monoclonal antibody which was first described in 1981 by the Milstein group [42]. Over the past three decades, the feasibility of antibody based tissue targeting has been clinically demonstrated (reviewed in [43, 44]). Monoclonal antibodies have been used as biological therapeutics and as targeters. The first FDA approval for therapeutic monoclonal antibodies for the treatment of cancer came in 1997 where rituximab (Rituxan™) was approved for treating patients with a form of non-Hodgkin's lymphoma [45]. A year later, in 1998 Trastuzumab (Herceptin™), a humanized anti-HER2 (against EGFR) monoclonal antibody, has been approved for the treatment of breast cancer [46]. Over the past couple of decades more than two dozen of different monoclonal antibodies have already approved by the FDA to treat certain cancers [47]. Among them, the first angiogenesis inhibitor to treat colorectal cancer, Bevacizumab (Avastin™) an anti-VEGF mAb, was approved in 2004 [48–50]. Emerging approaches include delivery of a chemoimmunoconjugate containing anti-CD33 antibody and calicheamicin which is in clinical Phase III development for the treatment of acute myelogenous leukemia [51]. Recently, a targeting angiogenic agent was approved by the FDA—ramucirumab (Cyramza, Eli Lilly and Company), a mAb that binds to vascular endothelial growth factor receptor-2 for the treatment of advanced gastric cancer [52].

Despite the recent success of monoclonal antibodies as targeting moieties, this class of molecule continues to exhibit important suboptimal properties. Foremost, the biological production of monoclonal antibodies can be difficult and unpredictable. In addition, the performance of antibodies can vary from batch to batch, in particular when production is scaled up. However, despite these disadvantages, antibodies represent the most common and potentially effective strategy for targeting cancer. A relatively new targeting technology that addresses many of the shortcomings of antibody based approaches involves use of high affinity oligonucleotides called aptamers.

7.5 Aptamers

Aptamers are short single stranded DNA or RNA oligonucleotides that have been selected *in vitro* from a large number of random sequences ($\sim 10^{14}$ – 10^{15}) and have a molecular weight (10–15 kDa), which is one order of magnitude lower than that of antibodies (~ 150 kDa) [53]. Aptamers are derived from an *in vitro* iterative protocol called systematic evolution of ligands by exponential enrichment (SELEX) (Fig. 7.3) that was first described independently by Tuerk and Gold [54] and by Ellington and Szostak [55] in 1990.

Since their original description, aptamers have been isolated to a wide variety of targets, including intracellular proteins, transmembrane proteins, soluble proteins,

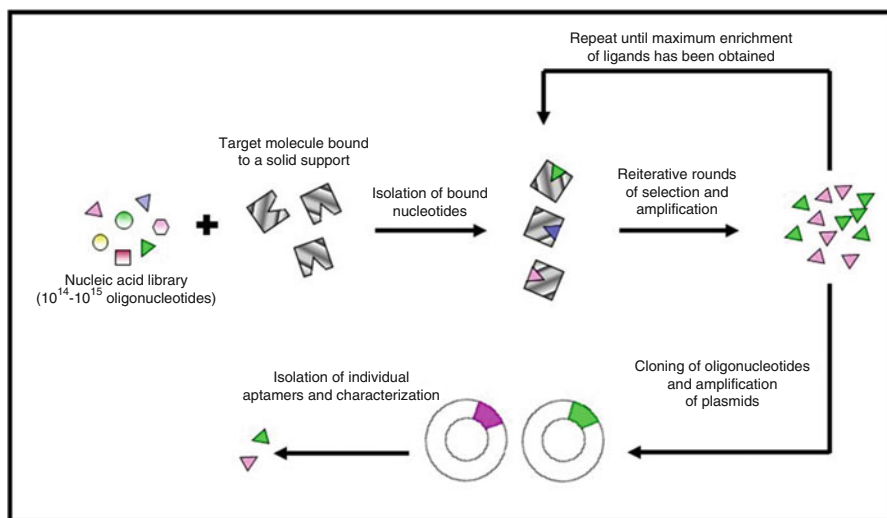


Fig. 7.3 Outline of the SELEX process. Target molecules are initially incubated with a combinatorial library of nucleic acids. Bound nucleotides are isolated followed by reiterative rounds of selection and amplification. This process is repeated with increased stringency with each round. Next, oligonucleotides are cloned and plasmids are amplified. Finally, through this enrichment process, a small number of tightly binding aptamers are obtained

carbohydrates, and small molecule drugs. Unlike antibodies, aptamers can be chemically synthesized at microgram or kilogram scales, and due to their small size and similarity to endogenous molecules, are believed to be non-immunogenic. It is possible to incorporate selection by intact biological entities such as cancer cells or tissues to identify an array of highly specific aptamers [56, 57]. Unlike antisense small interfering RNA compounds, which are single-stranded nucleic acids that affect the synthesis of a targeted protein by hybridizing to the mRNAs that encode it, aptamers may inhibit a protein's function through directly binding to it.

Due to their composition (nucleotides) and relative small size, aptamers may be degraded by nucleases or rapidly cleared from the blood. Unlike antibodies, aptamers require sheltering from degradation, which can be achieved through a variety of techniques [58–61]. A number of aptamers have been developed to bind specifically to receptors on cancer cells. For example, as guided delivery of nanoparticles for treatment of over expressing EGF receptor tumor cells [62] or as a selective delivery agent of chemotherapy based liposomes to prostate cancer cells [3, 63].

To date, there is one FDA approved aptamer based therapeutics called Macugen (Pegatanib) in treatment for age-related macular degeneration (AMD) and many more aptamers under clinical trials, one antitumor DNA based aptamer targeting nucleolin which is drawing attention according to its clinical trials results is the AS1411 [64].

7.6 Incorporation of Drugs into Nanocarriers

Traditional drug administration such as intravenous injection or multiple daily oral dosing has proven to be suboptimal or unsuitable for a variety of medical conditions. Aside from requiring high doses to reach target tissues, these types of therapeutics typically produce peaks in drug concentration above the therapeutic range followed by a rapid decrease in concentration to a level below the therapeutic range. During the past five decades delivery strategies have evolved and methods are now available for achieving a constant or pulsatile release, within the therapeutic range, over long periods of time [65–69].

Chemotherapeutic anti-cancer drugs and their carrier may be joined via two major techniques. One way is to entrap drug molecules into delivery vehicles for controlled release of the drugs. Controlled release occurs when a natural or synthetic polymer is combined with a drug in such a way that the drug is encapsulated within the polymer system for subsequent release in a predetermined manner. Drug delivery vehicles that are designed as particles can range in size from <10 nm (dendrimers) and 100 nm (nanoparticles) to over 10 μm (microparticles) and can release the encapsulated drugs via surface or bulk erosion, diffusion, swelling followed by diffusion, or response to a stimulus (e.g. pH, temperature, ultrasound, electrical stimuli). The release of the active agent may be constant or cyclic over a long period, or may be triggered by the environment or other external events [70]. In general, controlled-release polymer systems can provide drug levels in the

optimum range for an extended period of time than other drug delivery methods, thus, increasing the efficacy of the drug and maximizing patient compliance. The primary consideration of drug delivery is to achieve more effective therapies while eliminating the potential for both under- and overdosing. Other advantages of using controlled release delivery systems can include the maintenance of drug levels within a desired range, the need for fewer administrations, higher diffusion rates of the drug into the tumor due to a continuous-controlled release of it in the tumor proximity and optimal use of the drug in question. While these advantages can be significant, the potential toxicity of the materials used in these delivery systems may limit their translation into clinical practice in some cases [71].

Another approach is to conjugate drugs covalently to the bulk of drug carriers. carrier-drug conjugation usually inhibits metabolic activity until the conjugation bond is cleaved, resulting in activation of the drug at the target site [72]. The covalent conjugation between carriers and drugs has a number of advantages including: (1) Limiting the mechanism of drug release from the carriers and (2) reducing potential burst of drug as it is typically seen in entrapment approaches. A further detailed review on the polymer-drug conjugates and examples thereof can be found elsewhere [73].

7.7 Choosing a Carrier

In choosing an appropriate nanocarrier construct, one should consider the following criteria:

1. A carrier should be made from a material that is biocompatible, easily functionalizable, and well characterized.
2. A nanocarrier with targeting molecules should exhibit high uptake efficiency by the target cells unless they are required to accumulate on the cell surface and act as site specific drug depots.
3. The nanocarriers should be either soluble or colloidal in aqueous phase and exhibit an extended circulating half-life to increase the likelihood of their effectiveness.
4. The nanocarriers should have a low rate of aggregation and preferably a long shelf life.

A variety of potential nanocarrier compositions are available including liposomes, polymeric nanoparticles, inorganic nanoparticles, and dendrimers and these are reviewed below (Fig. 7.4).

7.8 Liposomes

Liposomes are the first nanocarriers described more than 50 years ago by Bangham [74–76], and the first that have been clinically approved by the FDA to carry chemotherapy (DaunoXome™ (50–80 nm) in 1996 [77], Doxil™ (100 nm) in 1997,

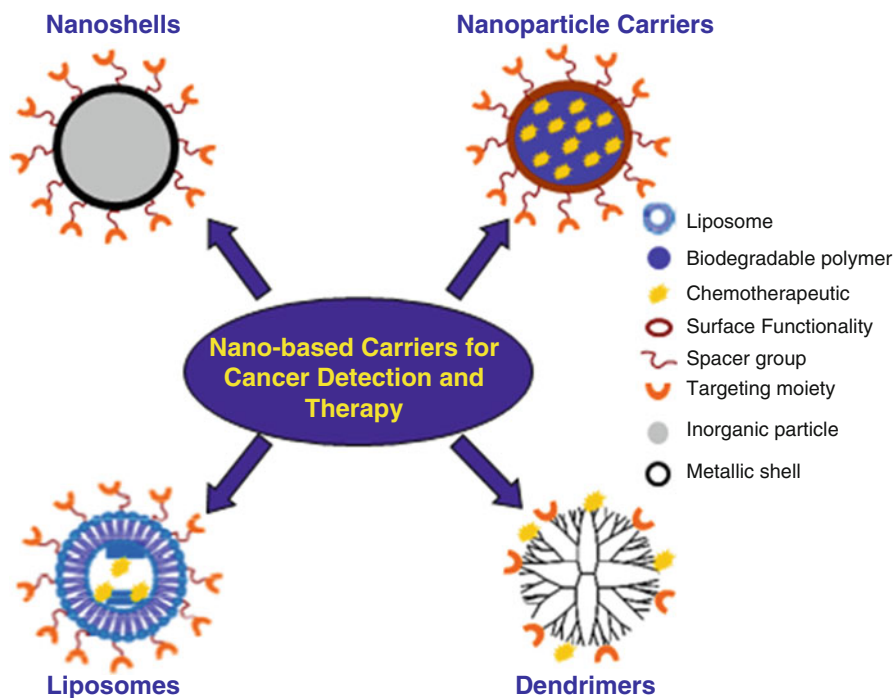


Fig. 7.4 The arsenal of nano-based carriers of targeting cancer. Liposomes (50–800 nm), polymeric nanoparticles (5–150 nm), dendrimers (2–10 nm) and nanoshells (5–100 nm)

Myocet™ (190 nm) in 1997 [78] [79–81]. Liposomes are spherical, self-closed structures formed by one or several concentric lipid bilayers with an aqueous phase inside and between the lipid bilayers. The most common way to prepare liposomes is the lipid film method [82] where the lipids are dissolved in organic solvents and evaporated to dryness under reduced pressure. Then they are hydrated by a swelling solution with the desired drug to form a multilamellar vesicle (MLV). An extruder is used to downsize the MLV to a nano-scale, unilamellar vesicle (ULV) or to small ULV (SUV) having progressively smaller pore-size membranes, with several cycles per pore-size and the result is a uniform nano-scale population of liposomes with a narrow size distribution. There are today more than 20 different liposomal formulations in different stages of clinical trials [83] (Table 7.1). Among the active chemotherapeutic drugs entrapped in the liposomes are: daunorubicin, doxorubicin, cytarabine, vincristine, lurtotecan, platinum based compounds and more [84, 85]. Liposomes have many attractive biological properties including: biocompatibility, biodegradability, the ability to entrap both hydrophilic drugs (through their interwater compartment) and hydrophobic drugs (through the membrane), and to protect the drugs from the biological environment. Their size, charge and surface could be easily modified by adding more agents into the lipid membrane or by surface chemistry modifications.

To achieve improved selectivity, a targeting moiety is attached to the liposomal surface via a PEG spacer arm to reduce steric hindrance of binding to the target [86].

Table 7.1 Representative FDA approved nanomedicines or in clinical evaluation

Compound	Commercial name TM	Indications	Status
<i>Liposomes</i>			
Daunorubicin	DaunoXome	Kaposi's sarcoma	Market
Doxorubicin (PEG-liposomes)	Doxil/Caelyx	Refractory Kaposi's sarcoma; recurrent breast cancer; ovarian cancer	Market
Doxorubicin (non PEGylated liposomes)	Myocet	Combinational therapy of recurrent breast cancer	Market
liposomal vincristine sulphate	Marqibo	lymphoblastic leukemia	Market
Albumin NPs encapsulate paclitaxel	Abraxane [®]	Breast cancer NSCLC	Market
Liposome encapsulated with semi-synthetic doxorubicin analogue	Annamycin	Acute lymphoblastic leukemia (ALL) and acute myelogenous leukemia (AML)	Phase I/II
Doxorubicin (thermally sensitive liposomes)	ThermoDox [®]	Primary Liver cancer RCW breast cancer	Phase III Phase II
Pegylated liposomes encapsulated Irinotecan	Nektar-102 IHL-305	Breast, colorectal cancer Advanced solid tumors	Phase III Phase I
Nanoliposomal irinotecan	PEP02 (MM-398)	Gastric and pancreatic cancer	Phase II
Liposome entrapped irinotecan and HCl:floxuridine	CPX-1	Colorectal neoplasms	Phase II
Liposome enrapped paclitaxel	LEP-ETU	Ovarian, breast and lung cancers	Phase I
Liposome entrapped with cisplatin	Aroplatin	Colorectal cancer	Phase I/II
Vinorelbine liposomes	Vinorelbine	Advanced solid tumors Hodgkins disease Non-Hodgkins lymphoma	Phase I
Liposome entapped with cisplatin	Lipoplatin	Pacreatic, head and neck, breast cancer	Phase III
Stealth liposomal cisplatin	STEALTH [®]	Head and neck, lung cancers	Phase III
Targeted liposomal cisplatin formulation (based on tumor triggered release mechanism)	LiPlaCis	Advanced or refractory solid tumors	Phase I
Vincristine	Onco TCS	Relapsed aggressive non-Hodgkin's lymphoma (NHL)	Market
Liposome entrapped cytarabine and daunorubicin	CPX-351	AML	Phase II
Mitoxantrone		Prostate cancer	Phase I/II
Liposomal SN-38 (DNA topoisomerase I inhibitor)	NK102	Colorectal cancer	Phase II
Liposomal Lurtotecan (Liposome entapped topoisomerase I inhibitor)	OSI-211	Ovarian cancer	Phase II/ III

(continued)

Table 7.1 (continued)

Compound	Commercial name TM	Indications	Status
Cytarabine liposome	DepoCyt	Esophageal cancer Lymphomatous meningitis	Market
Plasmid DNA encoding HLA-B7	Allovectin-7 [®]	Metastatic melanoma	Phase III
<i>Polymeric micelles</i>			
Cisplatin-incorporating polymeric micelles	NC-6004	Pancreatic cancer NSCLC	Phase III Phase I/II
Paclitaxel encapsulated micelle	NK105	Breast cancer	Phase III
Liposome encapsulated with (S)-camptothecin (aerosolized)	9-Nitrocamptothecin liposome-L9NC	Lung, metastatic endometrial cancer	Phase II
<i>Peptides/aptamers</i>			
Peptide conjugate based on PLGA-LHRH	Lupron [®] (leuprolide)	Prostate cancer	Market
LHRH synthetic decapeptide	Goserelin (Zoladex)	Breast cancer	Market
Peg-L-Asparaginase	Oncaspar	Acute lymphoblastic leukemia	Market
Pegylated recombinant human granulocyte colony-stimulating factor (rhG-CSF)	HHPG-19K	Breast cancer	Phase II
Paclitaxel poly-L-glutamic acid conjugate (PG-TXL)	CT-2103 (XYOTAX)	Metastatic breast cancer Ovarian cancer	Phase I Phase III
Doxorubicin-N-(2-hydroxypropyl) methacrylamide HPMA copolymer	HPMA copolymer	Breast cancer	Phase II

Some examples of selective targeting include use of anti-CD19 long-circulating liposomes that deliver doxorubicin to B cell lymphoma [87] and anti-Her2-liposomes that showed effective targeting to breast cancer cells overexpressing erbB2 receptor [88, 89]. Folate-targeted-liposomes have been proposed also as vehicles for chemotherapy to improve therapeutic activity [90].

Transferrin (Tf)-mediated liposome delivery is also an emerging strategy to target tumors overexpressing the transferring receptors. Some examples include Tf-coated liposomes entrapping doxorubicin and paclitaxel targeted to U-87brain glioma cells [24, 91] and to solid tumors [92].

Other approaches include the combination of fusogenic peptides, targeting moieties and pH sensitive lipids that will cause the liposome disruption in the endosome [93–98].

One of the potential challenges with drug containing liposomes is the inability to effectively reconstitute them after lyophilization and to maintain their dimensions. Successful attempts to address these challenges involve covalent immobilization of high molecular weight HA on unilamellar liposomes. This serves to cryoprotect against lyophilization [32, 99]. In addition, the HA coatings improves circulation time

and enhances targeting to tumors expressing HA receptors (CD44 and RHAMM) [4, 100, 101]. We developed antibody coated HA-liposomes (Integrin beta 7 targeted) that can be lyophilized and stored as a dry powder which preserves the nano dimensions and the binding capacity of the immobilized mAb on its surface [102].

Chitosan based NPs hold several intrinsic advantages of biocompatibility, biodegradability and a prepared infrastructure for chemical reaction, made them a suitable vehicle for encapsulating a variety of anti cancer drugs and surface readily modified for chemical cross link conjugation of different targeted moieties [103], such as folic acid [104], poly lactic-co-glycolic acid (PLGA) [105], cell-targeting peptides (for example CP15) [106] or provascular agent bradykinin potentiating peptide (BPP) [107]. The main advantage of these targeted NPs is longer circulation time, that enables significantly prolonged survival rates of tumor-bearing mice.

7.9 Polymeric Nanoparticles

One of the earliest reports of using nanoparticles for drug delivery was introduced in 1978 [108]. Since then, polymeric nanoparticles for clinical applications have been attempted using a wide variety of biodegradable polymers including: poly (lactic acid) (PLA) [109], poly (glycolic acid) (PGA), poly (lactic co-glycolic acid) (PLGA) [110], poly (orthoesters) [111], poly(caprolactone) [112], poly(butyl cyanoacrylate) [113], polyanhydrides [114], and poly-N-isopropylacrylamide [115]. Although many fabrication methods exist, polymeric nanoparticles are frequently made using an oil-in-water emulsion [116] which involves dissolving a polymer and drug in an organic solvent. Next, the organic phase is vigorously mixed with an aqueous phase, followed by removal of the organic solvent by evaporation which forces the polymer to precipitate as nanoparticles. The particles are then recovered by centrifugation and lyophilization. To prevent the aggregation of the nanoparticles after they are reconstituted, nanoparticles can be made from amphiphilic copolymers composed of two biocompatible blocks [65]. For example, copolymers containing both lipophilic (i.e. PLGA) and hydrophilic (i.e. PEG) polymers have been constructed where the PEG migrates to the surface of the nanoparticles which comes in contact with aqueous solution.

Although polymeric nanoparticles offer great potential for targeted drug delivery approaches, one key limitation is their polydispersity with respect to size which may increase non-specific uptake. Another class of nanoparticles made from inorganic materials represents an emerging class of nanocarriers that may be prepared with near monodispersity.

7.10 Inorganic Nanoparticles

One of the first uses of inorganic nanoparticles for cancer therapy was reported by Roy et al. in 2003 [117]. Aside from potential monodispersity, a unique and highly advantageous feature of inorganic nanoparticles is their ability to be non-invasively

imaged by magnetic resonance imaging (MRI) and high resolution superconducting quantum interference device (SQUID) [118]. Additionally, Inorganic particles offer advantages including ease of size control, potential functionalization to introduce targeting molecules and drugs, and the ability to induce cell apoptosis via hyperthermia. Examples of cancer cells' targeting by inorganic nanoparticles include galactose-PEG modified gold nanoparticles [119] and superparamagnetic iron oxide nanoparticles conjugated with methotrexate (as a targeted drug) [120].

To demonstrate the potential of the dual imaging/therapeutic feature, immunotargeted nanoshells were used to detect and destroy breast carcinoma cells that over-express HER2, a clinically relevant cancer biomarker [121]. 'Nanoshells' are composed of a core of silica and a metallic outer layer. Nanoshells have optical resonances that can be adjusted to absorb or scatter essentially anywhere in the electromagnetic spectrum, including the near infrared region (820 nm, 4 W/cm²) where transmission of light through tissue is optimal. Absorbing nanoshells are suitable for hyperthermia-based therapeutics, while scattering nanoshells are desired for imaging applications. In addition to utilizing a passive approach where Nanoshells preferentially accumulate in tumors due to EPR, gold nanoshells can be readily conjugated to antibodies, peptides or aptamers for targeting tumor cells. Recently, a cancer therapy based on absorption of near infrared light by gold nanoshells leading to rapid localized heating was developed [121, 122]. Tissues heated above the thermal damage threshold displayed coagulation, cell shrinkage, and loss of nuclear staining, which are indicators of irreversible thermal damage, whereas control tissues appeared undamaged [122]. Specifically, exposure to near infrared light caused the tumors to increase temperature by approximately 40 °C, while controls without nanoshells was heated by less than 10 °C.

One of the challenges of those inorganic nanoparticles is the inability to achieve high drug loading. Other than utilizing thiol groups, the conjugation chemistries are limited.

7.11 Dendrimers

Dendrimers are synthetic monodispersed branched macromolecules that form a tree-like structure and their synthesis represents a relatively new field of polymer chemistry. Particularly, polyamidoamine (PAMAM) dendrimers, since first synthesized by Tomalia and co-workers using the divergent synthetic route [123], have shown great promise to be used in biomedical applications due to: (1) ease of functionalization (conjugation with biomolecules such as targeting molecules, imaging agents, and drugs), (2) high water solubility, (3) well-defined chemical structure, (4) biocompatibility, (5) low production of toxic metabolites due to the dominance of renal filtration as the main route of clearance, and (6) the rate of clearance may be controlled by chemical modification [124]. Although cationic PAMAM dendrimers are biologically active and, thus, have been used as a non-viral gene delivery vectors [125], they lack specificity leading to substantial non-specific interactions [126]. Consequently, charge neutral dendrimers have been prepared by acetylation using

acetic anhydride and play as a platform to which the multiplicity of their functional end groups can be covalently attached. Although dendrimers have not yet progressed to clinical trials, they show great promise for medical applications based on the ability to potentially synthesize high densities of accessible functional groups in a well-defined and standardized manner and based on their non-immunogenic and minimally toxic properties [127]. A successful example of in vivo targeted delivery approach using dendrimers was demonstrated in 2005 where the Baker group used methotrexate conjugated to PAMAM dendrimer which led to a tenfold reduction in tumor size compared to the same molar concentration of free methotrexate [6]. PAMAM dendrimers have also been complexed with MRI detectable metal nanoparticles and in vivo biodistribution of the dendrimer nanocomposite was studied [128]. Other potential dendrimers for targeting cancer include polypeptide and polyester dendrimers [129]. Other examples of dendrimers are summarized in literature [130, 131]. Although promising, there are still many obstacle currently preventing progression to clinical trials and markets. Dendrimers are relatively expensive as compared to other nanoparticles and require many repetitive steps to be synthesized, posing a challenge for large-scale synthesis.

7.12 Polymer Conjugates

A versatile group composed from both natural and synthetic polymer conjugate molecules that can carry and efficiently deliver drugs via covalent or non-covalent carrier-drug bonds. The most important features of this group that make it a suitable drug delivery agent, based on its triangular features of safety, quality and efficacy.

FDA approved delivery formulation is a conjugate of neocarzinostatin and poly(styrene-comaleic acid), SMANCS, that have shown a potential for a reduction of toxicity and immunogenicity in addition to an extended half-time [132].

Peptide conjugate formulation based on pegylation using covalent or non-covalent attachment of PEG are very common drug delivery approach for many anti cancer drugs. For example, *Oncaspar*, a pegylated version of asparaginase, a bacterial enzyme, use to treat Leukemia [133]. Neulasta, another pegylated based formulation of human G-CSF, use to treat cancer patients in order to reduce chemotherapy side effects [134].

PEGylation coupling polymer strategy has been proven as a successful and powerful delivery carrier especially in the field of protein polymers conjugates [135]. The PEGylation backbone enhances the instability and short half time besides improving therapeutic effect of the free drug [136].

However there is no good without bad. Certain PEG conjugated formulations may cause toxicity side effects. Excessive toxicity signs were observed during phase II clinical trials in patients with advanced ovarian cancer treated with PEG-L-asparaginase (Oncaspar®): Pancreatitis, fatigue, neutropenia and hypoalbuminemia,

weight loss, dehydration, decreased fibrinogen and hypersensitivity were identified and the trial was ended before completing [134, 137].

Other polymer conjugates which is under preclinical trials is N-(2-hydroxypropyl) methacrylamide (HPMA) copolymer [138]. The first example of an anti-cancer agent coupled to HPMA is that of Duncan et al., that have demonstrated, although less effective than free daunomycin, an inhibitory activity of daunomycin-HPMA conjugates in vitro [139]. A more recent study by Satchi-Fainaro et al., have targeted angiogenesis with a conjugate of HPMA copolymer and caplostatin in order to overcome short circulation time in plasma and improve the therapeutic effect of the free drug [140].

Drug based polymer conjugates can also potentially improve cancer chemotherapy by preferentially target the tumor. Examples for chemotherapeutic drug conjugates which are under clinical investigation are the (N-(2-hydroxypropyl) ethacrylamide) (HPMA) copolymer-Dox [141] for the treatment of breast, lung and colon cancers. Poliglumex (Xyotax) is a paclitaxel-poly-L-glutamic conjugate [142] and is in phase II clinical trials in ovarian cancer patients [143] and phase III for advanced non-small cell lung cancer patients [144].

In order to increase treatment effectiveness, a combination therapy for blocking both cancer cells and cancer stem cells based on HPMA copolymer PI3K/mTOR inhibitor and docetaxel was developed by Zhou et al. [47].

Main advantages of HPMA as drug carrier include increasing solubility, prolonging the circulation time, and improving pharmacokinetic/biodistribution profiles of small molecule drugs.

Trojan peptides known as cell-penetrating peptides (CPPs) composed from short cationic arginine-rich amino acid sequence capable to traverse cell membrane efficiently. CPPs can be conjugated to chemotherapy drugs and used as a non-receptor-mediated delivery platform [145, 146]. However, CPPs-based delivery platforms suffer from low specificity and a limited therapeutic effect.

Other proposed polymeric conjugates for delivery are: poly(ethyleneimine) (PEI), polyvinylpyrrolidone (PVP), polydimethylacrylamide (PDMA), polyacrylamide (PAM), polyvinyl alcohol (PVA), chitosan and dextran [135].

Polymers can be designed as passive drug delivery conjugates and its penetration to the cancer cells is mainly attributed by the EPR mechanism, or when including a targeting moiety to increase specificity to cancer cells microenvironment via an active cellular targeting strategy [98].

Using folic acid as a targeting agent which can be easily conjugated to the polymer may facilitate drug uptake by the cancer cells via receptor-mediated manner and improve therapeutic effect as was demonstrated by pullulan conjugated doxorubicin delivery system [11].

Another tumor targeting conjugate is the arginine-glycine-aspartic acid (RGD) peptide, selectively target the cellular receptors for $\alpha v \beta 3$ and $\alpha v \beta 5$ integrins expressed on tumor vasculature cells [147].

HA can be conjugated directly to chemotherapy drugs to target highly metastatic CD44-positive tumor [148, 149].

7.12.1 *siRNA Conjugate Polymers*

A novel treatment modality could utilize RNA interference (RNAi) for therapeutic intervention in variety of diseases including cancer. Small interfering RNAs (siRNA) are a class of short double-stranded RNA molecules that can knockdown the expression of genes that has the complementary nucleotide sequence, yet siRNA delivery is a major abstacle due to its limited biological stability and negative charge characteristics. Polymer conjugates use for improving siRNA stability in circulation. In order to inhibit the potential for lung metastasis, Bonnet et al. intravenously injected TSA-Luc mammary tumor-bearing mice with a stable complex of siRNA against survivin and cyclin B1, key cell cycle regulators, both involved in cell proliferation and survival processes. A strong inhibition of lung tumor metastases was demonstrated. An insignificant inhibition (<20 %) was observed with the classic siRNA-PEI complex [150]. No additive inhibitory effect was demonstrated when the two genes were simultaneously targeted. The therapeutic effect of Cis was increased (up to 90 % tumor inhibition) when injected after cell cycle blockage with the ssiRNA-PEI complex treatment, compared with the 40 % inhibition when cisplatin was injected alone. Mouse survival rate following co-treatment demonstrated a similar trend [150].

Shen et al. presented simultaneous inhibition of both metastasis and tumor cells by co-delivery of shRNA together with Ptx. The master regulator Twist mediated tumor metastasis by promoting epithelial to mesenchymal transition (EMT), which occurs in metastatic sites. In order to block EMT, Twist shRNA (TshRNA) and Ptx were both conjugated with pluronic P85 and polyethyleneimine (PEI) polymer to form the D- α -Tocopherylpolyethyleneglycol 1000 succinate complex. In the pulmonary metastasis mouse model, a significant synergistic inhibitory effect of both tumor growth and pulmonary metastasis formation was demonstrated. The IC₅₀ of free Ptx was 63-fold higher in comparison with the TshRNA-Ptx complex. In addition, the complex had a prolonged circulation time and promoted increased accumulation of Ptx and TshRNA in lung and tumor tissues [151]. Finally, an elegant approach to conjugate siRNA to triantennary N-acetylgalactosamine (GalNAc) induce robust RNAi-mediated gene silencing in the liver, most probably due to an uptake mechanism mediated by the asialoglycoprotein receptor (ASGPR) on hepatocytes. This promising strategy is currently under clinical investigation [152]. It will be interesting to see if this strategy could also be applied to lung tumors and their metastases.

7.13 Conjugate Chemistry

Covalent conjugation has a number of advantages over physical adsorption. Specifically, (1) increased stability until the carrier reaches the targeting site, (2) control over the density of the targeting molecule on the nanocarrier surface, (3) linkers for improved availability. Conjugation of targeting moieties to nano-based

carriers can be achieved through a variety of chemistries including simple succinimidyl ester chemistry at room temperature [63]. For example, through reactive NH_2 terminated aptamers with polymers containing carboxyl groups (COOH) in the presence of Ethyl-3-(3-dimethylaminopropyl)-carbodiimide (EDC) and N-Hydroxysuccinimide (NHS) under aqueous conditions, amide linkage is formed and a product yield of 90 % should be routinely achieved [153]. The conjugation methodology can be categorized into five major reaction types: reaction between activated carboxyl groups and amino groups, which yields amide bond; reaction between thiols, which yields disulfide bonds; reaction between maleimide derivatives and thiols, which yields thioether bonds; reaction between hydroxyl and carboxyl, which yields ester bonds; and reaction between *p*-nitrophenylcarbonyl- and amino-group, which yields carbamate (urethane) bonds (Fig. 7.5).

Chemotherapeutic drugs can be also covalently conjugated to drug carriers. A preferred linkage for drugs would be ester bond since ester groups can be immediately cleaved by cytosolic enzyme esterases after endocytosis and, thus, can exploit the prodrug conversion mechanism. Ester linked drug carriers can be achieved by the EDC conjugation chemistry using hydroxyl group containing polymers and carboxylic acids or NHS attached drugs such as methotrexate and Taxol [154, 155]. The readers are referred to excellent reviews on conjugation chemistry in [84, 156].

Characterization of the binding of antibodies or aptamers to the nanoparticles can be achieved through conjugation of a fluorescent probe such as Fluorescein isothiocyanate (FITC), AlexaFluors, rhodamine or any other fluorophors and examining the particles under a fluorescent microscope or with flow cytometry. A more precise way is conjugation of ^{111}In or ^{125}I directly to the targeting moiety (usually antibodies) prior to their conjugation to the particles and an accurate measurement of the radioactivity prior and post conjugation to the particle. This can give also an estimation of the number of mAb per particle with a known diameter [84, 156–158].

7.14 Conclusion and Future Prospective

The development of nano-based carriers for targeting cancer is currently a vibrant area of research which promises the state of the art in cancer diagnostics and treatment. Although numerous nano-carrier based strategies exist, choosing the appropriate nano-carrier is still not obvious and comparative studies, of which there are few, are difficult to interpret and a suitable system which demonstrates optimal characteristics remains elusive. Determining the optimal nano-carrier system is especially difficult given the numerous factors which effect biodistribution and targeting. Therefore, successful targeting strategies at present must be determined experimentally on case by case basis. In addition, systemic therapies employing nano-based carriers require methods for overcoming few factors, such as: (1) non-specific uptake of nanoparticles by the mononuclear phagocytic cells and non-targeted cells (2) lack of the EPR effect in some cancers (3) the presence of necrotic processes in malignant masses [159].

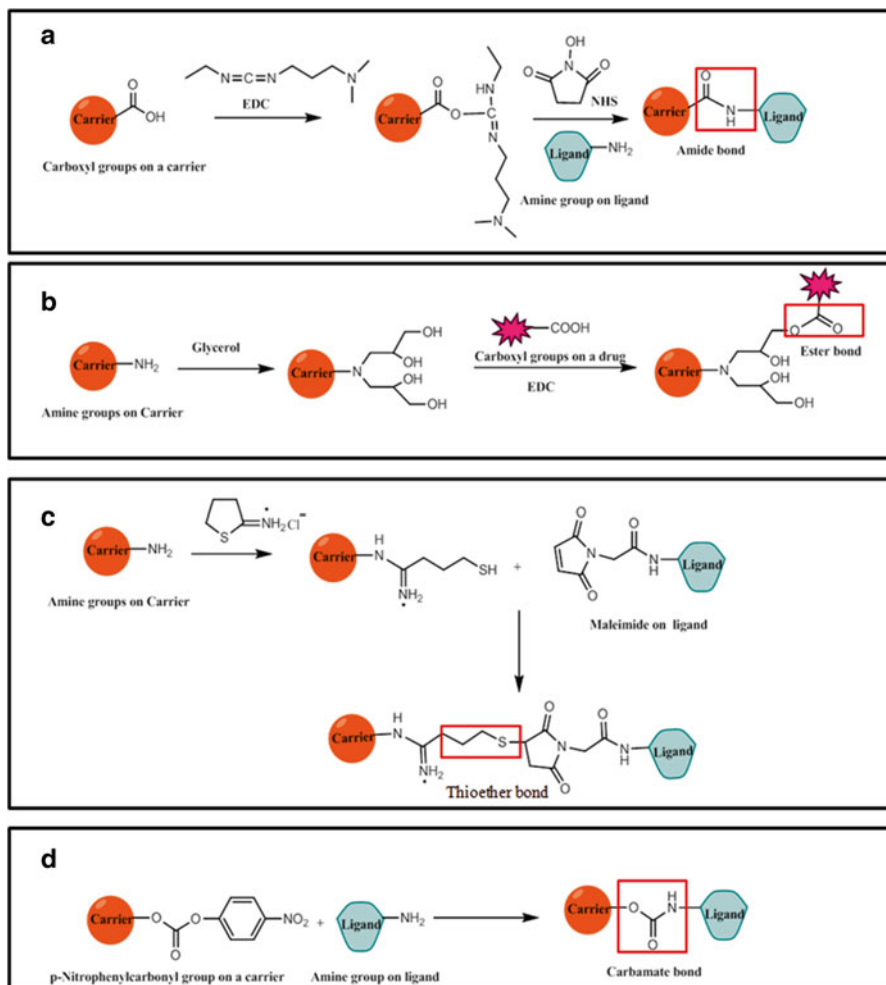


Fig. 7.5 Examples of common conjugation chemistry forming five major linkages. (a) Amide bond formation facilitated by EDC/NHS chemistry (two step reaction), (b) ester bond formation using glycerol as a linker (two step reaction), (c) thioether bond formation (two step reaction), and (e) formation of carbamate (urethane) linkage (two step reaction). Note that all the conjugation shown here, fall under the definition of Click chemistry due to the simplicity of their conditions, and may be achieved under aqueous conditions at room temperature

From a successful pharmaceutical delivery system perspective, there are few successful liposomal formulations (13) and a fewer number of conjugated drugs (3) on the market and above 300 different formulations in different stages of clinical trials—representatives can be seen from Table 7.1.

As this field matures, there will undoubtedly be a great demand for simple scale-up methods and high throughput assays for quality assurance and this is certain to

Table 7.2 Examples for targeting ligands and their receptors

Ligand	Receptor
Transferrin	Transferrin receptor [160]
Hyaluronan	CD44 [161]
Folic acid	Folate receptor
RGD peptide	Integrin $\alpha\beta3$ [162]
Somastatin analogues	Somatostatin receptor

be an active area of development. With the fulfillment of these requirements, we are likely to enter an era where nano-carriers will play a fundamental role in the oncology field in the upcoming years (Table 7.2).

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Chapter 8

The Importance of Particle Geometry in Design of Therapeutic and Imaging Nanovectors

Matthew J. Ware, Jenolyn F. Alexander, Huw D. Summers, and Biana Godin

Abstract One of the main objectives in nanomedicine is to enable specific delivery of therapeutics and imaging agents to the disease loci. While previously the main tool for targeting was incorporating entities with biological recognition on the surface of the nanovector, recently, the focus has shifted to the ability of the physical characteristics of particles to guide them crossing numerous barriers to the site of action. In this chapter we will focus on how the geometry of nanovectors affects their interaction with biological milieu, and as a result their therapeutic and diagnostic potential. The chapter is divided into sections describing interactions of various particles with extracellular components, whole cells and various cell organelles. In the intracellular interactions subsection, we focus on the initial interaction between particle and cell membrane, the uptake mechanism and intracellular trafficking of particles with various geometries. Furthermore, the importance of charge density and the zeta potential parameter is discussed. The chapter concludes with a discussion of challenges and outlook on future developments of theranostic particle delivery using particle geometry as the rational design feature.

Keywords Particle size • Particle shape • Zeta potential • Surface • Interactions with cells • In vivo • In vitro • Geometry

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8.1 Introduction

Due to the numerous physical and biological barriers to drug and contrast agent transport in the body, the minimization of side-effects with simultaneous targeted delivery of active agents to specific tissues of interest continues to pose a challenge. The use of nanoscale vectors to deliver drugs, genetic material, and imaging contrast agents has been an intensively investigated niche in medicine, as achieving desirable bioavailability of therapeutic agents been especially challenging in the fields of gene therapy [1] and oncology [2–4]. Furthermore, nanovectors may support the transport of hydrophobic drugs [5], as dissolution of such drugs in the bloodstream is difficult, and in cases where the drug or imaging agent is too fragile, insoluble, or toxic for direct in-vivo administration.

It is becoming increasingly apparent that the design of contrast agent and drug nanovectors involves the manipulation of several critical factors. These include their chemical functionality and mechanical properties, the degree of dispersion or encapsulation of the drug, on or within, the nanovector, the size and shape of the particle, and the surface charge.

The ability to easily fabricate drug or contrast agent-loaded spherical particles with a narrow size distribution by a number of bottom-up methods including emulsion, dispersion, suspension, polymerization, precipitation and spraying techniques has meant that the role of particle size and composition play in biodistribution [6], cellular binding [7], cell entry pathways [8], cell uptake [9–11], and tumor bed penetration [12, 13] have been extensively studied for spherical particles. However, the majority of naturally occurring (and blood-born) biological entities are non-spherical and biological processes typically occur under dynamic conditions in which the motion of spherical and non-spherical objects will differ. For example, Decuzzi and Ferrari's theoretical work predicts that, under conditions of linear shear flow such as that in the bloodstream, oblate particles adhere more strongly to biological substrate than spherical particles, and hence, the use of non-spherical particles for delivery is predicted to improve therapeutic and imaging efficacy [14]. Additionally, shape is crucial to the particle's mechanism of cell entry [15–17], and the release rate of the therapeutic cargo [18, 19]. Past limitations in the control of particle fabrication have been overcome as novel methods such as photolithography have been developed. This means researchers are now able to access a multitude of different shapes at the sub-micron size range on a sufficient volume scale to allow for extensive in vitro and in vivo biological studies (Fig. 8.1). This will allow a more complete understanding of how the combined effects of particle size, shape, charge, chemical composition and mechanical properties affect biodistribution, target tissue interaction and cell entry which will lead to the development of increasingly efficient drug and image contrast based nanovectors. While numerous studies further discussed in this chapter have shown that the geometry of the particle is an important parameter when the particles are administered intravenously, interactions with mucosal barriers have been shown not to depend on the shape of particulates [20]. In contrast, the size of nano- and micro-particles does play a role in permeation across the mucosal barrier. As an example, Hanes's group has shown that even when nanoparticles are larger than the mucus mesh size (e.g. 500 nm), they can

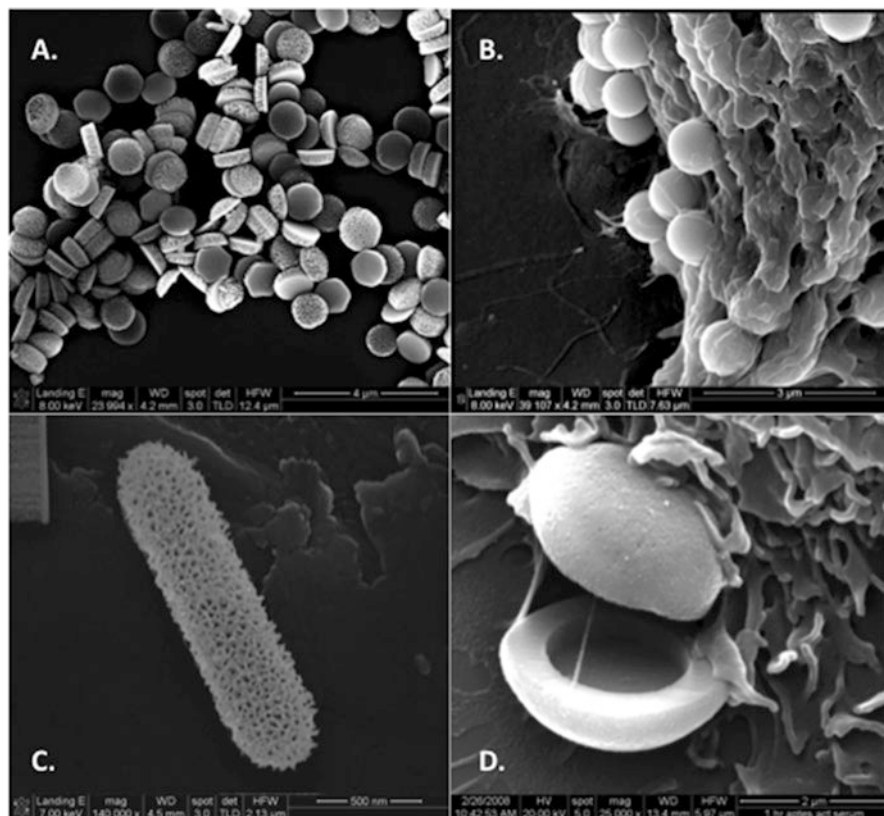


Fig. 8.1 Scanning electron micrographs of silicon particles fabricated by top-down approach using photolithography and electrochemical etching: (a) Discoidal porous silicon particles. (b) Spherical silica beads (fabricated by bottom-up synthesis) on a pancreatic cancer cell (PANC-1, ATCC). (c) Porous silicon rods. (d) Quasi-hemispherical porous silicon particles

be mobile in the mucus if coated with hydrophilic polymers [21]. Further in this chapter we will focus on systems that are intended for intravenous administration.

This review aims to summarize the various fabrication approaches, including ‘Top Down’, ‘Bottom Up’, and novel combinational fabrication techniques, which are able to create nanoscale particles with various intricate geometries tailored towards a specific biological function. We attempt to relate aspects of particle design with their systemic behavior *in vivo* and also their action at the target cell. We discuss the importance of the surface charge of nanoscale vectors and common assumptions regarding its measurement on non-spherical particulates, which affect understanding of the initial particle-cell membrane interaction as well as with other biological entities *in vivo*. We summarize the approaches to improve efficiency in particle function, including geometry and surface charge manipulation, the use of targeting moieties, ligand binding, PEG conjugation and other functional groups. The chapter concludes with an overview of future perspectives relating to the improved use of nanoscale vectors in active agent transport.

8.2 Particle Fabrication Approaches

8.2.1 ‘Top Down’ Fabrication

8.2.1.1 Photolithography and Electrochemical Etching

Top down fabrication of therapeutic and diagnostic particles of various shapes and sizes uses the same principles developed and refined in the microelectronics industry throughout the past several decades: Photolithography and electrochemical etching. These techniques are characterized by high reproducibility and yield, and allow particles to be fabricated with homogenous and precise dimensions “by-design”. Rationally designed particles have an increased ability to negotiate the numerous bio-barriers associated with the delivery of active agents in the body. For instance, in recent works multistage vectors (MSV) are presented [22, 23]. These vectors are comprised of several distinct nano-elements or ‘stages’; each designed to negotiate one or more bio-barriers from the site of administration to the target lesion. The discoidal stage 1 mesoporous silicon particles (S1MP) not only hold and protect the cargo drug or image contrast agent (Fig. 8.2), but also enable increased margination and interaction with vasculature walls and cell surface adhesion properties during their transportation in the blood [24, 25]. Porous silicon was used as the material for S1MP fabrication due to its biodegradability [26, 27], biocompatibility [27, 28] and versatile fabrication methods which allow the control of shape, size and porosity [23]. S1MPs are fabricated by an integration of two top-down approaches; photolithography and electrochemical etching for particle porosification. Concentration of the etching solution, doping, electrical current and etching time affect the porous structure, while photolithography is used to control the pattern and geometry of the particles via silicon micro-fabrication, which is well known in the silicon micromachining industry [23, 29, 30]. These techniques enable precision, reproducibility and scalability.

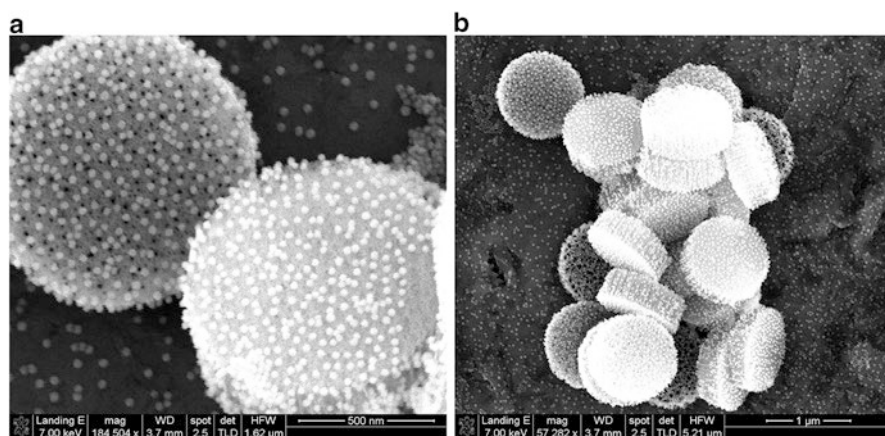


Fig. 8.2 Scanning electron micrographs of silicon mesoporous particles (S1MP) loaded with iron oxide nanoparticles for image contrast applications. Scale bar: (a) 500nm; (b) 1 micron

Chiappini et al. [29] outlined the fabrication protocol of quasi-hemispherical S1MPs yielding monodisperse particles 900 nm–3.2 μm in diameter, 30–65 % porosity and 3–55 nm pores. Briefly, starting with a silicon wafer coated with a dielectric silicon nitride (SiN) film, an array of particle patterns is first transferred to a dielectric layer by a photolithography process and Reactive Ion Etch. A silicon etch is then performed which defines the trenches that nucleate S1MPs with different profiles [29]. A successive, two-step electrochemical etch process is applied to yield the S1MP of desired porosity, pore size and thickness. A highly porous release layer is formed at the wafer-S1MP interface retaining S1MP on the substrate and allowing for differential chemical modifications of the two sides of S1MP if required (Fig. 8.3).

Discoidal vectors are fabricated by photolithographic patterning of a porous silicon film, a process that eliminates potential distortion effects introduced by non-uniform current distribution during electrochemical etch, simultaneously granting independent control of the porous structure. A novel sealing strategy to pattern pSi films using low-pressure chemical vapor deposition of LTO was implemented to overcome the problem of photoresist adsorption within the pores. After electrochemical etch of double layered porous silicon films, the porous films are sealed by LTO, and an array of circles is patterned on LTO film. A CF_4 Reactive ion etch is then performed to etch through LTO and two-layered porous silicon producing monodisperse discoidal S1MPs with diameter as small as 400 nm and pore size ranging 3–150 nm. The S1MP fabrication process is robust and well characterized. Using 1000 nm diameter disks as an example, more than 2 billion S1MP patterned on a 4 wafer can be obtained. Recent studies [22] point towards the possibility for a multi-layer fabrication method of discoidal particles, which should increase the production yield by tens times.

8.2.1.2 Step and Flash Imprint Lithography

Step-and-flash imprint lithography is a technique that replicates the topography of a rigid mold using a photocurable prepolymer solution [31]. In this technique a low viscosity, photocurable liquid or solution fills the void spaces of the mold. The solution consists of a low-molecular-weight monomer and a photo-initiator. UV light exposure polymerizes and, therefore, solidifies the precursor while in contact with the mold. Removing the mold leaves a topographically patterned inverse replica on the substrate.

Step and flash imprint lithography has recently been used to fabricate a range of non-spherical particles such as cubic, triangular and pentagonal cylindrical particles. From the numerous scientific studies involving enzymatically degradable peptide sequences for drug delivery [32–35], Glangchai et al. [36] were successful in using cathepsins to initiate the release of DNA plasmids from cubic, triangular and pentagonal cylindrical particles *in vitro*, by combining the superior shape control provided by lithographic methods with novel drug loading and release approaches.

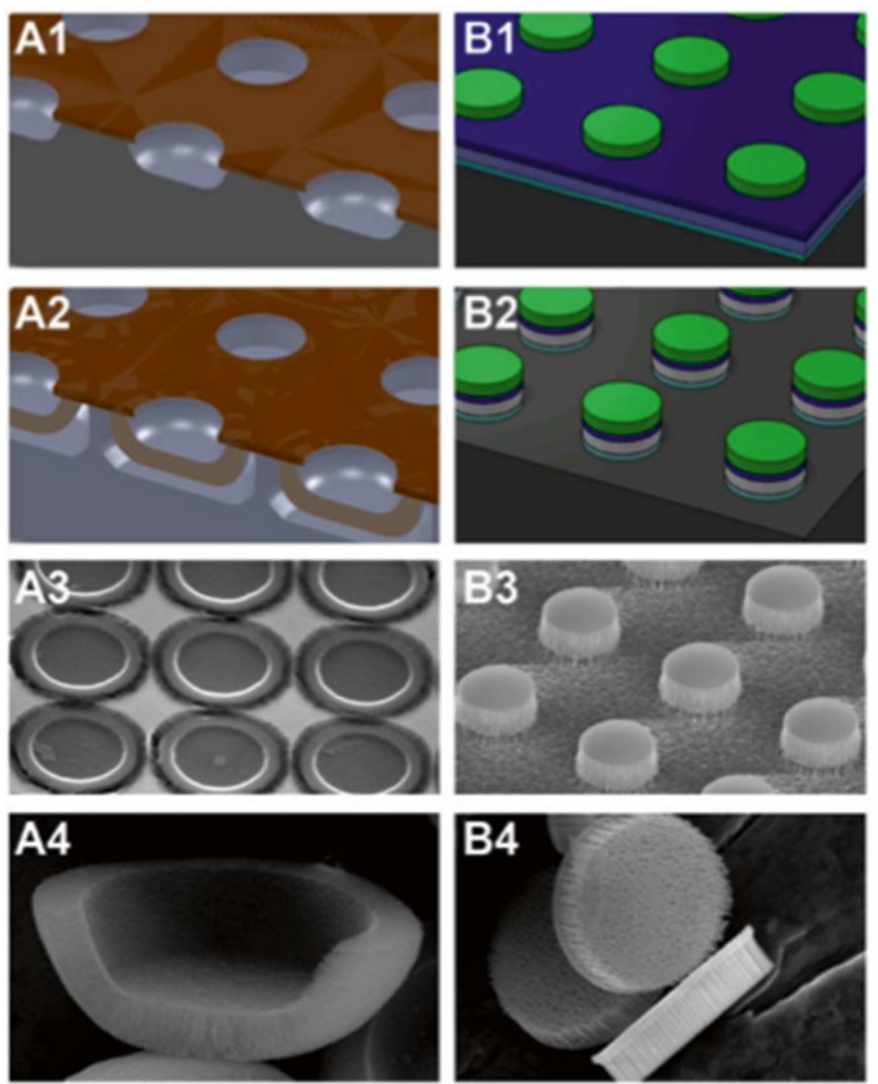


Fig. 8.3 Fabrication of silicon mesoporous particles (SIMP). (a1) Patterned silicon nitride (SiN) layer and trenches etched into silicon. (a2) Electrochemically etched SIMP with release layer. (a3) SIMP array on wafer after removal of SiN. (a4) Cross-section of hemispherical SIMP. (b1) Photoresist pattern on Low Temperature Oxide (LTO) capped porous silicon film with release layer. (b2) Particle array on wafer after RIE. (b3) Discoidal SIMP array on wafer after LTO removal. (b4) Released discoidal SIMP [22]

8.2.1.3 Particle Replication in Non-Wetting Templates (PRINT[®])

The PRINT[®] (Liquidia Technologies) process [15, 37–45] is a modification of traditional lithographic methods and can produce polymeric micro and nano vectors. PRINT[®] provides precision, control and flexibility in the fabrication of rationally

designed particles and can regulate particle size and shape independently of composition. Since PRINT[®] uses elastomeric fluoropolymers, instead of traditionally used silicones it provides a selective filling of nano-scale cavities in the mold using an organic liquid whilst avoiding the formation of interconnecting “flash layers” which are a common problem in traditional imprint lithography [46]. PRINT[®] particles can be produced with diverse surface chemistries, degradation characteristics and elastic moduli because organic liquids and sol-gel metal oxide precursors do not swell fluoropolymers in the same fashion as they swell silicones. Furthermore, photocurable perfluoropolyethers (PFPEs) used in the PRINT[®] process maintain the desirable flexibility of Polydimethylsiloxane (PDMS) whilst exhibiting an improved resistance to solvents [47, 48]. The PFPE molding materials are able to accurately pattern fragile or soft objects at the nanoscale, such as block copolymer micelles and naturally occurring entities, such as viruses [38]. The PRINT[®] process has constructed particles from natural matrices such as biologically relevant proteins [44] and synthetic matrix materials, such as highly cross-linked hydrogels and linear polyesters [37, 39] and has fabricated particles which have been shown to be effective delivery agents for therapeutic drugs and contrast imaging agents [42, 43], such as for the delivery of doxorubicin to cervical HeLa cancer cells [43].

Despite its uses, fluorinated molding matrices are more expensive than silicone materials. However, cost has been mitigated by the development of continuous mold manufacturing processes that use inexpensive backing materials coated with a fine film of the more expensive fluorinated molding material [49]. This strategy has enabled the PRINT[®] process to become economically more feasible and has led it to become a scalable particle manufacturing process.

8.2.1.4 PDMS-Molding Processes

Lightly cross-linked PDMS, also known as silicone rubber, is one of the most widely studied molding material due to its low toxicity, low modulus, high gas permeability, which allows dead-end filling and low surface energy which enables facile release from master pattern templates as well as replication of nanoscale objects [50]. However, for some applications, the swelling of PDMS molds in organic solvents or PDMS-fragment transfer to the sample are major weaknesses. Other silicone-based materials and filled molding material formulations have been developed that limit the mechanical distortion of the stamp [51]. Additionally, several mold surface treatments [52, 53], such as oxygen plasma and fluorosilane grafting, have been investigated to ensure the mold surface is more hydrophilic or hydrophobic, respectively, which effects wetting during mold filling and the release of the particles from the mold. Lastly, cleaning procedures have been developed which limit sample contamination from small fragments originating from the PDMS-stamp [54].

Hansford et al. have used PDMS stamps to produce both thermoset and thermoplastic microparticles and medical devices with various geometries [55–57] for drug delivery applications. Furthermore, Guan and colleagues highlighted the numerous possibilities created by the precise geometric control on the micro and

nanometer scale by the design of unique geometries with extensions that can self-fold to imbed in intestinal tissue [57] after oral delivery. The same group also fabricated microcapsules for intravenous delivery, which contained sucrose as the cargo and a thin poly(vinyl alcohol) (PVA) or poly(lactic-co-glycolic acid) (PLGA) based outer layer which swells in the presence of water due to osmosis. This releases the drug via diffusion through the outer membrane rather than dissolution of the thermoplastic matrix, and, thus, creating a novel drug transport mechanism with higher drug loading capacities than conventionally prepared PGLA microcapsules [58]. Additionally, Oudshoorn et al. have explored various molding techniques including rigid micro-molding, soft micro-molding, and photolithography to fabricate uniform methacrylated hyperbranched polyglycerol microparticles [52]. Square and hexagonal microparticles particles were obtained in high yields, however, PDMS-based molding techniques produced a higher number of less well defined particles in relation to shape when compared to the rigid, epoxy-based micro-mold.

In summary of this section, the ability to prepare particles in a manner than allows independent manipulation of one geometric parameter at a time has insights into shape dependent biological processes such as phagocytosis to be developed. Many techniques, such as stretching thermoplastic particles [16] or flash imprint lithography preparation of cross-linked particles as previously described [36], can be used to fabricate small amounts of shape-specific particles. Practical, large scale manufacturing techniques for fabrication of geometrically precise particles constitute a major development gained by the application of the PRINT[®] process to biomedical and other disciplines. Traditional soft lithography using PDMS-based molding materials offers an additional technique in the design of micron-sized particles with specific geometry. However, for applications such as intracellular drug and imaging agent delivery, where nanoscale vectors are often most efficacious, opportunities exist to improve on PDMS-based soft lithography using various other molding techniques.

8.2.2 ‘Bottom Up’ Design of Biomimetic Assemblies

Bottom up fabrication is a more classic method to produce nanoscale and microscale particles starting from molecules, which are the building blocks in such case. Since spherical shape is, generally, the most thermodynamically favorable one, most of the bottom-up produced particles are spheres. It is difficult to develop synthetic systems that mimic biological macromolecules and objects, as their structures and geometries are created with highly specific interactions that work at extremely small dimensions. However, using various building blocks and fabrication techniques, it is possible to manipulate the geometry of bottom-up produced particles to some extent. The use of self-assembling molecules, where the structure contains at least two distinctive moieties, a biologically active portion and a fraction that directs assembly, presents one particular design strategy that does emulate nature. This

approach fits into a larger framework known as ‘bottom up’ design, where complex functional structures are generated by assembling a group of molecularly interlocking parts [59]. Such systems mimic biological architectures by using a particular set of primary molecules that are capable of associating into secondary and tertiary structures. This process is initiated by incorporating particular peptides, nucleotides, sugars or binding synthetic molecules to the primary molecule to provide it with the appropriate biologically specific information. Secondly, a molecular region must be included to direct the assembly of the molecules into a highly controlled tertiary structure, such as micelles, vesicles, tubules, and other more complex architectures. The assembly provides potency to the biological information it carries by several possible mechanisms, which include the control of spatial relationships, and concentrating locally the bio-functional entities, as well as providing opportunities for combinatorial selectivity. However, the ability to precisely design for a specific function remains a significant challenge. Currently, an ever more-growing number of interesting new structures can be designed with function in mind; however, accurate structure function relationships in this type of assembly are warranted to further accelerate the effective design process. Additionally, many of these new biomimetic particle designs are comprised of relatively new molecules and the balance between therapeutic efficacy and toxicity is poorly understood. Several unknown parameters for these nascent molecular assemblies can result in the restriction of use, primarily toxic side-effects, but also the sterility, stability and scale-up of the complicated drug carrier systems [60].

Bottom-up strategies are frequently used by nature to combine proteins into quaternary structures to create systems with a specific function. Nature is adept at maximizing utility whilst minimizing the use of resources and economizing synthesis by assembling molecules in a variety of ways to give a multitude of binding targets by the integration of heteromeric assemblies. Dimers such as antibodies, integrins, and leucine zippers, trimers such as collagen, and larger multimeric assemblies such as viral capsids are examples of molecules that act in this fashion. Leucine zippers show that homodimers and heterodimers give quaternary structures where several molecules are coupled together in a combinatorial fashion to give greater target diversity and specificity. Another mechanism of control is provided by the coupling reaction itself, where the monomeric uncoupled assembly does not possess the desired binding pattern or function and is, therefore, in a dormant phase. Cellular networks comprised of these modular components of a structurally comparable class may selectively self-assemble to produce subtle functional correlations. This concept is found in a viral capsid, where the process of self-assembly has formed unique protein architecture to create a functional icosahedral cage and would be extremely useful to mimic for drug delivery applications. Several attempts have been made to imitate biological structure and function by using bottom-up strategies to design molecules that are capable of self-assembly. However, successful synthetic emulation of this type of assembly would require a comprehensive knowledge of the precise forces that facilitate self-assembly in a variety of nanoscale molecules that can be tailored to facilitate the delivery of drugs by emulating the characteristics of natural biological systems.

In summary, several physical parameters dictate the assembly of macro-molecular structures and nano-crystals/metallic particles, and predicting and controlling these aggregates is extremely complex. The synthesis of simple molecular building blocks can lead to intricate self-assembled architectures much more complex than the original target structure [61–63], which can be subsequently rationalized based on the physical parameters and chemical principles of self-assembly, which include, but are not limited to, intermolecular forces and thermodynamics which dictate the mechanisms of assembly [64].

8.2.2.1 Hydrophobic Interactions

Hydrophobic interactions play a major role in the rational design of self-assembling molecular aggregates. Amphiphilic molecules, which comprise both hydrophilic and hydrophobic moieties, use the hydrophobic interaction to form a variety of aggregated structures. The hydrophobic moiety provides a driving force for assembly, however, assembly can stem from several factors that make aggregation either favourable or unfavourable. Understanding these forces will help describe the general driving forces for aggregate formation [65, 66]. These forces include; the transfer of hydrophobic tails from an aqueous environment into the aggregate, the assembly and interaction of the alkyl groups with the solvent as a function of the micelle area, hydrophobic packing, which causes an entropic decrease that is associated with less conformational freedom in the aggregate and finally, forces involved with the attractive, or more commonly, the repulsive interactions of the head-groups, such as steric interactions of polymeric head-groups, or electrostatic repulsions in charged head-groups.

Self-assembling amphiphilic molecules have attracted attention for the novel design of complex structures with biological specific functionalities. Luk and Abbott [67] review a variety of head-group interactions that are biologically relevant, whereas, several other groups have explored the conjugation of carbohydrates [68], peptides [63, 69], polymerizable elements in the hydrophilic head-groups [70] and hydrophobic cores [71] for stability that affect the structure and function of amphiphiles. Nature's use of amphiphiles follows two specific design goals; the formation of organelles and other compartmentalized environments and simultaneously the precise presentation of specific signals at the interface of these enclosures. This is comparable to the applications pursued by chemists, who control assembly by the subtle manipulation of chemical structures and hydrophobic interactions which leads to the production of a variety of nanoscale assemblies, including vesicles, micelles, sheets, tubes and bundles [67].

8.2.2.2 Electrostatic Interactions

Electrostatic interactions provide a useful tool to manipulate the dynamics of self-assembly as they provide both attractive and repulsive forces, which can be controlled by adjusting salt concentrations within the system. For instance, coiled-coil

heterotrimeric assemblies have recently been designed via the manipulation of both hydrophobic and electrostatic interactions to drive assembly and to precisely match primary structures, respectively [72]. The specificity of assembly can further be controlled via the manipulation of charge-to-charge interaction, and provides a design that is selective for trimeric α -helical coiled-coils as building blocks. This coiled-coils design strategy demonstrates that specificity can be included in the assembly process via electrostatic interactions and these types of biologically relevant assemblies can be used to develop therapeutics with tightly controlled geometries.

Hydrogen bonding originates between positively charged hydrogen atoms and negatively charged electronegative atoms and, thus, may be considered a subset of electrostatic interaction [65]. The positive charge in hydrogen atoms is a result of a covalent bond to electronegative atoms, and quantification of this interaction is difficult as it is dependent on the chemical components and the angle of bonding between the atoms involved [73].

To conclude this sub-section, self-assembling systems must be designed for particular applications based on the class of molecules used and the forces that dominate their assembly. A combination of forces that describe self-assembly as a complex milieu with several local energy minima, much like protein folding, should be considered when making realistic predictions of self-assembled structure. This will enable evermore-complex drug and contrast agent delivery vectors with biologically relevant dimensions and functions to be designed, which take advantage of the elegant and intricate process of self-assembly.

8.2.2.3 Metallic and Carbon Nanoparticles

In addition to molecular assemblies, other nanoscale materials of different shapes, such as metallic and carbon nanoparticles can be synthesized via bottom up methods. Metallic nanoparticles are unique materials for optical, electronic, catalytic, and sensing applications. There is a great deal of interest in using metallic nanoparticles as building blocks in the development of more complex nanostructures through the use of a 'bottom-up' approach due to the flexibility in controlling the surface chemistry of these particles through functionalization. Using self-assembly techniques, one can exploit spontaneous chemical interactions to build complex nanoconstructs. Gold nanorods have been synthesized and modified with various polymers, inorganic oxides and organic ligands to establish principles for self-assembly of these unique nanomaterials, and are of great interest due to their strong optical absorption in the visible and near infrared regions, which can be tuned through material preparation and modification of the surrounding environment. Gold nanorods can be prepared via: (1) the template method, where gold is electrochemically deposited within the pores of either a polycarbonate or alumina template membrane; (2) the electrochemical method where an electrical potential is placed across a gold anode and a platinum cathode. The two electrodes are placed in an electrolytic solution containing cationic surfactants, which form cylindrical micelles. As gold ions from the anode are deposited onto the platinum cathode, the

surfactant stabilizes the rod allowing it to grow in a cylindrical manner and (3) the seed-mediated growth method, where a small amount of spherical particles, which act as seeds, are added to a growth solution containing additional metal salt and a structure-direction agent and is one of the most reliable and versatile methods to control the shapes of noble metal nanocrystals.

Several research groups have succeeded in fabricating organized CNT arrays by growing carbon nanotubes on patterned catalyst printed substrates or by post-synthesis alignment [74–76]. Controlled CNT arrays are usually fabricated either by growing CNTs on catalyst pre-patterned substrates, such as iron, nickel or cobalt substrates, in a chemical vapour deposition (CVD) reactor [75–77] or by self-assembling functionalized CNTs on the substrate [78, 79]. Aligned CNT structures by self-assembly approaches [78–80] present some solutions for the limitations of organized CNT structures prepared by existing approaches, including the ability to fabricate CNTs at room temperature and flexible substrate selections. Additionally, CNT arrays have been manipulated with functionalized CNTs by self-assembly. Shortened CNTs by chemical oxidation were self-assembled on silver coated silicon substrate [80], and Fe ion coated silicon substrates by interactions with carboxylic groups of CNTs and substrates [78]. Thiol functionalized shortened CNTs have been used for self-assembled arrays on gold-coated Si substrates [79]. For bulk production of aligned CNTs, self-assembly pyrolytic methods [81], thermolytic processes with sprayed solutions [82], and pyrolyze systems with sophisticated aerosol generators [83] have been suggested. Though self-assembly approaches demonstrate possibilities for fabricating CNT arrays, it is still challenging to precisely control nano-size dimension CNT arrays during their fabrication. Ultimately the combination of well-defined self-assembly techniques and nano-size patterning processes to obtain nano-size well organized CNT patterns for applications remains the goal.

8.2.3 *Combination Fabrication Approaches*

Several types of nanovectors with various geometries combining the top-down and bottom-up methods of fabrication have been recently developed with a view to combine the possibilities of both and/or overcome the limitations of either one of the independent approaches as discussed in this section [84, 85]. Examples include particles fabricated by photolithography followed by a self-assembling coverage of polymers or biological entities like phage [86–94].

Recent particles evolving in the world of nanotherapeutics are the hollow microcapsules composed of polymeric film, which have shown to possess extended circulation time [95], controllable degradation [96] and tunable drug release [97–99], capable of surface functionalization for enhanced cellular association, and function as suitable drug delivery agents to target cells [100–110]. These microcapsules can be generally produced using a self-assembly of hydrophobic-hydrophilic block copolymers resulting in polyerosomes or by combining top-down and bottom up

approaches. Microcapsules formed by self-assembly are commonly poly-disperse and limited to be spherical or worm-like in shape [111]. These two limitations are overcome by fabricating microcapsules by combining top-down and bottom-up methods; Rigid core-templates which are homogeneous in size and of desired shape are manufactured by methods that involve photolithography (top-down) and then are deposited with multiple layers of a polymeric monolayer or polymeric bilayers (bottom-up), depending on the functionality, on a rigid core template, following by dissolution of the core which forms hollow elastic microcapsules [90, 91, 93, 112, 113].

Using non-porous silicon discs as the rigid, core templates, erythrocyte-mimicking elastic capsule could be fabricated [114]. The sacrificial silicon discoid cores were made by photolithography combined with a reactive ion etching of Polycrystalline Silicon (PolySi). The particles upon detachment from the wafer were subject to PEI adsorption and then hydrogen bonded, pH sensitive copolymer layers of poly(N-vinylpyrrolidone) or poly(methacrylic acid) were assembled in a bottom-up fashion and cross-linked with carbodiimide. The rigid core-templates were dissolved in an aqueous solution of hydrofluoric acid, and dialyzed for 2–3 days. These cubical and discoid (erythrocyte-mimicking) capsules were hence programmed to exhibit change in dimension and volume based on environmental pH. Several such hollow polymeric particles have been engineered in the last decade, comprising of close-to-monodisperse, rigid sacrificial core-templates chemically synthesized from nano-seed solutions ammonium bicarbonate and manganese sulfate. By controlling the time, temperature and stirring conditions the shape of the rigid cores were manipulated to be spherical or cubical. The manganese sulfate concentration determined the size of the particles [93].

Polyelectrolyte microcapsules are capable of multiple functionalities by incorporation of quantum dots for imaging, superparamagnetic nanoparticles for application using magnetic fields, enzymes for pro-drug activation and intracellular sensing, pharmaceutical agents for therapy, metallic nanoparticles for remote drug release and surface functionalization for targeting purposes. The remarkable advantage of such systems is that all these functional properties can be potentially incorporated in a single nanoparticle system [108].

A recent study showed the design and therapeutic capability of a Multistage Nanovector based on Bacteriophage Associated Silicon Particles (BASP) [86]. This system, consisted of porous silicon particles manufactured by photolithography to produce quasi-hemispherical particles, and electrochemically etched in a hydrofluoric acid-ethanol mixture, to impart porosity. The particles were oxidized, and then functionalized with 2 % 3-aminopropyltriethoxysilane to confer a positive charge. Approximately 45 nm gold nanoparticles (AuNP) were prepared from Gold(III) chloride and trisodium citrate as per the Turkevich method [115]. Bacteriophages with suitable peptides were used to form AuNP-bacteriophage networks by overnight self-assembly [116]. The final BASP were obtained by mixing AuNP-bacteriophage networks with silicon particles for 1 h. By integrating the top-down and self-assembly approaches, it was possible to develop a unique theranostic multistage nanovector. Porous silicon particles, loaded with SPIONS coated with chitosan using combinational fabrication technology promise intercellular and intracellular targeting as well as transport of drugs and imaging agents [94].

8.3 Biological Interactions of Active Particles

In general, any drug molecule or imaging payload must reach its intended site of action to exert an efficient therapeutic or diagnostic action and to prevent adverse side-effects [117]. The delivery of such particles to the target organ or diseased loci generally requires a systemic administration through the circulatory system. Consequently, several biological barriers must be overcome to achieve efficient drug or contrast agent delivery [118]. These barriers involve systemic hurdles, including interactions with extracellular matrix and non-target cells, mononuclear phagocytic capture (MPC) by the mononuclear phagocytic system (MPS), and non-specific biodistribution, as well as cellular hurdles such as the target cell membrane, avoiding complete lysosomal degradation, and in some cases, overcoming the nuclear envelope. Particles must also remain stable and soluble, whilst not aggregating in the blood, or degrading which may expose the payload to degradation enzymes within the blood or inter-tissue fluid [119].

The manipulation of particle design is important for successful delivery to the target loci, and affects particles transport in the bloodstream, their extravasation out of the blood vessels, the diffusion through tissue, and finally, the degree of cellular internalization and subsequent uptake pathway and long-term fate. The appropriate design of a therapeutic/diagnostic particle requires a comprehensive understanding of all the characteristics of the vectors, as well as the mechanisms by which they interact within the biological environment and targeted cells.

8.3.1 Systemic Interactions

Regardless of whether the drug delivery vector is being administered via inhalation, intramuscular injection, gavage, or intravascular injection, it will unavoidably come into contact with the extracellular environment. Within this environment, multiple barriers exist which can result in rapid clearance and/or degradation of the nanovector before it ever reaches the intended site [120]. Furthermore, therapeutic efficacy of intravenously administered particles may be compromised by anatomical constraints and non-specific interactions with biological membranes, such as vasculature endothelium and connective tissue. Particle aggregation and disassembly may be caused by unwanted interactions with blood components, biological tissue and immune cells which are specialized in inactivation and removal of foreign particles, can lead to non-specific organ accumulation and low targeting efficiency and adverse side-effects [121, 122]. For example, blood circulatory half-life is highest for neutral particles, whereas cationic nanoparticles (NPs) possess the lowest half-life in the bloodstream and cause several complications such as hemolysis and platelet aggregation [123].

8.3.2 *Interactions with Immune Cells*

Generally, opsonisation of serum proteins, such as IgM, fibronectin and C-reactive proteins, to the surface of particle systems stimulates recognition by receptors on the surface of macrophages which enhances mononuclear phagocyte system (MPS) capture [124]. This leads to rapid removal from the blood and accumulation within non-target organs which contain a high population of resident tissue macrophages such as the liver [124], spleen and lungs. Most nanomaterials, when administered into the blood, are taken up within minutes or hours by phagocytic cells and are seen to possess a general predisposition for phagocytic clearance by the MPS. This is probably due to the fact that particles smaller than 5 μm in their largest dimension are comparable in size to the pathogens combated during the evolution of the immune system [125]. Micro-scale sized particles (0.5 μm and larger) have been described as invoking “bacteria-like responses” primarily being removed from circulation via phagocytosis, while NPs up to 200 nm in diameter are more likely to invoke “virus-like responses” [125]. Exploiting this trend, NPs have been intentionally designed to invoke, suppress, or avoid the immune response [126]. The use of nanovectors to target MPS organs has proven relevant in major therapeutic areas such as oncology, for the treatment of tumors, especially those that are situated in an organ of the MPS, such as hepatocarcinoma and liver metastasis and secondly in the treatment of infectious diseases, as many of pathogens that need to be eliminated are located in the MPS macrophages. Still the therapeutic moiety should escape phagocytic capture to exert therapeutic effect in other parts of the liver. Invocation of the immune response has driven size-dependence research in particulate vaccine vectors [127–129], suppression of an immune response could also make particulate carriers useful during organ transplant therapy, while avoidance of the immune system and reduced systemic toxicity using targeted particulate carriers is most often desirable in the case of imaging and therapeutic agents.

The emerging understanding of the importance of particle shape in phagocytosis by macrophages has been described and carefully studied by Champion et al. [16, 17]. The recognition that naturally occurring immunological targets vary widely in both size and shape provided motivation for this work. By carefully varying shape at constant size, the authors concluded that it is indeed particle shape, rather than size, which plays a dominant role for determining the complexity of the local actin structure, and ultimately whether phagocytic processes occur. Lipidic disks have been developed as alternatives to liposomes, having a diameter of 120 nm [130] to 250 nm [131], and a thickness of only a few nanometers, showing efficient uptake by MPS macrophages. Other lipidic assemblies such as cube-shaped so-called cubosomes [132] have been also proposed as drug nanocarriers. However, the impact of the shape of such systems on phagocytosis as compared to liposomes remains to be fully elucidated. As for polymeric systems, a recent study performed with polystyrene ellipsoids and discoidal particles has shown that, independently of opsonization, the local particle shape at the point of contact dictates whether macrophages initiate phagocytosis or simply spread on particles [16]. For example, a

macrophage attached to an ellipse at the pointed end will internalize it in a few minutes, while a macrophage attached to a flat region of the same ellipse will not internalize it for over 12 h. Interestingly, the same study has also shown that particle shape, but not size, plays a dominant role in phagocytosis. While initiation of phagocytosis in at least one orientation happened for particles of all shapes, the local particle shape, measured by tangent angles, at the point of initial contact is responsible for an ability of phagocytic cells to initiate/complete the phagocytosis or to “spread” the particle. On the other hand, particle size is sculpturing the ability of phagocytes to complete the process of the particle uptake [16].

Deformability, or rigidity, can also play a significant role in a particle phagocytic predisposition. As far as interaction with the cell membrane is concerned, macrophages tend to show a strong preference for rigid particles. One study showed that soft polyacrylamide particles were unable to stimulate the assembly of actin filaments required for the proper formation phagosomes, as opposed to rigid particles which had the same total polymer mass and surface properties [133]. Macrophage’s increased sensitivity to bead stiffness can be connected to the fact that bacteria and other pathogens have cell walls usually more rigid than the surrounding tissue they invade. On the other hand, particle rigidity can have an opposite effect on opsonization. Rigid liposomal membranes, composed of cholesterol and saturated phospholipids with a high melting point, are known to decrease complement activation and hence decrease phagocytosis [134]. Similarly, core-shell nanoparticles having a rigid polystyrene core were significantly less prone to uptake by MPS than NPs made of a more flexible core, based on fluid-like poly(methyl acrylate) PMA [6]. Particles with low elastic modulus are thought to provide a greater number of surface interactions with the biological environment. Though the research in this area is ongoing, there is not a well-defined relationship between particle rigidity and phagocytosis as of yet.

Clearance by phagocytes, a high blood perfusion and discontinuous sinusoidal endothelium are features that contribute to a general tendency for non-modified NPs to accumulate in the liver. Early work by Fidler et al. [135] reported clearance of liposomes to the liver, whilst Illum and Davis [136] showed a short circulatory half-life and liver accumulation in rabbits of non-modified polystyrene particles which showed the biodistribution could be changed with poloxamer 388 functionalization. This rapid clearance can be avoided by adding poly(ethylene)glycol (PEG) to the surface of NPs. Dunn et al. [136] showed removal within 1 h and sequestration of 100 nm polystyrene particles in the resident Kupffer cells of the liver in rats that could be overcome by particle PEGylation. The prevention of opsonisation with the addition of PEG drastically increases the blood half-life of all nanomaterials regardless of surface charge. Generally, the blood half-life of gold NPs is also increased by increasing the length of PEG, which causes the protective layer to thicken [137]. The synthesis of these long circulating “Stealth” NPs improves accumulation in target tissue. In addition to PEGylation of NP surfaces, blood half-life also depends on a NP’s shape and size and surface chemistry. For example, rod-shaped micelles have a circulation lifetime ten times longer than that of spherical micelles [138].

8.3.3 *Particle Dynamics in the Blood Stream*

NPs encounter numerous physical and biological barriers created by normal and abnormal physiology en route to their target site. Regardless of the targeting strategy, it is desirable for blood-borne particles to navigate within the vasculature, identify the abnormal blood vessels and there, either adhere firmly to the walls withstanding the dislodging hydrodynamic forces (vascular targeting) or extravasate, leaving the vessels through inter-endothelial openings, and seek the diseased cells within the surrounding tissue. Interestingly, and perhaps not surprisingly, this is the strategy used by circulating cells such as leukocytes, to marginate and adhere to the inflamed endothelium as well as platelets which are also mostly found in close proximity to the vessel walls, where they can periodically sense and adhere to sites of vascular injury. This characteristic platelet behaviour is ascribed to their small size ($3\times$ smaller than RBCs) and quasi-discoidal shape [139, 140]. Conversely, healthy red blood cells (RBCs) can continuously navigate the circulatory system for several weeks while delivering oxygen to the tissues. The size, shape, and physiochemical properties of the RBC membrane enable these cells to avoid entrapment within the complex microvasculature and, in larger vessels, to repel from the endothelial walls and preferentially stay within the vessel center [141, 142]. Despite these intricate design strategies exhibited by nature, limited investigations have focused on predicting the transport behaviour of synthetic particles with different sizes and shapes within the vascular system. Such a feature can be exploited in the rational design of particle-based intravascular and pulmonary delivery systems. Ferrari and Decuzzi [143] presented a general model to predict the dynamics of nano-/micro-particles immersed in a linear laminar flow. They showed that non-spherical particles under the simultaneous action of inertial and hydrodynamic forces marginate and, therefore, increase the interaction with the wall surface. Among the shapes considered, discoidal particles with low aspect ratio exhibit the largest propensity to marginate. The effect of shape has also been shown to play a critical role in the margination of NPs [143, 144]. It has also been predicted [14] that particle shape can dramatically affect firm adhesion under flow, proposing the use of oblate spheroidal rather than classical spherical particles, as oblate-shaped particles are subjected to torques resulting in tumbling and rotation [143, 145]. These complex dynamics of non-spherical particles cause translational as well as rotational motions. These theoretical predictions have been supported by both in vitro experiments with silica particles [144] and by in vivo studies with silicon discoidal particles [146] and discoidal polystyrene particles targeted against ICAM-1 molecules, that both exhibited higher tumor accumulation/targeting specificity compared to spherical particles [147]. Furthermore, the particle density also has a strong influence on margination, as generally, lighter particles marginate further than denser counterparts [148].

8.3.4 Biodistribution

For intravenously administered NPs, diameter is an important determinant of pharmacokinetics and biodistribution owing to the variable size of fenestrations lining blood vessels. Hydrophilic NPs with diameters smaller than 6 nm, can be eliminated from the body as they can be excreted by the kidneys [149]. However, the majority of injected nanomaterials are larger in size and, thus, accumulate in the organs of the MPS, namely liver, spleen and lungs. For instance, Godin et al. [146] described quantitative analysis of organ distribution of different discoidal particle sizes as compared to spherical particles of the same diameter (Fig. 8.4). They found that nanovectors with diameter of 600 nm accumulated more prominently in the liver, but to a lesser extent in the lungs when compared to their 1000 nm in diameter counterparts ($p < 0.05$). For 1700 nm particles the amount of Si increased dramatically in the lungs, liver and spleen, the MPS organs, as well as the heart [150]. A total of 66.4 %, 63.1 % and 68.2 % ID were recovered in the above five major organs and tumor for 600 nm \times 400 nm, 1000 nm \times 400 nm, and 1700 nm \times 400 nm in diameter nanovectors after 4 h. They attributed these numbers to a partial degradation of the discoidal Si particles leading to elevated levels of Si in kidneys without

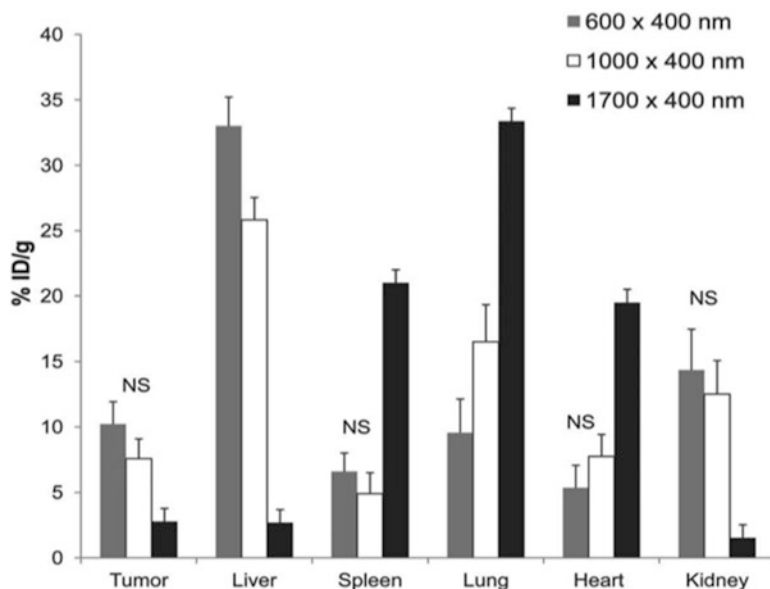


Fig. 8.4 Biodistribution of 600, 1000, and 1700 nm in diameter and 400 nm in thickness discoidal silicon particles 4 h following intravenous administration at the dose of 7.5–9 mg/kg to the tail vein of 4T1 orthotopic breast tumor bearing mice. Silicon analysis normalized to basal silicon levels in the tissues as evaluated by Inductive Coupled Plasma Atomic Emission Spectroscopy (ICP-AES). The data is presented as % of injected dose/g organ. NS, $p < 0.05$; NS— not significant [146]

apparent presence of the nanovectors. Another possibility is that a distribution of the nanovectors to other organs/systems such as gastrointestinal tract and pancreas (through entero-hepatic circulation), and lymphatic system occurred [146].

However, the long term fate of non-degradable nanomaterial that are too large to be excreted by the kidneys, remain in the tissues inside the body for up to 8 months [151] and may cause toxicity. Resident macrophages in MPS organs will phagocytose NPs, degrade a small fraction of them, and eventually exocytose both the degraded and intact NPs [152]. The long-term distribution and toxicity (>1 year) of non-degradable particles have not been evaluated, raising concern about toxicity. Cadmium-Selenium (CdSe) quantum dots exhibit hepatotoxicity in cell cultures but do not appear to be toxic in Sprague-Dawley rats on the basis of histopathology as well as liver and kidney marker analysis. In addition, pristine and PEGylated single-walled carbon nanotubes, persisted within liver and spleen macrophages for 4 months without apparent toxicity in rats and rabbits despite their reported toxicity within in vitro culture models. The differential toxicity may be related to a localized concentration of NPs [153]. In vivo, NPs seem to be transported from organ to organ, whereas in vitro, the NPs are confined within a limited cell population. Therefore, the differential toxicity between in vivo and in vitro may be due to exposure concentration.

A study of the biodistribution and pharmacokinetic study of [125I]-labeled non-targeted, cylindrical particles prepared by the PRINT[®] process in the 200 nm size range in healthy mice showed the expected uptake primarily in the liver and spleen [40, 41]. The conventional strategies to reduce the rapid clearance from the bloodstream and uptake by the liver and spleen have been to increase hydrophilicity of the particle surface and reduce particle size. In an alternative approach, however, Geng and coworkers have compared soft spherical assemblies with flexible filaments and found that the in vivo circulation time for non-spherical filomicelles was about ten times longer than the analogous spherical counterparts [138]. They extended their study to the delivery of paclitaxel and showed significant tumor shrinkage in a xenograft mouse tumor model, and showed that an increase in the filomicelle length had the same relative therapeutic effect as a similar increase in the paclitaxel dosage. These results show that, in applications where a prolonged circulation time is desired, a long, worm-like structure can be more effective than a sphere.

Though particle size has long been considered dominant in determining in vivo behaviours such as circulation time and biodistribution profile, it stands to reason that physical filtration barriers in the body could be navigated by larger, but more deformable particles [154, 155]. Deformable particles that resembled RBCs in size and shape have been shown to deform in restricted channels [156] or capillaries [112] that were smaller than the particle diameter, though the modulus of these particles was not characterized [156] or was poorly matched to RBCs [112] and was restricted to in vitro testing in both cases. In a simulation of renal filtration of soft particles, microgels translocated through pores that were 1/10th of the particle diameter under physiologically relevant pressures [157]. Merkel et al. [158] recently reported the fabrication of a series of discoidal, monodisperse, low-modulus hydrogel PRINT particles with diameters ranging from 0.8 to

8.9 μm , spanning sizes smaller than and larger than RBCs, which were injected into healthy mice. They tracked their concentration in the blood and their distribution into major organs and found that the particles demonstrated some hold up in filtration tissues like the lungs and spleen, followed by release back into the circulation, characterized by decreases in particles in these tissues with concomitant increases in particle concentration in blood. They concluded that particles similar to RBC in size demonstrated longer circulation times, suggesting that this size and shape of deformable particle is uniquely suited to avoid clearance. Another study [138] found filamentous worm-like micelles to have increased circulation times with respect to their spherical counterparts, though this effect was attributed to the size and shape of these particles as well as the deformability. Inspired by nature's example of long-circulating microparticles, Merkel et al., sought to mimic the size and shape of RBCs while varying the modulus across a physiologically relevant range to probe the physical barriers encountered in vivo by soft microparticles. They showed that increasing the deformability of RBC-sized particles increased their circulation times beyond that of conventional microparticles and significantly altered their biodistributions. As a consequence of their low modulus, these discoid microparticles bypassed several in vivo filtration mechanisms, illustrated by animal survival and dramatic increases in elimination half-lives of particles with decreasing modulus [95]. Such deformable and long-circulating particles may find utility in the fields of drug delivery and medical imaging, where long-circulation times and varied biodistributions are often desirable characteristics [159–161] and, hence, particle deformability must be considered as a vital design parameter during the fabrication of therapeutic or diagnostic particles.

8.3.5 *Tumor Accumulation*

Passive accumulation and physiological effects have been exploited for disease treatment in preclinical models for inflammation [162] and cancer [163] without surface modification or active targeting by ligand attachment. This may reflect the neovasculature associated with these conditions and the suggested enhanced permeability and retention effect (EPR) in tumors in which nanoscale particles migrate and accumulate across leaky vasculature in disease tissue. NPs can accumulate in tumors and be used to deliver therapeutic compounds and contrast agents for imaging. Tumors possess large fenestrations between the endothelial cells of blood vessels produced by angiogenesis and can retain particulates found in the blood. EPR effect allows NPs to accumulate inside the tumor if they are not cleared by the liver or spleen or excreted through the kidney. Longer circulation time in the bloodstream allows repeated passes of nanomaterials through tumor blood vasculatures and favors passive tumor targeting via an EPR effect [164]. To produce long-circulating NPs that can accumulate inside tumor tissues a diameter between 30 nm and 200 nm is desired [165]. Using this approach, researchers may induce

passive accumulation of nanomaterials inside a tumor. In fact, it is possible to control the overall accumulation and penetration depth into the tumor by changing the NPs diameter [137] and shape [166]. Researchers determined that the NPs capacity to navigate between the tumor interstitium after extravasation increased with decreasing size. By contrast, larger NPs (100 nm) do not extravasate far beyond the blood vessel because they remain trapped in the extracellular matrix between cells. Thus, the smallest NPs (20 nm) penetrate deep into the tumor tissue but are not retained beyond 24 h.

Surprisingly, adding a targeting moiety on the surface of the NPs does not appear to increase accumulation inside the tumor or change biodistribution. Active targeting of NPs changes the intra-tissue localization of NPs and their increased internalization into cancer cells [167, 168]. However, some studies have shown active targeting increases tumor accumulation of NPs. For instance, Godin et al. [22] discuss how their MSV particle is able to release ‘Stage 2’ NP (S2NP), such as cytotoxic agents, contrast enhancing imaging agents, bioactives or metal nanoparticles after the S1MP has adhered to the diseased vasculature. In the case of metal S2NP (e.g. gold, silver and iron oxide), the activation of the system can be triggered with an external energy source, such as radio-frequency [169] or near-infrared [170] energy. Another possible mode of action includes targeting specific cell elements, such as macrophages and myofibroblasts, in the disease location to achieve preferential localization. The intracellular payload can then be released to destinations influenced by S1MP and S2NP surface chemistry. However, owing to the many contradictory findings in the literature, it is not certain how active targeting affects NP accumulation in the target tissue [168], however, Frieboes et al. [171] recently presented a computational model which predicts vascular accumulation of blood-borne NPs. The fraction of injected NPs depositing within the diseased vasculature and their spatial distribution was computed as a function of tumor stage, from 0 to 24 days post-tumor inception. They explain that NP vascular affinity, interpreted as the likelihood for a NP in the blood to firmly adhere to the vessel walls, is a fundamental parameter and depends on NP size and ligand density, as well as vascular receptor expression. They go on to show that for high vascular affinities, NPs tend to accumulate mostly at the inlet tumor vessels leaving the inner and outer vasculature depleted of NPs, whereas, for low vascular affinities, NPs distribute quite uniformly intra-tumorally but exhibit low accumulation doses.

Unfortunately, tumor accumulation generally represents a relatively small fraction of total injected dose, due to a number of factors, such as high intra-tumoral pressures and the architecture of vasculature networks. For instance Frieboes et al. [171] have recently demonstrated that for insufficiently developed vascular networks, such as those seen within tumors, NPs are transported preferentially through the healthy, pre-existing vessels, thus, bypassing the tumor mass. However, a few recent studies with discoidal particles have shown enhanced accumulation in the tumor based on particle geometry [146, 166]. For instance, a high accumulation of discoidal porous silicon (pSi) nanovectors into orthotopically grown breast tumors in mice has been recently reported [146]. For 600 nm × 400 nm pSi particles, 10.2 % ID/g was found in the tumor, which is five times higher than their spherical

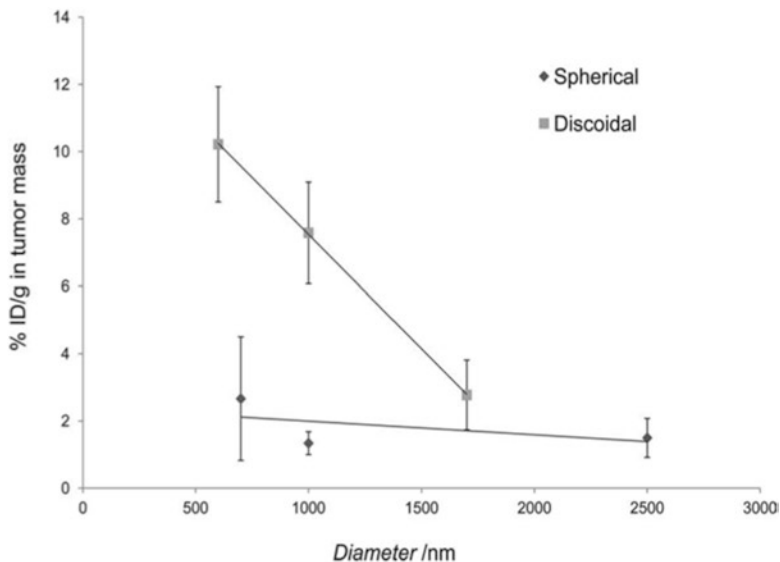


Fig. 8.5 Effect of particle diameter on tumor accumulation of discoidal particles as compared to spherical particles. For discoidal particles, the height is 400 nm [146]

counterparts (Fig. 8.5). Their results were achieved without the use of any active targeting strategy, and, therefore, emphasized the effect of geometry on in vivo behaviour of therapeutic particles.

In an additional study Van de Ven et al. [166] used intravital microscopy and elemental analysis to compare the in vivo localization of particles with different geometries and demonstrated that discoidal (plateloid) particles preferentially accumulate within the tumor vasculature at unprecedented levels, independent of the EPR effect. They found that in melanoma-bearing mice, 1000×400 nm plateloid particles adhered to the tumor vasculature at about 5 % and 10 % of the injected dose per gram organ (ID/g) for untargeted and RGD-targeted particles respectively, and exhibited the highest tumor-to-liver accumulation ratios (0.22 and 0.35), however smaller and larger plateloid particles, as well as cylindroid particles, were more extensively sequestered by the liver, spleen, and lungs. The authors concluded that plateloid particles appeared well-suited for taking advantage of hydrodynamic forces and interfacial interactions required for efficient tumoritropic accumulation, even without using specific targeting ligands [166].

Improvements in fabrication protocols have allowed to create a potential toolbox for increased targeting efficacy. For instance, in vitro and in vivo studies have demonstrated that liposomal carriers slightly improved the delivery of siRNA to melanoma, lung cancer, breast cancer, and ovarian cancer [172, 173]. The efficacy of hemispherical 1.6 μm pSi particles loaded with neutral dioleoyl phosphatidylcholine nanoliposomes (DOPC) containing EphA2-specific siRNA was evaluated by Tanaka and colleagues in two independent orthotopic mouse models of ovarian cancer [174]. EphA2 is an oncoprotein overexpressed in most malignancies

including ovarian tumors. After a single treatment with EphA2-targeted-MSV and without simultaneous chemotherapy, gene silencing and decrease in tumor burden was observed, evaluated by cell proliferation and angiogenesis. A dose twice as high and administered twice a week for 3 weeks was required to achieve a similar effect with siRNA-DOPC, alone. The mechanism of the efficient and sustained liposomal siRNA delivery was due to surface modification, tissue distribution, and slow biodegradation of the pSi particles (S1MP). S1MP not only served as storage for liposomal siRNA, but also shielded siRNA oligos extensive and prompt degradation by nucleases in the bloodstream. This novel approach enables new avenues in personalization of siRNA therapeutics through controlled delivery of synergistic payloads in a time-dependent manner.

8.4 Interactions with Cells

8.4.1 *Binding to the Cell Membrane*

Once the theranostic particle has reached the target cell, it faces the plasma membrane, which is the first barrier for the intracellular delivery of drug or imaging payloads. NPs were initially considered as benign carriers, but multiple studies have demonstrated that NPs' interaction with serum proteins and cell membrane receptors is determined by their design, in effect, influencing cell uptake, gene expression, and toxicity. Interactions between NP-bound ligands and cellular receptor depend on the engineered geometry, NP arrival rate, surface charge and the ligand density of a nano or micro-material (Fig. 8.6).

The particle size and shape dictates the number of ligands that interact with the receptor target. A multivalent effect occurs when multiple ligands on the NPs interact with the multiple receptors on the cells. The binding strength of complexed ligands is more than the sum of the individual affinities and is measured as the avidity for the whole complex. Similarly, on NP surfaces, the presence of ligands at a given density over a specific curvature will contribute to the overall avidity of the NP-linked ligands for available receptors. To illustrate this, the binding affinity of Herceptin to the ErbB2 receptor is 10^{-10} M in solution, 5.5×10^{-12} M on a 10 nm NP, and 1.5×10^{-13} M on a 70 nm NP [175]. This example shows how a ligand's binding affinity increases proportionally to the size of a NP owing to a higher protein density on the NP surface. However, when viewed in terms of the downstream signalling via the ErbB2 receptor, 40–50 nm gold NPs induced the strongest effect, suggesting other factors beyond binding affinity must play a role. Nevertheless, several studies have shown that NP design can cause differential cell signalling when compared with free ligand in solution. For example, the previously mentioned 40–50 nm Herceptin-coated gold NPs altered cellular apoptosis by influencing the activation of caspase enzymes [175]. Similarly, receptor-specific peptides improved their ability to induce angiogenesis

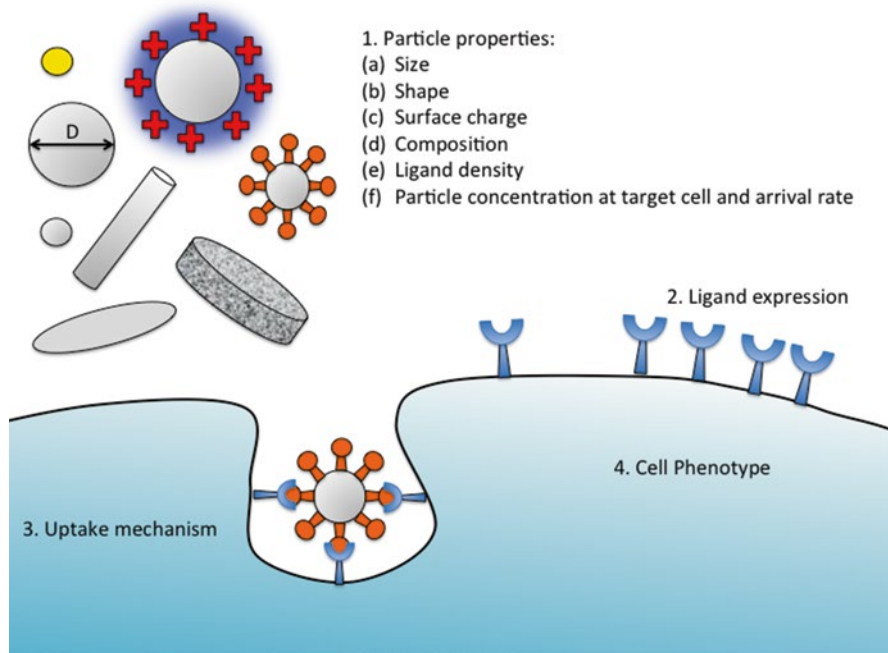


Fig. 8.6 Schematic presentation of the effect of physicochemical features of the particles on cellular uptake. Size, shape, surface charge, composition, ligand density and arrival rate influences a particles interaction with the surface of a specific cell

when conjugated to a NPs surface [176]. These findings highlight the advantage of having a ligand bound to a NP as opposed to it being free in solution. The NP surface creates a region of highly concentrated ligand, which increases avidity and, potentially, alters cell signalling.

Despite their advantages, a general concern with NP-ligand complexes is the potential denaturation of proteins when bound to the engineered NP surface. The denaturation of a protein can affect binding to its receptor, increase nonspecific interactions or provoke inflammation. For instance, when lysosome is bound to gold NPs it denatures and interacts with other lysozyme molecules and induces the formation of protein-NP aggregates [177]. Fibrinogen also unfolds when bound to the surface of polyacrylic acid-coated gold NPs, which means it is able to bind to the integrin receptor Mac-I which leads to an inflammatory response [178].

Surface charge is also important in dictating membrane interaction and subsequent cellular fate. Compared with NPs with a neutral or negative charge, positively charged NPs are taken up at a faster rate [179, 180]. It has been suggested that the cell membrane possesses a slight negative charge and cell uptake is driven by electrostatic interactions [181, 182]. A recent study demonstrated that this electrostatic attraction between the cell membrane and positively charged NPs favours adhesion onto a cell's surface, leading to uptake. Negatively charged heparan sulfate proteoglycans and integrins play a role in the cellular binding of positively charged parti-

cles. For example, Mislick and Baldeschweiler showed that the binding of cationic particles to the cell membrane was significantly reduced in proteoglycan-deficient mutant cells [122]. Even so, heparan sulfate proteoglycans may induce non-specific binding but the exact role they play in cellular uptake is not clear, and also non-adherent cells do not contain such molecules, which seems to result in lower uptake [183]. The clustering of trans-membrane proteins and syndecans at the plasma membrane during binding to cationic particles may stimulate interaction with the actin cytoskeleton and result in the formation of tension fibers. This tension provides the energy required for internalization to take place [184].

For small NPs (2 nm), a positive charge can perturb the cell membrane potential, causing Ca^{2+} influx into cells and the inhibition of cell proliferation [185]. For larger NPs (4–20 nm), surface charge induces the reconstruction of lipid bilayers [186]. It has been reported that NPs conjugated with polycationic molecules interact with fluid-phase domains of lipid bilayers [187] and can induce the development of plasma membrane pores leading to membrane disruption. Binding of negatively charged NPs to a lipid bilayer causes local gelation, whereas binding positively charged NPs induces fluidity. Several studies have confirmed the pivotal role surface charge plays in downstream biological responses to NPs.

Additionally, Particle-membrane interactions can be altered by the development of a protein corona which is caused by particle interactions with serum proteins such as lipoproteins, immunoglobulin, complement and coagulation factors, acute phase proteins and metal-binding and binding proteins [188] during circulation in the blood. This is particularly relevant for NPs with high surface charge density. As an example, Dawson and colleagues have demonstrated that positively charged polystyrene particles adsorbed a layer of plasma proteins changing the biological fate of the particles and treated cells. Namely, the corona enabled the protection of the cells from the damage induced by the highly positive NP surface, thus, significantly altering any particle-cell interaction [189].

Therefore, the measurement of surface charge plays a key role in the design of novel theranostic NP approaches. The Zeta-potential (ζ) is one of the most important micro or nano scale parameters in the interaction between particle and cell. Whilst ζ is relatively straightforward to measure, its relation to the structure and charge content of an agent can be complex and so care must be taken when interpreting ζ readings. Zeta potential is an energy potential that arises due to electrical charge on the particle. In particular the particle charge produces a distance dependent potential, ϕ that is described in 1-D by the Poisson equation:

$$\frac{d^2\phi}{dx^2} = -\frac{\rho}{\varepsilon} \quad (8.1)$$

where ρ is the charge density on the particle and ε the permittivity of the medium. Solution of the Poisson equation describes a potential, whose value is dependent upon the charge density on the particle and which decays exponentially with increasing distance, x from the particle. In the majority of practical situations there is further complexity due to interactions of ions within the local environment with

the particle surface charge. Ions of opposite charge to that of the particle will firmly bind due to electrostatic attraction to form a layer known as the Stern layer. Beyond this there are further electrostatic interactions between ions and particle but these are mediated by the presence of the bound ions within the Stern layer. It is the electric potential at the outer edge of the Stern layer that is defined as the ζ -potential. These interactions of solvent ions with particle make the ζ value dependent upon the properties of the medium (and hence experiment specific) as well as those of the particle.

ζ is usually measured indirectly via the electrophoretic mobility, μ , which quantifies the motion of the particle due to forces induced by an applied electric field. In practice a voltage is applied and particle motion measured, usually by optical means. The value of ζ is then given by:

$$\xi = \frac{\mu n}{\epsilon_0 \epsilon_r} \quad (8.2)$$

where η is the viscosity of the medium.

Often ζ is used as a ready 'look up' value to assess the effects of particle charge, however this can be misleading. As we have already seen the medium is intrinsically linked to the ζ value and so realistic conditions must be used, e.g. in in-vitro studies measurements in water will not accurately reflect the value pertaining to exposure within a cell culture medium containing a multitude of charged molecules. The size of the charged particle must also be carefully considered. Equation 1 shows that the value of ζ is dependent upon charge density and so different sized particles with the same ζ value will have different total charge. This may be important if considering effects where the sum charge amount is the determining factor. For example, if cationic particles of 2 nm and 10 nm diameter and the same ζ are used to create a proton sponge effect for endosomal escape, very different results should be expected as the total charge of the 10 nm particles is 25 times greater than that of the 2 nm.

The interaction of charged particles is known to be size dependent and so again the zeta potential is not the only parameter of importance. Consider the attraction energy per unit area due to Van der Waal's forces, for two identical spheres of radius, r and charge density, ρ_s this is:

$$W = -\frac{\pi^2 C \rho_s^2}{12d} \cdot r \quad (8.3)$$

where d is the separation distance of the spheres and C is a constant of proportionality. Thus the strength of interaction is determined by r , irrespective of the charge (and hence ζ) properties of the particle. This is true for a range of geometries, for instance consider the case for a sphere interacting with a charged plate, i.e. a physical model akin to membrane-particle interaction. Here the attraction energy is again dependent upon r and given by:

$$W = -\frac{\pi^2 C \rho_s \rho_p}{6d} \cdot r \quad (8.4)$$

In summary, ζ is a useful, and hence widely used, parameter that characterizes the electrical potential of charged particles. However care must be used when using ζ measurements in studies involving multiple particle types for they relate to charge *density* on the surface, i.e. particle size is factored out through this *per unit area* measure. The particle radius will still determine the total charge and the spatial arrangement of that charge in respect to structural components of the cell. Thus different sized particles may exhibit very different effects on cells despite having similar ζ values.

A complete understanding the measurement of ζ will foster a more a comprehensive understanding of the initial interaction between the particle and the cell membrane and will enable us to reduce cytotoxic effects associated with cationic polymers and simultaneously increase therapeutic efficacy. Furthermore, it is generally assumed that the initial interaction of the particle with the cell membrane is closely linked with the uptake pathway and subsequent intra-cellular trafficking and, hence, its therapeutic efficacy within the cell. Therefore, a promising strategy for increasing the efficiency of a drug or gene delivery vector is to target specific internalization pathways that improve the intra-cellular fate of the particle vectors.

8.4.2 Cellular Uptake

The plasma membrane presents a substantial barrier to particles successfully gaining entry to the cell, as it is a dynamic and a relatively lipophilic structure that restricts the admittance of large, hydrophilic, or charged molecules. As is well documented, NPs can be transported into cells through a process called endocytosis [190]. In particular, endocytosis occurs in all cell types and is mediated by at least four basic mechanisms: macropinocytosis (Fig. 8.7), clathrin-mediated endocytosis, caveolae-mediated endocytosis, and clathrin–caveolae and dynamin-independent endocytosis [191] (Fig. 8.8).

It is generally accepted that systems which are less than 200 nm in size [8, 192] and that have not been functionalized with cell surface receptors, such as transferrin or low-density lipoprotein receptors for specific binding [193–196], are internalized via clathrin-mediated endocytosis (CME) [8, 192]. CME is considered kinetically the most effective [197, 198] endocytic uptake pathway for various particles. However, a limitation of utilizing this pathway for theranostic delivery is that the internalized therapeutic drug or imagine contrast agent is typically destined for acidic and/or enzymatic degradation in the late endosome or lysosome in the final stages of the process [199, 200]. Thus, when nanovectors are taken up via CME, it is advantageous for them to be used in conjunction with a polymer material, such as PEI [201], polyamidoamine (PAMAM) dendrimers [201] or an imidazole-containing

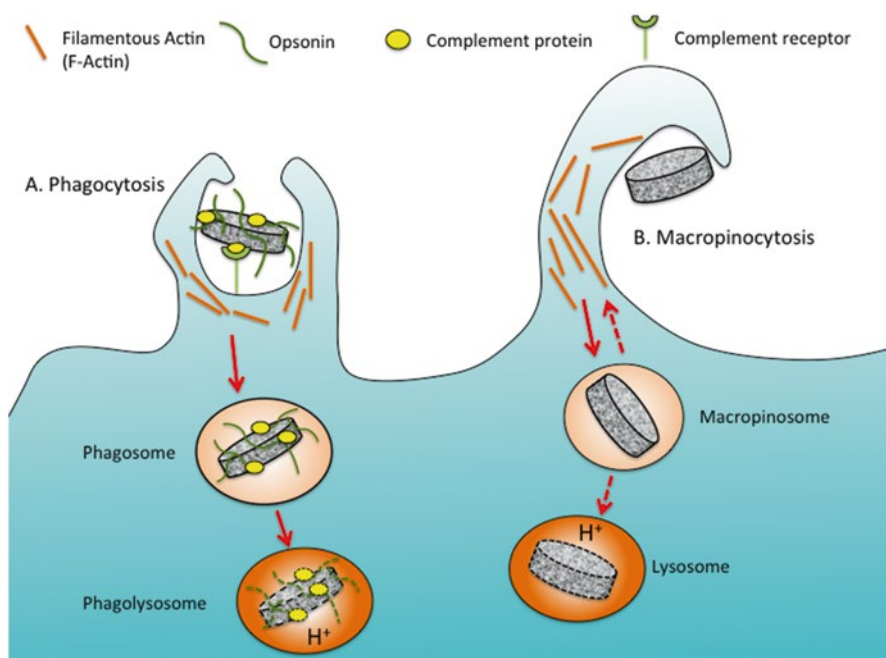


Fig. 8.7 Intracellular uptake of particles. (a) Phagocytosis is an actin-based mechanism occurring primarily in cells associated with Mononuclear Phagocytic System (MPS), such as macrophages, and is closely associated with opsonisation of complement proteins and other opsonins. However, fibroblasts, epithelial and endothelial cells may also display some lower phagocytic activity [235]. (b) Macropinocytosis is an actin-based pathway, engulfing NPs and other biological constituents with poor selectivity

polymers [202], that are capable of inducing endosomal escape of the polyplex into the cytoplasm of the cell before significant degradation occurs. Other desirable internalization pathways include macropinocytosis [201, 203], micropinocytosis [203] and caveolae-mediated endocytosis. Caveolae-mediated endocytosis maybe an advantageous internalization mechanism for gene transfection strategies as the vesicles that result from the internalization in this pathway do not develop into lysosomes and, therefore, avoid significant degradation [204].

The endocytic process occurs through either specific or nonspecific cellular uptake depending on the surface properties of NPs. In many cases, NPs enter the cell after binding to the receptor target. Once bound, the internalization of particle systems can vary greatly depending on a number of factors such as particle chemistry, cell type, cell polarization state and stage of cell cycle [205]. For instance, a NPs shape can directly influence uptake into cells, Gratton et al. [15] determined that rods were taken up the most, followed by spheres, cylinders, and cubes in synthesized NPs larger than 100 nm. In studies with sub-100 nm particles, spheres show an appreciable advantage over rods [206, 207]. In fact, at this size range, increasing the aspect ratio of nanorods seems to decrease total cell uptake. The effects of shape

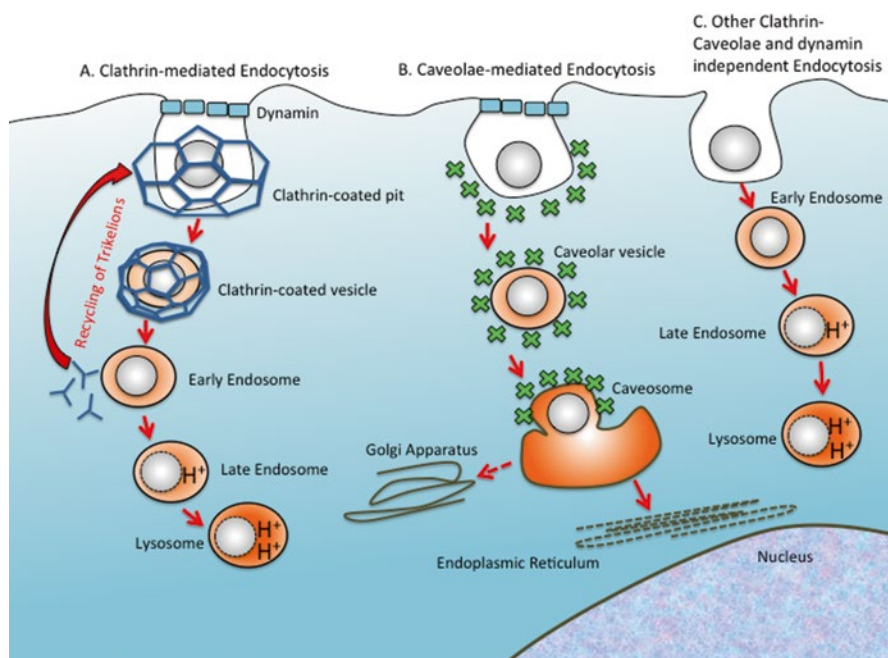


Fig. 8.8 Intracellular trafficking of nanovectors following internalization via clathrin-mediated endocytosis, caveolae-mediated endocytosis and other clathrin-caveolae and dynamin independent endocytosis. **(a)** Clathrin mediated endocytosis. The assembly of clathrin triskelions into a polygonal matrix helps deform the overlying plasma membrane to form a clathrin coated pit. After assembly of the cup-like clathrin matrix, dynamin is employed at the neck of the pit to mediate membranal fission. This leads to the release of the clathrin-coated vesicle to the cytoplasm. The following un-coating of the vesicle allows clathrin triskelion recycling. Clathrin-coated vesicles then form early endosomes, which are acidified and fuse with pre-lysosomal vesicles containing enzymes to give rise to late endosomes and eventually lysosomes. This pathway leads to substantial nanovector and drug degradation mediated by the acidic environment and enzyme action. **(b)** Caveolae-mediated endocytosis of a nanocarrier gives rise to a caveolar vesicle that is delivered to a caveosome, which does not have a degradative acidic and enzyme-rich environment

and size of metallic particles on cellular internalization were also reported by Xu et al., using layered double hydroxide Mg₆Al₂ nanoparticles with two distinct particle morphologies: (1) rods that were 30–60 nm in width and 100–200 nm in length, and (2) hexagonal sheets that were 50–150 nm wide (laterally) and 10–20 nm thick [208, 209]. The group found that both shapes are quickly taken up by CHO-K1, NIH 3T3, and HEK 293T cell and that rod-like particles specifically target the nucleus while the sheet-like particles are retained in the cytoplasm [208, 209]. In a study focusing on the uptake rates of length-fractionated single-walled carbon nanotubes (SWNT) 130–660 nm in diameter, it was suggested that nanoparticulate aggregates on the cell membrane form a cluster which is sufficient in size to generate a large enough enthalpic contribution for overcoming the elastic and entropic energy barriers associated with membrane vesicle formation. Endocytosis rate for

nanotubes was 1000 times higher than for spherical gold nanoparticles, while similar exocytosis kinetics was measured for poly(D,L-lactide-co-glycolide), SWNT, and Au nanoparticles across distinct cells [182]. Furthermore, Calderera-Moore et al. [210] focused on the dynamic manipulation in particles geometry as a tool to precisely control particle-cell interactions [210]. As an example, Yoo and Mitragotri [211] have designed polymeric particles able to switch shape in a stimulus-responsive manner. The shape-switching behavior was a result of a fine balance between polymer viscosity and interfacial tension and could be tuned based on the external stimuli (temperature, pH, or chemicals in the medium), enabling the modulation of phagocytosis of elliptical particles that were previously not internalized by the cells [211]. This indicates that non-spherical particle-cell interactions may be much more complex. For instance, ligand coated rod shaped NPs may present to the cell with two different orientations. Compared with the short axis, the long axis will interact with many more cell surface receptors [212] (Fig. 8.9). For spiky nanostructures such as gold nanourchins, whether the ligand is located on or between the spikes affects how it is presented to the target cell receptors [213].

Serda et al. have shown that discoidal SIMP vectors are internalized by endothelial cells through a combination of phagocytosis and macropinocytosis [214, 215] (Fig. 8.6). Both mechanisms are actin-driven processes that involve extensive membrane reorganization with the formation of pseudopodia and extension of the cell membrane to surround and actively engulf the particles.

Within a given geometric shape, a nanomaterial's dimensions are a strong determinant of total cell uptake. For gold spherical NPs, silica particles, single walled carbon nanotubes, and quantum dots, a 50 nm diameter is optimal to maximize the rate of uptake and intracellular concentration in certain mammalian cells [182, 212, 213]. In addition to size and shape, the composition of the nanomaterials also affects the uptake as single-walled carbon nanotubes and gold NPs, each 50 nm in diameter, possess endocytosis rates of 10^{-3} min^{-1} and 10^{-6} min^{-1} , respectively. This 1000-fold difference may be due to the intrinsic properties of carbon versus gold. Which ligand is used to coat the NP will also affect the downstream biological responses. For example, the uptake and cytotoxicity of NPs were significantly altered when the NPs were coated with two different proteins targeting the same receptor [181].

If a NP is bound to a receptor on the cell membrane then it will most likely enter the cell via receptor-mediated endocytosis [175, 207, 216]. The binding of the NPs-ligand conjugate to the receptor produces a localized decrease in Gibbs free energy, which induces the membrane to wrap around the NP to form a closed vesicle structure [217]. The vesicle eventually buds off the membrane and fuses with other vesicles to form endosomes, which fuse with lysosomes where degradation occurs. The size dependent uptake of NPs is likely related to the membrane wrapping processes. Small NPs coated with 50 kDa proteins may interact with only one or two cell receptors. By contrast, a 100 nm NP has many more ligand-receptor interactions per particle. Several small-ligand coated NPs must bind to receptors in close proximity to one another to produce enough free energy to drive membrane wrapping. Larger NPs can act as a cross-linking agent to cluster receptors and induce uptake. Thermodynamically, a 40–50 nm NP is capable of recruiting and binding enough

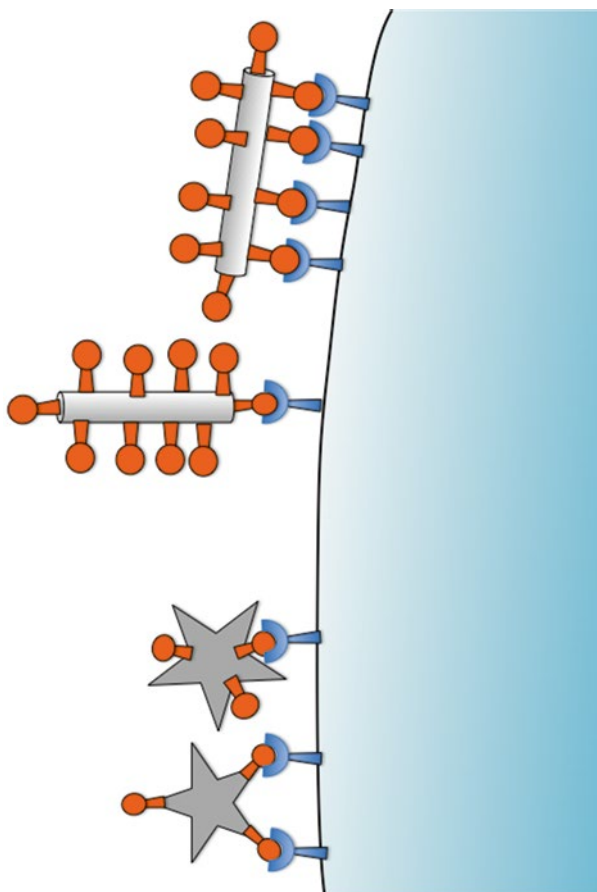


Fig. 8.9 Geometry of the nano- and micro-particles has a dramatic impact on the presentation of the NP at the cell interface, which affects cell-membrane interaction, uptake and long-term fate of the particle

receptors to successfully produce membrane wrapping. Above, 50 nm, NPs bind such a large number of receptors that uptake is limited by the redistribution of receptors on the cells surface via diffusion to compensate for local depletion. NPs larger than 50 nm bind with high affinity to a great number of receptors and may limit binding of additional particles. Mathematically modeling this phenomenon has demonstrated that optimal endocytosis occurs when there is no ligand shortage on the NP, and no localized receptor shortage on the cell surface [218], which occurs in NPs 30–50 nm in diameter where ligand density is optimal.

To conclude this sub-section, the design of smart functional nanosystems for intracellular imaging and therapeutic applications requires a thorough understanding of the mechanisms by which NPs enter cells. For the engineering process, asymmetrical NPs may provide another level of control in presenting ligands to the target receptors. In biological and clinical applications, the ability to control and manipulate the accumulation of NPs inside a cell by specific cellular uptake makes it possible to improve diagnostic sensitivity and therapeutic efficiency [219]. However, there is still limited information regarding the effects of particle geometry on uptake mechanisms and downstream cellular responses, such as cell proliferation, apoptosis, adhesion, migration and cytoskeleton formation. Therefore, investigating how the uptake of different shaped particles affects cell functions will assist in the design of nanoscale delivery systems and open up new NP bio-applications. Additionally, most studies elucidating the effect of NP diameter on uptake have been conducted primarily on immortalized cell lines. Furthermore, cells under culture conditions are constantly dividing, however, this may not be the case in vivo and could potentially be a further limiting factor for efficient delivery of therapeutic and diagnostic nanovectors. Therefore, the nuclear envelope cannot be neglected within in vivo situations. Optimal NP uptake size may depend on the cell being assayed because each cell type possesses a unique phenotype. Thus, it will be necessary to expand NP studies to include both immortalized and primary cells in different culture configurations to identify broad-scope design parameters for optimal uptake and accumulation in cells. Furthermore, it is not unreasonable to generalize that if nanovectors could be specifically modified to suit a particular internalization pathway, such as caveolae-mediated endocytosis, more improved efficiencies in the endosome escape processes regardless of the particle type could be achieved.

In the past decade, cell interaction studies of diverse variety of hollow microcapsules show that the nature of the capsule wall influences the shape, elasticity, deformability and thickness of the cell wall, which in turn affects the cell recognition and uptake. Examples of some of the factors influencing microcapsule-cell interaction are depicted here.

Spherical poly(methacrylic acid) capsules were internalized by HeLa cells at a faster rate than rod-shaped capsules with a higher aspect ratio, but both capsule types were subject to lysosomal compartmentalization [220]. It must be noted that in case of sized rigid particles, rod-shaped particles with higher aspect ratio showed a higher internalization rate [15].

In terms of particle elasticity, theoretical analysis and molecular modeling of cell uptake suggested that rigid particles are enfolded by membrane more entirely and more easily than softer particles implying that elasticity of the particles play a

crucial role in cell internalization [221]. Polystyrenesulfonate/polyallylamine hydrochloride multilayer capsules were shown to have no impact on normal rat kidney (NRK) cells at low concentrations of about 5 particles/cell. At higher concentrations, however, the capsules were found to settle over the cells and affect their viability. When these capsules were embedded with inorganic ions such as cadmium telluride (CdTe), toxic ions were released and impacted the cell viability drastically [222].

Humanized A33 monoclonal antibody (huA33 mAb) functionalized poly(allylamine hydrochloride)/poly(sodium 4-styrenesulfonate) capsules of ~500 nm in diameter, have shown to specifically target colorectal cancer cells and trigger internalization via receptor-mediated endocytosis [105]. In case of mixed cell populations, huA33 mAb functionalized PVPON capsules of diameter around 800 nm, showed significant uptake suggesting antibody-independent internalization such as macropinocytosis [104]. Micron sized capsules were taken up by cancer cells by means of phagocytosis, while the nature of the polymer decided the integrity and intracellular localization of the capsules—non-biodegradable, synthetic PSS/PAH microcapsules remained intact and in the cytoplasmic region after 3 h incubation with cells, while the biodegradable DXS/PRM microcapsules were found to be degraded and excluded from nuclei in the perinuclear region [106].

Hollow microcapsules fabricated with layer-by-layer assemblies of poly-(sodium 4-styrene sulfonate) [PSS] and poly-(allylamine hydrochloride) [PAH] have shown to be biocompatible and applicable for 3D tumor tissue studies comprising of cells such as C6 glioma and 3T3 fibroblasts [223]. Polyelectrolyte microcapsules consisting of layer-by-layer coatings of dextran sulfate and poly-L-arginine layers, with the number of bilayers, contributing to the integrity of the capsules in vivo, have been subcutaneously injected into mice eliciting a moderate immune reaction, with a majority of the injected capsules internalized within 16 days and also initiate degradation [96]. Multilayer hollow capsules thus show promising potential as drug delivery agents capable of cell-specific targeting, by manipulating their polymeric configurations and geometry.

8.4.3 Intracellular Trafficking

After internalization via endocytosis, the particles are predominately localized within endosomes which either fuse with lysosomes for degradation or recycle their contents back to the cell surface. The behavior of the particles in endo-lysosomal vesicles is not fully elucidated. Whenever the particles are decorated with targeting ligands, these can be cleaved by the protease cathepsin L inside endo-lysosomal vesicles [224]. In some cases (e.g. quantum dots), there is an evidence of the slow particle core structure decomposition by the lysosomal enzymes [152]. In general, the escape from endosomes is favorable, since the therapeutic payload is intended to act in the cell compartments such as nucleus and cytoplasm and the accumulation of particles in endosomes followed by the degradation in endo-lysosomes strongly limits the efficiency of a payload transfer.

If a NP is engineered to escape the endo-lysosomal system, it can enter the cytoplasm, where it may interact with intracellular organelles potentially affecting cell behavior. Cytoplasm is not enzymatically-inert and contains nucleases and proteases that can neutralize the active compound. For instance, it was shown that plasmid DNA is degraded in the cytoplasm of HeLa and COS cells with a half-life of 50–90 min [225]. Additionally, the cytoplasm is a viscous environment crowded with molecules, characterized by the decreased mobility of macromolecules [226–228].

There are controversial reports on the effect of the particle size on the intracellular trafficking. As an example, compartmentalization of cadmium-telluride (CdTe) quantum dots into subcellular organelles or their ability to transverse the nucleus was reported to be affected by the particle size [229]. On contrary, the uptake of silver NPs and quantum dots into macrophages was shown to induce the expression of inflammatory mediators such as TNF- α , MIP-2 and IL-1 β independent on size [152, 230]. In addition to modulating NPs size for controlling their intra-cellular fate, coating a NP with cationic polymers, such as polyethylenimine (PEI), may initiate endosomal escape via so-called “proton sponge effect” due to the buffering inside endosomes. PEI causes additional pumping of protons into the endosome, along with a concurrent influx of chloride ions to neutralize the net charge, this causes an increase in ionic strength inside the endosome and leads to osmotic swelling, physical rupture of the endosome, and the escape of the vesicle material into the cytosol avoiding the degradative lysosomal trafficking pathway [231]. This ability for osmolytic endosomolysis has been shown to be dependent on the particle size, with small PEI800 or PEI25/DNA complexes formed in water having lower transfection efficacy compared to larger complexes formed at higher ionic strength. The localization of NPs in the intracellular space may also be directed to mitochondria. When in the cytosol, NP can elicit biological responses by disrupting mitochondrial function, eliciting production of reactive oxygen species and activation of the oxidative stress mediates signaling cascade [232]. The production of reactive oxygen species can have detrimental effects on the mitochondrial genome, induce oxidative DNA damage, and promote micronuclei formation [233]. Furthermore, certain types of NPs can induce nuclear DNA damage, leading to gene mutations, cell cycle arrest, cell death, or carcinogenesis. Demonstrating the later, hydrophilic titanium oxide NPs will persist unless they are sorted back into the endolysosomal system where they can be exocytosed. NPs that persist in the cytosol during mitosis will be distributed within the daughter cells [234]. However, the effect of the NPs on subsequent cell generations remains unclear. To date, there is no consensus on the toxicity and properties on NPs inside the cytoplasm. Additional work is still required before researchers fully understand the fate and toxicity of internalized NPs.

8.5 Concluding Remarks

Whilst the biologically driven interactions of particulates based on specific recognition moieties are being investigated for the past few decades, effects of physical parameters and especially geometry of the nanovectors still pose more questions

than answers. Recent advances in fabrication of particles with various geometries, based on above described top-down and bottom-up approaches, allowed to isolate these parameters for careful systematic analysis. For example, lithography methods used in microelectronics industry are becoming valuable tools for imparting highly reproducible size, shape, and composition of the produced particulates. It is commonly agreed that the expected use of novel nano and micro-particles will lead to a substantial improvement in the delivery of therapeutic and diagnostic payloads to diseased loci. Though a large number physicochemical compositions have been identified and studied for this purpose, there are still questions about how the physical and chemical structure of the particle affects its interaction with biological milieu. Namely, the particle shape determines the kinetics it displays within the bloodstream and its uptake at the target cell. Studies have shown that nanovectors can be efficiently rationally designed based on theoretical models of flow in the bloodstream and interactions with cells at the target site. This rational design is based on changes in the physical factors such as flow, pressure, porosity in the tissue of interest. Moreover, a combination of physical and biological features of the particles can synergistically attenuate their accumulation in the desired organ. In this regard, the thorough advanced characterization of the particles to determine surface properties such as graft density of coatings or ligands (e.g. with X-ray photoelectron spectroscopy) is very important to determine the elemental composition as the surface properties have to be standardised between particles of different geometry. A detailed understanding of particle-cell interaction from molecular to tissue level as a result of the particle's geometry will allow improved particle-based drug and imaging payload delivery systems to be developed.

To conclude, recent developments in micro and nano-fabrication of rationally designed shape and size-specific vectors, show great promise in overcoming bio-barriers en route to the disease loci. Further incorporation of stimuli-responsive biomaterials into the rationally designed vectors will enable the development of conceptually novel drug delivery systems.

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Chapter 9

Delivery of Peptides and Proteins to the Brain Using Nano-Drug Delivery Systems and Other Formulations

David Stepensky

Abstract Peptides and proteins can potently modulate processes in the central nervous system (CNS) and can be used for management of CNS disorders and other diseases. However, peptides and proteins permeate to a limited extent into the CNS, due to the blood-brain and blood-cerebrospinal fluid barriers, and undergo rapid clearance by endogenous proteases and via other pathways. These pharmacokinetic drawbacks prevent accumulation of peptide and protein drugs in the CNS and render them pharmacologically inefficient. Different approaches for enhanced delivery of peptide/protein drugs to the brain have been developed in order to overcome these drawbacks and to increase their therapeutic efficiency. These approaches include: (a) focal administration into the brain tissue or fluids (via intracerebral, intrathecal, or intracerebroventricular injection), (b) incorporation of drugs into specialized nano-drug delivery systems (DDSs) and other formulations, (c) disruption of CNS barriers, (d) use of immune cells as ‘Trojan horses’ for peptide/protein brain deliver, and others. In this chapter, the major approaches used to deliver peptide and protein drugs to the brain are described and factors that affect the systemic and local drug disposition (distribution and elimination) are summarized. Current problems and limitations in the field of brain delivery of the peptide and protein drugs are presented, and recommendations for further brain delivery enhancement of peptide/protein drugs are given.

Keywords Peptide and protein drugs • Central nervous system disorders • Blood-brain barrier • Drug targeting to the brain • Drug delivery systems • Nano-formulations

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Abbreviations

BBB	Blood-brain barrier
B-CSF-B	Blood-cerebrospinal fluid barrier
BDNF	Brain-derived neurotrophic factor
bFGF	Basic fibroblast growth factor
CSF	Cerebrospinal fluid
DDS	Drug delivery system
EGF	Epidermal growth factor
MPS	Mononuclear phagocyte system
MRI	Magnetic resonance imaging
NGF	Nerve growth factor
PLGA	Poly(lactic-co-glycolic) acid
PEG	Polyethylene glycol
STL	Solanum tuberosum lectin
TNF	Tumor necrosis factor
VIP	Vasoactive intestinal peptide
WGA	Wheat germ agglutinin

9.1 Peptides and Proteins for Treatment of Brain Diseases and Barriers for Their Use as Drugs

Central nervous system (CNS) disorders comprise a group of diseases (neurodegenerative diseases, epilepsy, neoplastic and inflammatory CNS diseases, acute and chronic pain, and others) with complex etiology and pathophysiology that have a major impact on human life duration and quality. Despite many efforts to develop drugs for management of CNS disorders, the incidence of these diseases is constantly increasing during the last decades, in part due to prolongation of human life span, and new treatments are required for their better management. Due to advances in the understanding of the regulation of the CNS functions in physiological and pathophysiological conditions, peptides and proteins have emerged as promising therapeutic modalities for management of CNS disorders. Moreover, via their pharmacological activities in the CNS, peptides and proteins can modify activity of other organs and tissues and can be used for management of non-CNS diseases, such as gastrointestinal, metabolic, and hormonal disorders. Selected examples of peptides and proteins that can be used for management of CNS and non-CNS diseases are listed in Table 9.1.

In spite of their clear therapeutic potential, many peptides and proteins are clinically ineffective and cannot be applied for management of patients with specific diseases. This lack of efficiency stems from the pharmacokinetic limitations of peptides and proteins; low permeability to the CNS tissue and fluids, and high clearance (by endogenous proteases and via other pathways, see below) that prevent their accumulation in the CNS and render them pharmacologically inefficient. Permeability of drugs from the site of administration to the CNS is limited by the blood-brain barrier (BBB) that is formed by several types of cells and is strengthened by the tight junctions (see Fig. 9.1b).

Table 9.1 Examples of peptides and proteins that can be used for management of central nervous system disorders and of other diseases

Group	Examples
Neuropeptides	Neurotensin, melanocortins, brain natriuretic peptide
Opioid peptides	Enkephalin, kyotorphin, dynorphin
Ingestive peptides	Leptin, ghrelin, orexin
Cytokines	Interleukins, tumor necrosis factor α , granulocyte/monocyte colony stimulating factor, macrophage inflammatory protein-1
Neurotropic factors	Brain derived neurotrophic factor, nerve growth factor, epidermal growth factor, insulin-like growth factor I, neurotrophins
Proteins	Therapeutic antibodies, insulin, transferrin
Hormones	Oxytocin, luteinizing hormone-releasing hormone, corticotropin releasing hormone, somatostatin, vasopressin

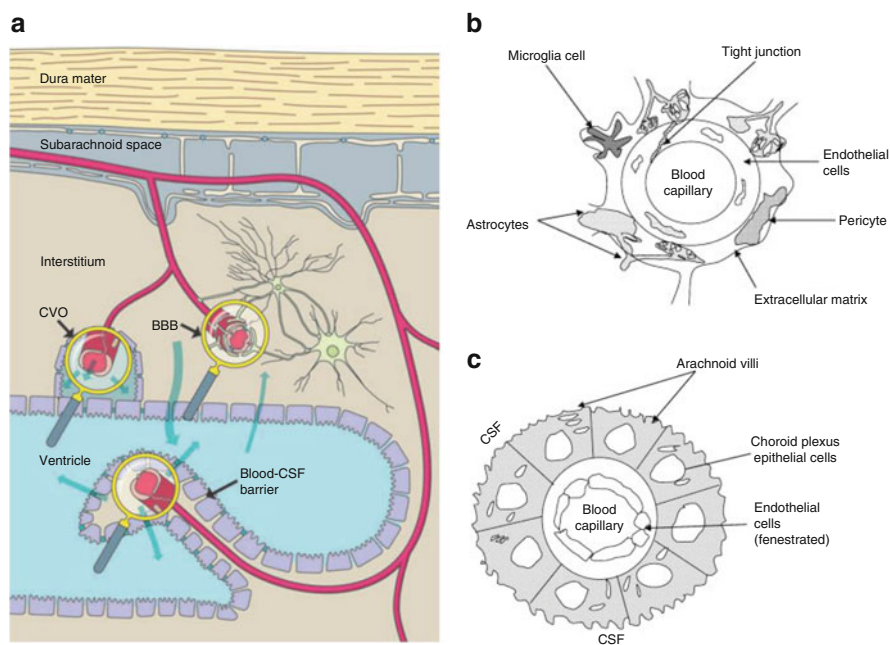


Fig. 9.1 Barriers for delivery of peptide and protein drugs to the brain. **A.** Brain fluid turnover and barrier system. Cerebral interstitial fluid is secreted at the cerebral capillaries (at the blood brain barrier) and flows from brain to CSF via perivascular spaces. The brain-cerebrospinal fluid barrier is formed by choroid epithelium where CSF is formed. In certain regions of the brain (circumventricular organs, CVO) capillaries are leaky and neuronal projections have direct access to solutes of the plasma. Reprinted from [70] with permission from John Wiley and Sons. **B.** Morphology of the blood-brain barrier (BBB) that is formed by the brain capillary endothelium, pericytes and astrocytes and strengthened by the tight junctions between the endothelial cells. Reprinted from [2] with permission from Elsevier. **C.** Morphology of the blood-cerebrospinal fluid barrier (B-CSF-B) situated at the choroid plexus and meninges. The barrier is formed by the choroid plexus epithelial cells and is strengthened by the tight junctions between these cells. Reprinted from [2] with permission from Elsevier

Table 9.2 Quantitative parameters for analysis of brain delivery of peptide and protein drugs [46, 71–74]

Parameter	Description
Brain uptake index	Relative uptake of a drug compared to a reference substance (compound freely diffusible across the BBB, such as ^{14}C -butanol)
Relative concentration	Concentration of the drug in the brain vs. other organs or body fluids; e.g., brain/plasma brain-plasma partition coefficient, K_p
Influx rate constant, K_{in}	The unidirectional influx constant from blood to brain, is most commonly determined after intravenous injection or after in situ perfusion of the compound
Permeability-surface area product, PS product	The unidirectional clearance from blood to brain, is most commonly determined after intravenous injection or after in situ perfusion of the compound
Concentration above threshold value	Concentration of the drug in the brain in comparison to a specific threshold value (e.g., the lower boundary of therapeutic window)
Fraction of dose	% of drug dose that reached the brain
Drug targeting efficiency percentage, %DTE	Quantifies the overall tendency of the drug accumulate in the brain following local (e.g., intranasal) administration vs. systemic administration
Direct transport percentage, %DTP	Relative amount of the drug that reaches the brain via direct routes, as compared to overall brain drug delivery

The structure of the BBB, including description of efflux and influx transport systems that affect the CNS permeability of specific peptides and proteins, is explained in a detailed fashion in several recent reviews [1–3].

Despite that the BBB is usually mentioned as the major barrier for CNS permeability of drugs, including peptide and protein compounds, it should be noted that the CNS is characterized by a complex fluid turnover and barrier system (see Fig. 9.1a). Specifically, permeability of a specific peptide or protein via the blood-cerebrospinal fluid barrier (B-CSF-B, see Fig. 9.1c) and the circumventricular organs, along with its BBB permeability, affects the extent of peptide/protein accumulation in the brain. Moreover, due to the directional flow (convection) of the interstitial and cerebrovascular fluids in the CNS, concentrations of the permeating compounds e.g., peptide/protein drugs are unequal in different parts of the brain. These BBB permeability and CNS convection pathways are altered in many pathological conditions (brain trauma or stroke, inflammation, tumor, epilepsy, neurodegenerative diseases, etc.) [4–6], affecting the patterns of peptides/proteins accumulation and distribution in the CNS [7–9].

Several parameters can be used to quantify the brain delivery efficiency of peptides/proteins, and of other types of drugs (see Table 9.2). There is no general consensus on the best quantitative parameter for analysis of brain drug permeability, and some of these parameters are better suited for clinical needs, while others are used predominantly in preclinical settings (for analysis of data obtained in animal

experiments). Nevertheless, it is commonly accepted that peptides and proteins are largely unable to reach the CNS following systemic administration [2, 10] due to inability to permeate the CNS barriers and rapid clearance, leading to low values of all the parameters listed in Table 9.2. Subsequently, numerous approaches have been proposed to increase the CNS accumulation of peptides and proteins for enhancement of their therapeutic efficiency.

9.2 Drug Delivery Systems for Enhanced Delivery of Peptide and Protein Drugs to the Brain

Different approaches for enhanced delivery of peptide/protein drugs to the brain have been developed over the last several decades (see Table 9.3). The drug can be focally injected into the brain tissue or fluids (as intracerebral, intrathecal, or intracerebroventricular injection) to bypass the BBB and B-CSF-B barriers. This form of drug administration is invasive and, due to safety concerns, is usually restricted to severe clinical conditions that require single drug dosing or rare drug administrations.

Alternatively, for enhanced delivery of peptide/protein drugs to the brain, the drugs' physicochemical properties can be masked by its encapsulation, adsorption or chemical conjugation to different types of nano-drug delivery systems and other formulations (e.g., nanoparticles, liposomes, polymer conjugates etc., see Table 9.3). As a result, the clearance of peptide/protein drug can be decreased by its protection from proteolytic cleavage. Indeed, the half-life of the drug/DDS in the central circulation is

Table 9.3 Approaches that can be applied for enhancement of brain delivery of peptides and proteins

Formulation types	Administration routes	Barriers disruption	Other approaches
Free drug (solution)	Intravenous	Hyperosmolar agents	Decoration with residues that can enhance BBB crossing via:
Nanoparticles	Intraarterial	Proinflammatory agents	• adsorptive transcytosis
Liposomes	Intranasal	Permeation enhancers	• transporters and pumps
Micelles	Intracerebral	Ultrasound	• receptor-mediated transcytosis
Bolavesicles	Intrathecal	Local heating	Inhibition of efflux pumps and transporters
Polymer conjugates	Intracerebroventricular		Use of immune cells (monocytes, macrophages, etc.) as 'Trojan horses'
Dendrimers			Magnetic field for magnetite-containing formulations
Pasty polymers			
Hydrogels			
...			

usually much longer than that of the free drug, especially if the DDS is decorated with polyethylene glycol (PEG) residues or other 'inert' protective groups [11, 12].

As a result of incorporation of peptide/protein drugs into the DDSs, the dimensions of the drug-containing particles are larger than those of the original free drug. Consequently, the permeability of drug encapsulated into the DDS via the CNS barriers is usually decreased. Several approaches have been proposed to overcome this problem and to enhance the brain permeability and accumulation of systemically administered DDSs (see Table 9.3). These approaches can be based on disruption of the CNS barriers by chemical compounds (e.g., mannitol or inflammatory cytokines) or external signals (e.g., ultrasound or local heating), inhibition of CNS drug efflux mechanisms, use of specific targeting residues or cell-mediated brain delivery pathways.

Nowadays, the most intensively investigated approaches for increased drug/DDS permeability to the CNS are based on their decoration with specific targeting residues. For instance, decoration with transferrin or with anti-transferrin receptor antibodies (such as OX26 murine monoclonal antibody) was proposed for enhanced transport of the nerve growth factor across the BBB into the CNS [13, 14]. Less frequently investigated approach involves 'feeding' drug/DDS into monocytes, macrophages or other immune cells that can subsequently permeate via the CNS barriers and deliver the drug into the brain (the 'Trojan horses' approach) [15–18].

It should be noted that some of the above-mentioned approaches can be used in combination e.g., DDSs decorated specific surface residues can be administered together with barrier disrupting agent or treatment. However, use of a specific approach, or combination of approaches, does not guarantee more efficient drug/DDS brain accumulation (see Sect. 9.3). Apparently, only in some cases the approaches listed in Table 9.3 are efficient enough to increase the values of the quantitative brain delivery parameters (see Table 9.2) for peptide/protein drugs. Only in rare cases, brain targeting of peptide/protein drugs is attained (i.e., increased drug accumulation in the brain, as compared to other organs and tissues). In part, these outcomes stem from the complex pathways of drug/DDS disposition (i.e., distribution and elimination) in the body (following systemic or local administration) and diverse effects of the specific brain targeting approaches on these pathways. For instance, application of focused ultrasound to disrupt the CNS barriers increases local permeability of the drug/DDS into the brain, but also enhances the drug transport in the opposite direction, from the brain into the systemic circulation, and can induce local release of the drug from the DDS (i.e., release of the drug with its subsequent degradation, not at the target site). As a result of these processes, focused ultrasound will not necessarily enhance the brain drug accumulation efficiency. Therefore, detailed analysis of the drug disposition pathways and of the resulting targeting efficiency of the individual targeting approaches and of their combinations is required for rational development and successful application of peptide/protein drugs intended for brain delivery. In the next sections of this chapter, detailed analysis of the major factors that affect the disposition and efficiency of systemically- and focally-administered peptide and protein drugs is presented and current limitations in the brain delivery of these drugs are summarized.

9.3 Disposition Pathways of Peptide and Protein Drugs Intended for Brain Delivery

9.3.1 Systemic Administration

Systemic parenteral administration (intravenous, subcutaneous, intraperitoneal, etc.) is the most commonly applied way to administer peptide/protein drugs intended for brain delivery. Enteral administration routes, oral, buccal, rectal, etc. are usually not applied for these drugs due to their rapid degradation in the gastrointestinal tract that leads to low bioavailability. Approaches for brain drug delivery via systemic parenteral administration routes are extensively described in several recent reviews [3, 19–21]. In addition, several reviews focused specifically on brain delivery of peptide/protein drugs following systemic administration [2, 10, 22].

Many strategies have been proposed for the purpose of brain delivery of peptide/protein drugs. Most commonly, these strategies are based on liposomes, nanoparticles, polymer conjugates, dendrimers, and other DDSs that encapsulate peptide/protein drugs, usually applied in combination with targeting approaches (based on CNS barrier disruption or transport modulation, see Table 9.3). For the purpose of brain-targeted drug delivery, these DDSs should encapsulate substantial amounts of peptide/protein drug, should be stable in the systemic circulation for sufficient amount of time (usually several hours or longer), should efficiently reach the target tissue (the brain), and eventually release the encapsulated drug in the brain.

The choice of the DDS type and the targeting approaches may significantly affect the systemic and local (brain) drug/DDS pharmacokinetics and pharmacodynamics. Following systemic administration and permeation into the bloodstream, peptide/protein drugs and their formulations are exposed to a plethora of processes which can interfere with their targeting efficiency (see Table 9.4). Specifically, drug/DDS can undergo elimination by proteases or via hepatic and renal elimination pathways, endocytosis and metabolism by the cells that are circulating in the blood, or the cells that are lining the blood vessels. DDSs that encapsulate the peptide/protein drugs can aggregate in the bloodstream (with subsequent deposition in capillaries or uptake by the phagocytic cells in the bloodstream), and can release the encapsulated drugs (decapsulate) in the systemic circulation, prior to reaching the target tissue. The remaining fraction of the free and encapsulated peptide/protein drugs that managed to avoid the above-mentioned processes can extravasate to reach the brain or other organs/tissues. For detailed analysis of the factors that affect the disposition pathways of free and encapsulated drugs, the readers are referred to several excellent recent reviews [23, 24].

It can be seen that the chances of the individual drug/DDS to reach the target tissue are low and drug targeting to the site of action in the brain requires blocking or avoiding the above-mentioned interfering pathways. Efficiency of the individual disposition pathways listed in Table 9.4 can be substantially affected by the dimensions (size and shape) and the surface properties (charge, surface residues) of the drug/DDS. For instance, the cells of the mononuclear phagocyte system (MPS)

Table 9.4 Major pathways of peptide and protein drugs disposition (distribution and elimination) following systemic (intravenous) administration (in a free form or encapsulated into drug delivery systems)

Pathway
Degradation in the systemic circulation
Elimination by liver and kidneys
Aggregation in the bloodstream
Interaction with the non-immune endogenous compounds (formation of ‘corona’)
Interaction with the soluble components of the immune system (antibodies, complement)
Uptake by mononuclear phagocyte system (MPS) cells
Uptake by the endothelial cells
Uptake by the red blood cells
Permeation to the target organ (the brain)
Permeation to other organs and tissues
Release of the encapsulated drug from the DDS

cells have their ‘preferences’ for the size, shape, and charge of the DDSs and will preferably endocytose charged (with positive or negative surface charge) drugs/DDSs with diameter bigger than 200 nm [23, 25]. Therefore, for the purpose of preferential drug/DDS delivery to the brain, systemically-administered formulations with the diameter of less than 200 nm should preferably be used.

It should be noted that during their presence in the systemic circulation, drugs/DDSs can interact with and adsorb endogenous compounds such as albumin, transferrin, lipoproteins, and other plasma components [23]. As a result of this interaction, coating or ‘corona’ is formed on the surface of drug/DDS which affects its surface properties and efficiency of its disposition pathways (see Table 9.4), and can further reduce the chances for efficient drug targeting to the brain. A popular strategy to block these pathways and to avoid ‘corona’ formation is drug/DDS decoration (via non-covalent adsorption or covalent conjugation) with polyethylene glycol (PEG) residues. PEGylated drugs/DDSs are often referred to as ‘stealth’ formulations, and PEGylation technology that forms ‘inert’ steric barrier on the drug/DDSs’ surface can indeed be used to prolong their residence in the bloodstream [11, 12, 26]. However, repeated administration of PEGylated drug/DDS can provoke immune reactions of the host against the PEG residues, via formation of anti-PEG IgM antibodies and complement activation, that can increase the drug/DDS clearance and diminish its brain accumulation [27, 28]. These immunogenic aspects of PEGylation need more detailed investigation, and combination of PEGylation with ‘active’ targeting approaches is recommended for enhanced drug/DDS accumulation in the brain.

Despite that brain drug delivery using systemic administration routes was analyzed in many studies, only in few of them quantitative data on efficiency of drug targeting to the brain was reported [29]. Examples of the studies that analyzed brain

delivery of peptide/protein drugs following systemic administration and reported quantitative brain targeting data are presented below.

Radiolabeled brain-derived neurotrophic factor (BDNF) was attached to the murine OX26 monoclonal antibody (that recognizes the rat transferrin receptor) using avidin-biotin technology [30]. Brain accumulation of this conjugate in experimental rats was threefold higher than that of the free peptide (0.49 brain/plasma ratio for the conjugate, as compared to the 0.16 ratio for the BDNF peptide) [29]. Subsequently, brain accumulation of PEGylated form of this conjugate was analyzed and was found to be approximately 17-fold higher than that of the free BDNF peptide [31].

Effect of decoration with the same OX26 antibody on brain accumulation of amyloid beta 1-40 peptide [32] and of PEGylated epidermal growth factor (EGF) [33] was investigated by the same research group. Brain accumulation of the amyloid beta 1-40 peptide decorated with the OX26 antibody was approximately 16-fold higher than that of the free peptide (based on the calculated values of % of the dose that reached the brain) [29]. Brain accumulation of the EGF-PEG3400-OX26 conjugate was 1.4-fold higher than that of the free peptide (0.31 brain/plasma ratio for the conjugate, as compared to the 0.22 ratio for the BDNF peptide) [29]. In a different study, decoration with the same OX26 antibody increased by eightfold the brain accumulation of chimeric opioid peptide DALDA (Tyr-DArg-Phe-Lys; 0.057 brain/plasma ratio for the conjugate, as compared to the 0.0063 ratio for the free DALDA peptide) [34].

Increased brain accumulation of a nerve growth factor (NGF) using two different approaches has been reported. In one study, NGF was encapsulated into PEGylated liposomes decorated with RMP-7, a ligand to the B2 receptor, leading to 12.5-fold its higher accumulation in the rat brain (0.30 brain/blood ratio for the decorated liposomes, as compared to the 0.024 ratio for the free NGF peptide) [35]. In another study, NGF encapsulation into poly(butyl cyanoacrylate) nanoparticles coated with polysorbate 80 induced a 2.8-fold increase in its brain accumulation, as compared to the free peptide [29], and significantly reduced the basic symptoms of Parkinsonism in a rat disease model [36].

The above-described examples summarize the findings in selected publications only, and for description of other studies and the major research trends in the field of brain delivery of peptide/protein drugs following systemic administration the readers are referred to several recent reviews [2, 10, 22]. Results of recent quantitative analysis indicate that the efficiency of brain drug delivery is dependent on the drug formulation: nanoparticle-based DDSs appear to be more efficient than the liposome- or conjugate-based DDSs [29]. Incorporation of specialized reagents, such as PEG residues and surface-active agents, decoration with specific targeting residues, and use of external targeting signals apparently can increase the efficiency of drug/DDS delivery to the brain and can lead to brain-targeted drug delivery. However, more detailed quantitative analysis of the drug/DDS disposition and its mechanisms is required to identify the limiting factors for drug brain delivery and the drug formulations that are the most suitable for this purpose [29].

9.3.2 Intranasal Administration

Nasal administration of drugs has been increasingly applied in recent years in an attempt to deliver drugs to the brain. The reasons for this are: (a) relative inefficiency of drug delivery to the brain via systemic routes [29], (b) non-invasive nature of nasal drug administration, as compared to the intravenous and other systemic parenteral administration routes, (c) possibility of direct drug delivery to the brain following intranasal administration, bypassing the BBB and avoiding the gastrointestinal and hepatic first-pass metabolism, (d) intranasal delivery may fasten the onset of action of CNS drugs, which can be of special clinical importance for the anti-pain and anti-migraine medications [37]. Detailed description of the brain drug delivery approaches via nasal route is provided in several recent reviews [38–42]. Additional publications describe brain delivery of peptide and protein drugs using nasal route [43–45].

The mechanisms of drug delivery to the brain following intranasal administration are understood only partially. It appears that following intranasal administration drugs/DDSs can reach the brain via several direct and indirect pathways (see Fig. 9.2). Specifically, some drugs/DDSs can apparently reach the CNS via the olfactory pathway, that consists of the olfactory epithelium, olfactory tract, anterior olfactory nucleus, piriform cortex, amygdala, and hypothalamus [40, 44]. An additional direct pathway that can mediate drug/DDS delivery to the brain from the

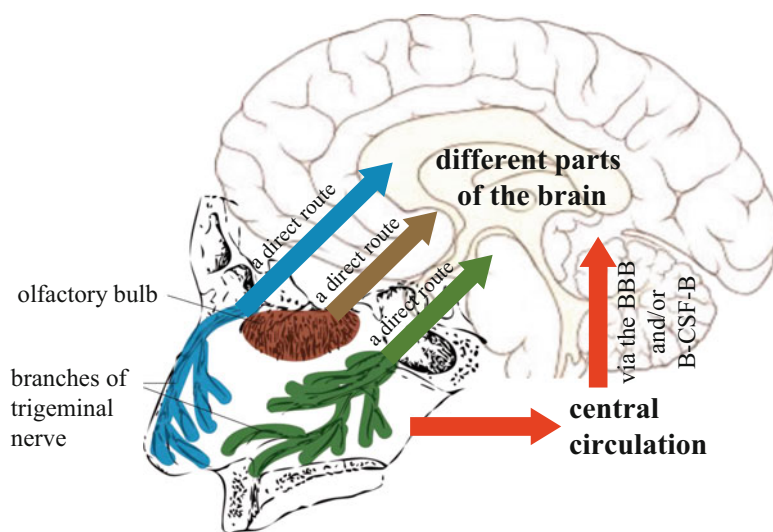


Fig. 9.2 Pathways of peptide and protein drug delivery to the brain following intranasal administration. Following intranasal administration, the drug (in a free form, or encapsulated into a drug delivery system) can reach the brain via one of the direct routes (via the olfactory and/or trigeminal nerves), or indirectly, by absorption to the central circulation, entry to the brain blood vessels, and passage through the blood-brain barrier and/or blood-cerebrospinal fluid barrier. Reprinted from [46] with permission from Elsevier

nasal cavity is via the three branches of the trigeminal nerve (ophthalmic division, maxillary division, and mandibular division) which merge at the trigeminal ganglion, enter the CNS in the pons, and terminate in the spinal trigeminal nuclei in the brainstem [40].

In addition to these direct pathways, drug/DDS can enter the brain indirectly, via the central circulation (and the BBB and B-CSF-B barriers, refer to Fig. 9.2), following its absorption from the nasal cavity to the systemic circulation. Indeed, drugs and DDSs can undergo efficient absorption into the local blood and lymph vessels of the highly vascularized nasal mucosa. This absorption can be increased in some pathological conditions e.g., flu, allergic diseases and other diseases accompanied by local inflammation and vessel permeability in the nasal cavity, or if drug formulation contains permeation enhancers such as penetrating peptides, surface-active agents, pro-inflammatory compounds, etc. In addition, efficiency of both direct and indirect nose-to-brain pathways can increase if the drug or its formulation induces local toxic effects that compromise the function of physiological barriers. This factor is overlooked by many researchers and insufficient amount of efforts is directed towards safety characterization of intranasally-administered drugs and DDSs.

Direct and indirect nose-to-brain routes apparently comprise minor routes of drug/DDS disposition following intranasal administration [46], especially for the peptide and protein drugs. The absolute bioavailability of the nasal desmopressin (nine amino acid peptide), buserelin (nine amino acid peptide) and calcitonin (32 amino acid peptide) in human subjects was reported to be lower than 10 % [47–49], reflecting low efficiency of drug absorption into the central blood circulation (directly and via the brain). Majority of the peptide/protein drug/DDS that is applied intranasally is not reaching the brain, but undergoes other disposition processes (see Table 9.5). Specifically, part of the administered drug/DDS is inhaled or/and ingested. Other part is undergoing enzymatic degradation and clearance by the mucociliary system (the self-clearing mechanism of the nasal cavity and of the airways). DDSs that are deposited in the nasal cavity may release the encapsulated drug which becomes available for enzymatic degradation in the nasal cavity or/and acquire coating or ‘corona’ which reduces the efficiency of the direct and indirect brain delivery pathways.

Table 9.5 Major pathways of peptide and protein drugs disposition (distribution and elimination) following intranasal administration (in a free form or encapsulated into drug delivery systems)

Pathway
Degradation in the nasal cavity
Discharge with mucus (e.g., due to ciliary movement)
Interaction with the endogenous compounds (formation of ‘corona’)
Release of the encapsulated drug from the DDS
Passage into the airways and lungs (inhalation)
Passage into the gastrointestinal system (ingestion)
Absorption into the central circulation
Brain permeation via the branches of the trigeminal nerve (ophthalmic division, maxillary division, and mandibular division) in the nasal cavity
Brain permeation via the olfactory bulb and olfactory nerve

Overall, it appears that only a small fraction of the intranasal-administered drug reaches the brain and exerts pharmacological effects in the CNS. Efficiency of drug delivery to the brain following intranasal administration is usually reported using the relative targeting indexes (such as brain/plasma concentrations ratio, drug targeting efficiency percentage and direct transport percentage, see Table 9.2) since it is difficult to measure directly the fraction of the dose that reached the brain in preclinical and clinical studies. In our recent publication, we analyzed the available quantitative data on drug delivery to the brain via nasal route and the relative brain targeting efficiency of different types of intranasal-administered DDSs [46]. Based on quantitative experimental data from 73 publications (that reported data on 82 compounds, only part of them from the peptide/protein group), we analyzed the efficiency of brain drug delivery following intranasal drug/DDS administration. Results of studies that investigated brain delivery of peptide/protein drugs following intranasal administration and reported quantitative brain targeting data are described below.

Brain-targeted delivery of peptide/protein drugs that were administered into the nasal cavity in the form of drug solution was investigated quantitatively in several studies. Following intranasal administration of hypocretin-1 neuropeptide (orexin-A, a 33 amino acid peptide) to rats, drug targeting to the brain and spinal cord increased five- to eightfold, as compared to the intravenous drug administration [50]. It was estimated that 77 % of hypocretin-1 reached the brain following intranasal administration via direct (olfactory and trigeminal) transport routes. The brain and cerebrospinal fluid levels achieved following intranasal administration of hexarelin (a 9 amino acid growth hormone releasing neuropeptide) solution to rabbits were approximately 1.6 times greater than those attained after intravenous administration, despite that the plasma drug levels were significantly lower after intranasal drug administration, as compared to intravenous drug injection [51]. Intranasal administration resulted in significantly different spatial distribution patterns of hexarelin in various regions of brain consistent with substantial drug traffic via olfactory route, whereas, intravenous administration yielded nearly homogeneous drug distribution in the brain. Intranasal administration of a TNF-alpha inhibitory single-chain antibody fragment (scFv, molecular weight of 26.3 kDa) solution to mice resulted in 0.3 % systemic bioavailability [52]. Addition of a penetration enhancing peptide to the formulation enhanced delivery of this drug to the olfactory bulb and to the cerebrum (estimated 98 % of brain drug delivery via direct transport routes [46]), without increasing systemic exposure.

In two studies, brain delivery of peptides/proteins encapsulated in nanoparticle formulations was investigated in quantitative fashion. Brain accumulation of vasoactive intestinal peptide (VIP, a 28 amino acid peptide hormone) encapsulated into poly(ethylene glycol)-poly(lactic acid) nanoparticles modified with wheat germ agglutinin (WGA) was investigated in comparison to VIP-loaded unmodified nanoparticles and drug solution [53]. It was found that higher amounts of VIP have reached the brain following administration of the drug encapsulated into the WGA nanoparticles (estimated 84 % of brain drug delivery via direct transport routes [46]), as compared to the unmodified nanoparticles and drug solution. The same improvements in spatial memory in ethylcholine aziridium-treated rats were

observed following intranasal administration of VIP encapsulated into unmodified nanoparticles and twofold lower dose of VIP encapsulated into wheat germ agglutinin-modified nanoparticles (25 $\mu\text{g}/\text{kg}$ and 12.5 $\mu\text{g}/\text{kg}$, respectively). In another study, nasal delivery of basic fibroblast growth factor (bFGF) encapsulated in polyethylene glycol-poly(lactide-polyglycolide) (PEG-PLGA) nanoparticles conjugated with *Solanum tuberosum* lectin (STL, potato lectin), which selectively binds to N-acetylglucosamine on the nasal epithelial membrane, was investigated [54]. The overall brain exposure to the bFGF in the olfactory bulb, cerebrum, and cerebellum of rats following nasal application of STL modified nanoparticles was 1.79- to 5.17-fold of that of rats with intravenous administration, and 0.61- to 2.21-fold and 0.19- to 1.07-fold higher compared with intranasal solution and unmodified nanoparticles, respectively. These findings indicate substantial contribution of direct nose-to-brain routes to overall brain delivery of bFGF following nasal administration (estimated 72 % for the STL-decorated nanoparticle formulation [46]). The neuroprotective effects of different bFGF formulations were determined using a rat model of Alzheimer's disease, and it was found that the STL-decorated nanoparticle bFGF formulation significantly improved spatial learning and memory of diseased rats following intranasal administration, while effects of other bFGF formulations (undecorated nanoparticles and solution) were less prominent.

Overall, the experimental data indicate that nasal administration of peptide/protein drugs can be used for management of brain diseases, but the efficiency of brain drug delivery following intranasal drug/DDS administration appears to be low. We were not able to identify correlations between the drug physico-chemical characteristics, formulation properties, or their combination, and the values of the analyzed brain targeting indexes [46]. Outcomes of data analysis indicate that particle- and gel-based DDSs offer limited advantage for brain drug delivery in comparison to the intranasal administration of drug solution. On the other hand, incorporation of specialized reagents (e.g., absorption enhancers, mucoadhesive compounds, targeting residues, etc.) can increase the efficiency of drug delivery to the brain via nasal route [46]. Overall, more elaborate and detailed investigations of drug/DDSs disposition following intranasal administration (see Table 9.5) are needed to reveal the mechanisms of drug targeting to the brain and to identify the most suitable drugs and DDSs for the management of brain diseases via nasal route.

9.3.3 Local Administration

Drugs can be administered directly into the CNS via the intracerebral, intrathecal or intracerebroventricular routes, as drug solution or encapsulated into DDSs or specialized implants. Due to the invasive nature of these administration routes, and safety concerns, clinical application of these administration techniques is limited to delivery of selected anticancer drugs. Indeed, specialized approaches have been developed for brain injection of anticancer drug solutions via different routes [55, 56] and implantation of drug-eluting polymers during the brain tumor resection

Table 9.6 Disposition (distribution and elimination) pathways following intracerebral administration of peptide and protein drugs (in a free form or encapsulated into drug delivery systems)

Pathway
Diffusion
Convection
Degradation
Aggregation
Interaction with the endogenous compounds (formation of ‘corona’)
Release of the encapsulated drug from the DDS
Uptake by the target cells (e.g., the neurons)
Uptake by non-target cells

procedure [57, 58]. Drug-eluting polymer formulations can be designed to release the encapsulated drug over the desired period of time, usually several hours—up to several weeks or months, with the desired kinetics. For example, release of carmustine from the Gliadel® wafers in the brain exhibits two phases: the induction period (‘burst’ release by diffusion of ~60 % of the drug over the first ~10 h) followed by the erosion period (~5 days) [59]. In addition, convection-enhanced delivery technique can be used to modulate the local drug concentrations in the brain tissue (e.g., within the brain tumor and its vicinity) [60, 61].

Local administration of peptide and protein drugs into the brain tissue and fluids has been investigated much less extensively as compared to that of the anticancer drugs, and no such treatments have been approved for clinical use. The reader is referred to the selected experimental and review manuscripts that analyzed drug peptide/protein delivery to the brain via local administration routes [62, 63]. Due to the analytical limitations, local disposition of peptide and protein drugs in the brain following their local administration is seldom analyzed, and pharmacological responses are usually reported in the experimental research studies. Within the brain tissue, locally injected or released drug is exposed to several disposition pathways, which are summarized in Table 9.6. The drug/DDS can traffic in the brain via simple diffusion, and with flow of the interstitial fluid and of the CSF, direction and extent of which varies in the different parts of the brain [7, 64]. In addition, the drug/DDS can undergo degradation by local proteases, interaction with the endogenous compounds, aggregation, and uptake by the target and non-target cells within the brain. As a result of these disposition pathways, the drug/DDS is not staying at the site of local injection or implantation, and a gradient of drug/DDS concentrations is formed in the brain tissue.

From the studies with anti-cancer drugs administered locally into the brain, it is known that the major limitation of the local drug administration approaches is that therapeutic drug concentrations are attained only in a close vicinity to the drug/DDS administration site [65, 66]. The thickness of this layer of therapeutic drug concentrations in the brain is up to several millimeters [58], as revealed by the non-invasive imaging-based or destructive analytical techniques (e.g., fluorescence, MRI, radio-labeling, etc.) [65, 66], reflecting the balance between the release and disposition

kinetics of the specific drug/DDS combination. This rapid drug disposition from the administration site and inefficient exposure of the brain to the locally-administered drug poses a major limitation for the clinical effectiveness of peptide and protein drugs. More detailed analysis of the individual disposition pathways of peptide and protein drugs and their dependence on the formulation properties is required for devising efficient local brain administration strategies for these drugs. On the other hand, locally-administered drugs/DDSs are usually characterized by high safety (high therapeutic index) due to the rapid drug disposition from the administration site and dilution of the drug in the brain and other organs, tissues and fluids [58, 66].

9.4 Current Limitations in the Field of Brain Delivery of the Peptide and Protein Drugs

9.4.1 Limited Targeting Efficiency

Based on the available data, it appears that peptide and protein drugs do not reach efficiently the intended site of action following administration via systemic and intranasal routes, and are rapidly disposed from the injection or release site following local administration into the brain. Development of specialized drug formulations that incorporate specific reagents (or their combinations), such as PEG residues and surface-active agents, decoration with specific targeting residues, and use of external targeting signals, apparently can enhance the drug/DDS delivery to the brain, elevate and prolong the exposure of different parts of the brain to therapeutic drug concentrations, and increase brain drug targeting efficiency. However, rational development of such drug formulations should be based on detailed understanding of drug disposition pathways inside and outside the brain (in the systemic circulation, in the nasal cavity, etc.). For this purpose, sensitive and reliable analytical methods should be used to quantify the peptide and protein drugs and DDSs in organs, tissues, and body fluids.

Unfortunately, many publications that investigate brain delivery of peptide and protein drugs report qualitative data and/or pharmacological effects of the investigated drugs. In the majority of studies with quantitative data, only a short description of the analytical methods is provided and it is not clear whether the specific method had been validated, and whether it was able to quantify reliably and separately the drug at its different forms in the analyzed samples (free, aggregated, protein-bound, encapsulated in the DDS, etc.). This problem is especially acute for the imaging-based analytical methods that are becoming the common way for drug quantification in preclinical and clinical studies [67, 68]. It is possible that insufficient characterization of the analytical methods could have introduced bias in the data that were reported in the individual manuscripts, and lead to erroneous conclusions regarding the drug/DDS disposition [29, 46]. It is recommended to perform detailed assay validation, including quantitative analysis of the different forms of the analyzed drug, in the future studies that will assess the drug/DDSs brain delivery and targeting via different routes.

9.4.2 Heterogeneous Drug Distribution in the Brain

The drug that permeates the brain following systemic or intranasal administration, or that is locally administered into the brain, is exposed to complex disposition processes (see Table 9.6) that lead to heterogeneous drug distribution in different parts of the brain. For instance, uptake to the brain via olfactory route leads to increased drug concentrations in the amygdala and hypothalamus; uptake via trigeminal nerve leads to increased drug concentrations in the trigeminal nucleus and thalamus, etc. From these sites, the drug can subsequently permeate into other parts of the brain by simple diffusion and according to the directions of convection of the interstitial fluid and of the CSF flow [7, 64]. Indeed, in several publications, drug concentrations in different parts of the brain (cerebrum, hippocampus, etc.), and/or in the CSF were reported, and substantial differences in drug concentrations in the analyzed samples were found (see the raw data from the individual studies that are summarized in the Supplementary Tables of [29, 46]).

For selective and efficient treatment of specific CNS diseases, intra-brain drug/DDS targeting to the specific parts and structures of the brain may be required (e.g., substantia nigra in Parkinson's disease, lesions sites in Alzheimer's disease, etc.). Thus, brain drug delivery strategies should preferably take into account the three-dimensional structure of the brain, the directions of CSF flow and of the convection-mediated drug/DDS disposition pathways in the brain [7, 64]. Future drug/DDS brain delivery and targeting approaches will efficiently rely on the heterogeneous nature of drug distribution in the brain, but more detailed knowledge of brain drug/DDS disposition pathways will be required for this purpose.

9.4.3 Toxicity Risks

Drugs and DDS components can exert toxic effects on the physiological barriers (the nasal epithelium, the brain endothelium, and others). Specifically, the BBB and B-CSF-B barriers function can be compromised by specific formulation components and treatments (permeation enhancers, hyperosmolar agents, efflux pumps inhibitors, ultrasound, etc., see Table 9.3). These alterations in brain barrier functions can lead to permeation of undesired compounds into the brain and subsequent alterations of brain function or/and toxicity. Extent, duration and consequences of these alterations in brain barrier permeability should be determined for the individual drugs/DDSs to pave the way for their safe clinical application. To this end, specific markers (e.g., methylene blue) or imaging approaches [69] can be applied in preclinical and clinical settings.

Depending on the administration route and the type of DDS, additional patterns and mechanisms of toxicity, in addition to the toxicity at the BBB and B-CSF-B barriers, should be investigated. For instance, local drug/DDS injection or implantation into the brain can exert direct toxic effects on neurons and other types of cells. On the other hand, nasal-administered drugs/DDSs can be toxic to the nasal epithelium,

and/or to the neurons of the olfactory and trigeminal nerves. These types of toxicity have not been studied in sufficient fashion, and their dependence on the drug/DDS composition (e.g., presence and content of permeation enhancers), dosing frequency and application volume is largely unknown. For the purpose of safe and efficient application of brain delivery of peptide and protein drugs, all these potential toxicity pathways should be investigated and ruled out.

9.5 Summary

Peptide and protein compounds can be potentially used for management of CNS disorders and of some non-CNS diseases. However, pharmacological efficiency of drugs from this class is limited by their inefficient permeation into the CNS and rapid clearance by endogenous proteases and via other pathways. Different types of nano-drug delivery systems and other formulations, formulation components, targeting approaches and administration techniques have been developed to overcome these pharmacokinetic limitations of peptide and protein drugs. During the recent decades, substantial advances in development of peptide/protein drugs and drug delivery systems were made. However, the currently available delivery approaches apparently offer limited brain exposure to the administered drugs and are characterized by low pharmacological effectiveness. More detailed analysis of the individual drug/DDS disposition pathways in the brain and in other organs/tissues/fluids is required to devise more efficient brain drug delivery approaches. In addition, the mechanisms and magnitude of toxic effects induced by the peptide and protein drugs and drug delivery systems, in the brain and in other organs and tissues, should be investigated and ruled out to pave the way for efficient clinical application of these drugs.

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Chapter 10

Polymer-Based DNA Delivery Systems for Cancer Immunotherapy

Ayelet David and Adi Golani-Armon

Abstract The use of gene delivery systems for the expression of antigenic proteins is an established means for activating a patient's own immune system against the cancer they carry. Since tumor cells are poor antigen-presenting cells, cross-presentation of tumor antigens by dendritic cells (DCs) is essential for the generation of tumor-specific cytotoxic T-lymphocyte responses. A number of polymer-based nanomedicines have been developed to deliver genes into DCs, primarily by incorporating tumor-specific, antigen-encoding plasmid DNA with polycationic molecules to facilitate DNA loading and intracellular trafficking. Direct *in vivo* targeting of plasmid DNA to DC surface receptors can induce high transfection efficiency and long-term gene expression, essential for antigen loading onto major histocompatibility complex molecules and stimulation of T-cell responses. This chapter summarizes the physicochemical properties and biological information on polymer-based non-viral vectors used for targeting DCs, and discusses the main challenges for successful *in vivo* gene transfer into DCs.

Keywords Antigen presenting cells • Cancer immunotherapy • Cationic polymers • Chitosan • Dendrimers • Dendritic cells • DNA vaccine • Immunization • Gene delivery • Polyamidoamine • Polyethylenimine • Poly(lactic acid) • Poly(lactide-co-glycolide) • Poly(glycolic acid) • Poly(lysine)

Abbreviations

Ag	Antigen
APC	Antigen-presenting cell
BMDCs	Bone marrow-derived dendritic cells
CRD	Carbohydrate recognition domain
CLR	C-type lectin receptor

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CTL	Cytotoxic T-lymphocytes
DCs	Dendritic cells
ER	Endoplasmic reticulum
GM-CSF	Granulocyte-macrophage colony-stimulating factor
HLA	Human leukocyte antigen
iDCs	Immature DCs
MΦs	Macrophages
MHC	Major histocompatibility complex
MR	Mannose receptor
MRD	Minimal residual disease
NK	Natural killer
NLS	Nuclear localization signal
OVA	Ovalbumin
PAMAM	Polyamidoamine
PEG	Poly(ethylene glycol)
PEI	Poly(ethylenimine)
PIC	Polyion complexes
PLGA	Poly(lactide-co-glycolide)
PLL	Poly(L-lysine)
PS	Polystyrene
SiRNA	Small interfering RNA
TAA	Tumor-associated antigen
TAP	Transporter associated with antigen presentation
WTC	Whole tumor cells

10.1 Introduction

Cancer has long represented a major burden on health and longevity [1]. The high prevalence of the disease has made it the basis of a major research focus, with the disease being investigated in many contexts. Intensive research has yielded a large body of information that has served to uncover biological, biochemical and pathophysiological aspects of the disease, as well as their underlying mechanisms. Based on the information gathered by the efforts of numerous researchers, novel nanosized medicines (nanomedicines), that can be effectively localized to tumors and actively taken up by cancer cells have been developed, and are expected to replace traditional chemotherapy and possibly irradiation due to increased efficiencies and attenuated adverse side-effects [2].

Unfortunately, recent years have seen only minor improvements in cancer morbidity and mortality rates, with the most commonly used cancer therapies still being conventional chemotherapy, despite their serious side-effects. This has encouraged researchers to seek answers in the field of immunotherapy [3]. Whereas exogenous anti-cancer agents have failed to generate strong and effective responses against tumors and induce regression, the endogenous activity of the immune

system was hoped to elicit specific anti-tumor immunity that would result in tumor regression and elimination. In particular, dendritic cell (DC) vaccinations were expected to induce immunological memory that would enable recognition and destruction of residual tumor cells that evaded earlier treatment, thereby avoiding recurrence of the disease [4]. Previous studies employing mouse models indicated that the establishment of anti-tumor immunity required the presentation of tumor-associated antigens (TAA) by DCs [5, 6]. Two unique properties make DCs the most potent antigen-presenting cells (APCs), namely their ability to present exogenous antigens on major histocompatibility complex (MHC)-I molecules to prime CD8⁺ T-cells (i.e., cross-presentation), and their capability to initiate, activate and modulate the various arms of the immune system in a coordinated manner. Indeed, *in vivo* targeting of tumor antigens into DCs was shown to elicit strong TAA-specific DC4⁺ and CD8⁺ immune responses [7]. Therefore, DCs loaded with TAA or tumor antigen-encoding plasmid DNA may facilitate the development of new immunotherapies for cancer treatment. This chapter highlights the repertoire of non-viral, nanosized polymeric DNA delivery systems (polyplexes) available to achieve efficient gene transfer into DCs for immunotherapeutic applications in cancer therapy. The physicochemical characteristics and surface properties of polyplexes required for efficient gene transfer and gene expression in DCs are discussed. The influence of the ligand valency on the targeting of DCs via dendritic cell surface receptors is also described.

10.1.1 Cancer Immunotherapy

The field of immunotherapy aims at activating a patient's own immune system against a host disease. Immunotherapy includes the investigation of different mechanisms leading to activation of the immune response, and the development of methods to control such responses in a desired manner so as to combat disease, either as a single strategy or in combination with other treatments. In the specific case of cancer immunotherapy, it is speculated that the *in vivo* activity of endogenous immune cells would be far more sensitive and specific than would any exogenous treatment [8]. Anti-cancer vaccines have, thus been designed to induce both tumor-specific effector T cells and tumor-specific memory T-cells [4]. Such anti-cancer vaccination, it is hoped, will induce regression of established tumors, as well as prevent the onset of secondary tumors and metastases, possibly by addressing the issue of minimal residual disease (MRD). In this case, sparse cancerous cells that evaded irradiation or chemotherapy will be detected and destroyed by the immune system cells that, unlike the temporary exogenous treatment, are present in the body at all times. A major requirement for an efficient anti-cancer vaccination is the generation of a cytotoxic T-lymphocyte (CTL) response, but manipulation of the other arms of the immune response, mainly CD4⁺ T helper cells, is critical for a robust and long lasting CTL response [9–11]. CTL response generation depends on antigen (Ag) presentation on MHC to naïve DC8⁺ T cells. The poor Ag presentation

activity demonstrated by tumor cells [12] highlights the need for professional APCs to generate the desired CTL response. Among the several types of APCs in the body, DCs are the most suitable for this purpose.

10.2 Dendritic Cells

10.2.1 *Dendritic Cell Biology*

In 2011, Ralph Steinman was awarded the Nobel Prize for his 1973 discovery of DCs. DCs are a family of professional APCs. As the most powerful APCs, they are equipped with specialized machinery that enables them to regulate the initiation of a primary immune response [13–15]. Originally derived from bone marrow, DCs are seeded in all tissues, where they sample their environment in an attempt to detect tissue damage, pathogen entry, inflammation and malignantly transformed cells by taking up particles and molecules, processing them into short peptides and presenting the resulting peptides on MHC molecules. DCs use different routes to capture Ags, including phagocytosis, micro- and macropinocytosis, and receptor-mediated endocytosis [9, 15, 16]. Importantly, receptor-mediated endocytosis can be exploited for the targeting of TAAs into DCs.

Immature DCs in peripheral tissues are characterized by high Ag capture activity, low surface MHC levels and co-stimulatory molecules, as well as a tendency to induce tolerance to self-antigens. Upon recognition of “danger” signals, DCs undergo a maturation process characterized by a reduction of Ag capture activity and an increase of MHC and co-stimulatory molecule expression on the cell surface [17]. Mature DCs migrate to adjacent lymph nodes, where they present antigenic peptides to naïve T-cells [4]. Ag presentation, co-stimulatory molecules expression, and the secretion of an appropriate set of cytokines and chemokines by DCs facilitate the differentiation of naïve CD4⁺ and CD8⁺ T-cells into effector T-helper or T-cytotoxic cells, respectively [18, 19]. DCs can, thus, simultaneously activate both a CTL response to directly kill pathogens or affected cells and a T-helper response to recruit and enhance the activity of other immune cells in the right context. This unique ability of DCs to activate the various arms of the immune system to achieve a powerful and efficient immune response makes them the most promising candidates for immunotherapeutic manipulations.

10.2.2 *Antigen Presentation Pathways in Dendritic Cells*

Three distinct pathways are used by different cell types for the presentation of antigens, namely MHC class I and class II antigen presentation, and cross-presentation.

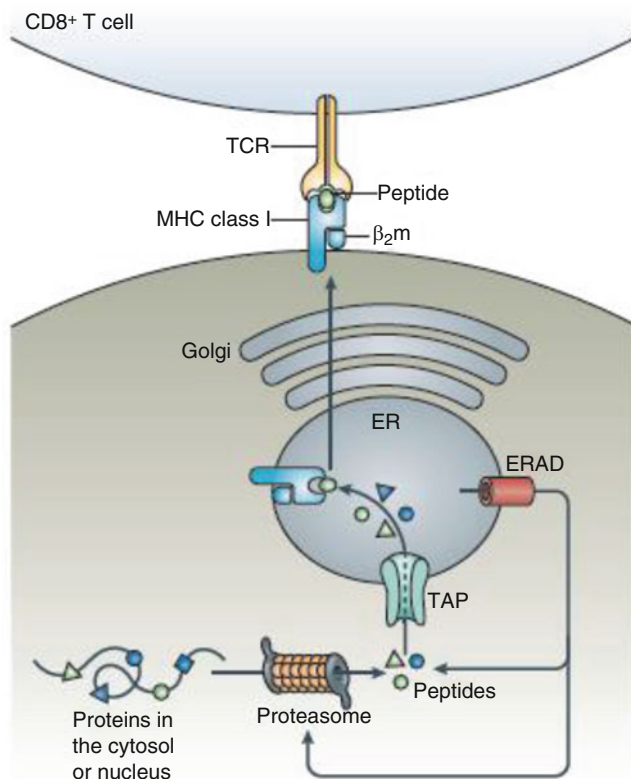


Fig. 10.1 MHC class I processing and presentation pathway. Taken with permission from Ref. 18

10.2.2.1 MHC Class I Antigen Presentation

This pathway is present in all nucleated cells, and is used to present endogenous antigens derived from the nucleus or the cytoplasm. MHC class I antigen presentation can, therefore, report intracellular bacterial or viral infections, as well as malignant transformation in the presenting cell. The resulting intracellular products are degraded in the cytoplasm by the proteasome, translocated into the endoplasmic reticulum (ER) via the transporter associated with antigen presentation (TAP), and loaded on newly formed MHC class I molecules. The MHC class I/peptide complexes are then exported along the constitutive secretory pathway to the cell membrane, where they are presented to naïve CD8⁺ T-cells. The interaction between MHC class I/peptide complexes on DCs and T-cell receptors (TCR) on CD8⁺ T-cells induce the differentiation of naïve CD8⁺ T-cells into cytotoxic T-cells that are capable of killing the infected or transformed cells (Fig. 10.1) [14, 18, 20].

10.2.2.2 MHC Class II Antigen Presentation

This pathway is restricted to APCs, including macrophages (MΦs), DCs and B lymphocytes, and is used to present exogenous antigens. The exogenous antigens are taken up into these cells by endocytosis, where they are found in early endosomes, a mildly acidic cell compartment containing a small amount of proteases. Early endosomes develop into late endosomes that are more acidic and present higher proteolytic activity, which process the captured Ag into short peptides suitable for MHC presentation [21]. The α and β chains of the MHC class II molecule are assembled in the ER and are associated with the Invariant chain (Ii) that

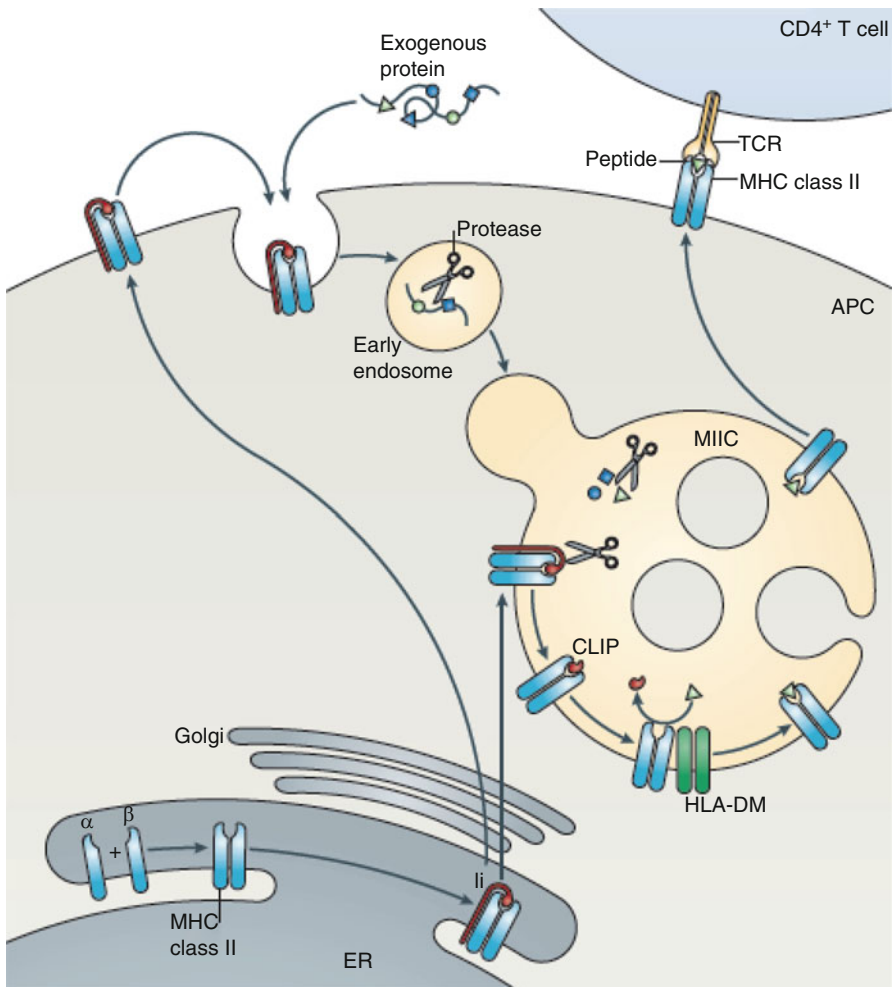


Fig. 10.2 MHC class II processing and presentation pathway. Taken with permission from Ref. 18

occupies the peptide-binding groove of the complex. The MHC class II molecule is then transported to the late endosome, also referred to as the MHC class II compartment, where, following partial digestion of Ii by cathepsins, it finally binds the exogenous peptide. The MHC class II/peptide complexes are transported to the cell membrane for peptide presentation to CD4⁺ T-cells, and induction of T-helper response to support and enhance the activity of other immune cells (Fig. 10.2) [18].

10.2.2.3 Cross-Presentation

The generation of a CTL response against virally infected or transformed cells is possible through the MHC class I pathway described above. However, other kinds of threats can be addressed by the immune system APCs through the MHC class II pathway, but this pathway exclusively generates a T-helper response. As it is obvious that not every viral infection or malignant transformation involves APCs, and that many pathogens can impair antigen presentation by their host cells, an alternative mechanism must exist that would enable APCs to generate a cytotoxic T-cell response against the various exogenous threats they encounter [22]. Such a pathway indeed exists and is a unique property of DCs. In addition to the MHC class I and MHC class II antigen presentation pathways, DCs can present exogenous antigens on MHC class I molecules to CD8⁺ T-cells. This pathway is termed cross-presentation, implying that it involves mechanisms from both the MHC class I and the MHC class II pathways [23]. Although first described more than three decades ago, the details of this pathway remain only poorly understood, with two main theories attempting to explain the mechanisms leading to MHC class I loading of endocytosed exogenous antigens. The cytosolic track suggests that the antigen is translocated from the endosome into the cytosol, where it enters the cytosolic processing pathway. The endocytic track, on the other hand, suggest that MHC I molecules are recycled from the cell membrane to the endosome, where they are loaded with antigenic peptides processed by endosomal proteases [21, 24].

10.2.3 *The Role of DCs in Anti-Tumor CTL-Response Generation*

As mentioned earlier, anti-cancer immunity requires the generation of a specific anti-tumor CTL response that, in turn, requires proper antigen presentation. As part of the many techniques tumor cells use to avoid or weaken the immune system, they can substantially reduce the expression of MHC class I/peptide complexes on their cell surface. Since tumor cells are, thus, poor Ag-presenting cells, cross-presentation of tumor antigens by DCs is essential for the generation of tumor-specific CTL responses [4]. In addition, given the unique ability of DCs to initiate and orchestrate

the various arms of the immune system, including recruitment and activation of macrophages, natural killer (NK) cells, T-helper cells and B-cells, a strong and comprehensive anti-cancer immune response can hopefully be achieved by inducing such cells to present an appropriate tumor antigen [10].

10.3 Dendritic Cell-Based Cancer Vaccination

10.3.1 DC Immunization

DC immunization requires that an appropriate antigen be presented to DCs in a proper manner and in the right immune context so as to allow Ag uptake, processing and presentation on MHC molecules, parallel to DC maturation and T-cell priming. The first method developed for DC immunization involved the use of whole tumor cells (WTC) or tumor cell lysates, either alone, mixed with an adjuvant, or genetically modified to express an adjuvant. The major advantage of this method is that no Ag identification is required, and multiple Ags are being delivered simultaneously. Nevertheless, all WTC vaccine methods showed limited efficiency in clinical trials, probably due to insufficient interaction between the tumor antigens and the DCs [25]. Pulsing DCs with antigenic proteins or peptides corresponds to another approach. Protein-pulsed DCs are capable of presenting Ags on both MHC class I and MHC class II molecules, with a long half-life of MHC class I presentation and no HLA restriction (i.e., no need for patient selection), but they require an appropriate processing of the protein by DCs, which may be difficult to achieve. Peptide-pulsed DCs, on the other hand, require no processing. Still, they show limited MHC class II presentation, a short half-life of MHC class I presentation, and a HLA restriction. Specific or total tumor mRNA can be processed and presented on both MHC class I and class II molecules. This, however, requires mRNA extraction from a tumor sample and is, thus, patient-specific [15, 26]. Lately, the delivery of tumor Ag-encoding plasmid DNA has emerged as a promising method for DC immunization [19, 27].

10.3.2 DNA Vaccines

DNA vaccines are DCs that were genetically modified to express TAAs. DNA vaccines are advantageous for several reasons. They enable the presentation of multiple epitopes of full-length TAAs on MHC class I and class II molecules, and since processing occurs within the cell, they are not HLA-restricted. In addition, efficient gene transfer allows for a continuous supply of peptides in the modified DC [10]. Genetic modification of DCs can be achieved either by ex vivo gene delivery methods, including use of a gene gun, electroporation, ultrasound and microinjection, or by in vivo approaches, including naked DNA delivery, and viral or synthetic

vectors [8, 27]. Ex vivo delivery offers many advantages. The extracted cells are cultured and undergo maturation under controlled conditions and in the absence of inhibitory signals provided by the tumor cells, such that maturation status can be determined before re-administration. Since transfection occurs in culture, high specificity is achieved as only the selected cells are transfected. Nevertheless, ex vivo gene delivery carries substantial limitations. The process is laborious, costly and time-consuming [12]. Patients must undergo cytophoresis, followed by culturing and maturation of the acquired cells, steps that must be performed for each patient separately. Reproducibility is very low and different quality control methods are used. Thus, only a limited number of patients would be expected to benefit from this approach. Hence, ex vivo gene delivery is not likely to become widely marketable. Moreover, transfected mature DCs can show poor distribution from the injection site and ex vivo maturation can impair DC trafficking to lymph nodes, a critical requirement for cross-priming [28, 29]. In contrast, in vivo gene delivery may be expected to become “off the shelf” therapy. As a single product suitable for all patients, in vivo gene delivery can be produced on a large scale with lower costs. Simple and uniform manufacturing and product control procedures will enable reproducibility and control over product quality. In addition, DCs can be targeted at different sites and in their natural environment, thereby not impairing their natural course of maturation and activation [25, 29]. For these reasons, in vivo gene delivery is considered by many to be the best strategy for Ag delivery into DCs [8, 10, 19].

10.3.3 In Vivo Gene Delivery

In vivo gene carriers must meet important requirements. First, they should be able to incorporate their plasmid DNA cargo into the core of the nanoparticle and be stable enough to carry such cargo in the circulation and protect it from degradation. Second, they should be able to selectively target the desired cell type and be properly internalized. Third, they must facilitate escape from the endosome, cytoplasm trafficking, nuclear transport and DNA unpacking [30]. To date, the most efficient gene delivery systems are viral vectors. Using small amounts of DNA, viral vectors can induce high transfection efficiency and stable, long-term gene expression. Unfortunately, viral vectors possess some serious safety issues, including toxicity, immunogenicity and oncogenicity, with numerous clinical trials having been terminated because of this [15, 31, 32]. Restricted gene size is another major limitation of viral vectors. Synthetic vectors correspond to cationic lipids or cationic polymers, respectively termed lipoplexes or polyplexes, which electrostatically bind the negatively charged DNA. The main drawback of such vectors is their low transfection efficiency, as compared to viral vectors. Yet, synthetic vectors are simple, safe, easy to manufacture on a large scale and can carry plasmids of unrestricted size. Moreover, they can be easily modified to possess desirable properties, including targetability, serum stability, reactivity to external signals and an ability to stimulate

an immune response. In fact, cationic polymers themselves may provide immune stimulation and mimic nuclear localization signals (NLS), facilitating nuclear transport of their cargo [9, 30].

10.4 Polymer-Based Systems for DC-Targeted Gene Delivery

Various polymeric carriers for DNA and RNA delivery have been developed in the last decade as alternatives to viral vectors. Cationic polymers with free primary, secondary and/or tertiary amines were the earliest DNA carriers investigated as transfection reagents in mammalian cells. Such polymers can condense large genes into smaller structures, protect DNA from enzymatic degradation, and can mask the negative charges of DNA, a prerequisite for successful transfection of most types of cells. Some cationic polymers, such as poly-(L-lysine) (PLL), L-polyethylenimine (L-PEI) and chitosan, are linear, while others, like B-PEI and polyamidoamine (PAMAM) dendrimers, are highly branched chains [33]. Branched cationic polymers, such as PEI and PAMAM, exhibit high transfection efficiency in mammalian tissues. Micro- and nanoparticulate systems which adsorb or encapsulate oligonucleotides or genes, based on, for example, poly(lactide-co-glycolide) (PLGA), polycyanoacrylate, polyorthoesters, gelatin, alginate or chitosan, are also under investigation as sustained release matrices for genetic drugs. Some of the most important polymer-based systems utilized for DNA delivery into APCs are discussed below.

10.4.1 Poly(L-Lysine)

Poly(L-lysine) (PLL) was the first cationic polymer developed for gene delivery. PLL is a linear polypeptide presenting L-lysine residues in repeat units (Fig. 10.3c). This first generation cationic polymer bears ϵ -amino groups that electrostatically bind the negatively charged nucleic acids, and interact with the negatively charged cell membrane. The large number of active functional amine groups allows for easy modification with targeting ligands [34]. However, the transfection efficiency of PLL is very low because it cannot mediate escape from the endosomal compartment and release into the cytosol. To overcome this barrier, PLL is usually used in combination with chloroquine, although use of this agent is limited due to its cytotoxicity [35]. Another disadvantage of PLL polyplexes is that transfection efficiency is significantly influenced by serum, probably due to the rapid binding to negatively-charged serum components [36]. Furthermore, in vivo applications of PLL polyplexes are complicated by a high level of cytotoxicity [37] and lack of in vivo stability [38, 39]. Because of low transfection efficiency, imidazole groups ($pK_a \sim 6.5$) were introduced into PLL to improve buffer capacity, thereby, enhancing transfection efficiency [40]. PLL with high imidazole content mediated high

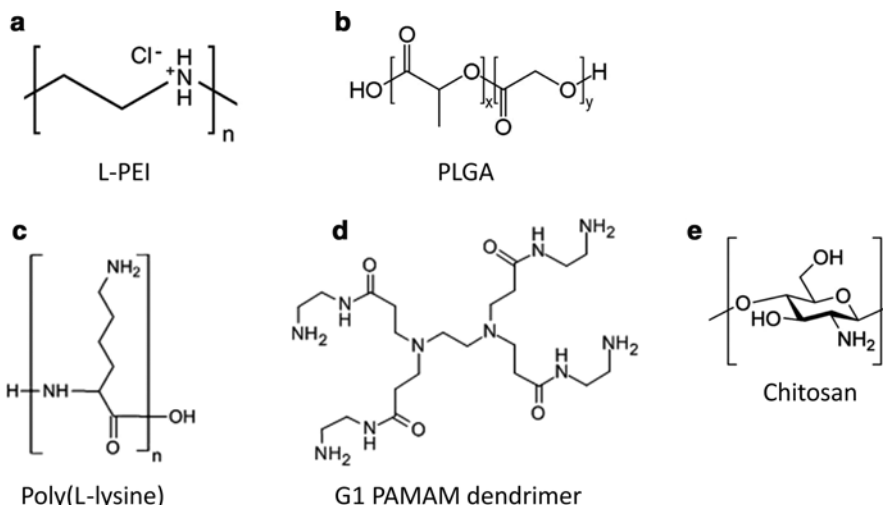


Fig. 10.3 Structures of (a) linear PEI, (b) PLGA, (c) poly(L-lysine), (d) first generation (G1) PAMAM dendrimer and (e) chitosan

gene transfection efficiency, with gene expression levels close to those of PEI (discussed below), and exhibited low cytotoxicity. The PEGylation of PLL can also improve its cytotoxicity profile [36].

Finally, whereas it has been found that PLL/DNA complexes can be taken up by DCs, PLL/DNA did not alter DC phenotype through surface marker expression [41]. Complexation of plasmid DNA encoding for chicken egg ovalbumin (OVA) with PLL-coated polystyrene (PS) particles induced high levels of CD8⁺ T-cells as well as OVA-specific antibodies in C57BL/6 mice, and further inhibited tumor growth after challenge with OVA expressing tumor cells [42]. PLL-based microspheres displaying mannan or mannoside-modified surfaces (for targeting the mannose receptor, discussed below) were readily phagocytosed by both DCs and macrophages, however neither surface-assembled mannan- nor mannoside-modified microspheres could stimulate DC maturation [43]. Thus, despite the promising results shown in early studies, PLL is not likely to find clinical applications [30].

10.4.2 Poly(ethylenimine)

Poly(ethylenimine) (PEI) is a second generation and one of the most useful polycations for gene delivery [44]. Linear PEI has only secondary amino groups that are almost all protonated under physiological conditions. Branched PEI presents not only primary and secondary amines but also tertiary amines. As such, only about

two-thirds of the amino groups in branched PEI are protonable under physiological conditions. The transfection efficiency of PEIs depends on the molecule weight, the PEI nitrogen/DNA phosphate charge ratio (N/P) and the cell type. As every third position on the polymer backbone is occupied by a protonable amino group (Fig. 10.3a), its cationic charge density is very high, allowing for condensation of the negatively charged DNA. Under physiological conditions, only about 20 % of the PEI nitrogen atoms are protonated, leaving the other 80 % to facilitate the important step of endosomal escape by the so-called “proton sponge effect” [30, 45, 46]. Thus, PEI is the most popular and most effective polymeric transfection reagent cited to date [47]. However, the positive charge of PEI/DNA polyplexes cause some serious problems, including adsorption to cells and negatively charged blood components, recognition by the immune system components, resulting in rapid clearance from the circulation, and cytotoxicity to non-target cells [44, 48]. To overcome these limitations, PEI has been conjugated to hydrophilic polymers (i.e., poly(ethylene glycol) (PEG) [49], or hyaluronic acid (HA)) of different molecular weights [50]. PEG is widely used in drug and gene delivery systems to shield charged, immunogenic or toxic segments, resulting in less toxic “stealth” particles that can evade the immune system and, thus, avoid rapid clearance from the circulation [51, 52]. The conjugation of PEG to PEI was previously shown to increase polyplex solubility and serum stability and reduce cytotoxicity by shielding the high positive charge of PEI [49]. PEGylated PEI, however, showed lower transfection efficiency, as compared to PEI/DNA complexes, and also lacked cell-specificity [53]. Ligation of PEI/DNA complexes to molecules targeting DC uptake receptors can significantly increase transfection efficiency.

Approaches for targeting DC receptors have generally involved either natural receptor ligands or the use of antibodies raised against specific receptors. Targeting DC cell surface receptors may also provide cell activation signals [4, 54]. Different DC receptors have been used to facilitate targeted delivery of gene and drug carriers into DCs, including β -integrins (CD11b and CD11c) [55], CD40 [56] and the Fc receptor [57], but the most studied proteins for this purpose are members of the C-type lectin receptor (CLR) family [58]. CLR is a family of receptors sharing structural homology in their carbohydrate recognition domain (CRD), where specific sugar residues are bound in a calcium-dependent manner. One of the CLRs, the mannose receptor (MR, CD206), is the most widely used receptor for targeting DCs, using both mannose and mannan [59]. While several studies demonstrated that mannosylated-PEI conjugates are effective in gene delivery via MR [60–62], their transfection potential for primary human and mouse DC was found to be rather low [60]. This might be due to the low affinity of carbohydrate ligands to their receptors, which is usually in the low millimolar range [63], thus, limiting in vivo transfection efficiency. The eight adjacent CRDs in MR may help to increase the binding affinity and specificity of polyplexes containing mannosylated glycans in a multivalent display [59, 64]. We have recently described the design of multivalent mannosylated PEI/DNA complexes bearing mono- and trivalent mannose as a ligand for targeting MR-positive DCs [65]. Complexes bearing mono- and trivalent mannose (Man-PEG-b-PEI/DNA and Man₃-PEG-b-PEI/DNA, respectively) were safe and demonstrated significantly higher in vitro transfection efficiency in DCs. The mannosylated

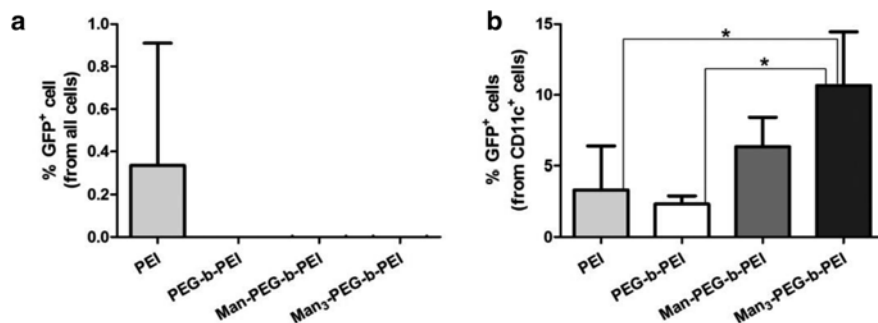


Fig. 10.4 In vivo uptake of mannosylated-PEI/DNA complexes by cells in the draining lymph nodes of C57/BL6 mice 24 h after subcutaneous injection. Results represent the percentage of GFP⁺ cells in the entire cell population (a) and in the CD11c⁺ cell population (b). Taken from Ref. 65

complexes were injected into the tail base of C57/BL6 mice, and 24 h post-injection, the percentages of GFP positive cells in the entire cell population and in the CD11c⁺ cells taken from the inguinal lymph nodes were measured. When examining the entire cells population extracted from the lymph node, only PEI/DNA complexes showed detectable GFP activity, but no significant change was observed between the treatments (Fig. 10.4a). However, Man₃-PEG-*b*-PEI/DNA was significantly more efficient in transfecting CD11c⁺ DCs collected from inguinal lymph nodes, as compared to polyplexes prepared with PEG-*b*-PEI/DNA or PEI/DNA (Fig. 10.4b).

One challenge of targeting the MR, as holds true for all of the other above-mentioned receptors, is that they are not exclusively expressed on DCs, and are also on several other cell types [57]. MR, for example, is also expressed on monocytes, subsets of endothelial cells and tumor-associated macrophages [9]. Furthermore, the synthesis of carbohydrate ligands and analogs, especially multivalent or complex carbohydrates, often requires many time-consuming, low yielding steps [66]. The stereochemistry of such carbohydrate ligands is difficult to control, and the products are difficult to purify. A frequently used alternative for carbohydrate ligands are antibodies raised against DC receptors (i.e., integrin CD11c/CD18 [55], Fc receptors [67], DEC-205 [7], DC-SIGN [68] and MR [69]), but these, even when humanized, may still elicit adverse immune responses that decrease the efficiency of treatment and induce auto-immune side-effects [70].

The use of peptide ligands remains, nonetheless, a promising method to target DC receptors. Such ligands are easily synthesized, do not possess immunogenicity, and show high selectivity to DCs (Table 10.1). Accordingly, peptide ligands that could serve as DC-targeting moieties have been sought.

A DC3-nona-arginine fusion peptide (DC3-9dR) that binds nucleic acids by electrostatic interactions was previously exploited for the specific delivery of siRNA to DCs in vitro and in vivo [78] and was also used to silence immunosuppressive molecules in DCs so as to induce strong human T-cell immune responses [79]. The DC3 targeting peptide (Table 10.1) was recently studied by our group to mediate the specific delivery of PEGylated-PEI/DNA polyplexes into DCs [80]. Polyplexes

Table 10.1 Examples for peptide ligands used for targeting DCs

Peptide name	Amino acid sequence	Characteristics	Reference
C-GRWSGWPADL-C	C-GRWSGWPADL-C	A circular peptide that binds to human CD11c/CD18, which shares homology with the D4 domain of intracellular adhesion molecule (ICAM)-1	[71]
APEDNGRSFS	APEDNGRSFS	Derived from ICAM-1, and shares homology with C-GRWSGWPADL-C	[72]
DC3	FYPSYHSTPQRP	Identified using a phage display peptide library. Binds specifically to CD11C+ cells but not monocytes, T and B lymphocytes, or NK, endothelial or fibroblast cells. The cognate receptor for DC3 on the DC surface is unknown	[73]
P-D2	VTLTYEFAAGPRD	Derived from the Ig-like domain 2 of intercellular adhesion molecule 4 (ICAM-4)	[74]
TP	TPAFRYS	Identified by phage display method. The counterpart receptor for TP on BMDC surface is unknown	[75]
NW	NWYLPWLGTDNDW	Identified by phage libraries on monocyte-derived immature DCs (iDCs)	[76]
Pan HLA DR-binding peptide	PADRE	MHC class II-binding pan-DR peptide, acts directly on iDCs to induce differentiation into mature DCs, with potent T cell-stimulating capacity	[77]

show significant transfection efficiency in DCs when decorated with DC3 peptide but not in endothelial cells. The transfection efficiency observed was higher than that of PEI/DNA, signifying the potential use of DC3-bearing polyplexes for immunotherapy via DCs.

10.4.3 Polyamidoamine (PAMAM) Dendrimers

Polyamidoamine (PAMAM) dendrimers are hyper-branched, symmetrical, flexible and monodisperse polymeric molecules (Fig. 10.3d). Szoka et al. firstly investigated PAMAM cascade polymers as non-viral gene delivery vectors [81]. These polymers have an ammonia initiator core and amido-amine repeat units of different

generations (G2–G10). Their stepwise synthesis results in products that are uniform in terms of structure and size [82] and bear a well-defined number of primary and tertiary amines. The surface primary amines are protonated, resulting in an extremely positively charged surface that enables DNA binding in stable complexes, and interaction with the cell membrane. The tertiary amines in the interior of the polyplexes are protonable, endowing the dendrimers with pH buffering capacity that enables endosomal escape of the polyplexes following cellular uptake [83]. All these properties, in addition to their non-immunogenic nature, make PAMAM dendrimers an alternative to the highly efficient and highly immunogenic viral vectors. However, a major drawback of PAMAM dendrimers is that low generation dendrimers (G5 or lower) show poor transfection efficiencies, while dendrimers of high generation (G6 and higher) are efficient transfection agents, yet possess serious cytotoxicity [82]. In addition, synthesis of high generation dendrimers is a high cost, labor-consuming process that last several days, making them less likely to be designed for large scale production [83].

Mannosylated G4-PAMAM dendrimers conjugated to OVA specifically targeted DCs and induced cross-presentation in vivo [84]. Moreover, pre-immunization with mannosylated PAMAM-OVA leads to delayed onset of B16-OVA melanoma development, slower kinetics of tumor growth and increased survival of OVA-immunized mice. G4-PAMAM dendrimers bearing DC-SIGN ligand in multivalent presentation also achieved efficient DC targeting properties, however, did not affect DCs maturation [20]. It has been postulated that mannosylated dendrimers, as opposed to DC-SIGN-modified dendrimers, may trigger not only DC-SIGN, but also other mannose-specific CLRs that contributes to DC maturation and activation. G5-PAMAM dendrimers conjugated to MHC class II-targeting peptide (PADRE, Table 10.1) and surface-loaded DNA have been shown to effectively transfect murine and human APCs in vitro [85]. When applied subcutaneously, this conjugate preferentially transfected DCs in draining lymph nodes, promoted generation of high affinity T-cells, and elicited rejection of established B16 tumors.

10.4.4 Chitosan

Chitosan is a linear polysaccharide (Fig. 10.3e) obtained by deacetylation of its parent polymer chitin, a compound that is widely distributed in nature. Chitosan is composed of randomly distributed β -(1-4)-linked D-glucosamines (deacetylated units) and N-acetyl-D-glucosamines (acetylated units) and has an apparent pKa value of 6.5 [86]. Chitosan is a biodegradable and biocompatible polymer, with low or no immunogenicity and antibacterial activity [86, 87]. It was previously shown to be non-toxic in both test animals [88] and humans [89]. Due to these properties, chitosan is attractive for drug and gene delivery. The high density of positive charges along the polymeric chain contributes to the condensation of the negatively charged DNA, and more frequently, siRNA molecules, into compact structures, thus protecting them from degradation by blood nucleases, promoting

their cellular uptake [90]. Chitosan also possess adhesive properties by interacting with glycoproteins in the mucus [91], and is thus used in muco-adhesive drug delivery systems as an adsorption enhancer [92]. DNA plasmid incorporated into chitosan nanoparticles was able to induce DCs maturation and increase IFN- γ secretion from T cells after pulmonary mucosal immunization [93]. However, unless modified, chitosan has low solubility at physiological pH, which significantly limits its use in many applications [87, 92]. Chemical modifications of chitosan have been performed to improve its solubility at physiological pH, including PEGylation or quaternization of the amine groups of chitosan. However, the transfection efficiency of chitosan-based derivatives reported so far is generally not superior to that of PEI [94].

Various DC-targeted chitosan-based DNA delivery formulations has been designed to overcome the major obstacles facing the clinical development of chitosan, namely, the lack of cell-specificity and low transfection efficiency. Mannosylated chitosan/DNA complexes were more efficient in transfecting DCs, as compared to water-soluble chitosan/DNA, and induced better IFN- γ production from DCs [95]. Mannosylated-chitosan-entrapping PEI/HBV-DNA complexes induced significantly enhanced serum antibody production and CTL levels after intramuscular immunization [96]. Biotinylated chitosan nanoparticles were modified with bifunctional fusion protein (bfFp) vectors for achieving DC-selective targeting. bfFp is a recombinant fusion protein consisting of truncated core-streptavidin fused to an anti-DEC-205 single chain antibody (scFv). Intranasal administration of plasmid DNA-loaded bfFp/chitosan nanoparticles, along with anti-CD40 DC maturation stimuli, enhanced the amount of mucosal IgAs, as well as systemic IgGs, against nucleocapsid (N) protein of severe acute respiratory syndrome coronavirus (SARS-CoV) [97]. Finally, to improve low transfection efficiency, chitosan-linked-PEI/DNA complexes have been designed, and showed high transfection efficiency and low cytotoxicity towards DCs [98]. Vaccination with DCs transfected with chitosan-linked-PEI/DNA encoding gp100 (melanoma-associated antigen) slightly improved resistance to the B16BL6 melanoma challenge.

10.4.5 Micro- and Nano-Particulate DNA Delivery Vectors

Polymeric particulates have been shown to be efficient in delivering plasmid DNA into APCs (reviewed in [99–101]). A key advantage of particulate vectors relative to other non-viral gene delivery systems is their superior in vivo stability. The principal types of polymers studied in this context include those made of poly(lactide) (PLA; reviewed in [99]), poly(lactide-co-glycolide) (PLGA; reviewed in [99, 100, 102]), polyorthoesters [103], polystyrene (PS) [104, 105] and poly(ϵ -caprolactone) [106]. Of these, PLGA has been studied most extensively in terms of its capacity to stimulate APCs.

10.4.5.1 Poly(lactide-co-glycolide)

Poly(lactide-co-glycolide) (Fig. 10.3b) is an FDA- and European Medicine Agency-approved polymer for drug delivery systems for parenteral administration [107]. This biocompatible and biodegradable polymer slowly degrades *in vivo* by hydrolysis, and its byproducts (lactic and glycolic acid) are easily metabolized and excreted. DCs and macrophages appear to have high affinity for PLGA particles, as a high level of internalization of antigen-loaded particles has been demonstrated in both *in vitro* and *in vivo* settings [108, 109]. PLGA microspheres carrying protein antigens or antigen-encoding plasmid DNA are capable of eliciting potent antigen-specific immune responses [100, 110, 111]. PLGA also affected expression maturation markers and cytokine production in DCs [112]. This capacity appears to be driven by unique physical features of PLGA particles (i.e., surface charge, shape and the rate of polymer degradation [100]). The mechanisms leading to maturation induction in DCs, however, remain unclear. With respect to their stability in PLGA particles, antigen-encoding plasmid DNA offers advantages over protein-based immuno-modulators, which can lose their biological activity in response to small changes in their tertiary and quaternary structures during formulation.

DNA delivery into DCs can be improved by using ligand-decorated PLGA particles. Mannose-grafted PLGA nanoparticles lead to a significant enhancement of OVA accumulation in the inflamed colon compared to the healthy one, underlining the benefit of active targeting of macrophages and dendritic cells in diseased tissues [113]. Mannan-decorated OVA-loaded PLGA nanoparticles simultaneously enhanced antigen-specific CD4+ and CD8+ T-cell responses in vaccinated mice [114]. Modification of PLGA micro-particles with P-D2 peptide (Table 10.1) significantly improved DC antigen presentation *in vitro*, and increased the rate and extent of microsphere translocation by DCs and macrophages *in vivo* [110].

PLGA has been also utilized in combination with other cationic polymers for gene expression in DCs, with [115] or without targeting ligands [116, 117]. PLGA scaffolds encapsulating PEI/DNA and granulocyte-macrophage colony-stimulating factor (GM-CSF) led to a significant increase in gene expression, and high levels of expression that persisted for a period of time [116]. Yet, since PLGA nanoparticles are hydrophobic in nature, they tend to form aggregates that reduce the efficiency of the system [108]. In addition, PLGA particles are opsonized by the immune system components, and degraded before reaching their destination. Another drawback of PLGA for delivery of genetic vaccines stems from their sustained release property [102]. PLGA particles release their cargo very slowly, over days or even weeks, but *in vivo* they may be exocytosed from the cell or degraded in the lysosome over much shorter periods of time, before sufficient amounts of cargo are released [35]. Moreover, most DCs die within 7 days after activation and migration to draining lymph nodes [118], hence even fast degrading PLGA systems, which fully release the encapsulated DNA within few weeks, cannot meet with the rapid release kinetic criteria and thus fail to induce high levels of target gene expression [108]. Finally, hydrolysis of PLGA leads to low pH within the particle and thus to DNA degradation [102]. For these reasons, despite the promising results obtained with animal models [114] and clinical trials [119, 120], PLGA has been of limited use in this sense [121].

10.5 Conclusion

The delivery of antigens into DCs carries tremendous potential for immune modulation, and specifically, for cancer immunotherapy manipulations. Various polymeric nanomedicines for DNA delivery, ranging from linear homopolymers to block copolymers, branched polymers, as well as combinations of different types of polymers, selected to reduce cytotoxicity and facilitate endocytosis of particles into DCs, are now being routinely examined for their ability to modulate DCs and macrophages. Ultimately, DNA delivery systems that can also induce DC maturation and activation would be advantageous. The main challenges for successful *in vivo* gene transfer into DCs are the cytotoxicity associated with many cationic polymer gene carriers, the lack of cell specificity and relatively low transfection efficiency, when compared to viral vectors. Together with the fact that DCs compose only 1–3 % of cells in peripheral tissues, such gene systems will have limited success unless targeted. Given the number of targeting molecules, immune-modulatory agents, chemokines, growth factors and antigens that can be considered for DC-specific delivery, large numbers of potentially useful formulations for DC manipulation are available. Targeting DCs *in vivo* with tumor antigen-encoding plasmid DNA can elicit effective and long-lasting tumor antigen-specific immunity, with minimal inconvenience to the patient. Furthermore, DNA delivery formulations that are stable for extended periods of time and which can enhance antigen presentation on both MHC-I and MHC-II molecules are preferable. With respect to clinical translation, efficacious non-viral gene delivery into DCs will depend on the combination of intelligent material design, the appropriate tumor specific antigen-encoding DNA and immuno-stimulatory molecules to promote DC maturation and activation.

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Chapter 11

The Use of Silk in Nanomedicine Applications

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Abstract Biopolymers made up of silk proteins have been used in numerous drug delivery applications and represent an excellent source of natural biomaterials. In particular silk fibroin has proved valuable as a building block for nanomedicines and drug delivery implants, owing to its favorable biocompatibility, degradation, stabilization and controllability. In this chapter we will discuss the various sources of silk biomaterial and how this naturally occurring biopolymer has been utilized in the development of nanomedicines and implantable drug delivery systems, demonstrating how silk is a unique biological template which has opened up many possibilities for the generation of functional biomaterials and drug delivery systems in a green and cost-effective manner.

Keywords Silk • Polymers • Silk fibroin • Nanomedicine • Drug delivery • Silk biomaterials • Nanoparticle

11.1 Introduction

Since the lucrative trade of silk began along the Silk Road during the Chinese Han Dynasty (206 BC–220 AD), it has been regarded as a valuable commodity. Spiders and silkworms produce silk protein fibers that have unmatched mechanical strengths in comparison to synthetic biomaterials, and in addition to wide-ranging applications in textiles, silk proteins and fibers have also been utilized in biomedical research for the development of nanomedicines and drug delivery scaffolds. Silk is a non-toxic biomaterial that can degrade at favorable rates for drug delivery

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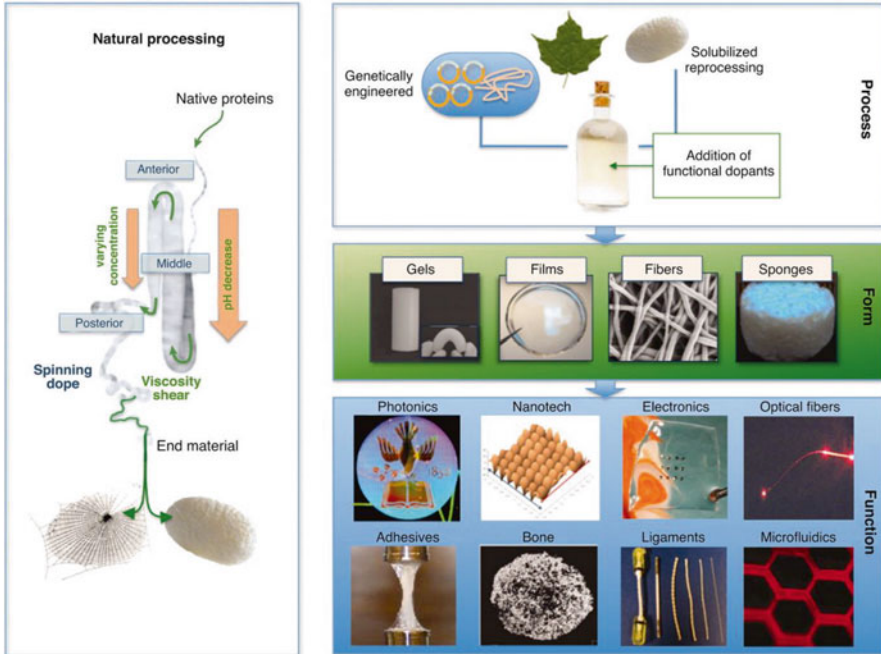


Fig. 11.1 Silk generation and materials development. *Left*: process of natural silk production. *Right*: silk produced from reconstituted native silk proteins or genetically engineered silk, and the variety of silk materials and technology platforms generated from silk. Figure taken from Omenetto et al. [12]

applications, and has ideal mechanical and chemical properties [1], which have enabled its use in a range of wound dressings and sutures [2–11]. Silk proteins have led to the development of new materials, technology platforms and functional devices (Fig. 11.1) [12]. In addition to the development of macroscale products, silk polymers have also been used as building blocks for the formulation of a range of nanomedicines with biodegradability and controlled release properties. In this chapter we will firstly discuss the origins and types of silk biopolymers, and then explore the uses of silk and silk-like products for the development of nanomedicines and healthcare products such as drug depots and implants.

11.1.1 Spider Silk

Orb weaving spiders, such as *N. calvipes* and *E. australis* spin complex webs to catch flying prey [13]. These spiders have seven different spinning glands with each producing a silk with different physical properties and functions; such as pray wrapping, web construction, and coating of egg cocoons [13]. The major ampullate gland produces dragline silk which is important in web construction, has the highest mechanical strength and has been the most extensively studied [14].

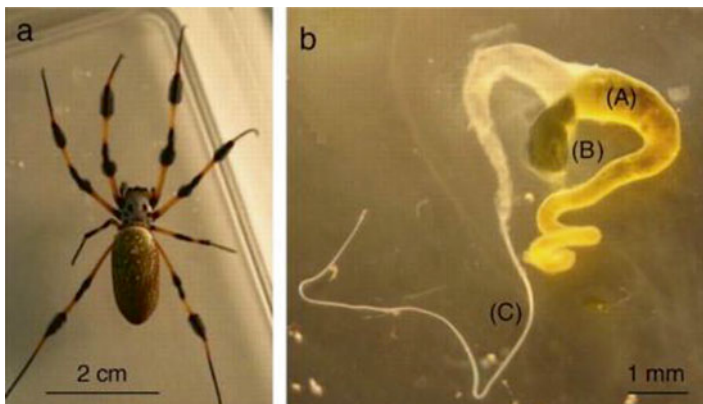


Fig. 11.2 Orb weaving spider and its ampullate. (a) Adult female *N. clavipes* (golden-orb) spider, (b) dissected major ampullate gland (A) tail (B) sac (C) duct. Figure taken from Tokareva et al. [18]

The spider silk gland contains a tail, sac and duct region (Fig. 11.2). The two major proteins produced in this gland are known as major ampullate silk protein 1 and 2 (MaSp1 and MaSp2) [14]. These are relatively large proteins with molecular masses between 260 and 350 kDa [15], with both proteins possessing a polar N- and C-termini which flank long repeating hydrophobic sequences with small interspersed hydrophilic sequences [16]. The central region of MaSps contain many repeats of certain short amino acid sequences such as; alanine and glycine repeats (poly(A), poly(GA) and GGX), proline containing regions (GPGGX/PGGQQ), and short hydrophilic spacers [17–19].

The central region of MaSps contain many sequences of short alanine and glycine repeats (poly(A), poly(GA) and GGX), proline containing regions (GPGGX/PGGQQ), and short hydrophilic spacers [17–19]. The alanine and glycine repeats have the potential to form anti-parallel beta-sheets, while the proline-containing motifs form beta-spirals [17, 20]. The N- and C-termini form dimers which link individual fibroins [20–24]. The secondary structure is determined not only by the amino acid sequences, but also by their location and the pH and ionic environment of the major ampullate gland [25]. Silk proteins are synthesized in the tail and stored in the sac. During spinning, silk is pulled through the three limbs of the duct and exits at the spigot [26]. Silk is stored in the sac as a concentrated aqueous solution (30–50 % w/v) [25], here, it lacks ordered secondary structure and appears to be randomly coiled and oriented [27]. The pH of the aqueous silk solution drops from about 6.9 in the ampulla to 6.3 by the third limb of the duct, and the sodium and chloride ion concentrations decrease while potassium, phosphorus and sulfur ion concentrations increase [25, 28, 29]. The combination of changes in pH and ionic strength and the shear forces in the narrow ducts, cause the conversion of disordered turns and helices of the liquid silk into the ordered beta-sheet conformation of spun silk fiber [25]. The dimerization of C-termini and the pH-dependant dimerization of N-termini is also an essential step in the formation of fibers [21–24]. The secondary structure of spun major ampullate silk is approximately 40 % beta-sheets which are

oriented parallel to the fiber axis [30, 31]. This structure is responsible for the exceptional mechanical strength of spider silk fibers [20, 26, 32].

11.1.2 Silkworm Silk

Silkworm larvae form a protective cocoon during their transformation into an adult moth. These cocoons contain a mixture of silk proteins and have been harvested for hundreds of years. Silk is produced commercially from the cocoons of *Bombyx mori* (*B. mori*) [25]. Silk fibroins from *B. mori* are composed primarily of two peptides made of a heavy chain (300–400 KDa) covalently attached to a light chain (25 KDa) [33–35]. The structure of silkworm fibroins and the associated spinning process is similar to that of spiders [36]. The heavy chain fibroin contains 12 hydrophobic blocks of around 400 amino acids made up of small glycine and alanine-rich repeats (GAGAGS, GX, GAAS) [11, 37]. Between these blocks are hydrophilic spacers of about 30 amino acids and hydrophilic C- and N-termini flank the sequence (Fig. 11.3) [11, 37]. The heavy chain-light-chain dimer also associates non-covalently with a protein called P25 (30 kDa), forming a complex in a 6:1 ratio (dimer: P25) [33, 35, 38]. These hydrophobic proteins form insoluble fibers which make up 70–80 % of the protein in the cocoon, while the rest of the cocoon is made up of a group of proteins known as sericins [11, 39]. These are hydrophilic proteins which act as glue holding the fibroin fibers together, and have molecular masses up to 400 kDa [16, 44]. Similar to spider silk, the secondary structure of silkworm-silk is dependent on both the amino acid sequence and the pH and ionic environment of the silk gland [41, 43].

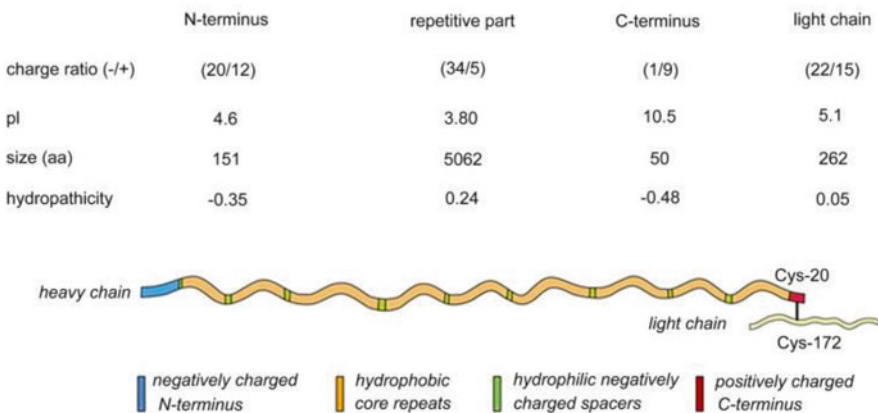


Fig. 11.3 Illustration of the size, charge and hydropathicity of silk fibroin from *B. mori*. The heavy and light chains are linked by one disulfide bond between Cys-20 and Cys-172. Figure modified from Lammel et al. [40]

Silk fibroin is synthesized in the posterior part of the silk gland and transported to the middle of the gland where sericin is added [41, 43], and as silk travels through the anterior part of the silk gland, silk fibers are formed [36, 38]. Much like spider silk, the fibroins are stored in a concentrated, aqueous state with randomly coiled secondary structure in the posterior and middle part of the gland [41, 45]. As they are released through the anterior part of the gland and spinning duct, the fibroins are transformed into an insoluble fiber dominated by beta-sheets [36, 41]. The shear stress caused by the decreasing diameter of the duct, the changes in ion composition and the drop in pH are thought to be responsible for the formation of silk fibers [42, 43]. These factors contribute to the removal of water and contraction of the structure which is necessary for the alignment of anti-parallel beta-sheets that lead to the formation of silk fibers [41–43].

11.1.3 Isolation of Silk Proteins

The majority of silk fibroin comes from silkworms and is typically isolated from the cocoons of *B. mori* [44, 45]. The cocoons are boiled in a dilute aqueous solution of NaCO_3 (around 0.02–0.05 M) for at least 15 min [44, 46–51]. This dissolves the hydrophilic sericins and leaves undissolved silk fibroins which are rinsed with deionized water and dried, followed by solubilisation in an aqueous solution, often while being heated, using a concentrated lithium salt (9–10 M) or a concentrated mixture of CaCl_2 , ethanol (or methanol) and water (1:2:8 molar ratio). The salts stabilize the silk in aqueous solution which is then dialyzed into deionized water to remove excess salts [44, 46–51]. The resulting aqueous silk is made up of disordered and partially digested silk fibroin proteins with molecular weights ranging from 8 to 70 kDa [50, 52].

Other methods of isolating silkworm-silk fibroins involve dissection of the silk gland prior to cocoon formation [48, 53]. The posterior silk glands are removed and rinsed with deionized water to remove traces of sericin. Tubes of the gland are squeezed to expel protein which is then dissolved in sodium dodecyl sulfate and dialyzed into deionized water [48, 53]. Generally, the farming and isolation of spider silk is challenging due to the cannibalistic nature of spiders and, therefore, the large-scale production of spider silk by natural means is not feasible [54]. Other attempts to produce spider silk involve the expression of recombinant silk genes in *E. coli*, yeast, mammalian cells, transgenic plants, and even transgenic silkworms [55–59].

11.2 Silk in Nanomedicine Applications

Nanomedicine involves the application of nanotechnology to medicine, whereby nanoscale structures are developed for medical applications such as drug delivery, controlled drug release, and imaging. The following section discusses recent work

regarding the formation and potential use of silk-based nanoparticles for biomedical applications. To-date, silkworm silk and recombinant spider silks have been used as the core of nano- and micro- particles, as coatings for nanoparticles, or blended with nanoparticles or other proteins and polymers due to their enhanced biophysical and mechanical strengths [46, 60–63]. Nanoparticles made from silk fibroin have the potential to increase the efficacy of a range of therapeutics and to minimize their adverse effects.

11.2.1 Preparation of Silk Nanoparticles

Many different techniques have been successfully used to form silk nanoparticles and microparticles with the simplest being nanoprecipitation (also known as solvent displacement) and desolvation [35, 46, 60]. These techniques change the solvent conditions around solvated silk proteins to induce the precipitation of silk as nanoparticles. Like in the formation of silk fibers, precipitation of aqueous silk into nanoparticles corresponds with an increase in the beta-sheet content and decrease in the random coiled content [52, 53]. Nanoprecipitation involves the drop-wise addition of aqueous silk solution into a larger volume of miscible organic solvent, such as acetone or ethanol [60]. Desolvation is the opposite process in which a miscible organic solvent or a high ionic strength solution, such as 2 M PO₄, is added to an aqueous silk solution, up to a certain concentration using dialysis or mixing [40, 46, 64]. Bulk nanoprecipitation can yield particles with diameters between 40 and 170 nm and desolvation with organic solvents has achieved diameters around 150 nm [46, 50, 53, 65, 66]. Desolvation with concentrated aqueous salts avoids the use of organic solvents, but leads to larger particles with the smallest so far reported being around 330 nm [40, 67].

Solution-enhanced dispersion by supercritical CO₂ is another notable method of preparing silk nanoparticles [68, 69]. Using this technique, nanoparticles with diameters as small as 50 nm have been made. In this method, isolated *B. mori* silk is dissolved in 1,1,1,3,3,3-hexafluoro-2-propanol and pumped under high pressure in combination with supercritical CO₂ into a high-pressure vessel forming nanoscale droplets leading to nanoparticles. Microfluidics is another promising new method of nanoparticle formation in a more controlled and reproducible manner [70, 71]. Mintopolus et al. were able to make silk fibroin spheres as small as 210 nm. Here, a stream of aqueous silk combines with a stream of poly(vinyl alcohol) to form tiny monodispersed drops, allowing for the silk to precipitate as nanoparticles [72]. Other methods of making silk nanoparticles and microparticles include spray drying, laminar jet break-up using sound waves and electrospinning dilute solutions under low voltage [35]. Loading of drugs into silk nanoparticles is typically accomplished by mixing the cargo with the silk protein prior to, or during the formation of nanoparticles [65, 66]. Particles can also be loaded by charge-based association [64]. Silk particles bear a negative charge at physiological pH, therefore, drug loading post formulation is limited to cationic species [35].

11.2.2 *Silkworm Silk-based Nanoparticles*

The majority of nanoscale silk research has involved nanoparticles made from the silk of silkworm cocoons. This section highlights particles loaded with drugs and small molecules, proteins and peptides, as well as briefly discussing the functionalization of silk particles with targeting ligands.

11.2.2.1 **Small Molecule Drug Delivery with Silk Nanoparticles**

Silk nanoparticles, similar to other nanomedicines, are capable of encapsulating and facilitating the controlled release of small molecules and drugs. For example, Chen et al. loaded paclitaxel into silk fibroin nanoparticles using desolvation [73]. Among different formulations, the smallest particles were 270 nm in diameter with zeta potentials between -20 and -27 mV. Drug loading was between 1 % and 7 % and the encapsulation efficiency was between 67 % and 100 %. Release of paclitaxel in vitro was relatively rapid, with the entire drug content being released in 2 h, however, the larger particles (diameters up to 520 nm), had a slower release lasting up to 300 h. Wu et al. prepared smaller paclitaxel-loaded nanoparticles, also by desolvation, with diameters between 160 and 210 nm [66]. In vitro release showed a rapid 40 % release in 8 h followed by a delayed release, totalling 47 % after 100 h. The particles were cytotoxic to two human gastric carcinoma cell lines (BGC-823 and SGC-7901) in vitro, while silk fibroin alone was not. In vivo testing on a subcutaneous human gastric cancer (BGC-823) nude mouse xenograft model showed decreased tumor volume and weight when treated with local injection of nanoparticles versus the equivalent dose of drug. Gupta et al. loaded curcumin, a potential anti-cancer drug, into silk particles prepared using the capillary microdot technique [74]. Silk and the drug mixture was dispensed on a slide, frozen, lyophilized and resuspended in methanol. Particle sizes ranged from 40 to 70 nm and in vitro release showed a burst phase followed by a slower sustained release over 8 days. The particles were shown to be taken up by and to inhibit growth in two human breast cancer cell lines (MCF-7 and MDA-MB-453) in vitro. Zheng et al. made 50 nm diameter particles using solution-enhanced dispersion by scCO₂ which were loaded with the anti-inflammatory drug indomethacin [69]. Drug loading was 6 % by mass with encapsulation efficiency of 31 %. In vitro release was relatively well sustained and stable: 61 % after 6 h and 87 % after 24 h.

Silk can also be blended with other polymers or macromolecules to modify properties such as drug loading [46, 75]. Subia et al. formed cross-linked albumin-silk fibroin nanoparticles loaded with the anti-cancer drug methotrexate [46]. Particles were formed by desolvation and cross-linking was done using glutaraldehyde. The particle diameters were between 100 and 200 nm with a zeta potential around -25 mV. The drug loading was 15–24 % by mass and the encapsulation which was 83–87 % was higher than that of particles made from only silk fibroin or albumin. Up to 90 % of the methotrexate was released from the various formulations after 10 days.

Both silk fibroin-albumin nanoparticles and pure silk fibroin nanoparticles inhibited the growth of the human breast cancer MDA-MB-231 cells, and were more effective than the free drug. The particles were also found to be non-toxic to human erythrocytes. They were taken up into the cytoplasm and perinuclear space in feline fibroblast cells (AH-927) as observed by confocal laser scanning microscopy after conjugation of fluorescein isothiocyanate to the nanoparticles.

Attempts have also been made to combine nanoparticles with other biocompatible structures. Numata et al. incorporated silk nanoparticles into a silk hydrogel to combine the fast release of molecules from the hydrogel with slower release from nanoparticles [76]. The particles were 175 nm in diameter with a zeta potential of about -12.5 mV. Dyes (Rhodamine B, Texas Red and fluorescein isothiocyanate) were incorporated into both the particles and the gel, and release was measured in the presence of protease. Within 1 h, 90 % of dye was released from the hydrogel while the nanoparticles had a relatively constant release rate over 5 days which was not influenced by enzymatic degradation. No cytotoxicity was observed against human mesenchymal stem cells.

Glue-like sericin proteins are normally removed from cocoons during the degumming process [1, 49–52], and appear to be non-toxic [3, 77]. Hanjin et al. created drug-loaded microparticles by electrohydrodynamic spraying using these proteins [75]. The resulting particles, however, were large; 150–300 μm . The anti-inflammatory drug diclofenac was loaded into the particles and 60–70 % was released in 7 h. Mandal and Kundu were able to make sericin nanoparticles from *A. mylitta* silk by mixing this with poloxamers (triblock copolymers with hydrophilic-poly(propyleneoxide)-hydrophilic sequences) [63]. Their diameters ranged from 60–130 nm and both hydrophobic paclitaxel and hydrophilic insulin were successfully loaded. The growth of MCF-7 breast cancer cells was reduced when they were treated with paclitaxel-loaded nanoparticles. The authors suggested that the positively charged sericin could aid cellular uptake by charge-based association with the plasma membrane, however, in this experiment the nanoparticles were not more effective than the drug alone. Sericins might have some utility for drug delivery, however this has yet to be fully demonstrated in vivo. Silk fibroin nanoparticles on the other hand, have greater potential for the loading and controlled release (via diffusion) of drugs and small molecules.

11.2.2.2 Peptide or Protein Drug Delivery with Silk Nanoparticles

Proteins and peptides represent a further important group of pharmaceuticals and constitute therapeutic hormones, growth factors, clotting factors, anticoagulants, drug-activating enzymes, and antibodies [78]. However, these types of biologic drugs suffer from degradation by serum proteases and clearance by the immune system [78, 79]. Incorporation into nanoparticles could potentially mitigate these effects. Hai-Bo et al. formulated particles of silk fibroin cross-linked to insulin by nanoprecipitation [50]. Silk-insulin particles were cross-linked using glutaraldehyde post-formation and their diameters ranged from 40 to 120 nm. Compared to

the native peptide, encapsulated peptide had improved resistance to degradation in human serum and to degradation by trypsin *in vitro*. Kundu et al. observed the endocytosis of particles made from *B. mori* and *A. mylitta* silk using nanoprecipitation [53]. The particles were between 150 and 180 nm in diameter with a zeta potential around -25 mV (Fig. 11.4).

The particles were endocytosed into murine squamous carcinoma cells (SCC7) and were detected in the cytoplasm and nuclear membrane. Cytotoxicity was measured using murine fibroblasts (L929) and the particles were found to be non-cytotoxic up to 100 $\mu\text{g}/\text{mL}$. The nanoparticles made from *A. mylitta* silk were loaded with vascular endothelial growth factor by charge-based association and *in vitro* release was approximately linear for 5 days followed by a slower, sustained release. Cao et al. prepared nanoparticles loaded with the enzyme beta-glucosidase by nanoprecipitation [65]. The particles were between 50 and 150 nm and the beta-glucosidase was incorporated by charge-based association. Wang et al. formulated nanoparticles of silk blended with poly(vinyl alcohol) using a different technique which involved drying of the silk-poly(vinyl alcohol) mixture to leave a thin film, followed by resuspension in water [51]. Rhodamine and tetramethylrhodamine conjugated to either bovine serum albumin or to dextran, was loaded into particles by mixing with silk prior to nanoparticle formation. Various silk-polymer ratios gave a large distribution of diameters with the smallest being 300–400 nm. Dextran was

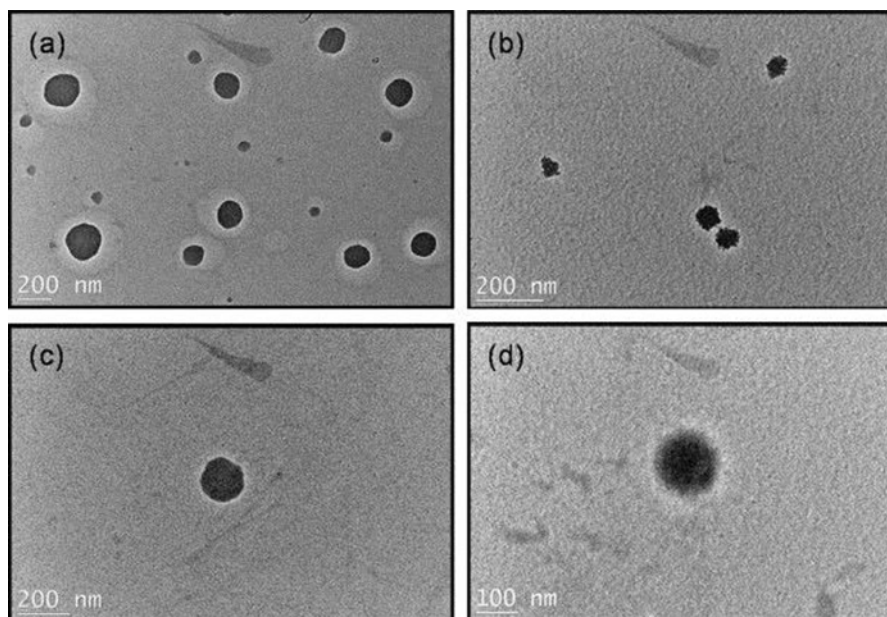


Fig. 11.4 TEM images of silk fibroin nanoparticles prepared from *A. mylitta* silk (a) and (c) and *B. mori* silk (b) and a single silk fibroin nanoparticles prepared from *B. mori* (d). Taken from Kundu et al. [53]

released from the nanoparticles at a reasonable rate: 60 % in 14 days, however, less than 5 % of the protein payload was released in the same amount of time.

Both proteins and peptides can be effectively incorporated into silk nanoparticles. Encapsulation can shield and modulate the release of proteins and peptides into their surrounding environment which would improve stability by reducing the rate of degradation by proteases [50]. More work is needed to demonstrate this in vivo and with other peptide and protein drugs.

11.2.2.3 Targeted Nanoparticles

The location and accumulation of nanoparticles in vivo is subject to blood hemodynamic forces and diffusive mechanisms [80]. Rather than relying on these passive means of accumulation, attachment of ligands, such as peptides, antibodies, or small molecules, to the surface of nanoparticles allows them to preferentially bind target cells to improve the efficiency of drug delivery [80]. Wang et al. functionalized the surface of silk microparticles (2–3 μm) with avidin using carboxydiimide coupling followed by the attachment of biotinylated antibodies to the particle surface [61]. Fluorescein isothiocyanate was bound to avidin to image the particles by confocal laser scanning microscopy. The targeted particles bound to CD3 positive human T-lymphocytic cells (ATCC[®]TIB-152[™]) with improved specificity. Another targeted silk nanoparticle was made by Subia et al. [48]. Folic acid was conjugated to nanoparticles, made from *A. mylitta* silk by nanoprecipitation, using carboxydiimide coupling. Folic acid is taken up by endocytosis and the folic acid receptor is overexpressed on many cancer cells, making it a useful tumor targeting ligand [81, 82]. The particles were around 200 nm in diameter and the anti-cancer drug doxorubicin was loaded by charge-based association. Uptake was tested in human breast cancer cells (MDA-MB-231) and was observed using confocal microscopy by conjugating rhodamine isothiocyanate to the particles. Immune response was tested by measuring cytokine expression in mouse bone marrow macrophages (RAW-264.7). Particles decorated with folic acid were taken up more effectively than non-targeted nanoparticles. Targeted nanoparticles had improved cytotoxicity to breast cancer cells and a relatively low immune response in vitro. They observed near-linear drug release over 7 days followed by slow release until 21 days [48]. Although research has been done in this area, these experiments show that it is possible to functionalize silk nanoparticles using standard carboxydiimide coupling. These experiments demonstrate that attachment of targeting ligands can improve cellular binding and drug delivery.

11.2.3 Spider Silk Nanoparticles

Spider silk proteins possessing similar chemical and physical properties as silkworm silk proteins have also been used to create a range of nanoparticles [64, 67]. Despite the many new methods of spider silk production, nanoparticles have so far been

developed mostly from silk produced in *E. coli* by expression of an artificial gene construct. Hofer et al. expressed a synthetic gene, containing elements from MaSp of the spider *A. diadematus* in *E. coli* which produced a 48 kDa protein [64]. The particles were made by desolvation in 2 M KPO_4 and lysozyme was incorporated into the particles by charge-based association, with 40 % loading efficiency. Release was observed to be pH sensitive, above pH 6, lysozyme remained associated with the particles and was not released and only between pH 4 and 2 lysozyme release was observed (70–90 % after 24 h). Lammel et al. prepared recombinant spider silk and nanoparticles in the same way [67]. The average diameter was 320 nm with a large distribution and a zeta potential of -22 mV. The cationic dye methyl violet was incorporated by charged-based association and the authors showed that the dye permeated the core of the particles rather than simply associating with the surface. A release study showed a modest burst release which was accelerated in the presence of the proteases; elastase and trypsin. Xia et al. created a different gene construct containing the beta-sheet forming repeat of the *B. mori* heavy chain interspaced with the elastic motif of mammalian elastin [83]. This hybrid silk-elastin protein attempts to combine the mechanical strength of silk with the flexibility of elastin [84]. Varying ratios of silk and elastin sequences produced proteins with molecular masses between 48 and 53 kDa. The resulting micelle-like particles formed in water, were 100–200 nm in diameter. Even smaller nanoparticles with diameters 23–38 nm, have been made from silk-elastin-like proteins using an electrospraying technique [85]. Numata et al. designed a unique spider silk construct for the purpose of gene therapy [86]. The recombinant protein made by Numata et al. contained hydrophobic repeats from *N. clavipes* MaSp, 30 lysine residues, and a tumor binding peptide (Fig. 11.5) [86]. The sequence was expressed and isolated from *E. coli*. The positively charged lysines were incorporated to hold the negatively charged DNA within the particle

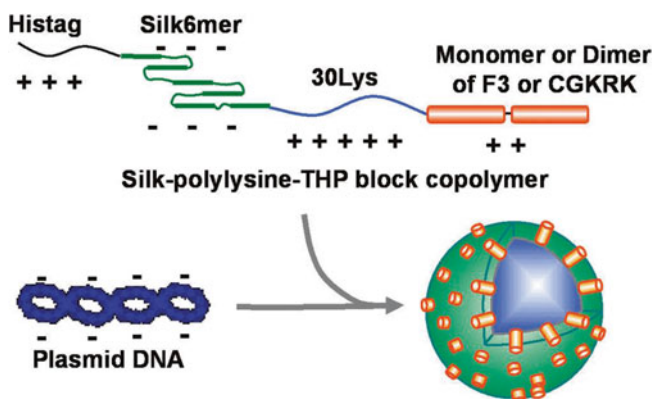


Fig. 11.5 Spider silk-based gene carriers for tumor cell-specific delivery. The recombinant peptide contains silk fibroin domains, lysine residues and one of two tumor binding peptides. The His-tag was incorporated to aid in isolation of the construct. Figure taken from Numata et al. [86]

which, in this case, was a plasmid containing the gene for luciferase. The smallest particles were 170–260 nm in diameter.

In vitro testing showed gene expression only in human melanoma cells (MDA-MB-435) and human breast cancer cells (MDA-MB-231) and not in non-cancerous mammary epithelial cells (MCF10A), indicating that the targeting peptides were effective. In vivo testing showed no toxicity in mice and significantly more luciferase expression in tumor-infected mice. Spider silk nanoparticles have yet to be studied as extensively as those produced from silkworm silk, however as evident from these preliminary investigations spider silk nanoparticles also have potential as drug delivery vehicles. Genetic engineering can further be used to incorporate other useful elements in lieu of chemical coupling [86]

11.2.4 Silk-Coated Nanoparticles

Rather than acting as the core of nanoparticles, silk can be used to coat other nanoparticles and improve their biocompatibility [62, 87]. Metallic nanoparticles have potential clinical applications in imaging and can be further stabilized by silk coatings. Liu et al. used silkworm-silk fibroin as a coat and scaffold for ZnFe_2O_4 nanoparticles [62]. Dissolved silk fibroin was present during the formation of the metal particles and the negative silk proteins are thought to act as a scaffold to hold Zn and Fe ions. The coated particles showed improved compatibility with P12 human brain astroglia cells versus non-coated particles. Silver and silver nanoparticles are also of interest in biomedical applications. Silver nanoparticles have been shown to be effective anti-bacterial agents against common bacteria [88, 89]. However, silver can cause significant toxic effects due to oxidative stress [90–93]. Silk has been used as a scaffold and coating for silver nanoparticles to help improve biocompatibility. Fei et al. prepared silver nanoparticles in the presence of silkworm silk fibroin [49]. The composite was effective against methicillin-resistant *Staphylococcus aureus*. Aramwit et al. used silkworm sericin as a reducing and stabilizing agent for silver nanoparticles [94]. The particles were 50–120 nm in diameter with a zeta potential around -20 mV. Increasing dosages of the particles inhibited the growth of 3 g positive bacteria and to a lesser extent, the growth of 3 g negative bacteria.

Quantum dots are another form of metallic nanoparticle which can be used as fluorescent probes for labelling and imaging proteins, DNA and cell structures, however, their cytotoxicity is a significant concern [95, 96]. In order to improve biocompatibility, Nathwani et al. coated CdSe/ZnS quantum dots with silkworm silk fibroin [87]. The resulting particles were 4–7 nm in diameter and were successfully used for fluorescent imaging of HeyA8 ovarian cancer cells. Uncoated particles could only be stabilized in organic solvent (chloroform) and caused cell death (possibly due to the solvent). Coating particles with silk fibroin improved biocompatibility, whereby the particles became soluble in water and had no observed cytotoxic effects. Instead of metallic nanoparticles, another study coated solid lipid

nanoparticles, made from steric acid and ceramide, with silkworm-silk fibroin [97]. Solid lipid nanoparticles have applications in delivery of hydrophobic drugs, for example, the dermal delivery of hydrocortisone [98]. With the silk coating, the average particle diameter was around 500 nm with a large distribution [97]. Coating of the lipid particles increased the zeta potential (although still negative) which increased the skin permeation of ceramide particles in the dorsal skins of female nude mice. These examples demonstrate that silk polymers have potential to act as a coating agent to improve the biocompatibility of metallic nanoparticles [99]. The coating of solid lipid nanoparticles demonstrates that silk fibroin is a biocompatible coating agent which can enhance in vivo distribution.

11.2.5 Nanofibers and Electrospinning

Nanofibers are nanoscale structures with potential applications in controlled drug delivery, tissue engineering and wound dressings [103, 104]. This section will introduce electrospinning, the method used to generate these fibers, and research using silk fibroin nanofibers. Electrospinning mimics the spinning process that occurs in spiders and silkworms by drawing dissolved silk, with disordered and coiled structure, through a very fine stream to form insoluble fibers with aligned beta-sheet structures [100–102]. Mechanical extrusion of silk through a very fine nozzle can make fibers with diameters between 10 and 500 μm [101]. In order to create nanoscale fibers, electrical forces are used to draw and spin silk [100–102]. Silk solution is added to a container which ends in a capillary. A high voltage electrode is inserted into the solution or connected to the capillary and then to a collection plate which is several centimeters from the edge of the capillary. As charge builds up in the solution, and the attraction of the solution to the collection plate by electrical forces eventually overcomes the surface tension holding it in the capillary. The aqueous silk leaves the capillary in a very fine stream causing the solvent to evaporate. The spun fibers are collected on a plate, spool or into a solution. The applied voltage, flow rate, collector distance, and polymer and solution properties all affect the size of spun fibers [100–102]. Thangaraju et al. loaded the hydrophobic molecule curcumin into silkworm silk fibroin nanofibers [103]. The spun fibers had diameters ranging from 50 to 200 nm, and release of curcumin was monitored in vitro which totaled 80–84 % over 10 days. Jingwen et al. prepared aspirin-loaded nanofibers from a mixture of poly(lactic acid) and silkworm silk fibroin [104]. Fiber diameter was between 80 and 210 nm. The ratio of silk to polymer affected the drug release by changing the swellability and corrosion of the fibers. Drug release was 9–16 % over 10 h and only 10–20 % over 72 h. Sheikh et al. made silk nanofibers which incorporated both silver and hydroxyapatite nanoparticles [105]. The electrospun mats had significant antibacterial activity on *E. coli* and *S. aureus* cultures. Unfortunately, toxicity and attachment inhibition was observed in fibroblast cells. Based on these experiments, it is evident that electrospun silk nanofibers have potential for the

encapsulation and controlled release of drugs, however, more testing is required to demonstrate their utility *in vivo*.

11.2.6 Potential Biomedical Applications of Silk Nanostructures

The development of improved medicines is the ultimate goal of research involving drug-loaded nanostructures. Table 11.1 summarizes promising biomedical applications of silk nanoparticles and nanofibers. These applications focus on the delivery and controlled release of anti-cancer drugs and are also used as biocompatible coatings. Although a wide range of silk nanostructures have been investigated in animal models to date, the potential utility of nanomedicines made using silk components remains to be evaluated in clinical trials [80, 106, 107].

11.3 Silk in other Drug Delivery Applications

Silk has extensively been used in biomedical applications and particularly in surgical applications. Silk sutures are mechanically superior materials with regards to yield point, toughness and elasticity moduli when compared with other types of biodegradable sutures (such as collagen, hyaluronins, alginates, absorbable synthetic polymers such as polydioxanone, polyglycolic acids and non-degrading like Teflon-coated polyester and nylon) [109]. Silk biomaterials exhibit slow biodegradation [110, 111], high biocompatibility, low incidence of acute inflammatory activity [4], and avoidance of organic solvents, surfactants and crosslinking agents in their design. Mild aqueous processing at ambient temperatures allows for the incorporation of sensitive biological cargo [10, 40], and silk encapsulation exerts a remarkable stabilizing effect on small molecules [112], protein [113], and enzyme payloads [114, 115]. Furthermore, silk produces non-toxic degradation products circumventing the need for post-surgical removal of local drug delivery implants [110]. Silk biomaterials can be processed to incorporate zero-order [116, 117], bi-phasic [118] or physiologically responsive release kinetics [119, 120], enabling desirable plasma drug levels with minimal interventions and effective long-term preventive care [121]. Drug release from silk based nanomedicines can be tuned by varying a range of parameters including; the degree of crystallinity, polymer composition and molecular weight. Silk biomaterials are mechanically robust, biocompatible high performance materials that provide an optimal platform for precise spatio-temporal control over the administration of bioactive drugs and biological signals for a variety of biomedical applications (Fig. 11.6) [35, 109, 122].

Table 11.1 Summary of biomedical applications of silk nanostructures

Nanostructure	Potential clinical application	Example	Experimental results	Particle/fiber diameter (nm)	Ref.
Nanoparticles loaded with small molecules	Delivery and controlled release of anti-tumor drugs	Anti-tumor drug paclitaxel-loaded nanoparticles	Reduced tumor size in mouse xenograft model more effectively than free drug when injected locally.	160–210	[66]
Nanoparticles loaded with proteins and peptides	Protection of peptide drugs from degradation	Insulin incorporated into silk particles	Improved resistance to protease degradation	40–120	[50]
Silk-coated nanoparticles to improve biocompatibility	Improve biocompatibility of silver nanoparticles	Sericin-coated and stabilized silver nanoparticles	Inhibited growth of six common bacteria	50–120	[94]
	Improve biocompatibility of metallic nanoparticles	Silk coated-ZnFe ₂ O ₄ nanoparticles for MRI contrast agents and brain imaging	Good biocompatibility and low cytotoxicity with human brain astroglia cells in vitro		[62]
Functionalized spider or silkworm nanoparticles	Target cancer cells for delivery of anti-tumor drugs	Nanoparticle with folate as targeting ligand on surface and loaded with doxorubicin	Improved inhibition of human breast cancer cell growth in vitro	~200	[48]
	Target cancer cells for delivery of anti-tumor genes	Recombinant spider silk construct containing tumor-homing sequence loaded with reporter gene	Increased reporter gene expression in mouse breast cancer model	170–260	[86]
Nanofibers loaded with small molecules	Localized, controlled release of anti-tumor drugs	Silk nanofibers loaded with curcumin	Delayed, 10 day in vitro release	50–200	[103]
	Anti-microbial mats	Silver and hydroxyapatite-incorporated nanofibers mats	Anti-bacterial activity against <i>E. coli</i> and <i>S. aureus</i>	100–300	[108]

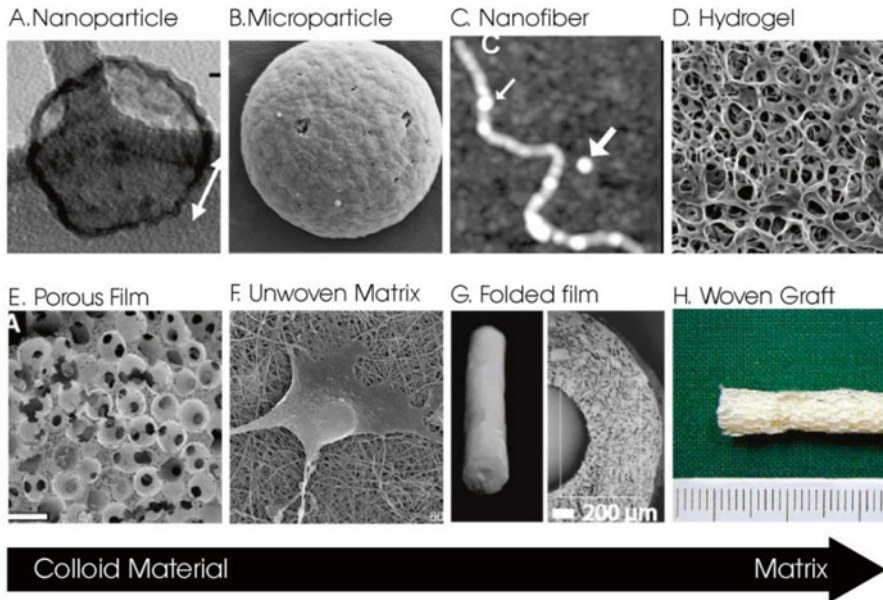


Fig. 11.6 Diverse morphologies of processed silk. (a) Silk elastin nanoparticle [85]. (b) PVA silk microparticle [51]. (c) AFM micrograph of dragline spider nano fiber [123]. (d) SEM micrograph of a pH induced *B. Mori* hydrogel [124]. (e) Porogen mediated porous scaffold [125]. (f) Osteoblast on electrospun silk matrix [126]. (g) Nerve conduit made from folded NGF loaded film. (h) Knitted micro-porous ACL scaffold [127].

11.3.1 Silk Processing

Natural silk is potentially immunogenic and can lead to IgE upregulation [128], and must firstly be degummed of hydrophilic sericin glycoproteins by boiling in alkaline sodium carbonate solution [4, 11]. Lindsay et al. demonstrated how boil times have a significant effect on silk thermal stability, decreased constituent fragment size distribution, pore-size, stiffness, porosity and in vivo degradation rate of the final fibroin, despite no appreciable disruption in β -crystallinity [129]. The extracted core silk protein can be used as adhesion matrices in tissue engineering applications [130, 131], or dissolved in concentrated salt solutions at 50–70 °C for processing into several material formats. Dialysis yields a pure silk solution that can be lyophilized for storage or alternatively reconstituted in the organic solvent HFIP for drug integration in depot and controlled release applications [44, 110]. Fabrication protocols for microparticles, films, and hydrogels take advantage of fibroin self-assembly in techniques such as freeze-drying, gelation, Electrospinning, laminar-jet break up, and phase-separations (Fig. 11.7) [11, 109, 132]. The mechanisms of microparticle and fiber assembly are physically identical but vary in nucleation and elongation kinetics and can be committed to either morphology at critical buffer

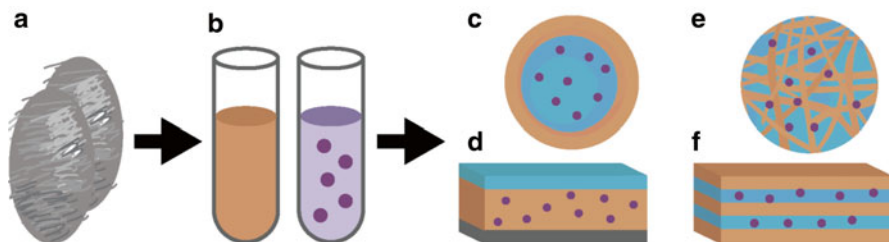


Fig. 11.7 Fabrication of silk materials. (a) Silk can be harvested from silk worm cocoon, recombinant cultures or from widely available textile manufacturers. It must be degummed before use. (b) Solubilized silk solutions (*brown*) are mixed with drug (*purple*) to be processed into; (c) microparticles, (d) films\membrane\coatings, (e) hydrogels, (f) 3D scaffolds

concentrations [133]. Template-assisted film-coating layer-by-layer deposition and film-casting techniques, at the cost of fabrication complexity, allow nanometer level control of material properties and facilitate integration of multiple drug delivery vehicles [10, 11, 35, 44].

A final step in silk processing of β -sheet induction is biomimetically adapted to control crystallinity-dependent material properties through changes in pH [124, 134], salt concentration [40, 135, 136], alcohol mediated dehydration [137], or the application of shear forces [138], and water annealing [139], for alcohol sensitive cargo [140]. Silk can be sterilized through autoclaving, γ -radiation [141], or 70 % ethanol without compromising beta-sheet crystallinity or induced gel properties preclusive to administration, however, sterilization treatment may cause leaching or loss of activity for biological cargo.

11.3.2 Drug Integration Techniques

Silk materials can be loaded with small molecules, peptides, proteins, and DNA constructs at high encapsulation efficiencies through non-specific surface adhesion, physical entrapment and covalent modifications at ambient temperatures. Silk is inherently negative at physiological pH levels as it has a theoretical pI of 4.2. Mixing and co-incubation of fibroin and drug solution results in positively charged molecules decorating hydrophilic spacer regions of the heavy chain silk proteins while non-polar molecules [4, 142], are captured in crystalline regions through hydrophobic interactions. The efficacy of this simple encapsulation technique has been successfully validated in loading electrospun matrices [143], films [118, 144, 145], particle [51, 140, 146], and tablets with optimal retention of bio-activity. Silk and Silk-Elastin like polymer(SELP) hydrogel solutions can be loaded with drug and physically stimulated to gel, immobilizing the cargo immediately prior to administration. Highly bioactive forms of the drugs are disseminated by avoiding possible exposures to harsh hydrogel processing

conditions [147–149]. Non-specific covalent coupling of amide linkages with N-hydroxysuccinimide (NHS) activated carbodiimide coupling have been used to immobilize BMP-2, linear gradients of HRP [150], recombinant PTH [151], RGD adhesion peptides [152], and biotin [61], to silk proteins. A 10 % addition of tyrosine in silk permits higher functionalization than diimide chemistry as demonstrated by Wenk et al. who achieved 99 % loading of FGF-2 [153] using sulfonic acid surface decoration in conjunction with previously described diazonium-tyrosine chemistry [154]. Yasusi et al. also successfully sulfated silk membranes using chlorosulfonic acid to exert heparin like anticoagulant activity [155]. Genetic modification of primary recombinant silk can be used to tailor silk mechanical properties and co-polymer chemistry; enriching amino acids that participate in the aforementioned coupling techniques, as well as the introduction of cysteine residues which allow disulfide coupling and sulfhydryl interfacing with colloid gold particles [156] and polylysine domains to sequester pDNA in silk-protein based gene delivery systems [157, 158].

11.3.3 Modulating Release Kinetics

Efflux of encapsulated molecules from silk fibroin occurs in a combination of extrusion through diffusion—a larger factor in small molecules, and degradation (via hydrolysis or enzymatic action)—a more relevant factor in high molecular weight drugs [10]. Primary sequence and crystallinity of the fibroin can be manipulated to dictate microenvironment factors such as porosity and thickness of the silk vehicle, in order to act as diffusional barriers, as well as to exert an effect on diffusion rates of encapsulated drugs based on electrostatic and Van der Waals forces [159]. Columbic interactions influence liberation of charged compounds from negatively charged fibroin as evident by effective entrapment and slow release for cationic alkaline model drugs such as rhodamine, methylene dye [160], and burst kinetics of negatively charged model drug such as azo-casien [160], loaded in comparable amounts [40, 161]. Entropically driven coacervation of silk with mAbs and a model drug protamine, facilitates exposure and subsequent attractive interactions between hydrophobic domains. Reduction in ionic shielding between the interacting non-polar proteins can cause gradual repulsion mediated release profiles demonstrated with a hydrophobic molecule such as propranolol [113, 159], and hydrophobic antibody domains [113, 162]. Higher concentrations of silk fibroin, crosslinked with EDC/NHS and film coating [144, 146] of drug reservoir reduces material diffusivity resulting in more ideal prolonged recovery of salicylic acid [161], propranolol, theophylline [117] and buprenorphine [152] amongst other drugs [35, 130, 163, 164]. Composite materials such as microspheres loaded into films have yet to be systematically studied but potentially enable two levels of control over release kinetics for additional precision in moderating complex cellular milieus by biological signaling and drug delivery [165].

11.3.4 Direct Drug Integrated Silk Microparticles

Silk microparticles are a versatile platform for sustained release and depot applications due to readily tunable drug release profiles, ease of fabrication, potential for active targeting and high bio-compatibility (Fig. 11.8). Common fabrication methods of monolithic single layer microparticle methods include phase separations, laminar jet breaks, oil/water emulsion and spray drying, all of which exploit silk self-assembly to form microparticles from mixtures of fibroin and drug solutions. Spray drying techniques subject biological materials to high temperatures and are not as therapeutically relevant [140, 166].

11.3.4.1 Phase Separations

Silk microparticles can be produced using salting-out methods. Lemmel et al. were able to generate microparticles by salting-out silk fibroin using 500 mM KPO_4 [167] in which by exchanging chaotropic sodium and chloride ions for kosmotropic ions, phosphate and potassium stimulates fibroin dehydration resembling salt exchange in distal spinning ducts of spiders [168]. Larger particles were generated with increased fibroin concentration and smaller particles were generated by changing the micro mixing intensity and potassium chloride concentrations in the range 250 nm–3 μ m [67]. Potassium chloride and silk concentrations can modulate sizes between 0.5–4 μ m, whereas lower pH levels and methanol treatments enhanced beta-sheets and encapsulation of cationic model drugs such as crystal violet alcian blue and rhodamine [40]. A mechanistic study of salt induced phase-separation in eADF4(C16) engineered silk microparticles assembly stipulates that the phase transition between nanofibril and microparticles occurs at a critical salt concentration, as a result of charge shielding that facilitates higher rates of spherical aggregation as compared to rates of elongation [133]. In a systematic investigation of charge-based release, release profiles were measured using eight cationic, three neutral and one negative model drug and loaded into salted-out particles sized between 0.17–0.7 μ m. Silk

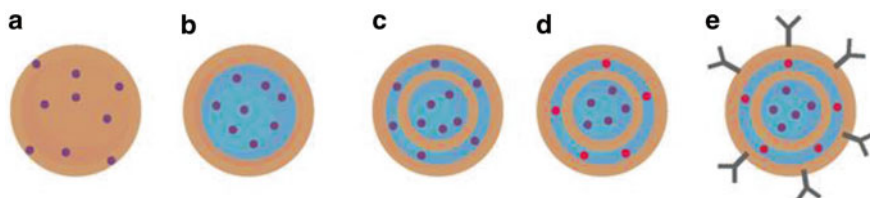


Fig. 11.8 Progression of silk microparticle designs. (a) Direct loaded silk microparticles. (b) Dissolved core, template based particles. (c) Template based particles with multiple fibroin coating containing drugs within and on the interfaces of the coating. (d) The encapsulation of sequential therapies with varying diffusional barriers. (e) Active targeting ligands for localization and up-take of microparticles

proteins were effective carriers of hydrophobic and basic molecules which were loaded at high efficiencies between 20.7–98.2 % and slowly released for up to 8 days, compared to 0.2–17 % encapsulation and burst release kinetics for acidic small molecules [67]. Xiaoqin et al. report an all aqueous microparticle technique based on the spontaneous phase separation of polyvinyl alcohol and drug loaded silk fibroin solutions [51]. The blended polymer solution is film cast in a petri dish and once dried, can be dissolved in water to produce silk microparticles in the range of 0.3–20 μm depending on concentration ratios of silk and PVA. The resultant microparticles doped with TMR-BSA, TMR-dextran and rhodamine-B demonstrated sustained release up to 4 weeks. This technique was adapted to co-flow microfluidics devices by Mitropoulos et al. [72], who were able to consistently generate microspheres with tunable diameters, by varying flow rates of the PVA over the silk solution, and it was demonstrated that smaller spheres exhibited larger cumulative burst release.

11.3.4.2 Laminar Jet break up

Wenk et al. outline a protocol to fabricate microparticles with sizes of 138–440 μm using a laminar jet break-up technique [161]. Consistently sized droplets force solutions of IGF-1/SF as a continuous flow through a vertically vibrating 200 μm nozzle in resonance with the Plateau-Rayleigh instability [161]. The particles were then dripped into liquid nitrogen and lyophilized. Post-treatment with methanol (or water vapor) produced diverse morphologies despite similar beta-sheet content [161]. This technique has produced impressive encapsulation efficiencies ($\sim 100\%$) of the model drugs salicylic acid, propranolol HCl and IGF-I loaded with a prolonged release of up to 7 weeks in vitro [140].

11.3.4.3 Oil/Water Emulsion

Nitchtl and Scheibel demonstrated that eADF4(C16) spider dragline silk will self-assemble at a water-toluene barrier as a permeable microcapsule that is both chemically stable to 2 % SDS, 8 M urea, and mechanically stable with a Young's modulus of 0.7–3.6 GPa [120]. Medical grade silica oil and subsequent ethanol treatments were used in the next rendition of this microcapsule to enhance beta-sheets (circumventing the need for toluene), a biological toxin. Active β -Galactosidase encapsulation is demonstrated through colorimetric progression of the O-Nitrophenyl-, D-galactopyranoside (ONPG) reaction even in the presence of endoproteinase AspN. Moreover, an α -complementation analysis showed a significant increase in ONPG cleavage activity after the addition of a terminal like α -donor protein EA22 to a terminal deficient enzyme. This displayed the ability to produce responsive micro-reactors for applications requiring tuned activation. Cheng et al. showed a similar pH responsive microcapsule in FITC-dextran loaded microcapsules that were produced using an oil/water emulsion technique. The pH-responsive capsules transition between open and lock states as lower solution pH causes swelling in the

shell walls, narrowing of the pores, and decreasing diffusion out of the capsule [169]. These microcapsule technologies form a semi-permeable enclosed reaction chamber which initiates reactions and diffuses reactants, while protecting cargo and allows for later activation, showing promise in drug delivery applications [115].

11.3.4.4 Template Based Techniques

A polyelectrolyte deposition process on a colloid template with subsequent core decomposition as described by Donath et al., has been effectively adapted to silk materials for developing precisely fabricated delivery systems [170]. This type of colloid synthesis provides structural control for encapsulation, diffusional control (addition of masking layers) of drug release and allows for the creation of sequential release therapies [35]. Wang et al. detail the use of horse radish peroxidase (HRP) loaded DOPC liposomes to generate homogeneously sized microparticles around 200 μm in diameter that could deliver detectable amounts of HRP activity in vitro for over a month, depending on the processing [140]. Once drug infused liposomes are formed in solution, they are emulsified with silk solution and subjected to freeze-thaw cycles, with homogenous liposome sizes which approximately doubled the HRP loading efficiency [171]. The removal of the lipid core and induction of crystallinity is achieved through incubation in methanol or 1.7–3.6 μM NaCl solution. Moreover PLGA, and Alginate microspheres coated in silk, delayed PLGA degradation, and maintained a diffusion barrier for loaded drugs; HRP, tetramethylrhodamine conjugated BSA encapsulated in these microspheres with and without silk coating. Drug release was significantly retarded by silk coating, further hindered through methanol treatment with the mechanically stable shell now posing as a diffusion barrier to the drugs [146]. Silk integrated microspheres therefore show an effective improvement on alginate and PLGA based particles alone by reducing potential harmful levels of drugs at the local or systematic levels, by controlling burst release, while also protecting labile cargo that may degrade once released into the plasma [172].

11.3.5 Hydrogels

Hydrogels are heavily hydrated solids composed of over 95 % water content and three-dimensional polymer networks that can physically swell as they absorb water [173]. Silk hydrogels are attractive drug delivery systems because gelation kinetics are temperature dependent and the structural composition can be tuned through polymer concentration, the presence of salts and pH [174]. Significant developments have been made in tissue production using hydrogel scaffolds by promoting morphogenesis and closely reproducing signaling micro-environments, however these are exhaustively reviewed elsewhere [131, 164]. Moreover, hydrogels exhibit solid-like mechanical stability with liquid-like elasticity through a combination of osmotic and entropic structural forces [136]. Hydrogels can be formed through the

addition of energy via ultra-sonication [173] and vortexing [46] the silk protein solution, which drives chain mobility that creates permanent, physical and intermolecular crosslinks. They can also be formed through decreased pH [124, 136] and surfactants [175], which facilitates the extravasation of water. Material properties such as stiffness and porosity are controlled through further dehydrating aqueous impurities from β -sheet block regions by post-treatment with methanol [136].

11.3.5.1 Physically Induced Gels

Xiaoqin et al. demonstrated the induction of liquid-gel transition in silk hydrogels in physiological potassium salt via ultrasonication [147]. The sonication-induced gel exceeded mechanical properties and displayed enzymatic degradation resistance congruous gelatin [176], PEG-fibrinogen [177] and alginate [178], gel systems without the use of hazardous crosslinking agents or solution conditions that may deteriorate labile bioactive cargo [147]. Moreover, the ultrasonication method allows for processing hydrogels at physiological pH and ionic strengths amicable to stabilizing sensitive cargo closely juxtaposed within beta-sheets [179]. Dialysis of supersaturated SF slowly leaches silk I stabilizing salts yielding metastable α -helical silk that anneals to hydrophobic β -sheet silk II on timescales dependent on the rate of salt diffusion [180]. Residual post-dialysis calcium concentrations in fibroin preparations reduce surface porosity of the silk material [136]. Furthermore, freezing at $-20\text{ }^{\circ}\text{C}$ as compared to freezing in liquid nitrogen allows slow nucleation of ice crystals that tend to grow larger and produce larger pores once sublimed in a lyophilization process which influences the permeation and diffusivity of drugs through the matrix [135]. Chemical cross linking with ammonium peroxodisulfate eADF4(C16) spider fibroin produce β -sheet rich gels with higher shear and elastic moduli which develop into sponge like networks rather than pleated sheet-like structures and significantly reduce pore size [174]. Jai-You et al. induced gelation in reconstituted fibroin-buprenorphine with citric acid mediated pH reductions and showed a nearly zero order release of buprenorphine, and were able to control rates by varying the proportion of low and high molecular weight silk polymers [124].

11.3.5.2 High Performance Gels

Ultrasonicated and vortexed [148] silk allow for milk aqueous processing but exhibit relatively low resilience—deforming at strains greater than 10 % due to the brittle nature of beta sheets in high displacement [181]. Silk electrogels withstand up to a 1000 % greater compression but rely on strong voltages applied to fibroin in high pH. This reduces helical propensities through a local proton enrichment [182]. The ease of administration via hand-held applicators and the reversible adhesive nature of e-gels warrant future investigation for topical and surgical applications. Benjamin et al. proposed HRP mediated oxidative tyrosine crosslinking to form robust flexible silk hydrogel networks akin to di-tyrpsine bridges in hyaluronins and

aligantes [149]. Enzymatically crosslinked silk is capable of recovering from cyclic applications of extremely compressive strain (up to 70 %) with a shear moduli of 10 Pa–10 kPa and a tangent moduli of 15–400 kPa similar to e-gels. The mild cross-linking process incorporate anti-bodies [162], growth-factors [183, 184] and antibiotics [179] for delivery as implants dressing of dynamic tissues.

11.3.5.3 SELPs

Silk-like and Elastin-like Polymers (SELP) are recombinant block copolymers in which physicochemical properties of the material are tailored by carrying composition, length and ratios of silk and elastin amino acid sequence blocks. X-ray diffraction and differential scanning calorimetry (DSC) are consistent with crystallization of silk-like β -sheet block thermal and chemical stability interrupted by flexible, more soluble elastin regions [84]. SELPs are liquid polymers that spontaneously undergo an irreversible liquid-gel transition in clinically useful timescales allowing the inclusion of dextrans, dansyl amino acids and other protein drugs shortly before administration as gels. SELPs can be processed in isotonic, physiologically compatible solvents allowing for the delivery of nucleic acids [185], microparticles, liposomes, live cells, and other bioactive compounds which may be sensitive to organic solvents. SELF-47 solution mixed with DNA solidified in an hour at 37 °C, released gene payloads for up to 28 days in a manner highly sensitive to ion strength of the medium that shield the charge of the phosphates in the DNA. Genetic engineering enables precise control over SELP charge densities facilitating internal control over charge-based release kinetics [186]. Controlled DNA cumulative release of only 5 % of total pFG-ERV plasmid load was seen after 28 % with preserved structure as shown by electrophoresis. Subcutaneous injection of luciferase plasmids loaded SELP gels proximal to tumours in MDA-MB-435 breast tumors in mice showed protection of DNA from nuclease activity and 3.5 times transfection of tumors as compared to naked DNA [186]. More recently silk-elastin gels loaded with adenovirus carrying β -galactosidase and luciferase reporter genes showed 4–8 times higher gene expression levels compared to the virus alone and minimal dissemination to the liver in the head-and-neck cancer nude mice models [185]. Moreover, xenograft head-and-neck cancer JHU-012 cells targeted by gels embedded with therapeutic adenovirus encoding thymidine kinase (Ad-Tk) and ganciclovir show significant increases of Ad-Tk mRNA eliciting reductions in tumor volumes compared to PBS and β -galactosidase controls at 7 and 14 days [185].

11.3.6 *Surface Coatings for Medical Devices and Regenerative Implants*

Silk coatings are an effective system as drug-eluting coatings, however silks micro-mechanical and biological properties has high utility in medical grade implants [187] and regenerative grafts as well [130]. For example, for cardiovascular surgery

applications, Wang et al. demonstrated silk surface coatings of luminal stents can alleviate the potential for restenosis through prolonged release applications of heparin, paclitaxel and clopidogrel in a swine model. Coated stents reduce excessive proliferation of vascular smooth muscle cells (SMCs), thrombosis, and chronic inflammation often aggravated by foreign metallic device implantation to facilitate post-angioplasty vascular repair [188]. Surface modification of silicone breast implants with micrometer-thin layer of recombinant spider silk proteins have also shown to reduce unspecific protein adhesion and post-operative inflammation through surface hydrophobicity reduction [189] and proving a great candidate to resolve periprosthetic capsular fibrosis in silicone breast implants with reasonable effort [190].

Silk based growth factor therapies have been purposed for regenerative therapies in a plethora of tissue types [164], but show poor clinical transition due to rapid degradation and clearance in vivo, in addition to lack of understanding of how materials interface with the complex cellular milieu [191, 192]. Epidermal growth factor and sulfadiazine loaded electrospun silk nanofiber mats exceed commercially available topical wound dressing Tegadern Hydrocolloid(3 M) in terms of re-epithelialization, dermis proliferation, collagen synthesis and scar formation [193]. Schneider et al. show how EGF incorporated into silk mats with slow release (25 % in 170 h), optimizes the wound healing response in facilitating the alleviation of infection, inflammation and promotion of growth factor release from surrounding tissues, observed in previous in vitro studies using electrospun mats on wounded human skin models [194]. Silk is a prime candidate for the development of growth factor delivery scaffolds as it is resilient under strains associated with structural tissue such as bone [152], tendons and ligaments [127]. Wenk et al. showed how IGF-1, an important molecule in chondrogenesis can be incorporated into microparticles embedded within scaffolds for better chondrogenic outcome [195]. Kaplan et al. further demonstrated the dual control of silk particles loaded with osteoconductive morphogenic factors in electrospun matrices. These hybrid systems allow for adjustable loading by the protection of the encapsulated drug through the matrix's processing and modulated the release of the drug through the compositions of the loaded microparticles and their diffusion of the fibers [183].

11.4 Conclusion

The versatility of silk and silk-like products in drug delivery and tissue engineering applications has been demonstrated as evident by the large body of research publications over the last two decades. Insect and spider silk produce natural biopolymers with remarkable strength and flexibility and along with their biodegradability, biocompatibility and ease of production, make silk an exceptional biomaterial. In this review we have seen how the relationships between the chemical and molecular composition, secondary structures and mechanical properties of silk biopolymers can be used to create controlled release drug delivery nano and microparticles for the release of a range of payloads with a variety of structures and sizes.

Our understanding of the genetic coding of silk biopolymers is continuously increasing, and with improved cloning and expression of silk, we are now in a better position to understand the complex self-assembly of silk polymers. This in turn serves as an inspiration for the design, and development of future materials based on the properties of silk biopolymers for more improved nanomedicines and technology tools for healthcare.

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Chapter 12

Nanotoxicology and Regulatory Affairs

Christiane Beer

Abstract Like other parts of the nanotechnology revolution nanomedicines hold great promise and in the case of nanomedicines the potential for more efficient therapies. Engineered nanomaterials that are used as nanomedicines for therapeutic and diagnostic purposes are often designed to specifically interact with cells of tissues and organs of the human body. However, the unique physicochemical properties of particles at the nanoscale may contribute to adverse effects requiring nanomaterial-specific safety considerations. Therefore, before nanomedicines can be approved by organisations such as the U.S. Food and Drug Administration (FDA) or the European Medicines Agency (EMA) and reach the market, safety, efficiency and efficacy have to be shown. Beginning with some short critical remarks, this chapter addresses the toxicology of nanomaterials referred to as nanotoxicology with special attention to nanomedical applications. The second part of this book chapter will briefly describe the general drug approval process, introduce risk assessment procedures and give an overview of safety and regulatory challenges for nanomedicines.

Keywords Nanotoxicology • Pharmacokinetics • Toxicokinetics • Drug safety • Risk assessment • Nanomedicine regulation

12.1 Introduction

The application of nanotechnology offers many advantages, and products based on engineered nanomaterials (ENMs) are used in nearly all parts of our daily live [1]. Nearly everyone in the developed world have been unintentionally or intentionally in contact with nanotechnology in one way or another, either as consumers, or as workers. Nanomedicines based on nanotechnology and ENMs offer great potential in the treatment of diseases. These new developments promise, for example, improved bioimaging properties and more efficient and targeted drug delivery with fewer side-effects [2]. Due to their small size nanomedicines can cross endothelia

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and epithelia biological barriers and enter organs, tissues and cells [2]. The advantageous properties of nanomedicines include a higher drug dissolution rate and enhanced drug adsorption, and increased bioavailability [2]. As much as ENMs offer the possibility to create new innovative products and new groundbreaking technologies there are, however, concerns that ENMs may pose a threat to human health and also to the environment. These concerns are based on the fact that ENMs may have unique physical, chemical and toxicological properties that differ from the parent material in a way that cannot be predicted by studying the larger-sized material. ENMs are known to be more reactive as their larger-sized counter parts, and it is, therefore, reasonable that ENMs may react with biological systems in new unpredicted ways that could lead to toxicity. Studies on natural occurring (e.g., forest fire) and unintentional (e.g., diesel exhaust) arising particles have shown that particle exposure leads to respiratory and cardiovascular diseases, cancer and possibly allergy in humans [3, 4]. Especially the particle fraction with a diameter less than 100 nm seem to play an important role for cardiovascular mortality and morbidity [3]. Due to the fast growth of nanotechnology and an increasing use of ENMs a whole new scientific branch, nanotoxicology, has emerged.

As nanomedicines consist of ENMs, the concerns regarding their toxicity due to their nano-particulate form lead to safety and regulatory challenges, especially as the field of nanotoxicology is still quite young and not all information that is needed for a comprehensive safety evaluation is available at this point.

12.2 Some Critical Remarks

Although the toxicity of particles in air pollution and their health effects have been investigated for some decades, the field of nanotoxicology with systematic toxicological investigations of ENMs is not more than 10 years old [5]. The term nanotoxicology was first mentioned in the scientific literature in a news article in *Science* by Robert F. Service in 2003 and was proposed as a new subcategory of toxicology by Donaldson and colleagues in 2004 [5]. Since then, the number of published nanotoxicology related articles has been increased from ~36 in 2004 to ~1919 in 2013. During the last 10 years, it has become clear that the unique physicochemical characteristics of ENMs not only make their toxicity towards biological systems difficult to predict, but that these characteristics also have a profound influence on their behavior in experimental settings. Even small changes of, e.g., their size or surface properties can have large effects on the observed toxicological outcome [6]. Over the years, it has become clear that the toxicological investigations of ENMs require high demands on the characterization of the particles and a well described experimental setting. The results from the first studies within nanotoxicology are hard to interpret due to incomplete or missing particle characterization. In addition, differences in the experimental setting due to studies in different cell line from different tissues and organs, exposure media, concentration and purity of the ENMs, dispersion protocols and media [7, 8] makes it difficult to compare the published

results and to draw definite conclusions on the toxicological potential of ENMs. For example, it became clear that physicochemical properties of ENMs can change considerably in different exposure media and that the results of genotoxicity studies are influenced by the preparation of ENMs, their concentration, used dispersion agents and impurities, cell type used, bioavailability and uptake of the ENMs [7]. These parameters most likely affect the toxicity of ENMs in general.

Today the demands on the researchers and the quality of their investigations are much greater. It is difficult to publish in peer-reviewed scientific journals without a thorough characterization of the used ENMs. But not surprisingly, there is still an apparent lack of consistent results for the toxic effects of ENMs.

Another problem that has been recognized during the past years is the interference of some ENMs with toxicity tests like the Comet assay (metal oxide based ENMs) and the MTT assay (carbon nanotubes) [9–12]. Therefore, appropriate controls are of utmost importance to exclude false positive and negative results. Furthermore, when reading and interpreting the scientific literature, a critical and objective review of the available information should include an assessment if a thorough characterization of the ENMs are made, if appropriate controls are included, and if the studied concentrations are meaningful and cover realistic exposure scenarios. In many studies, animals have been exposed to an unrealistic high dose of ENMs resulting in overload scenarios inducing health effects that are not observed at concentrations that cover realistic worst case scenarios [7, 8].

However, it has to be kept in mind that the vast majority of nanotoxicological studies have been performed *in vitro* and that there might be a difference to the *in vivo* situation. One point is that *in vitro* cell cultures mainly are cancer or transformed cell lines that might react more or less sensitive to ENMs. Another point is that ENMs in *in vitro* systems interact with and possibly bind to proteins that originate in the uppermost cases from bovine serum, whereas in the *in vivo* situation, the ENMs are in contact with human blood including all blood cell types as well as the proteins of the human serum. The interaction of ENMs with the different serum proteins and blood cells might influence the results of toxicological studies as well.

12.3 Toxicology of Nanomedicines—Pharmacokinetics and Toxicodynamics

If not directly used as imaging agent e.g., SPIONs or Quantum dots, ENMs can be used as carriers for therapeutic drugs. In either case ENMs are a main component of nanomedicines, which by themselves could pose a potential health threat [6, 13, 14]. Based on studies on the ultrafine nanoscaled particle fraction from air pollution it is known that exposure to these particles increases the risk to develop airway and cardiovascular diseases [13, 15]. Due to increased use and exposure of consumers and workers to ENMs has resulted in concerns about potential adverse health effects of ENMs and the development of a new toxicological field, nanotoxicology. Based on the definition of toxicology by The Society of Toxicology (SOT) [16] nanotoxicology

has been described by Oberdorster et al. “as the study of the adverse effects of engineered nanomaterials (ENMs) on living organisms and the ecosystems, including the prevention and amelioration of such adverse effects” [17]. In comparison to other ENMs, ENMs specifically used in nanomedicines have not been in as much focus of toxicological investigations yet, as the likelihood for an exposure of the general population by nanomedicines is considered to be low. The focus has been mainly on the development of efficient nanomedicines. However, as ENMs are a main component of nanomedicines the general nanotoxicological concepts for ENMs apply and general findings from other ENMs might be transferred to nanomedicines.

The toxicity of ENMs is crucially dependent on their physicochemical characteristics which play, therefore, a key role for the pharmacokinetics and toxicodynamics of nanomedicines. Pharmacokinetics, or toxicokinetics in the case of non-pharmaceutical substances, describes what the organism does with the nanomedicine, whereas, toxicodynamics describes what the nanomedicine does to the organism. Both, pharmacokinetics and toxicodynamics, therefore, reflect the inseparable interconnection of efficacy and toxicity of nanomedicines. This makes the task to develop nanomedicines that have the highest efficacy and, at the same time, the lowest possible toxicity not as simple as it might look at first glance. It is in fact a most difficult challenge as changes of the physicochemical properties can influence the toxicity, absorption, distribution, metabolism, and excretion of nanomedicines at the cellular but also organism level [18, 19]. Therefore, an early implementation of toxicological investigations is fundamental when developing new nanomedicines.

12.3.1 Physicochemical Properties of ENMs that Affect Toxicodynamics

Efficacy and toxicity of nanomedicines are both dependent on the physicochemical properties of the used ENMs. Through changes of their surface charge, shape, size and surface coating, one is able to control and target the drug load and delivery, influence the biodistribution and clearance from the bloodstream. However, a number of toxicological investigations of ENMs have shown that these physicochemical properties also affect the toxic potential of ENMs as they influence their interaction with the organism. The most important physicochemical properties in this connection are size including surface area, agglomeration and aggregation state and porosity; surface chemistry including surface charge and coating; shape; and chemical composition [8, 20–23].

12.3.1.1 Size, Surface Area and Shape

The physical behavior of particles changes dramatically when they reach sizes below 100 nm. Below this size and the smaller the particles are the rules of quantum physics apply more and more resulting in new chemical, mechanical, electrical,

optical and/or superparamagnetic characteristics of the particles. Due to these changes of the characteristics of the particles, it is nearly impossible to extrapolate the biological reactivity and toxicity of ENMs from their larger-sized counterparts [21].

The size is undoubtedly the most important property of ENMs from a toxicological point of view as it influences a number of particle characteristics. These characteristics are high surface to volume ratio, high surface reactivity, absorption of compounds, ability to cross cellular membranes and strong interparticle forces.

How much the reduction of the size to below 100 nm size can change the physicochemical properties of the parent material becomes clear when looking at the example of gold. Gold is known to be one of the least reactive chemical elements. Larger-sized gold particles do not react with oxygen of the air or water. However, when occurring as nanoparticles with less than 10 nm in diameter, gold will burn once it is in contact with oxygen. This example shows that nanoparticles indeed may behave completely different compared to their parent material and this might as well be true for their toxic potential. First of all, the reason for that nanoparticles behave different compared to their parent material lies in the dramatic increase of the surface to volume ratio the smaller the particles become. For example, if a 1 cm cube is divided into 1 million 1 nm cubes the volume is still the same (6 cm^3) but the surface area has dramatically increased from 6 cm^2 to $60,000,000 \text{ cm}^2$. In addition, the percentage of molecules that are at the surface of the particles increases exponentially when the particle size is below 100 nm [13]. This and the enormous increase in the surface area enhances the possibility of ENMs to react with biological systems, including binding to cells, proteins and other biological active molecules. This characteristic of ENMs might, on one hand, be very desirable and useful for their use as nanomedicines but gives, on the other hand, reasons for concern as these interactions might be detrimental as well as uncontrollable.

The size of particles influences where the particles accumulate, how and how fast the body is able to clear particles and presumably directly influence the mechanism and level of toxicity [24]. For titanium dioxide it has been shown that the reduction of their size from 250 nm to 20 nm increases the inflammatory response in lungs of rats and mice; at least when looking at the same mass dose of the particles. At the same surface area, which ultimately means a lesser mass dose for the small particles, the large and small particles induced the same toxicity [13]. Furthermore, the surface area as a direct function of particle size has been shown to be related to inflammation and genotoxicity for nanoscaled carbon black, carbonaceous nanoparticles and titanium dioxide nanoparticles, *in vitro* and *in vivo* [25, 26]. For porous or very rough ENMs the specific surface area (Brunauer Emmett Teller (BET) surface) should also be considered as this considers all surface molecules not only at the outer surface but also those in pores [21].

If the volume of the particles is kept as a constant, the surface area is dependent on the shape. Spheres have the highest volume to surface ratio, with the ratio decreasing for cubes and fibers. Therefore, the toxicity of ENMs is considered to be dependent on the shape of ENMs [27, 28]. Besides the influence on the surface area, shapes with sharp edges might be prone to faster degradation and in the case of

metal-based ENMs this might be leading to the release of toxic metal ions. However, how important the shape of ENMs is for toxicity is still not clear.

When considering shape and toxicity, nano-fibers are of special concern. The global exposure of workers but also of the general population to asbestos fibers resulted in disorders of the lung and pleura like lung cancer and mesothelioma, pulmonary fibrosis and plaques at a pandemic scale. In 1997 the World Health Organization (WHO) published a definition for the so-called fiber paradigm for high aspect ratio materials [29]. Fibers with a diameter smaller than 3 μm , length greater than 5 μm and an aspect ratio greater than 3:1 are considered as harmful. This is also true for ENMs that have a high aspect ratio. Due to the similarity in their appearance to asbestos and the severity of the asbestos pandemic, high aspect ratio nanomaterials (HARNs) are generally classified as harmful. The mechanism behind the fiber pathogenicity is an incomplete engulfment of the fibers by macrophages also called frustrated phagocytosis. Fibers that are longer than 20 μg in length cannot be fully engulfed and this incomplete uptake process results in pro-inflammatory responses by the macrophages [23]. Chronic exposure to the fibers and subsequent persistent inflammation has been associated with the deposition of scar tissue in the lung (fibrosis), tissue damages and carcinogenicity of high aspect ratio fibers. However, tangled fibers that appear more like a sphere than a fiber will be completely taken up by macrophages and degraded if the material is biodegradable.

One key question, not only for toxicological investigations but also for regulatory purposes is the investigation of the aggregation and agglomeration state of ENMs. Agglomeration is a reversible process as ENM agglomerates are formed by weak bonds whereas aggregates are formed by strong covalent bonds. Aggregation of particles can change profoundly the size and size associated properties like transport, deposition and material release. Therefore, if ENMs are present as aggregates that are within the micrometer range or several hundred nanometers in all directions the material will no longer be considered as a true ENM. Agglomeration and aggregation are of special concern for nanomedicines when applied intravenously directly into the body as this can lead to thrombosis, a potentially lethal obstruction of the blood flow. Agglomeration but eventually also aggregation can occur, e.g., through binding of plasma proteins to ENMs making the surface characteristics and coating of ENMs important factors for the toxicity and intracellular fate of ENMs.

12.3.1.2 Particle Coating and Protein Corona

At the moment ENMs enter the human body they can interact theoretically with any protein of the plasma proteome that consists of ~3700 different proteins. Therefore, it is safe to say that ENMs will never exist as uncoated particles in the body. The interaction between particles and proteins results in a protein corona covering the surface of the particles. Which proteins in particular bind to the ENMs and become a part of the protein corona is, however, dependent on the chemical composition and surface charge of the particles. Several of the proteins in the protein corona of carbon black, silica, titanium dioxide and acrylamide nanoparticles particles have been

identified and a number of these proteins are ligands to receptors at the cell surface [32–34]. Although binding of these proteins to the ENMs might lead to their unwanted uptake, this can also be exploited for targeted drug delivery when the ENMs are deliberately coated with ligands for receptors that are present at the cell surface of the target cells. However, unless the proteins are covalently bound to the surface of ENMs the protein corona is not a static structure and other proteins that are present in the surrounding body fluids can replace the original coating. The coating and the protein corona of ENMs have not only an influence on the clearance and targeting but also on the surface reactivity. If the surface atoms of the ENMs are covered by proteins, their surface reactivity is affected and the biological responses to the ENMs might be reduced [35]. This is exploited when adding a protein or polymer coat to the particles. For example, surface coating or surface functionalization has been shown to improve the therapeutic efficacy and minimizing adverse effects of mesoporous silica nanoparticles based nanocarriers [30]. Furthermore, coating of polystyrene nanoparticles with bovine serum albumin (BSA) has been shown to prolong the circulation time in the blood and significantly reduced the particle clearance compared to uncoated particles of the same size, although ~90 % of the particles were no longer present in the blood after 60 min [31]. However, as it takes approximately 1 min for a blood cell to circulate the body, even small increases of the circulation time will increase the likelihood that nanomedicines can reach their target organ before they are cleared by macrophages. The reason for the quite fast and efficient uptake of uncoated ENMs by macrophages lies in the binding of proteins like immunoglobulin G (IgG) and fibrinogen, so-called opsonins, to the particles. This opsonization of the particles marks the particles for destruction. Opsonins are recognized and bound by macrophages, thereby, enhancing the phagocytosis through these cells.

Taken together, coating of ENMs has several purposes: to avoid agglomeration/aggregation of the particles, to change the surface charge of the particles, target ENMs for uptake by specific cell types, change the bioavailability and degradation of the particles. All this leads to an improvement of the performance of nanomedicines as it diminishes the host defence mechanisms.

12.3.1.3 Surface Charge

Another physicochemical property that has been identified to play an important role for the toxicity of ENMs is their surface charge that can either be positive, negative or neutral. Studies showed that positively charged ENMs induce higher levels of toxicity compared to negatively charged particles of the same chemical composition. The induced toxic responses included cytotoxicity, disruption of cellular membrane integrity, apoptosis, necrosis, loss of mitochondrial membrane potential [36]. The surface charge of ENMs is dependent on their surface coating and surface functionalization. For example can the surface of carbon nanotubes (CNTs) be functionalized using acids. This treatment will result in carboxyl, carbonyl and hydroxyl groups at the surface of the nanotubes thereby leading to a negative charge of the

nanotubes. A negative surface charge has been linked to an increased cytotoxicity [37–42]. On the other hand was shown that a surface functionalization that increases the water solubility of the ENMs decreases the cytotoxicity of CNTs [43].

Interestingly, just by altering the size of ENMs their hydrophobic or hydrophilic properties can change. The reason for that lies in a changed curvature of the particle-water interface. Small-sized ENMs can, therefore, be hydrophobic whereas larger ENMs of the same chemical composition and same coating can be hydrophilic [44].

For the development of nanomedicines, it is important to notice that positively charged particles often form aggregates upon intravenous injection. This can cause, e.g., potentially lethal embolisms in the lung capillaries [45]. The surface charge has also indirectly an influence on the toxicity of ENMs as it influences the efficiency of the uptake of the particles by the cells, the uptake pathways and the cellular distribution [36, 39, 40].

12.3.2 Pharmacokinetics

The physicochemical characteristics of ENMs does not only has an influence on the toxicity of the particles but also on their pharmacokinetics. Pharmacokinetics, the knowledge and investigation of what the body does to a drug, follows the so-called ADME scheme studying the **A**dsorption, **D**istribution, **M**etabolism and **E**xcretion of a drug. Knowledge about pharmacokinetics is, therefore, very important for the development and fine-tuning of the efficacy, but also, toxicity of nanomedicines and plays a crucial role in health risk assessment for nanomedicines.

12.3.2.1 Absorption

In pharmacokinetics absorption is the process by which the drug crosses biological membranes and reaches the bloodstream. The absorption efficiency and involved absorption mechanisms of nanomedicines are greatly depending on how the nanomedicine is administered. Administration of nanomedicines can occur orally, via inhalation, dermally, intravenously, subcutaneously and intramuscularly. Obviously, the absorption process for a drug is bypassed if the drug is directly injected into the bloodstream. As most of the nanotoxicological studies focus on the unintentional exposure of workers and consumers to ENMs, absorption after subcutaneous and intramuscular administration is much less investigated and there is no conclusive information on the absorption process available at this point.

The effectiveness of the absorption process differs greatly with uptake through the skin (dermal) as the least effective and injection, either subcutaneously, intramuscular or intravenously as the most effective administration routes. Although research is still ongoing, most of the studies on dermal absorption of ENMs confirm that the human skin can normally not be penetrated by particles even when they are in the nanoscale range [23]. Neither titanium dioxide nanoparticles nor quantum

dots were able to reach the bloodstream [21]. Even if titanium dioxide nanoparticles were found in the stratum corneum the particles never reached the lower layers of the dermis. The stratum corneum consists of multiple layers of dead keratinized cells, which are difficult to penetrate for ENMs. However, this might be the case if the skin is damaged by wounds, sunburn or skin diseases like eczema. In addition, movement and stretching of the skin, particle charge, follicular openings, gender, and age might affect the barrier function of the skin [21]. But as there are reports showing the absorption or at least penetration of the stratum corneum there is currently no consensus about the ability of ENMs to be absorbed through the skin [46].

Absorption of ENMs via inhalation is probably the best investigated exposure route and will also be discussed in more detail in the section “Pulmonary toxicity”. The most important parameter for absorption of ENMs through inhalation is their size and aerodynamic diameter. Depending on these parameters particles will be deposited more or less deep in the respiratory tract. The main mechanism for particle deposition is diffusion due to displacement when the particles collide with the molecules of the air. Depending on the size of the ENMs, they will be deposited in the nasopharyngeal, tracheobronchial or alveolar region [13]. Alveoli are the deepest part of the lung and nanoscale particles have been found to reach this region. The approximate size limits for particle deposition are that particles with an aerodynamic diameter $>50\ \mu\text{m}$ do not enter the respiratory tract as they are filtered quite efficiently by the nose, particles $>10\ \mu\text{m}$ are deposited in the upper respiratory tract, particles between 2 and $10\ \mu\text{m}$ can reach trachea, bronchi and bronchioles, and particles smaller than $1\ \mu\text{m}$ can reach the alveoli [13]. ENMs that reach the alveoli are mainly cleared by alveolar macrophages [13]. However, if the particles persist in the alveoli they are able to access the pulmonary interstitium either through diffusion or, more likely, transcytosis through the alveolar epithelium. From there the particles can cross the endothelium of the capillary, enter the bloodstream and translocate to systemic sites. Alternatively, the particles could enter sensory nerve endings that are embedded in the airway epithelial, a mechanism that seems to be specific for nanoscaled material [23, 47].

Particles that have been inhaled may be cleared from the lungs via the mucociliary escalator and through this way reach the gastrointestinal tract (GIT) [13, 48]. In addition, the GIT is also an important absorption route for nanomedicines and nanocarriers are currently under development for a more effective oral uptake of drugs and vaccines [46]. Nanomedicines that are administered orally are ingested into the GIT and are absorbed by a process called persorption, the paracellular translocation through transitory leaks in the epithelial cell layer [49–51]. It is thought that loosened tight junctions in the mucosa allow undissolvable particles in nano- but also microscale to be transported in the epithelial cell layer. From there the transport occurs into the sub-epithelial region via the thoracic duct either through lymph tracts or through veins and reach the bloodstream. This process seems to be quite fast as within a few minutes particles are found in the peripheral blood [49–51]. However, newer studies on ENMs showed that the absorption through the GIT increases with decreasing particle sizes and that micro-sized particles are trapped within the Peyer’s patches, which are organized lymphoid nodules that are found in the lower regions of the small intestine. Particles trapped there does not seem to

reach the systemic circulation in high numbers [48]. Besides the size, the charge of the ENMs is important as positively charged particles are more efficiently absorbed than negative or neutral charged ENMs [46].

12.3.2.2 Distribution and Cellular Uptake

After absorption or direct administration by intravenous injection the nanomedicine is distributed in the body through the bloodstream. Within pharmacokinetics, distribution is the reversible transfer of drugs away from the site of absorption to other sites within the body, including the target site, into interstitial and intracellular fluids. Again, the physicochemical properties of the ENMs are important for the biodistribution as they influence the way ENMs interact with cells, body fluids and proteins. The binding of ENMs to proteins can influence the mobility of the particles. If these proteins promote cellular uptake of the ENMs in specific organs or immune cells the biodistribution of the particles might be limited.

A general rule is that smaller ENMs have a much greater biodistribution compared to larger ENMs. For example, intravenously administered 10 nm gold nanoparticles were found in liver, spleen, kidney, testis, thymus, heart, lung and brain whereas 50 and 250 nm gold nanoparticles were only found in liver and spleen. One explanation is that the 10 nm particles were too small to be efficiently recognized and internalized by professional phagocytes that normally will clear the blood for foreign particles. Therefore, the 10 nm particles were able to reach more organs compared to the larger particles [52]. Iron oxide nanoparticles with a size of 22 nm were shown to be quickly translocated to the bloodstream and distributed to liver, spleen, kidney and testis after intratracheally instillation of rats [53]. Although research is still ongoing it seems that in general particles with a small diameter (10–20 nm) or a positive charge are more easily translocated through the alveolar barrier of the lung [21]. However, the situation might be different when chronic exposure occurs or in the case of a pathological situation. For example, inflammation seems to increase the translocation of ENMs from the alveoli into the blood and has, thereby, an influence on the biodistribution of the particles [54, 55].

The cell membrane is the last barrier ENMs have to cross if they are used as carriers to transport the drug into the target cells. Due to the particulate or vesicular form of nanomedicines most of the cellular uptake of nanomedicines will occur via active transport mechanisms into the cell. These active transport mechanisms include internalization pathways like phagocytosis, (macro)pinocytosis and receptor-mediated endocytosis via clathrin coated pits or caveolae [21]. Which of these cellular uptake mechanisms apply is greatly dependent on the size of the particles and on their surface coating [21, 56–60]. If particles reach a size of larger than approximately 500 nm they are mainly taken up via phagocytosis by so-called professional phagocytes like neutrophils, monocytes, macrophages, dendritic and mast cells; smaller particles are primarily processed by endocytic pathways. An alternative for the uptake of larger aggregates (0.5–5 μm) might be micropinocytosis [61]. However, at this point the precise role and importance of this pathway for the uptake

of ENMs is not very well investigated. The surface coating can, e.g., allow binding to specific receptors on the cell surface. This is exploited for targeted drug delivery by interaction with receptors that are exclusively expressed at the surface of the target cells. In addition, targeting a specific receptor will also define by which cellular uptake routes the particles enter the cell as some receptors are exclusively found in clathrin-coated pits or caveolae [21]. After entry into the cells ENMs are present in intracellular membrane-coated vesicles. Depending on the uptake pathway, vesicles can be, e.g., endosomes, lysosomes or caveosomes. Furthermore, ENMs were also found in mitochondria, the nucleus or just free in the cytosol [21].

12.3.2.3 Metabolism

Metabolism of drugs covers the biochemical modification or biotransformation of pharmaceutical active substances or xenobiotics, substances that are foreign to the organism. The goal of these biochemical modifications is to convert lipophilic substances into more readily excreted hydrophilic products. The metabolic pathways are the same as for detoxification of poisons and include usually specialized enzymatic systems like the cytochrome P450 oxidase protein family. These enzymes are involved in the first of three metabolic phases where they introduce reactive or polar groups. In phase II, transferase enzymes such as glutathione S-transferases are catalyzing the conjugation of these modified substances to polar compounds that are in some cases further processed in phase III before they are pumped out of the cells by efflux transporters. These reactions are a defense mechanism of the cells to detoxify foreign substances, however, in some cases metabolic intermediates of normally non-toxic compounds can themselves be toxic.

Whereas the metabolism or biotransformation of the pharmaceutical active substance of the nanomedicine is likely to be well investigated and known, the metabolism of their carriers, the ENMs, is generally not very well investigated and understood. So far, ENMs were predominantly found not to be metabolized but that is, of course, very much dependent on the chemical composition of the ENMs. Qdots, e.g., seem to have a very long half-life in the body of several weeks or month. In contrast to nanoparticles, nanoscaled liposomes are likely to be much easier degraded and metabolized if they are able to fuse with cellular membranes. However, generally, there is still very little information available about what happens to ENMs after they have been taken up by cells. And there are concerns that breakdown of the nanostructures can again lead to unique unpredictable molecular responses.

Nevertheless, metabolization of ENMs has to be considered a very important step for clearance of the body from the particles. If ENMs are not metabolized or degraded they might not be excreted and, therefore, accumulate in the cells of the body. This might especially be a problem for repeated long-term administration of nanomedicines. In addition, if the particles are non-biodegradable even a short-term exposure and low toxicity of the administered ENMs might lead to a cumulative toxic effect over time. This is of special concern if an interaction with DNA occurs which could

result in carcinogenesis. Furthermore, it has to be kept in mind that theoretically all biological effects of ENMs can be enhanced if the particles persist within the body for several months, years or even through the entire residual lifetime.

Though, in some cases can the solubility of ENMs result in toxic effects. For example, some or all of the observed toxicity of ENMs consisting of ZnO, CuO or Ag has been attributed to the released metal ions [62, 63]. The dissolution of ENMs can either occur in body fluids but also intracellularly. Here, especially the acidic environment of endosomes and lysosomes are thought to contribute significantly to the degradation of ENMs and to the release of toxic metal ions [64]. However, the pH alone is not in all cases enough for the dissolution of particles. In case of silver nanoparticles the interaction with cellular proteins seem to play an important role for the degradation of the particles as well [65]. Nevertheless, it is important to notice that not all ENMs end up in endosomes and/or lysosomes and that there is a number of materials that either cannot be degraded in endosomes or lysosomes or are able to escape these compartments. Factors that are important for the dissolution rate of ENMs are the size of the particles, roughness, coating, and aggregation state [21, 66–69]. After dissolution of the ENMs the particle compounds might be available for biotransformation and subsequent excretion.

12.3.2.4 Excretion

The two major routes for excretion are through feces and urine and only to a lower degree via the lung and skin. When discussing the excretion of ENMs one has to differentiate between biodegradable and non-biodegradable particles. Biodegradable ENMs are digested and the metabolites excreted by the body through urine or feces, and does no longer pose a health threat. However, excretion of non-biodegradable ENMs might take very long or might even be not at all existing. In general, circulation of the particles in the blood is a prerequisite for their excretion. Also for non-biodegradable ENMs the major routes for excretion are via urine or feces.

For the excretion via urine the blood is filtered in the kidney through the renal glomerula and via this way particles with a size lower than 8 nm can be filtered out of the blood whereas particles that are larger in size will accumulate in the mononuclear phagocyte system (MPS) [21]. The MPS consists of phagocytic cells that are located in reticular connective tissue, which is found around the liver, kidney, spleen, and lymph nodes as well as in bone marrow. In the liver, this system is particularly well developed and the macrophages of the liver, the Kupffer cells, are responsible for clearance of the largest part of the particles. In the case of non-biodegradable ENMs, the particles accumulate and persist in the macrophages. In addition, hepatocytes are able to take up particles via endocytosis but if they can metabolize and secrete the particles into the bile is not known.

Although there is still a great demand for investigations on the fate of non-biodegradable ENMs it seems that particles that are administered intravenously are either rapidly cleared by the kidney or are taken up by the mononuclear phagocyte

system and persist in the body [21]. Water-soluble single-walled carbon nanotubes have been shown to be excreted via the renal route in rats and mice, whereas, titanium dioxide nanoparticles accumulate in the liver and spleen for several weeks [21]. Independent of the physicochemical properties of ENMs, the highest accumulation of particles is in general found in the liver [52, 70].

If particles or agglomerated ENMs reach a size of larger than approximately 500 nm they are mainly cleared from the blood via phagocytosis by so-called professional phagocytes like neutrophils, monocytes, macrophages, dendritic and mast cells. Especially macrophages have been in focus for pharmacokinetics and toxicity studies as they quite efficiently can clear the bloodstream from foreign particles. This might be a great problem for the efficacy of a drug but can also lead to toxicological complications and inflammatory responses if the particles persist within the cells. Positively charged nanomaterials are cleared fast from the blood and their aggregates accumulate in the liver and lung. Neutral ENMs have a decreased rate of uptake by macrophages of the liver or spleen. Neutral surface charge increases, therefore, the half-life of ENMs in the blood and the availability for uptake by other organs. In addition, binding of opsonins leads to enhanced phagocytosis and clearance of the particles from the bloodstream.

12.3.3 Mechanisms of Toxicity of Nanomaterials

Due to the small size of ENMs, the particles enter the organs, tissues and cells of the human body much easier than their larger counterparts. However, one of the most important questions is if the particles induce a toxicological response in the body once they are absorbed and what happens if non-degradable or slowly degradable ENMs accumulate in the body. If talking about toxicity of agents or drugs, a number of terms and definitions are used depending on what is the focus and aim of the toxicological study. Toxicity can be described based on the route, number and duration of exposure, primary toxic effects (target organ), and mechanism of toxicity. Terms like local and systemic toxic effects; acute, subchronic, chronic toxicity; transient, persistent, cumulative, latent toxicity are briefly described in the next section.

12.3.3.1 Toxicity Terms

The toxic effects of a drug, no matter if these are desired therapeutic effects or undesired side effects, can be either local or systemic. Local effects are those harmful effects that occur at the site of the initial contact, e.g., contact dermatitis. Systemic effects occur after absorption of the agent and include toxic effects in organs or tissues that are distant from the site of the original exposure [71]. Local and systemic effects can occur after acute (single) and repeated exposure where the repeated exposure can either be short-term (5 % of lifespan), subchronic (5–20 % of lifespan) or chronic (majority or entire lifespan) [71]. The vast majority of studies on

nanotoxicity are on short-term effects after acute exposure while long-term effects after chronic exposures are mainly unknown. Dependent on when and how long the toxic effects are arising, one distinguishes between transient, persistent and latent toxic effects. Transient effects are temporary and reversible whereas persistent effects are permanent and present during the complete residual lifetime. Latent toxic effects have a delayed onset and can appear days, weeks, month or even years after exposure. Latent toxic effects occur mainly after acute exposure, whereas, cumulative toxic effects are progressing effects after repeated exposure [71].

As nanomedicines are usually a mixture of different components, e.g., the pharmaceutically active drug and the carrier ENM, also other toxicological terms might be important that play a role especially for exposures to mixtures. The toxic effects of the different components of nanomedicines can be *additive* ($2+3=5$; the overall toxic effect is the sum of the toxicity of each component); *antagonistic* ($2+3<5$; at least one of the components antagonize the toxicity of the other); *potentiating* ($0+3>3$; one non-toxic component enhances the toxicity of another toxic component); or *synergistic* ($2+3\gg 5$; two toxic components are increasing the overall toxicity much more than the sum of the toxicity of each component). Potentiating and synergistic toxic effects can easily be confused. However, in case of a potentiating toxic effect one of the components has to be non-toxic, in case of a synergistic effect both components have to be toxic.

Although it is not very well studied how the different components of a nanomedicine are affecting each other's toxicity, there are a few examples where the co-exposure with two different kinds of ENMs leads to a potentiating or synergistic toxic effect. For example, pure cobalt and carbide particles have no toxic effect whereas the combination of both components leads to hard metal lung disease caused by the release of reactive oxygen species [72]. Oxidative effects are also observed after co-exposure with carbon black and iron oxide nanoparticles that are not observed for either particle type alone [73].

12.3.3.2 Reactive Oxygen Species, Oxidative Stress and Inflammation

The toxicological effects of ENMs on cells include cytotoxicity and genotoxicity. One if not the most important underlying mechanism for these effects is the induction of oxidative stress in the cells [74, 75]. Oxidative stress is caused by an imbalance between the formation of reactive oxygen species and the antioxidant capacity of the cells [13, 76]. Reactive oxygen species are chemically reactive molecules, and as the name suggests, do contain oxygen. Examples for reactive oxygen species are oxygen itself, superoxide anion, peroxide, hydroxyl radicals and ions, and hydrogen peroxide. These molecules are always present in cells as they are natural byproducts of the oxygen metabolism but, e.g., cellular stress, infection or other environmental factors can lead to an excessive formation of reactive oxygen species. In addition to reactive oxygen species, reactive nitrogen species containing nitric oxide can also be involved in the induction of oxidative stress [77].

There are different mechanisms of how exposure to ENMs might lead to an increased formation of reactive oxygen species. One possibility is the generation of free radicals by ENMs in aqueous suspensions *in vitro*. Another possibility is an increased production of reactive oxygen species in mitochondria but also the depletion of antioxidants and the subsequent impairment of the antioxidant capacity have been discussed as possible mechanisms. Reactive oxygen species and oxygen-free radicals are mainly produced in the mitochondria and thereby the mitochondria themselves are a major target for oxidative stress and injury.

The existence of too high concentrations of reactive oxygen species within the cell induces lipid peroxidation, mitochondrial damage, damages to DNA, RNA and proteins and lead to the induction of redox sensitive pathways that are involved in pro-inflammatory responses, cell cycle/proliferation as well as apoptosis (programmed and targeted cell death) and necrosis (non-programmed cell death) [74, 75]. The increased formation of reactive oxygen species is thought to be involved in the inactivation of protein functions that are important for cellular DNA repair. Reactive oxygen species might directly attack DNA leading to modified DNA bases like, e.g., 8-oxo-7,8-dihydroguanine (8-oxoG) as the most abundant and best investigated DNA alteration [75, 78]. Impaired repair of DNA alterations such as modified nucleotides is associated with mutagenesis and carcinogenesis. Furthermore, oxidative stress is thought to be involved in a number of different diseases like, e.g., Parkinson's and Alzheimer's disease, and cardiovascular diseases like atherosclerosis and myocardial infarction [74]. Therefore, the increased formation of oxidative stress by most of the investigated ENMs can be linked to a number of diseases and is the reason for concerns about the health effects of ENMs. However, as mentioned before, epidemiological human studies on ENMs are still very rare and many studies are using unrealistic high particle concentrations or purely characterized ENMs. Therefore, definite and especially general conclusions cannot be drawn on the induction of reactive oxygen species by ENMs and the toxicity of ENMs at this point.

12.3.3.3 Genotoxicity and Carcinogenicity

The high surface reactivity of ENMs and the induction of oxidative stress upon exposure to ENMs has raised the concern that they might be genotoxic and carcinogenic. An agent is classified as genotoxic when it has a DNA damaging capacity. Genotoxic events are normally very efficiently repaired by the cellular DNA repair system unless the DNA damage is too extensive. If the latter is the case programmed cell death, apoptosis, is induced. Mutagenesis is the permanent change of the original genetic information and occurs only if the DNA damage leads to persistent mutations within the genome. These persistent mutations can eventually lead to uncontrolled cell growth in form of neoplasms, which in the worst case can be malignant. The formation of malignant neoplasms, also better known as cancer, is called carcinogenesis. Carcinogenesis is a multistep process and requires not only an initiation stage but also a promotion stage. In the initiation stage the cell is exposed to a genotoxic agent whereas in the promotion stage the initiated cell is

exposed to a promoting agent. This multistep process is resulting in carcinogenic characteristics of the cell like evasion of apoptosis, uncontrolled cell growth and metastasis. The promoting agent does not need to be genotoxic itself but either persistent or repeated exposure is required for carcinogenesis. Whereas the initiating stage has no threshold (one DNA damaging event is in theory enough at this stage to initiate the cell), the promoting activity of an agent may have a threshold. In principle, the initiating and promoting agent could be the same [71].

The genotoxicity of ENMs and the underlying mechanisms are, at the point of writing this book chapter, not very well understood. In 2012, 4346 articles had been published on nanotoxicology whereof 94 described *in vitro* and 22 *in vivo* genotoxicity studies [7]. Although this number has tripled since then, it shows that there still is limited information on the genotoxicity of ENMs available considering the large amount of different types of nanomaterials that has been developed. However, *in vitro* studies suggest that several ENMs may have genotoxic potential, e.g., carbon nanotubes, C60 fullerenes, titanium dioxide and silver nanoparticles [7, 9, 79–81]. However, the results are somewhat conflicting and often due to limited information on the physicochemical properties of the investigated ENMs or variations in the experimental settings hardly to compare. Despite of these limitations, several mechanisms and factors are currently discussed that could lead to genotoxicity and carcinogenicity of ENMs. These could either be direct primary mechanisms (direct interaction of the particles with the genome), indirect primary mechanisms (interaction with proteins involved in cell cycle, binding to mitotic spindle components, inhibition of antioxidant defense and DNA repair activity or release of toxic ions from soluble ENMs, formation of reactive oxygen species by mitochondria) or secondary mechanisms (formation of reactive oxygen species by inflammatory cells) [7, 82]. These genotoxic mechanisms can result in oxidative modifications of DNA bases, bulky DNA adducts, single and double strand breaks, structural changes of the DNA (deletions, duplications, inversion and translocation of chromosome segments) or changes in the number of chromosomes [7].

Importantly, information on genotoxic and carcinogenic effects of ENMs on humans is even more limited at this point. Epidemiological studies on workers that were exposed to titanium dioxide nanoparticles were inconclusive and could not show an association between exposure to these particles and an increased cancer risk [83]. However, indications for a genotoxic potential of especially non-biodegradable persistent ENMs should of course be taken seriously and require further and more detailed investigations.

12.3.3.4 Neurotoxicity

Neurotoxicological health effects are adverse effects on the brain and the central nervous system. Normally, the blood-brain barrier protects the brain from entry of foreign particles or other unwanted compounds. The passage even of small molecules is tightly regulated and efficient translocation of drugs through the blood-brain barrier is hard to achieve. However, depending on their physicochemical

properties, especially their size and surface charge, nanoscale particles might be able to enter the brain or the central nervous system. ENMs that are 20–50 nm in size as well as hydrophobic particles might be able to enter the brain even if the blood-brain barrier is intact [23, 84–86]. However, the reports in this matter are conflicting and intravenous injection of ENMs like 40 nm gold nanoparticles did not lead to translocation of a detectable amount of particles into the brain [87]. In contrast, a recent study by Huang et al. showed an accumulation of intravenously administered lipid nanoparticles in the brain parenchyma of mice after 3 h. The particles persisted there for more than 24 weeks [88]. Furthermore, for polymeric nanoparticles the surfactants seem to be more important than the size of the particles [89]. Aging, injury or disease may limit the protective capacity of the blood-brain barrier and allow for an easier access [90]. Another possibility is the entry via the olfactory bulb where there is a connection between the nasal epithelium and olfactory neurons [47]. This has been shown for carbon nanotubes, gold nanoparticles, quantum dots and manganese oxide nanoparticles [13, 47, 91–94]. ENMs were found in the olfactory bulb but have also been found in the hippocampus [95]. The entry into the brain was shown to be associated with an inflammatory response [93, 95, 96]. Although animal studies have shown that ENMs can reach the brain through the olfactory bulb, it is not known which role this entry pathway might occur in humans, as humans have a significantly less developed olfactory bulb compared to rodents [21]. In addition, it is not known what the health effects of ENMs actually are after they reach the brain or central nervous system. Animal and *in vitro* studies suggest that the presence of ENMs in the brain can cause brain damage. Neutrophils and lymphocyte numbers as well as protein carbonyl levels were increased and oxidative stress, lipid peroxidation and inflammatory responses (glia activation) have been discussed to be induced as a result of the high surface area and reactivity of ENMs [47, 76, 88, 93, 97, 98]. In addition, it is more likely that neurotoxicity is of chronic nature than to be acute due to the difficulties of reaching the brain [90].

12.3.3.5 Pulmonary Toxicity

From studies on nanoscale particles in air pollution we know that their inhalation can have adverse health effects and is associated with increased risk to develop cardiovascular diseases, lung fibrosis and lung cancer. The adverse health effects that are associated with inhalation of particles are occurring due to the deposition of the particles in the lung. Although there has been a number of studies on nanoscale particles in air pollution on human health, there are still only limited data available on the health effects of ENMs. This is also due to the difficulties to separate exposure to ENMs from background exposure of ambient particles. In a study on workers that were exposed to polyacrylate nanoparticles it has been suggested that these particles induce pleural effusion, pulmonary fibrosis and granuloma [99]. The respiratory effects that have been described are mainly inflammation, oxidative stress and functional disturbances. The inflammatory response includes local invasion of

leukocytes and release of cytokines [21]. As described before, the toxic effect of ENMs are dependent on the size and shape of the particles. Nanoscale particles have been shown to reach the alveoli, the deepest part of the lung. Alveoli have an extreme large surface area (estimates are between 30 and 100 m²) but the distance between the surface of the alveoli and the bloodstream measures only 2 µm [21]. Therefore, this region is less protected against inhaled particles [100]. Macrophages are mainly responsible for clearance of particles in the lung via phagocytosis. However, the efficiency of this process is strongly dependent on the size of the particles [21, 101]. It seems that alveolar macrophages are unable to recognize particles as foreign and to phagocyte them when they are less than 70 nm [23]. In contrast, nanofibres with a length of more than 20 µm are too long for phagocytosis. In both cases, the particles are suspected to stay in the lung for month or even years resulting in nonspecific pulmonary inflammatory responses. These inflammatory processes might even spread systemically as, e.g., also an increased risk for cardiovascular diseases is associated with pulmonary exposure to ENMs [23].

12.3.3.6 Cardiovascular Toxicity

Cardiovascular toxicity of ENMs or other nanoscale particles, e.g., ultrafine particles in air pollution, has mainly been observed after exposure via inhalation. Several epidemiological studies have shown the association between particles in air pollution, especially the ultrafine fraction, and cardiovascular diseases [102]. The observed short-term health effects after exposure to nanoscale particles and particles in air pollution include arrhythmia, coagulation disturbances, thrombosis, blood pressure abnormalities and in the long perspective a generally increased risk for development of cardiovascular diseases. Although the reasons and mechanisms are still somewhat unclear, it is thought that deposited particles in the lung induce inflammatory responses and conditions that cause these health effects. The release of inflammatory and prothrombotic mediators from the site of exposure into the blood might cause the activation of immune cells leading to the development of these adverse conditions [103]. Especially if these inflammatory responses are chronic and become systemic they will, obviously, have stronger effects on the cardiovascular system. After uptake by alveolar macrophages ENMs might be translocated from the respiratory to the cardiovascular system where the particles can directly induce cardiovascular toxicity by induction of inflammatory responses through cellular stress and increased release of reactive oxygen species [21]. Although epidemiological data for ENMs are still rare, it is believed that ENMs underlie the same toxicological mechanisms and, thereby, have somewhat the same adverse effects on the cardiovascular system as ultrafine particles in air pollution. As several studies have shown that ENMs can induce pulmonary toxicity this is believed to be an indicator for potential cardiovascular damage due to the close association of pulmonary and cardiovascular toxicity [104].

12.3.3.7 Reproductive Toxicity

The reproductive toxicity, which includes adverse effects on the sexual function and the fertility of adult males and females as well as the developmental toxicity in the offspring, is probably the least investigated toxicological effects of ENMs. Some studies in mice have shown that titanium dioxide nanoparticles were able to cross the blood-testes barrier and reduced the sperm production in the offspring of the treated mice. In addition, it has been suggested that these particles might be able to affect the development of the central nervous system in the offspring as they affected the gene expression of genes involved in the development and function of the neural system [46]. However, since a possible accumulation of ENMs has been found, further studies are needed to exclude any reproductive and developmental effects due to an exposure to ENMs or nanomedicines. In addition, the potential of ENMs to cross the fetal-placental barrier has to be investigated as well [46].

12.4 Drug Safety Testing

Like conventional drugs, also nanomedicines have to be approved before used on patients. In the U.S.A. this is done by the U.S. Food and Drug Administration (FDA) and in countries of the European Union, the European Medicines Agency (EMA) is the responsible regulatory agency. The approval process involves several phases including preclinical studies as well as clinical trial phases. In these different phases the safety and efficacy of the drug is investigated by the applicant under the supervision of the responsible regulatory agency.

12.4.1 Preclinical Studies

Before entering the clinical trials drugs are tested in pre-clinical studies, normally by the drug-developing pharmaceutical company. The aim of these pre-clinical studies is to collect basic safety and efficacy data. Based on these data a plan for further testing of the drug on humans is developed and an application for clinical trials is submitted. Another important goal of these studies is to ensure that the drug is a promising candidate that justifies the enormous costs and efforts that are associated with a drug approval process. Therefore, these pre-clinical studies are quite extensive and include *in vitro* cell culture studies as well as *in vivo* animal studies to investigate the preliminary efficacy, toxicity and pharmacokinetics of the drug. In the case of nanomedicines, nanotoxicological aspects have to be considered in addition to standard toxicological investigations and evaluations.

12.4.2 Clinical Trials

The clinical trials involve three phases where the drug is tested on either healthy volunteers (phase I) or patients (phase II and III). The goal of phase I is the determination of the most frequent side effects of the drug and, frequently, the pharmacokinetics of the drug that gives information about how the drug is taken up, transported, metabolized and excreted. Typically, between 20 and 80 volunteers are involved in phase I and the safety of the drug is stressed. However, there are circumstances when patients have to be enrolled. This is the case when the drug is expected to cause severe side-effects in healthy individuals. After a successful phase I, approximately 100–300 patients are enrolled in phase II where the effectiveness of the drug is investigated. Normally, the effect of the drug on patients will be compared to patients receiving either a placebo or standard treatment. Phase III contain 1000 or more patients to further investigate the safety and effectiveness of the drug. Different dosages and the use of the drug in combination with other drugs are studied. Based on the results from the clinical trials the authorities decide whether the drug can be approved or not. However, as it is not possible to predict and determine all side-effects and especially long-term effects in the clinical trials, the drug will be further monitored to detect any adverse effects when on the market. Again, when investigating the safety of nanomedicines their specific nano-related characteristics and properties have to be taken into consideration in all phases of the drug approval process as special nanoscale related safety issues have to be addressed.

12.5 Risk Assessment of Engineered Nanomaterials

12.5.1 Risk Assessment

Toxicological investigations of ENMs are important not only for drug safety testing, but are also essential parts of the risk assessment of these agents. Nanomedicines and the used ENMs have to be manufactured and depending on the scale of this manufacturing process unintentional exposure of workers could occur. Risk assessment is also necessary to regulate the use of nanomedicines and ENMs properly. To cover all aspects of safety considerations for nanomedicines a short introduction into risk assessment will be given based on the WHO tool kit and IPCS harmonization project (WHO Human Health Risk Assessment Toolkit: Chemical Hazards; http://www.who.int/ipcs/methods/harmonization/areas/ra_toolkit/en/).

For the investigation of the adverse effects of a newly developed drug, a broad spectrum of methodologies are used that range from experimental in-vitro studies to animal studies and epidemiological investigations on whole populations. The identification of adverse effects form the basis for risk assessments of any given chemical,

physical or biological agent. *Risk Assessment* is a process where it is evaluated to which degree, and with which probability, these agents affect human health and the environment. It is the primary objective to identify and characterize potential hazards, estimate exposure and assess the overall risks for humans or the environment. The assessment if an agent poses a risk is the first component of a risk analysis that also includes risk management and risk communication.

When talking about risk it is important to keep in mind that a hazardous substance does only pose a risk to humans but also the environment if there is a likelihood for an exposure. Risk is therefore defined as a function of hazard and exposure:

$$\text{Risk} = f(\text{Hazard}; \text{Exposure}) \quad (12.1)$$

In Eq. (12.1) is risk = zero if either hazard or exposure equals zero. Equation (12.1) is probably the most important risk assessment paradigm and can be very illustratively explained using the tiger in a cage example. Everybody will probably agree that it is a risk to visit a living tiger inside its cage, especially a hungry one. In this situation, we are exposed to a hazardous biological agent. However, when the tiger is separated from the visitor by a cage there is no risk to the visitor (there is a hazard but no exposure) just as a mounted tiger outside a cage poses no risk to the visitor (there is an exposure but no hazard). This example shows that for a thorough risk assessment the exposure assessment is just as important as the identification of the potential hazards of chemical, physical or biological agents.

A risk assessment will always begin with a problem formulation to establish the scope and objective of the risk assessment. The risk assessment itself consists then of four steps including *hazard identification*, *hazard characterization*, *exposure assessment* and *risk characterization*. In the following chapters a short overview over these four steps is given.

12.5.1.1 Hazard Identification

The first step in risk assessment is the hazard identification that is mainly based on the results from toxicological studies. These toxicological studies include human studies (mostly epidemiological studies), animal-based and in vitro toxicology studies as well as structure-activity studies. Although risk assessment and toxicology also include the investigation of adverse effects on the environment, the following will focus on the health hazards and effects on humans as nanomedicines are primarily intended to be used in humans being well aware of that the production of nanomedicines could pose environmental hazards and risks.

The purpose of the hazard identification is to identify (1) the specific hazard, (2) the type and nature this hazard may have to an individual or (sub)population and (3) investigate if exposure to the agent has the potential to be harmful.

The hazard identification begins with the identification of the chemical composition and, as nanomedicines consists of nanomaterials, particle characteristics of the nanomedicine. Identifying the chemical composition of a nanomedicine will give

information if the parent material is already classified by the CLP regulation of the European Union (CLP stands for “Classification, Labelling and Packaging”) and if it is already known to be hazardous. The health hazards that are CLP classified include acute toxicity, sensitization of respiratory tract and skin, skin corrosion and irritation, serious eye irritation and eye damage, reproductive toxicity, germ cell mutagenicity, carcinogenicity, specific target organ toxicity after single and repeated exposure, and aspiration toxicity.

In addition, as previously mentioned high aspect ratio nanomaterials like, e.g., nanotubes, nanofibers, nanowires and nanorods are generally considered hazardous when they are at the same time biopersistent, able to pass ciliated airways and able to initiate frustrated phagocytosis, which leads to the release of pro-inflammatory molecules.

If the nanomaterial is not categorized as high aspect ratio nanomaterials (HARN) there has to be investigated if the nanomaterial induces acute or chronic toxicity including genotoxicity, neurotoxicity, carcinogenicity, pulmonary, cardiovascular, or reproductive toxicity or if the nanomaterial accumulates in organs. If there cannot be excluded that the nanomaterial is potentially hazardous one proceeds with the hazard characterization.

12.5.1.2 Hazard Characterization

Whereas the hazard identification recognizes the type and nature of the hazard, the objective of the hazard characterization is to obtain a qualitative or quantitative description of the inherent properties of the agent that is potentially hazardous when one is exposed to it. A quantitative description will, wherever possible, include a dose-response assessment, identification a no-observed-adverse-effect level (NOAEL), no-observed-effect level (NOEL) or cancer potency factor and take uncertainty factors into account. In addition, based on dose-response assessments no effect levels (NEL) of an agent are derived.

The information on NOAEL and an eventual cancer potency factor are used to establish tolerable daily intake (TDI), acceptable daily intake (ADI) value as guidance values while including uncertainty factors like, e.g., interspecies and intraspecies variability, and data quality. TDI and ADI are both referring to a dose that is safe to consume for humans during an entire lifetime. ADI is used for, e.g., food additives, whereas, TDI is used for agents we are unintentionally exposed to like, e.g., air pollution, contaminants of water.

Depending on the uncertainty level of these data, e.g., if there has to be extrapolated from *in vitro* or animal studies to humans, if it is necessary to include susceptible population groups etc., uncertainty factors in the range of 10–10,000 are applied to cover also worst case scenarios and population groups. By applying an uncertainty factor the acceptable concentration for the exposure to an agent is reduced to a value where also the most susceptible population groups are not experiencing adverse health effects. Thereby it is avoided that parts of the human population might be unprotected.

Like for the hazard identification, data are obtained from human studies (mostly epidemiological studies), are animal-based or *in vitro* toxicology studies as well as structure-activity studies or combinations of these studies.

12.5.1.3 Exposure Assessment

The exposure assessment does not only include the investigation whether there is a contact with a potentially toxic agent. It determines also the concentration, route and duration of exposure. It has also the goal to establish safety margins and thresholds by evaluating the likelihood and level of exposure. An exhaustive exposure assessment requires that all possible exposure scenarios are taken into account and that includes the identification of particular susceptible population groups like children, pregnant woman, elderly and predisposed people. Another important part of the exposure assessment is the identification of the route and duration of exposure. In addition, this knowledge is important for the regulation and legislation of toxic agents but also nanomedicines. As described before, the exposure can occur orally, via inhalation, dermally, intravenously, subcutaneously and intramuscularly and the toxicity of a substance may be dependent on the route of exposure. For the estimation of exposures, either measurement or modelling approaches are used. In most cases when unintentional exposures occur the exact measurement and determination of an exposure is not available and a worst case scenario is modelled.

For nanomedicines the administered dose is exactly known. However, the internal dose is dependent on the absorption and excretion of a drug and can therefore vary from the administered dose. In some cases, the internal dose can be estimated using biomarkers. Biomarkers are measurable indicators for the presence of a substance in the body and can be measured in tissues or body fluids like blood, urine but also feces. The duration of an exposure can be acute, subacute, subchronic or chronic. The acute exposure has a duration of less than 24 h and is often a single exposure. Subacute exposure refers to repeated exposures with a duration of up to a month and subchronic exposure lasts for 1–3 month. If the exposure duration exceeds 3 month chronic exposure occurs.

12.5.1.4 Risk Characterization

The last step in the risk assessment process is the risk characterization. The aim of the risk characterization is, if possible, quantitative determination of the probability that known potential adverse health effects occur under defined exposure conditions. These exposure conditions might be actual or predicted exposures. Risk characterization includes the results that have been obtained from hazard identification and characterization and exposure assessment. Based on these results, risk quotients or margins of safety are calculated and exposure and no effect levels are compared to estimate the risks. However, there are no

absolute measures of risks and the conclusion if and when a given agent comprises a risk might vary from scientist to scientist especially if they include implicit value judgments.

12.6 Regulatory Affairs

The regulation of the use of ENMs is still debated due to the relatively short time period ENMs have been in focus of toxicological investigations. Many toxicity and safety related uncertainties of ENMs have not been clarified and this gap in knowledge results of course in uncertainties about the safety of nanomedicines. Nevertheless, nanomedicines have been authorized by licensing agencies like the FDA and EMA for more than 30 years.

The primary regulatory bodies in the U.S. and European Union (EU) that are relevant for the regulation of nanomedicines are the FDA and EMA, respectively. The EMA is accompanied by several committees and groups, whereof, the Committee for Medicinal Products for Human Use (CHMP), the Innovation Task Force (ITF) and the New and Emerging Technologies (N&ET) Working group are relevant for the regulation of nanomedicine. In the case of the FDA the following centers and groups are relevant: Center for Drug Evaluation and Research (CDER), Center for Devices and Radiological Health (CDRH), Center for Biologics Evaluation and Research (NTIG) and Nanotechnology Task Force (NTF). In other countries, national agencies are responsible for the approval of nanomedicines.

12.6.1 *Definition of Engineered Nanomaterials for Regulatory Purposes*

For risk assessments and regulatory purposes a definition of the term nanomaterial is of utmost importance. However, the sheer number of different nanomaterials makes a definition much more complex than one perhaps first realizes. On one hand, the definition for nanomaterials has to be so comprehensive that it includes all nanomaterials but should, on the other hand, be also be simple and precise as possible. At the moment, there are no standardized definitions what a nanomaterial is and the definition varies between organizations and countries. One of these definitions was proposed by the International Organization for Standardization (ISO) in cooperation with the European Committee for Standardization (CEN). According to the ISO/TR 11360:2010 definition a nanomaterial is a material with any external dimension in the nanoscale or having internal or surface structure in the nanoscale with nanoscale (or nano range) defined as size range from approximately 1 nm to 100 nm (ISO/TR 11360:2010 Nanotechnologies—Methodology for the classification and categorization of nanomaterials <https://www.iso.org/obp/ui/#iso:std:iso:tr:11360:ed-1:v1:en>).

From a scientific point of view an implementation of a fixed size limit might not make sense and the approximate size range might be preferred. However, for regulatory purposes a fixed size limit is needed and, therefore, implemented by the EU Commission. This definition is based on the ISO definition, an opinion of the Scientific Committee on Emerging and Newly Identified Health Risks (SCENIHR) and a report of the Joint Research Centre (JRC). Nanomaterial means a “natural, incidental or manufactured material containing particles, in an unbound state or as an aggregate or as an agglomerate and where, for 50 % or more of the particles in the number size distribution, one or more external dimensions is in the size range 1–100 nm.

Fullerenes, graphene flakes and single wall carbon nanotubes with one or more external dimensions below 1 nm should be considered as nanomaterials.

In specific cases and where warranted by concerns for the environment, health, safety or competitiveness the number size distribution threshold of 50 % may be replaced by a threshold between 1 % and 50 %”. In addition, the European Commission has acknowledged that a upper limit of 100 nm might not always be scientifically justified and that there are special circumstances prevailing in the pharmaceutical sector (EU Scientific Committee on Emerging and Newly Identified Health Risks. Scientific basis for the definition of the term ‘Nanomaterial’. European Commission, Brussels, Belgium (2010)).

In the U.S.A. the FDA has another definition for nanomaterials and according to this a nanomaterial is defined to be any material with at least one dimension smaller than 1000 nm and a nanoparticle is an object with all three external dimensions in the size range from ~1 nm to 100 nm (<http://www.fda.gov/RegulatoryInformation/Guidances/ucm257698.htm>; accessed August 2014).

The lack of an adequate definition of nanomaterials becomes especially problematic when dealing with follow-up nanomedicines that are based on already approved medicines formerly not classified as nanomaterials or not registered to contain nanomaterials.

Another reason for the still ongoing debate on the definition of nanomaterials is the challenge of a comprehensive characterization of ENMs used as nanomedicines. The methods that are available for characterization are not necessarily applicable for ENMs in complex mixtures and sometimes only the primary material might be suitable for characterization. A too rigid definition and regulation might, therefore, lead to the reluctance of regulatory agencies to issue manufacturing licenses or marketing authorizations. Therefore, not only a definition for nanomaterials is needed but also a definition of standards for the characterization of nanomaterials [105].

12.6.2 Regulation of Nanomedicines

The exact definition of a nanomaterial is only one of many questions that have to be addressed for a proper regulative approach for nanomedicines. In addition, the situation might be even more complicated when it has to be decided if a nanomedicine that uses ENMs as carrier is a medicine (medical product) or a medical device? This

classification is not unimportant as medical products and medical devices are regulated in different ways. Furthermore, before a new nanomedicine can be accepted for the use on patients it has to be decided, which regulatory regime is applicable. A medical device fulfills its function by physical means like mechanical or chemical action, whereas, a medical product exclusively fulfills its function by pharmacological, immunological or metabolic means [105]. Clarification on these matters is of course of uppermost importance not only for the safety of patients but also for the pharmaceutical industry that demands a greater harmonization in current nanomedicine regulatory framework. In the EU, the decision on the classification as a medical device or medicine is based on the EU Directive 2001/83/EC on human medicines, as amended, and the EU Directive 93/42/EEC on medical devices, as amended. According to these directives the decision is been made according to the principal mode of action of the nanomedicine. Especially for nanomedicines, which have a complex mode of action, this may prove difficult as their mode of action might involve and combine physicochemical and pharmacological properties. In addition, in some situations, when the nanomedicines are based on viable cells or tissues, they might also be classified as advanced therapy medicinal products and fall under the Regulation (EC) 1394/2007 on advanced therapy medicinal products [105].

Toxicological investigations are the basis for the safety and risk assessment of nanomedicines and as such crucial for their approval. For conventional medicines a battery of OECD approved methods are available for the investigation of eventual toxic effects of a drug. However, it is not clear if these tests are applicable for ENM based nanomedicines. For example, bacteria-based genotoxicity assays like the Ames test may not be appropriate as ENMs may not be able to penetrate the bacteria [105].

Taken together, it becomes clear that the regulation of ENMs and nanomedicines is still under development. Just recently as from 21 June to 13 September 2013 the European Commission had launched a public consultation on the modification of the REACH Annexes (REACH—Registration, Evaluation, Authorisation and Restriction of Chemical substances) on nanomaterials with the aim to improve the clarity on how nanomaterials are defined and their safety demonstrated. This consultation was open for the public and interested stakeholders. The European Commission states that “The REACH legislation must ensure a high level of health, safety and environmental protection. At the same time it should permit access to innovative products and promote innovation and competitiveness.” (http://ec.europa.eu/nanotechnology/policies_en.html).

12.7 Conclusion

Efficacy and toxicity of nanomedicines are inseparable interconnected as changes of the physicochemical properties can influence absorption, distribution, metabolism, and excretion of nanomedicines at the cellular but also organism level. For example, lipid particles or biodegradable ENMs might be less harmful than non-

degradable inorganic ENMs and are most likely to be used in the clinic sooner. Therefore, an early implementation of toxicological investigations is fundamental when developing new nanomedicines.

Although there are no nano-specific directives and regulations at this time, it is important to point out that nanomedical products are not unregulated. Although it is not clear at this point if this procedure is adequate, the FDA and EMA are applying of course the existing legislation on medical products and devices, tissue engineering etc. that are relevant for nanomedicines. Despite the doubts, in 2013 as much as 247 nanomedicine products were listed in FDA registers as approved or to be in various stages of clinical trials most of them attended to be administered intravenously [106]. Hopefully, in time the safety and regulatory challenges that come with nanomedicines are solved to utilize the full potential of nanomedicines.

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Chapter 13

The Application of Nanotechnology for Implant Drug Release

Morten Østergaard Andersen

Abstract The use of medical implants is a cornerstone of modern medicine. All implants face, however, a number of challenges including infection and inflammation which cause many of them to fail. In addition, tissue engineering implants must also direct stem cell differentiation and tissue regeneration in order to work properly. These problems may be overcome using drugs that are delivered directly from the implant. For this to work the drugs have to be protected until they have performed their function, their release must be timed with when they are needed, they may have to affect specific regions of an implant only and some drugs must be delivered to specific sub-cellular locations in certain cell types. This chapter explores how various forms of nanotechnology may be employed to reach these goals and reviews many of the studies that have used nanotechnology for different implant mediated drug release applications.

Keywords Controlled drug release • Drug encapsulation • Drug adsorption • Local drug delivery • Stem cell differentiation • Implant associated infection • Inflammation control • Tissue engineering

13.1 Introduction

Nanotechnology plays an increasing role in improving drug release from medical implants, thereby, improving their ability to combat infections, modulate inflammation and promote tissue regeneration. Organ failure caused by trauma, congenital defects, infection or cancer has always been a major problem for human health. As a result, implants that support or replace our body parts have been in use for millennia as evidenced by ancient dental implants [1], wooden prosthetic limbs [2] and precious metal plates for use in cranioplasty after surgical trepanation [3]. Major

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new improvements in treating organ failure took place in the twentieth century including the first successful donor organ transplantations of kidneys in 1954 [4] and of hearts in 1967 [5] as well as the development of complex artificial electromechanical organs such as the Jarvik-7, the first working total artificial heart, implemented in 1982 [6]. Today the implantation of both donor organs and electromechanical implants form a critical part of modern medicine. However, the inherent limitations to both of these types of organ and tissue replacements have led to the development of a new research field: Tissue Engineering (TE). TE combines implant materials with cells and drugs to establish biological replacement organs. Unfortunately, donor organs, electromechanical implants and tissue engineered organs all face several major limitations which limit their clinical potential. Many of these problems arise from detrimental interactions between the implant and its biological surroundings such as implant associated infections, the immune system, and the peri-implant tissue. Various strategies have been devised to address these issues such as using different implant materials or changing surface chemistry or topography to prevent infection [7] or control stem cell differentiation [8]. The release of drugs from implants to modulate their surroundings has also been investigated extensively. The first of these devices such as drug eluting vascular metal stents [9] and antibiotics releasing orthopedic implants [10] have long been on the market and have helped reduce the occurrence of in-stent restenosis and implant associated infections, respectively. Yet many implant related challenges still exist. This chapter explores how nanotechnology is, and will be, used to improve implant functionality by modulating release of drugs. It is not possible to cover all studies on device-based drug delivery in this chapter; there are tens of thousands of publications already on this topic. Instead selected studies are reported that demonstrate different delivery/release principles, different nanotechnologies and different applications. Given the sheer number of studies that already exist, important studies may have been overlooked; however, the described studies do provide sufficient data to infer important generalized conclusions about device-based drug release. The following section introduces the three types of implants and some of the challenges they encounter, ending with an introduction into the concept of biocompatibility.

13.1.1 Donor Organ Transplantation

Organ transplants have successfully prolonged the life of millions of patients, with ~112.800 solid organ transplantations in 2012 carried out worldwide in order of frequency, kidney, liver, heart, lung, pancreas and intestinal transplants [11]. The benefits of living organ transplantation include patients receive a living implant that is fully capable of carrying out the multitude of functions that organs perform and that they can interact fully with the rest of the body, for example, through hormones and nerves. However, donor organs also have two severe limitations, the availability and their immunogenicity, as well as several other issues such as the risk of transferring undetected cancer or pathogens.

There are three main reasons for the lack of donor organs. First of all, a donor is needed, and with the exception of kidneys and small tissue grafts this means a recently deceased person who has consented to donate. Donor organs also need to be immunologically matched to the recipient otherwise the organ will be rejected [12]. Finally, donor organs must also have an appropriate size to fit into the patient which is a problem when either the donor or the recipient is a child [13]. The whole process is associated with several ethical issues such as the definition of being brain dead, the presumption of consent and the allocation of the organs [14]. As long term banking of living organs is not yet possible [15], patients have to wait until a suitable donor becomes available. In the USA alone [16], over 120,000 patients were on a waiting list for an organ at the start of 2014 and some 115,580 patients died while on a waiting list between 1995 and 2013. These numbers exclude even greater numbers of patients who could have benefited from an organ transplant but did not make it onto an official waiting list.

The second major limitation is the immunosuppressive medicine that organ recipients need to take for the rest of the life. Patients need to take these medicines to prevent organ rejection as perfect immunological matching of the donor is never achieved except in the rare cases where the organ is from an identical twin. Immunosuppression, unfortunately, leads to increased susceptibility to infections, diabetes, hypertension, cancer and several other diseases [17]. While implant survival and immunosuppressive therapy has improved, there is still a need for other treatment options where immunosuppression is not necessary and where the treatment can be carried out without waiting for a donor, until recently, that meant an artificial organ.

13.1.2 Artificial Electromechanical Implants

The main advantage of artificial electromechanical organs, such as total artificial hearts and ventricular assist devices, lies with their availability [18]; they are, thus, very useful in the cases where a suitable donor organ cannot be found in time. They also behave relatively reproducibly, compared to biological organs, even if the responses of different patients to identical implants differ. Devices also exist that can assist or replace tissue functions that are not amenable to organ transplantation; such devices include cochlear implants and deep brain stimulators. However, artificial implants also suffer from several major drawbacks and the clinical outcome of modern artificial implants are only beginning to match what is achieved with donor organs despite all their drawbacks [19]. Many human functions are too complex to completely emulate with current technology, examples include the signaling in the central nervous system, liver metabolism, kidney excretion and re-adsorption. Artificial organs also cannot communicate with the rest of the body for example via hormonal exchange, and, thus, cannot adapt to changing conditions else in the body, a total artificial heart is not regulated by thyroidal hormones nor does it secrete natriuretic hormones like a living tissue would. In contrast to many

biological tissues, artificial organs cannot repair themselves when damaged which leads to mechanical fatigue, functional failure and the shedding of wear particles. The materials that artificial organs are composed can also be non-biocompatible and induce inflammatory foreign body responses, thrombus formation and/or lead to the destruction of the surrounding tissue due to mechanical strength incompatibilities. Mass produced, one size fits all, implants may inadequately fit patients and do not grow with the rest of their body; this is especially a problem for pediatric patients. Finally, electrically active implants depend on a power source as they cannot utilize the body's own energy carriers. Some of these limitations may be overcome by delivering drugs from the implant to modulate e.g. the foreign body response.

13.1.3 Tissue Engineering

While both donor organ transplantation and artificial organs continue to improve, their inherent drawbacks have led to a parallel effort being pursued where the aim is to engineer patient compatible living replacement organs, this research field is known as tissue engineering. The overall concept for tissue engineering was laid out by Langer and Vacanti in a 1993 paper [20]. The general approach consists of extracting cells from a patient, placing these cells on a scaffold material and then applying stimuli that induce the cells to form the correct tissue on this scaffold. The resulting tissue can then be re-implanted into the patient to replace a failing organ. This approach is illustrated in Fig. 13.1.

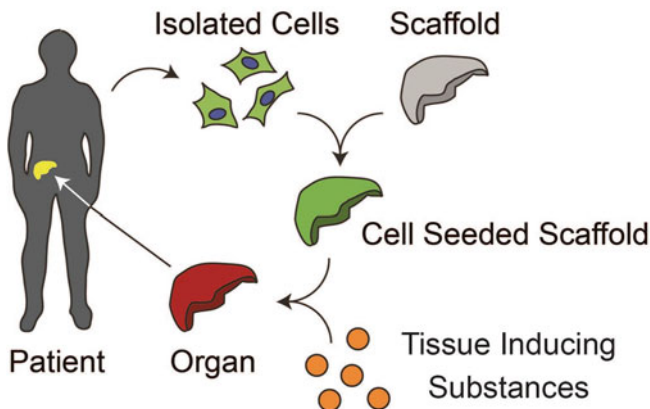


Fig. 13.1 Tissue engineering. Cells are first extracted from the patient and expanded in vitro, these are then seeded onto a scaffold where they are subjected to tissue inducing factors that may be delivered from the scaffold or an external source. When the desired tissue has formed it can be implanted into the patient

The first successful clinical transplantations have now been carried out using tissue engineered arteries in 2001 [21], heart valves in 2006 [22], bladders in 2006 [23] and trachea in 2008 [24]. Since then the long-term clinical success of tissue engineered organ transplantations has been proven [25]. The advantages of tissue engineering are plentiful. Ideally the organ becomes a living complete replacement organ fully capable of carrying out all normal functions and of communicating with the rest of the body. Since the tissue is derived from the patient's own cells no immunological rejection should take place. The organ shape and size is determined by the scaffold so they it can be made to fit the patient perfectly. No immediately deceased highly matched donor is required, and in the cases where a synthetic scaffold is used, no donor is required at all. Finally, since the graft is completely biological in origin it is ideally suited for pediatric patients [26]. The advantage in this case is that the transplant can grow with the patient obviating the need for repeated surgical interventions and reimplantations [27].

There are still many limitations to tissue engineering including cost, complexity, comparisons of different approaches and development of standardized best practices that enable studies from different groups to be compared. More animal studies are needed to determine the fate of the implants and potential problems they may cause. Clinical trials are also lacking. The lack of animal and clinical data causes concern [28] but that fact remains that some patients have been treated with tissue engineered organs with long-term success. This certainly indicates that tissue engineering will clearly be a treatment option in the future [29] even if the optimal approach has yet to be determined. However, the main tissues that have been developed so far have been relatively simple in their macro- and microstructure. To progress and develop more complex tissue engineered organs such as livers, kidneys, lungs and hearts (the organs that make up the majority of transplanted organs) there is a need to developed more complex scaffolds capable of inducing the formation of the intricate structures seen in these organs. This may potentially be achieved using scaffolds capable of advanced spatial and temporal control over drug release.

13.1.4 The Convergence of Artificial and Biological Implants

While tissue engineering offers limitless perfectly matched replacement organs that may be used to restore normal bodily function the possibility of merging artificial active implants with tissue engineered implants has been investigated. Mannoor et al., for example, additively manufactured a bionic ear composed of a chondrocyte filled alginate hydrogel and an inductive coil antenna made from conductive silver nanoparticles [30]. Such cyborg organs composed of biological and electronic components may provide capabilities far beyond what human tissues are normally capable of, and they could be the ultimate future implants. But even these implants will face problems that necessitate drug release to control infection, inflammation and differentiation.

13.1.5 Biocompatibility

Biocompatibility is one of the key issues in implant medicine [31]. Traditionally biocompatibility was seen as the absence of a host response as this led to the greatest success for implants. With the rise of tissue engineering where the ultimate goal is for the implant to merge completely with the body, this is no longer the case. Therefore, biocompatibility has been redefined as: “biocompatibility refers to the ability of a material to perform with an appropriate host response” [32]. Biocompatibility is clearly a complex and application dependent matter, but major determinants of biocompatibility and ultimately of implant success are implant associated inflammation and infection, as well as the interaction with the surrounding tissue. These are biological responses and, thus, amenable to pharmaceutical modulation.

Releasing drugs from implants has many advantages over systemic intervention such as greater local effect, lower side-effects at distant sites and smaller amounts of expensive drugs. The efficiency of drugs released from implants relies greatly on how they are interfaced with the implant in the first place. Drugs may, for example, be adsorbed or encapsulated into the implant, leading to fast and slow release, respectively. Poor drug stability and/or weak drug-implant interactions may prevent inclusion of drugs and in these cases, the drugs may be embedded in polymeric layers that are coated onto implants or in drug delivery particles that can be adsorbed onto or encapsulated into the implant surface. These then protect the drugs and determine how fast they are released. Nanotechnology is used in various ways to improve the functionality of these drug release system. Nanoparticles may, for example, be used to ensure maximum dispersion of the drug in the implant leading to a constant release; they may through their large surface area promote greater interaction with the implant matrix and they may facilitate delivery of bio-molecular drugs such as DNA and RNA to the intracellular sites of action. The following section describes how drug release strategies have been applied to various implants to improve their functionality in specific applications.

13.2 Implant Drug Release and Applications

13.2.1 Nanotechnology, Drug Release and Implant Associated Infections

It has long been known that implants provide extremely good breeding grounds for fungal and bacterial pathogens and that they may increase the virulence of bacteria more than 10^5 -fold [33]. Implant associated infections are still a major problem [34]. Infections have traditionally been prevented by maintaining standard anti-septic operating conditions, thoroughly sterilizing implants and by

giving patients antibiotics systemically after the surgery. However, even when sterile surgical conditions and implants are employed, a low number of pathogens may still find their way into the body. Furthermore, post-implantation, bacteria may arrive via the bloodstream and colonize otherwise sterile implants [35]. Interestingly, the implant material seems to have little effect on the infection rate, one study found that different metals gave roughly the same infection rate in vivo [36] and another that biological donor materials behaved like mechanical implants with regards to infection rates [37]. This is thought to be a consequence of the rapid formation of a serum protein film on the surface of all implant materials that subsequently masks the underlying material and provide good adhesion for microbial organisms [38]. When implant associated infection take place, the bacteria start to form biofilms on the implant surface [39]. These provide protection against the systemic treatment with antibiotics which cannot penetrate into the biofilm; they also deactivate the host's immune response by preventing the attachment of IgG and C3a [40]. As a result the implant usually has to be removed. To address this problem a wide range of new approaches have been developed [41, 42] these include changing the surface nano- and microstructure and its chemistry [7] as well as developing entirely new polymers [43]. However, immobilization [44] and/or release [45] of antimicrobial compounds such as organic molecules and silver are also being investigated often using nanotechnology to improve functionality. The progression of biofilm formation and treatment is illustrated on Fig. 13.2.

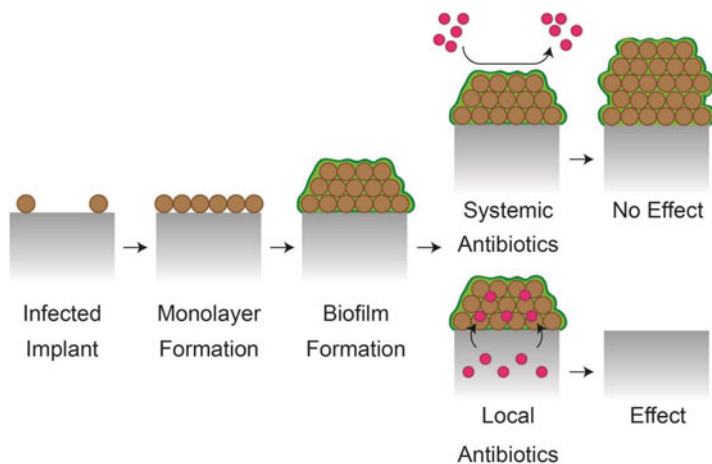


Fig. 13.2 Implant associated infections. Initially bacteria, implanted together with the implant either as free planktonic bacteria or already be attached to the implant, colonize the surface inside the body as a monolayer and then start to form a 3D multilayer biofilm. Systemic antibiotics cannot penetrate this biofilm but antibiotics released from the implant can due to their local high concentration

13.2.1.1 Controlled Release of Antibiotics from Implants

Chemical antibiotics have been in clinical use since the discovery of sulfonamides in the beginning of the twentieth century [46]. As antibiotics perform poorly when given systemically against implant associated infection due to biofilm formation, the local release of antibiotics from implants has been explored extensively and several products have reached the market [10]. The idea is that by delivering the drug beneath the biofilm it can locally accumulate to a much higher concentration and better penetrate the film. Liu et al. developed ethylcellulose particles encapsulating the antibiotic ceftazidime, which were encapsulated into a hydroxyapatite/polyurethane scaffold for bone tissue engineering [47]. The resulting scaffold released ceftazidime and could inhibit grow of *S. aureus*, additionally, the use of adsorbed ceftazidime encapsulating particles provided a longer release and bacterial inhibition than the simple inclusion of a “naked” ceftazidime in the scaffolds. Ma et al. demonstrated that nanosized cationic liposomes containing the antibiotic vancomycin could be incorporated into porous nano-hydroxyapatite/chitosan/konjac glucomannan tissue engineering scaffolds [48] which could then release liposomal vancomycin which destroyed *S. aureus*, the release rate could be tailored by the amount of konjac glucomannan that was incorporated into the scaffold. Similar to the study by Liu et al., this study also compared the release and antimicrobial activity of “naked” antibiotic with the formulated antibiotic and also found that encapsulating a formulated antibiotic led to prolonged release and anti-bacterial effect. Feng et al. showed that the antibiotic doxycycline could be incorporated into poly (lactic-co-glycolic) acid (PLGA) nanospheres than could then be placed on a porous nanofiber polylactic acid (PLA) scaffold [49]. This paper also found that doxycycline nanospheres provided a longer release and anti-bacterial activity than “naked” adsorbed doxycycline, with high molecular weight and lactic acid content in the PLGA indicative of longer release. Hong et al. developed an electrospun scaffold composed of polyurethane and PLGA fibers, where the PLGA fibers contained the antibiotic tetracycline which was then released destroying the *E. coli* [50]. Kim et al. demonstrated that the antibiotic cefoxitin could be incorporated into electrospun PLGA fibers where from it could be released and destroy *S. aureus*, introducing a polyethylene glycol (PEG)-PLA copolymer enabled better drug retention and prolonged the release [51]. Teo et al. blended the antibiotic gentamicin with polycaprolactone (PCL) and tricalcium phosphate (TCP) and deposited the blending into a wound healing mesh using fused deposition modelling, the mesh then released gentamicin and inhibited bacterial growth in vitro and in vivo where it promoted faster wound healing [52]. As an alternative to incorporating the antibiotic into the implant material, Li et al. demonstrated that the antibiotic cefazolin could be coated onto orthopedic implants using a layer-by-layer (LbL) technique; the resulting surfaces released the antibiotic and killed *S. aureus* [53]. These concepts are not limited to anti-bacterial compounds, Verreck et al. showed that the anti-fungal compound itraconazole could be encapsulated and released from electrospun polyurethane fibers [54].

These studies clearly indicate that by formulating antibiotics using nanotechnology, their activity, local retention and duration of effect can be improved. A major problem with using antibiotics to combat implant associated infection is, however, resistance to these drugs in bacterial communities including those found in implant associated infections [55–57]. This development has prompted a search for alternative means to control infection including implants that release non-organic compounds such as nitric oxide [58] and silver [59], amongst these approaches, silver has received by far the greatest interest.

13.2.1.2 Controlled Release of Silver from Implants

Silver has long been used as a bactericidal implant coating [60], and several commercial anti-infection products based on silver are on the market including Acticoat and Aquacel AG by Smith & Nephews and ConvaTec, respectively. Their effect is thought to be caused solely by the silver ion Ag^+ which silver surfaces release when oxidized [61]. Due to their large surface area which facilitates Ag^+ release and the ease of incorporating them into materials, silver nanoparticles are now used extensively to coat implants [62], where they seem to give better results than traditional silver formulations [63]. Those tested clinically include wound dressings [64], burn dressings [65], urinary catheters [66], ventricular drain catheters [67] and central venous catheters [68, 69] although the clinical effect varies. However, great caution must be exercised when using any drug for combating infections as they may have adverse effects on neighboring cells. Silver nanoparticles are known to be cytotoxic to a wide variety of cells [70] including stem cells commonly used for tissue engineering [71] and have been shown to be cytotoxic to osteoblasts and osteoclasts at concentrations lower than those needed for their antibacterial effect [72]. The use of nanotechnology such as silver nanoparticles to overcome one problem such as infection may, therefore, cause another problem such as local tissue damage. It is realistic; however, that further nanotechnological progress will enable us to use such potentially toxic particles in a controlled manner so that they do not cause harm, the key is to keep the local concentration of silver within the therapeutic window by controlling its release rate and local retention that can be achieved with enabling nanotechnologies.

This can be done by encapsulating the silver into a polymer or by attaching it onto the surface of the implant. Taglietti et al. used (3-aminopropyl)triethoxysilane to bind silver nanoparticles onto the surface of glass substrate where they inhibited microbial biofilm formation [73]. Gordon et al. developed a nanostructured coordination network composed of silver coordinating polymers and silver nanoparticles that could be coated onto titanium cages [74]. These cages effectively killed *S. epidermis* while the only damage to the mouse host was transient and minor damage to leukocytes. Wang et al. encapsulated silver nanoparticles into a PLGA electrospon coating that could be spun onto, and around, implants and showed that this coating was capable of killing *S. aureus* and *E. coli* while releasing concentrations far below those that caused acute toxicity [75]. Mohiti-Asli et al. showed that an

electrospun nanofibrous PLA scaffold could be coated with proprietary formulation of AgNO_3 and polymer binder where after the scaffold became bactericidal [76]. Sheikh et al. showed that silk nanofibers containing hydroxyapatite and silver nanoparticles could be electrospun into a mat that inhibited *S. aureus* and *E. coli* but also induced cytotoxicity in fibroblasts [77]. Samberg et al. developed an electrospun PLCL skin tissue engineering scaffold that incorporated silver nanoparticles and showed that it could both support a confluent layer of fibroblasts as well as kill *S. aureus* and *E. coli* [78]. The use of silver is often promoted as a solution to drug resistance and does indeed kill bacteria that are resistant to antibiotics as shown by Cheng et al. who formed silver nanoparticles inside titania rods that had been created on the surface of titanium implants by anodization, these silver nanoparticles could subsequently kill multi resistant *S. aureus* (MRSA) [79]. However, it is important to note that microorganisms may also evolve genes that confer resistance to silver and the increasing use of silver in infection control may, therefore, trigger the evolution and spread of these genes rendering the silver approach no better than organic antibiotics [80].

13.2.2 Nanotechnology, Drug Release and Implant Associated Inflammation

13.2.2.1 The Foreign Body Response

When any implant is placed within the body a sequence of inflammatory events occurs that either resolves itself or leads to a foreign body response that causes implant failure [81]. During surgery, skin, connective tissue, blood vessels and possibly other tissue types are damaged and blood accumulates around the implant. The inflammation sequence starts when blood proteins adhere to the surface of the implant forming a provisional matrix composed of coagulated fibrin which encapsulates a range of blood borne bioactive proteins including mitogens, cytokines, complement factors, growth factors and chemokines. This matrix then acts as both a controlled release device shedding these bioactive proteins and as a cellular attachment scaffold. Neutrophils are among the first cells that attach to this matrix and they trigger the innate immune response. Mast cells are recruited and degranulate releasing histamine which, among other things, attracts macrophages. This acute inflammatory phase then either resolves itself typically within a week following normal wound healing or it progresses towards chronic inflammation. Chronic inflammation is characterized by the continual presence of lymphocytes and macrophages on the implant surface. If this persists, a granulation tissue is formed around the adhering immune cells by infiltrating fibroblasts. This granulation tissue may eventually progress and become a fibrous capsule that encapsulates and isolates the implant from the body causing the implant to fail. At the same time macrophages on the implant surface may fuse and become foreign body giant cells that form a large isolated pocket between themselves and the implant surface into which they secrete

protons, reactive oxygen species and degradative enzymes which destroy the implant. These end results are known as the foreign body response. The direction that the inflammatory response takes (normal healing versus foreign body response) depends on the amount and type of proteins adsorbed onto the surface of the implant which depends on the surface chemistry and topography. A simplified overview of the foreign body response is shown in Fig. 13.3.

13.2.2.2 Wear Debris

On a longer term some implants shed nano- and micrometer sized debris particles [82]. This is the case when non-biodegradable metals, ceramics and polymers break down due to mechanical fatigue in areas where they are subjected to movement or loading e.g. in joint replacements. Wear particles then disperse into the surrounding tissue and can even travel to distant sites in the body. This is a major problem as

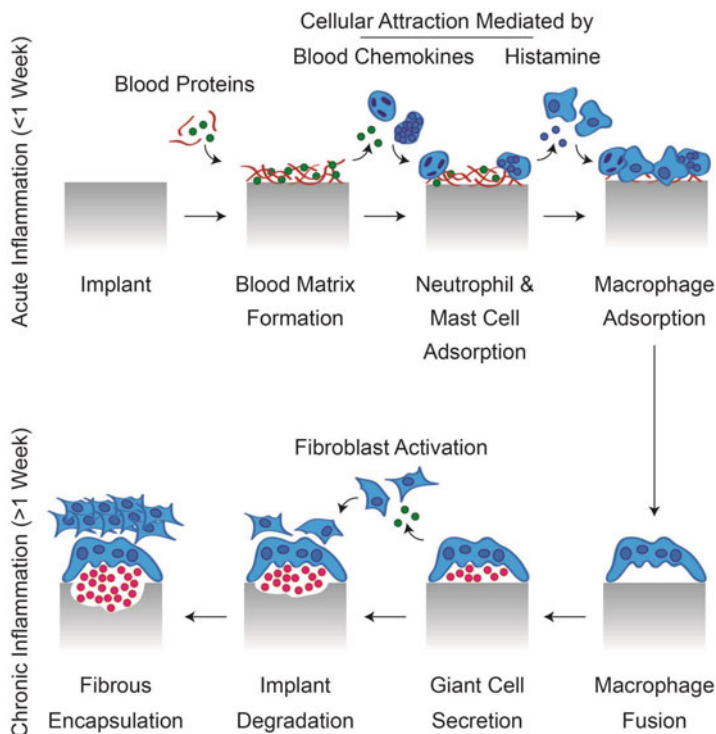


Fig. 13.3 The foreign body response. When an implant is inserted into the body a blood derived matrix forms attracting neutrophils, masts cells which then recruit macrophages. This acute inflammatory phase then either resolves itself, typically within 1 week, or it becomes chronic when macrophages start to fuse and degrade the implant while the implant is, in parallel, isolated by the formation of a fibrous capsule

macrophages engage these non-degradable particles. In the case of microparticles that are too big to be internalized, a frustrated phagocytosis response is triggered leading to giant cell formation and the production of reactive oxygen species. Some non-degradable wear debris nanoparticles are readily internalized by macrophages but since they cannot be broken down in the endo- and lysosomes, they end up destabilizing these [83]. Nanoparticles released by the wear of non-degradable polymer implants, such as those made of polyethylene, may also trigger inflammation through the TLR 1/2 receptor. In these cases the end result is the activation of the NALP3 inflammasome and the release of the pro-inflammatory cytokines IL-1, IL-6, IL-10, IL-12, TNF- α and IFN- γ . Additional pathways such as metal allergies may also contribute to the inflammatory response. The result is continual myelomonocyte infiltration and generation of additional active macrophages [84]. Eventually this cellular response causes peri-implant osteolysis where the bone tissue surrounding the implant is broken down due to three reasons: (1) Formation of bone-degrading osteoclasts triggered by inflammatory cytokines, (2) wear particle-induced apoptosis of bone-building osteoblasts and (3) degradation of the bone tissue which is caused by acidification of the tissue by bursting endosomes and the release of degradative enzymes such as collagenases and matrix metalloproteinases by the activated immune cells and osteoclasts. The end result is the degradation of the surrounding tissue and the loosening and eventual implant failure necessitating major surgical intervention.

13.2.2.3 Drug Release and Inflammation Control

Implant associated inflammation can be controlled by releasing drugs from the implants. However, there are two major problems with using drug release from to modulate inflammation. The first problem is that a certain degree of inflammatory response is absolutely critical to the success of most implants. The immune system controls implant associated infection, the immune system is involved in healing surgical tissue damage [85] and in the case of biodegradable implants, it is involved in degrading and clearing the implant and remodeling the resulting tissue to work together with the surrounding tissue. The second problem is that the inflammatory response may last as long as the material persists in the body. Non-biodegradable metal or polymer implants may, for example, trigger immune responses many years after their implantation by shedding wear debris that activates surrounding macrophages. Retaining a release of functional anti-inflammatory drugs for such a prolonged period is difficult. Despite these concerns, drug release has been studied to counteract implant associated inflammation in many different studies. These are increasingly making use of various nanotechnologies to improve functionality, for example, by achieving greater control over drug release.

To reduce early inflammation anti-inflammatory drugs may be released. Jayant et al. demonstrated that alginate microparticles containing and releasing the anti-inflammatory drugs dexamethasone or diclofenac could reduce the inflammatory response to a glucose sensor [86], in this case an LbL nanofilm was deposited on the

surface of the particles to control drug release. Vacanti et al. incorporated dexamethasone into electrospun PCL or PLA nanofibers and demonstrated that drug loaded PLA nanofibers could inhibit inflammation and fibrous capsule formation *in vivo* [87]. Interestingly, dexamethasone release differed greatly between the two polymers, it was burst released from PCL but not PLA fibers and only drug loaded PLA fibers were capable of reducing the foreign body response.

Another approach is to target the late stages of inflammation such as the fibrous capsule formation, this leaves the acute response unaffected and able to clear infections and damaged tissue. Rujitanaroj et al. have developed a biodegradable electrospun polymer nanofiber scaffold that encapsulated different siRNA nanoparticle formulations targeted against Collagen 1 A1, a major component in the fibrous capsule [88]. These scaffolds were capable of silencing Collagen 1 A1 in cells and when placed *in vivo* fibrous capsule formation was inhibited. Additionally, release rates and duration of silencing could be altered by choosing different nanoparticle formulations. Takahashi et al. used a similar strategy, encapsulating in PEG based hydrogels, polyethylenimine formulated siRNA nanoparticles targeted towards mTOR, a gene involved in fibroblast proliferation and collagen production [89]. While they observed effective results *in vitro* no effect was observed *in vivo*.

Implant associated pain is another area where drug release may be used to benefit a patient. The implantation of an implant is a serious intervention which is often associated with great pain for the patient. Weldon et al. has, therefore, developed an electrospun PLGA fiber suture which incorporated bupivacaine, a local anesthetic which caused anesthesia around the implant in an animal model [90].

13.2.2.4 Inflammation and Neuroprosthetics

Tissue inflammation is also a major problem for neural implants [91] which typically consist of metal electrodes such as those used for deep brain stimulation. In the CNS, inflammation is mediated by reactive astrocytes and activated microglial cells and results in glial scar formation which removes the nerves from the implanted electrodes which cease to function as the impedance rises. Device based drug release has been pursued to inhibit the inflammatory process in some cases using nanotechnology [92]. Mercanzini et al. developed dexamethasone encapsulating polypropylene sulfide nanoparticles that could be coated onto electrode neuroprosthetics using a dissolvable polyethylene oxide (PEO) based coating [93]. When implanted in the motor cortex of rats the nanoparticles were released from the electrodes but stayed in the vicinity of the implant which resulted in lower inflammation and impedance. Similarly, Kim and Martin developed dexamethasone encapsulating PLGA nanoparticles which were coated onto neuroprosthetic implants using alginate hydrogels, these coatings also reduced inflammation and impedance when the implants were implanted in the auditory cortex of guinea pigs [94]. Interestingly, this study also showed that the release rate was much lower when the particles were embedded in an alginate hydrogel than when they were in a PBS solution.

13.2.2.5 Immunostimulatory Nanocoats

Interestingly, the opposite approach, local immunostimulation, has also been investigated with the aim of boosting inflammation to reduce implant associated infections [95]. Li et al. developed an immunostimulatory nanolayer that was deposited onto surgical steel K-wires by electrostatic LbL deposition of IL-12, BSA and PLL. When implanted the deposition of IL-12 led to a marked reduction in infection rates [96]. The concept has later been expanded by including chemokines such as MCP-1 in the coating to attract immune cells to the implant, this also led to a reduction in implant associated infection [97]. While this strategy may prove effective against bacteria that are resistant to antibiotics and silver, its use must be carefully balanced against increases in inflammation and the foreign body response.

13.2.3 *Nanotechnology, Drug Release and Stem Cell Differentiation*

Promoting tissue development is important in tissue engineering where a new tissue has to be created de novo within the implant. But it is also relevant for non-biodegradable artificial implants such as orthopedic and dental implants where a strong fixation in the bone is desired and bone development on the implant surface is, therefore, required, known as osseointegration. During embryonic development as well as in adult tissue repair, tissue development typically takes place when stem cells proliferate to generate more specialized cell types by asymmetric cell division [98]. These specialized cell types then undergo a succession of differentiation steps which increasingly specialize the cell. Embryonic stem cells, for example, differentiate into endodermal, mesodermal and ectodermal stem cells during embryonic gastrulation, these then specialize further to produce the tissues our bodies contain. Likewise, mesenchymal stem cells can undergo differentiation into structural tissues such as bone, fat, cartilage and muscle in a step wise manner by first producing precursor cells typically suffixed “blasts” which can then form the final cell types typically suffixed “cytes” as in osteoblasts and osteocytes, respectively. Often additional steps are present, commonly indicated by prefix “pre” as in a pre-osteoblast. During each of these differentiation steps the cell types perform different functions, pre-osteoblasts, for example, lay down the protein components of bone matrix, osteoblasts mineralize the matrix and osteocytes maintain the developed bone. Each differentiation step is typically started by exogenous factors such as the extracellular matrix or endocrine or paracrine signaling through hormones (often steroids and vitamins) and protein growth factors. These factors then act on transcription factors and microRNA which combine to down regulate the set of genes that define the current stage of differentiation as well as upregulate the expression of the set of genes that are required for the next stage of development.

13.2.3.1 Drug Release and Stem Cell Differentiation

Different strategies exist to control the development of tissue on the surface of implants including changing the surface chemistry and topography [99]. However, since the body itself to a large extent controls stem cell differentiation through extra- and intracellular biomolecules it is only natural that much focus has been placed on delivering these biomolecules as well as synthetic molecules from implants to promote tissue development. Molecules delivered from implants to influence stem cell differentiation include proteins [100], vitamins [101], plasmid DNA [102] as well as various RNA molecules such as microRNA [103] and siRNA [104]. These molecules have been incorporated into scaffolds using a wide variety of methods. Wadagaki et al., for example, demonstrated that the synthetic molecule simvastatin could be incorporated into electrospun nanofibers that were deposited into a scaffold and that the subsequent release of simvastatin enhanced the formation of bone *in vivo* [105]. Shah et al. used a chitosan/poly aspartic acid LbL nanofilm system to load the osteoinductive bone morphogenetic protein 2 (BMP2) and hydroxyapatite onto durable implants made of PEEK or titanium [106]. The inclusion of a hydrolytically cleavable molecule in the films enabled tuning the BMP2 release and the coatings promoted bone regeneration *in vitro* and *in vivo*. Nie and Wang formed chitosan nanoparticles containing plasmid DNA encoding BMP2, these were then either encapsulated or adsorbed into PLGA scaffolds, and a sample with adsorbed “naked” plasmid was also prepared [107]. When placed in phosphate buffered saline the “naked” DNA was released fastest and the encapsulated plasmid nanoparticles slowest, the encapsulation of the plasmid led to the best combination of transgene expression and cell viability.

13.2.3.2 Drug Release and Other Tissue Engineering Applications

Adequate cell seeding on a tissue engineering scaffold is of great importance for the formation of functional tissues once the scaffolds has broken down. While most tissue engineering strategies rely on cell seeding prior to implantation in order to achieve sufficient cell densities, other approaches utilize implant mediated release of chemoattractive molecules to stimulate the mobilization and invasion of desired cell types. Zhao et al. developed a hydrogel that released the chemoattractive protein hepatocyte growth factor (HGF) [108]. This hydrogel was then capable of attracting mesenchymal stem cells in an *in vitro* transwell invasion assay whereas non-loaded hydrogels were not. Similarly, Li et al., demonstrated that a HGF loaded hydrogel could attract neural stem cells in a transwell assay [109].

Diffusion of oxygen and nutrients from the bloodstream into tissues and is another area of critical importance to tissue function and where insufficient transport takes place necrosis occurs [110]. Diffusion is distance limited and can typically only supply cells that are within 100 μm of a capillary. Neovascularization is promoted automatically when cells within a scaffold suffers from a lack of nutrient or oxygen [111], but may take place too slowly to prevent necrosis in large implants. Releasing drugs

that promote vascularization from scaffolds has, therefore, received much attention in tissue engineering. Nelson et al. formed polymeric nanoparticles containing siRNA against prolyl hydroxylase domain 2 (PHD2), an anti-angiogenic factor, and incorporated these into a biodegradable polymeric foam scaffold by mixing the nanoparticles with the scaffold foam prior to foaming [112]. These scaffolds were then capable of silencing PHD2 and promoting the ingrowth of more and denser vessels.

13.2.4 Other Applications of Implant Drug Release

Besides controlling implant associated infection, inflammation and tissue engineering, implant drug release is also being employed for other applications such as combating residual cancer cells around implants. When cancer tumors are removed a surgeon often has to balance conservative resection that retains tissue function but risk incomplete removal of all cancer cells against aggressive resection that more likely removes all cancer cells but which may destroy tissue function. Implants are often placed where tumors are removed to fill the void, therefore, the possibility of release drugs to kill residual cancer cells have been explored extensively [113]. Wie and Wang developed an electrospun PLGA fiber scaffold that encapsulated the chemotherapeutic paclitaxel and showed that its release rate could be tailored by controlling the fiber diameter with nanofibers releasing faster than microfibers [114], released paclitaxel was capable of killing glioma cells. Yohe et al. showed that the anti-cancer drugs CPT-11 and SN-38, camptothecin analogues, could be incorporated into an electrospun PCL/ PGC-C18 meshes designed to bridge colorectal anastomosis, these meshes could the kill colorectal cancer cells [115]. Intriguingly this study uses included air as a means to control drug release.

13.3 Nanotechnology Strategies in the Development of Drug Releasing Implants

Drugs can be interfaced with implants using a variety of means such as encapsulation and adsorption which affects amongst other things, drug accessibility and release rate. These parameters can be additionally tuned by formulating the drugs in nanoparticles. Figure 13.4 shows some simple modes for drug/implant interfacing.

13.3.1 Drug Adsorption

The simplest method for incorporating drugs is to adsorb them onto the surface of the formed implant by drying or lyophilizing a solution of the drug onto the surface. In this case the release of the drug from the surface depends on the binding strength

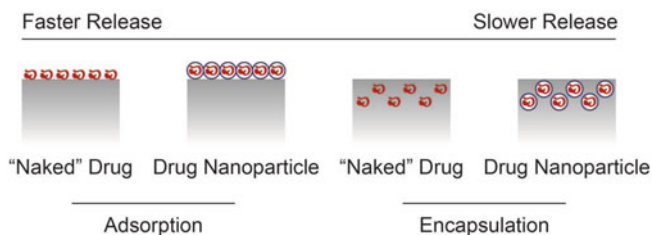


Fig. 13.4 Drugs can be adsorbed onto the surface of a polymeric implant after that surface has been formed (*far left*). If the drug is incorporated into a drug delivery vehicle before adsorption and that vehicle promote adhesion the release rate may be lowered (*center left*). If the drug is added to the polymer melt or solution before forming the implant surface the drug may be encapsulated which delays its release (*center right*). By incorporating the drug into a drug delivery system before its encapsulation the release may be further slowed if the system limits drug diffusion within the polymeric matrix (*far right*)

between the surface and the drug relative to the flow and content of the medium surrounding the implant. If this medium contains molecules that bind to the surface these may displace the drug. If slower release is desired, the drug may be incorporated into a drug delivery system that exhibits greater binding to the implant surface. A drug can also be covalently tethered to the surface, either permanently if the drug is only to affect adhering cells or temporarily if the drug is to be released slowly to the implant surroundings. To increase surface loading LbL depositions may be used, in this case several alternately charged polymers are deposited as layers onto the implant surface, this provides a matrix wherein drugs may be loaded. Alternatively the implant surface can be modified so it provides a larger surface area by which the drug may bind stronger and to which more drug may be bound. Additionally surface cavities or tubes may be used to protect adsorbed drugs from the flow of the surrounding medium slowing their release. Çalışkan et al. recently showed that nanotubes could be formed on titanium surfaces using anodic oxidation and that these improved the loading and delayed the release of gentamicin; these surfaces, but not plain titanium, could kill *S. aureus* [116]. It is worth noting that adsorbed drugs are exposed to the surrounding aqueous medium and that many labile biological drugs may rapidly degrade in this release mode if unprotected. The typical adsorption modes are illustrated in Fig. 13.5.

13.3.2 Drug Encapsulation

If a slower release or greater drug protection is desired, the drug may be incorporated into the implant surface or the entire implant. This can be achieved by dissolving the drug in the polymer melt or solution prior to forming the implant, the drug then becomes encapsulated as the polymer matrix solidifies. Using this

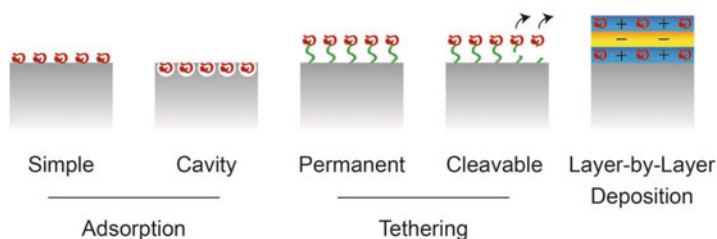


Fig. 13.5 Adsorption modes. There exist several different methods for adsorbing drugs onto an implant surface. Drugs may simply be adsorbed onto a flat surface but will in this case be susceptible to fast release. To slower the release rate, the drugs may be placed in cavities that reduce fluid motion and increase the surface area with which the drug may bind. Alternatively the drug may be covalently attached to the surface either permanently or by a linker that degrades slowly or fast in response to stimuli. Finally, drugs may be adsorbed as part of a layer-by-layer coating

method requires that the drug can tolerate the solvent or temperature used in the implant formation process which may not be the case for labile biological drugs, in this case a solvent/water emulsion may be used so the drugs can be kept in a aqueous phase prior to dehydration. Several studies show that compared to absorption, encapsulated drugs are released slower. The release rate depends on the drug diffusion rate through the polymer matrix and in the case of a biodegradable matrix, on the degraded volume as a function of time. Therefore, the desired release rate may be achieved by embedding the drug in a matrix with a tailored thickness or degradation rate. Alternatively the drug may be encapsulated into particles that are in turn encapsulated into the implant matrix, these particles may then retain the drug within the matrix and can also function as delivery vectors for drugs such as nucleic acids that need to reach the intracellular space of their target cells.

13.3.2.1 Drug Encapsulation by Co-Axial Electrospinning

Drugs can be incorporated onto or into electrospun polymeric implant the same ways as a normal polymeric implant by adsorption or encapsulation. Electrospinning, however, offers an additional mode of drug incorporation known as co-axial electrospinning, this method deposits a core-shell fiber containing an internal core phase and an outer shell phase which may be chemically different [117]. This allows the deposition of functional drugs into a drug compatible core phase even when harsh solvents are employed for the outer shell phase. In one example, Korehei and Kadla used co-axial electrospinning to deposit a T4 bacteriophage into hollow poly(ethylene oxide)/cellulose diacetate fibers and demonstrated that these fibers could subsequently release bacteriophages capable of killing *E. coli* in vitro [118]. The different delivery modes in electrospinning can be seen in Fig. 13.6.

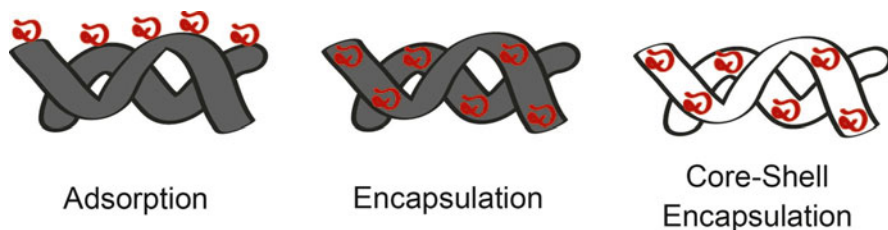


Fig. 13.6 The drug incorporation modes commonly applied in electrospun implants. Adsorption provides faster release than encapsulation. Core-shell encapsulation may be used if the polymer processing solvent is incompatible with the drug to be encapsulated

13.3.2.2 Drug Release from Metal and Ceramic Implants

Metal and ceramic implants can be coated with drugs by adsorption the same way as with polymeric implants, indeed dip coating metal prosthesis in solutions of antibiotics prior to implantation is a simple solution that has been employed [119]. Unfortunately, such coated drugs are released fast. Drug encapsulation as a means to control the release is not as easy as with polymeric implants as most drugs will not survive forging or sintering temperatures or pressures. In these cases other approaches have to be tried. A solution is to coat the metal or ceramic surface with a polymeric or hydrogel layer encapsulating the drug for example using the LbL method [53]. Achieving a uniform layer using this method is, however, not simple on three dimensional implants. A different solution is to electrospin polymeric fibers around an implant, as was demonstrated by Wang et al. These three methods of drug functionalizing metal and ceramic implants are shown on Fig. 13.7.

13.3.3 Co-Release of Multiple Drugs

In many cases it would be beneficial to release more than one drug from an implant. In some cases it could be that different drugs accomplish different functions such as combating infection while promoting stem cell differentiation [120], whereas in other cases, drugs act synergistic to induce a desired phenomenon. Each step in stem cell differentiation, for example, is typically promoted by more than one growth factor and co-delivering several growth factors is, thus, necessary to recapitulate natural development completely [121].

We have shown one application where the anti-cancer drug doxorubicin was co-released with chitosan/siRNA nanoparticles from tissue engineering scaffolds to adjacent cancer cells wherein gene silencing and cell death was induced in lung cancer cells with the aim of their susceptibility to the chemotherapeutic [122]. Zheng et al. demonstrated that antimicrobial silver nanoparticles and the osteogenic growth factor BMP-2 could be co-released from a PLGA scaffold killing *S. aureus*

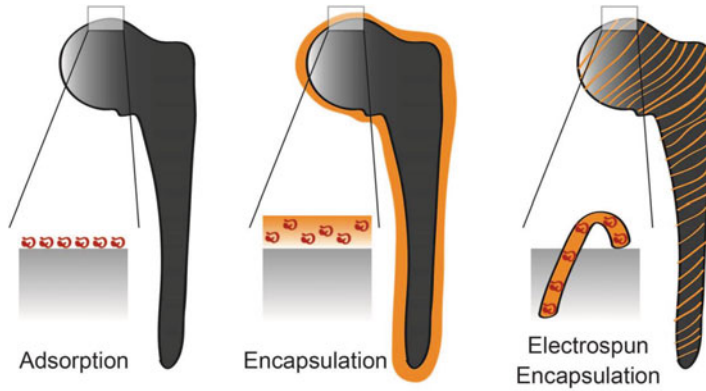


Fig. 13.7 Strategies for functionalizing metal and ceramic implants with drugs, illustrated with a hip prosthesis. The drug may either be adsorbed onto the implant surface (*left*) or be encapsulated in a polymeric layer that is coated onto the implant (*center*). A novel way of drug functionalizing metal and ceramic implants may be to wrap it in electrospun fibers that encapsulated drugs (*right*)

while promoting bone healing [123]. Similarly, Stevanovic et al. showed that silver nanoparticles and the anti-oxidant/osteogenic factor ascorbic acid could be co-incorporated into PLGA nanoparticles which could then kill *E. coli* and MRSA as well as promote bone formation [124], although in this case incorporation into medical devices was only suggested not tried. He et al. showed that polyethylenimine nanopolyplexes containing plasmids encoding vascular endothelial growth factor (VEGF) and fibroblast growth factor (FGF) could be incorporated into fibers scaffolds made of PEG-PLA [125]. When implanted into mice these plasmids synergized to promote ingrowth of denser mature vessels than either plasmid alone. Interestingly, the authors compared polyplex encapsulation with adsorption and showed that encapsulation led to greater expression of the encoded proteins as well as greater vessel density than adsorption.

13.3.4 Temporal Controlled Drug Release

Many implant related phenomenon are time dependent. Implant associated infections and acute inflammation take place immediately upon implantation, whereas, fibrous encapsulation occurs weeks after implantation and wear debris induced inflammation may not occur until years following implantation. The same is true in stem cell differentiation in tissue engineering where each differentiation step occurs in successions. Controlling drug release temporally is, thus, of great importance and can be achieved by choosing an appropriate delivery strategy that has suitable release kinetics for the biological function that needs to be modulated.

Drug adsorption, for example, leads to rapid release and may be suitable for combating infections or acute inflammation early on, whereas drug encapsulation leads to a slower release and may be more suited for modulating chronic inflammation or stem cell differentiation. By combining various delivery methods it is even possible to deliver different drugs from the same implant at different times. Clark et al. incorporated anti-inflammatory ketoprofen into a poly (β -amino ester) based hydrogel and BMP2 into PLGA microspheres, these two components were then sintered together to provide a composite scaffold [126]. When hydrated, the ketoprofen was released rapidly, whereas, the BMP2 was released slowly. Min et al. used an LbL technique to place BMP2 beneath a layer of the gentamicin on an implant [127]. Gentamicin was released fastest and was capable of killing *S. aureus*, subsequently BMP2 was released and was capable of promoting osteogenesis in pre-osteoblasts. Interestingly, the addition of clay layers in the layer by layer film was able to slow the release. In a different study by the same group, BMP2 was deposited underneath the angiogenic growth factor VEGF and demonstrated that VEGF was released first, followed by BMP2, and that their co-release led to denser bone formation than delivering BMP2 alone [128]. This was speculated to be due to the inclusion of fast releasing VEGF mediated the formation of an early vascular network through which cells penetrated the implant before dense bone formation was induced by slower releasing BMP2. Other strategies may also be used to control temporally controlled delivery of multiple drugs. Basmanav et al. incorporated BMP2 and BMP7 (BMP7 is another later acting osteoinductive growth factor) into crosslinked alginate microspheres with different alginate content and showed that the alginate determined the release rate [129]. When fast releasing BMP2 microspheres and slow releasing BMP7 microspheres were incorporated into a PLGA scaffold subsequently seeded with BMSCs bone formation was promoted. In a series of studies by the same group [130–132], Yilgor et al. incorporated BMP2 and BMP7 into nanocapsules composed of fast degrading PLGA or slow degrading PHBV, these particles were then either adsorbed onto or encapsulated into chitosan scaffolds where encapsulation led to a slower release. The authors then used these delivery modes to deliver BMP2 and BMP7 sequentially which led to greater bone formation than delivering them simultaneous or singularly.

13.3.5 Spatial Restricted Drug Release

Spatial restriction by nanofunctionalization is useful in the cases where a differential response is desired in different parts of an implant. This is relevant, for example, when complex tissue composed of multiple cell types has to be grown. Traditionally, such tissues have been grown by seeding different cell types onto different parts of an implant [133, 134] or by using different materials for each phase [135–138]. Sheehy et al., for example, created an osteochondral tissue by

encapsulating chondrocytes and mesenchymal stem cells in separate regions of an alginate hydrogel implant whereas Filardo et al. repaired the knees in patients using a cell-free osteochondral implant consisting of collagen I layers with or with hydroxyapatite. Another strategy is to load each phase with different drugs typically loaded in drug vehicles, these drugs can then be released in a spatially restricted manner and induce the desired local effects [139, 140]. Dormer et al., for example, generated an osteochondral tissue using a bi-phasic scaffold composed of TGF- β and BMP2 loaded regions. Mohan et al. generated an osteochondral tissue by combining a hydroxyapatite gradient and a BMP2/TGF- β particle gradient in a biphasic scaffold [141]. Using a similar strategy [142], we have previously grown a bone/fat tissue using siRNA nanoparticles. In this case we formed a macroporous PCL scaffold with additional nanopores which we filled with nanoparticles containing siRNAs against either TRIB2, an inhibitor of adipogenesis, or BCL2L2, an inhibitor of osteogenesis. The nanoparticles adhered to the scaffold for days, presumably because of ionic interaction within the pores, and were able to silence genes locally when mesenchymal stem cells were seeded into the scaffold. When scaffolds containing two compartments with either of the siRNAs were implanted into mice, a spatially differentiated tissue development took place (Fig. 13.8).

Using the aforementioned techniques is only relevant to the engineering of “simpler” tissues where the different cells, materials or drugs can be arranged in geometries amenable to separate cell/drug seeding or material joining by hand. Most

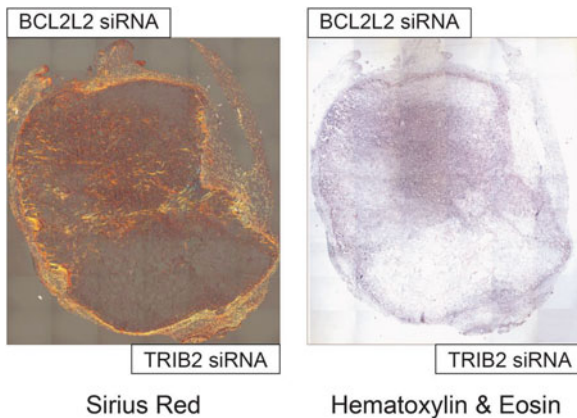


Fig. 13.8 A two-sided cylindrical scaffold containing adipogenic TRIB2 siRNA nanoparticles on one side and osteogenic BCL2L2 siRNA nanoparticles on the other side was seeded with mesenchymal stem cells and implanted sub cutaneously in a mouse. After 8 weeks pre-adipogenic tissue had developed in the TRIB2 side whereas pre-osteogenic tissue had developed in the BCL2L2 side. Figure modified from Andersen et al. [142]

internal organs and tissues, such as the kidney, liver, spleen, lungs and pancreas have more complex microstructures that necessitate different strategies. Towards engineering such tissues, various additive manufacturing strategies are being explored [143]. Many approaches rely neither on nanotechnology nor on drug release, instead, most research groups focus on printing different cell types into different volumes of an implant [144–147]. However, a different approach would be to 3D print different drugs into different compartments that correspond to different cell types in the desired tissue similar to the strategies that have worked well with simpler tissues [139–142]. Towards this, we have recently shown that nanoparticles containing different siRNAs can be 3D printed into different regions of a three component carbohydrate hydrogel using a patient derived CT-scan as the deposition guide [148], our approach is shown on Fig. 13.9. Once MSCs were seeded the siRNAs could then induce spatially restricted gene silencing. Co-printing cells and drug formulations can be combined, Xu et al. co-printed plasmid nanoparticles and endothelial cells into a collagen gel implants, during subsequent culture the endothelial cells became transfected by the plasmids [149].

13.4 Conclusion

The release of drugs from implants has clearly shown its clinical efficacy in fighting implant associated infections, modulating inflammation and in promoting tissue healing. It is, therefore, likely that most future implants will utilize drugs and that they will use nanotechnology to modulate their release and function. For artificial implants there will likely be a move towards employing multiple drugs to control infection, inflammation and other tissue responses. Nanotechnologies such as controlled release nanoparticles, responsive tethering and various layer encapsulation strategies will be used increasingly to ensure that effective but non-toxic concentrations of the different drugs are released at the specific times when each of these drugs are needed. For biological implants, there will likely be increased use of multiple drugs that promote different events in tissue regeneration. An implant will thus likely contain different stimulators of the progressive stages of stem cell differentiation as well as modulators of vascularization, reinnervation and cell survival, proliferation and migration. Many of these drugs will be labile biological macromolecules that are needed only at specific time points and locations. Nanotechnology will surely play a role in protecting the drugs until they are needed and in delivering the drugs to the right sub-cellular compartment in the targeted cells at the right time in the correct regions of the implant. This will be achieved by combining advanced targeted drug delivery systems with the advances in new scaffold fabrication technologies such as additive manufacturing and electrospinning.

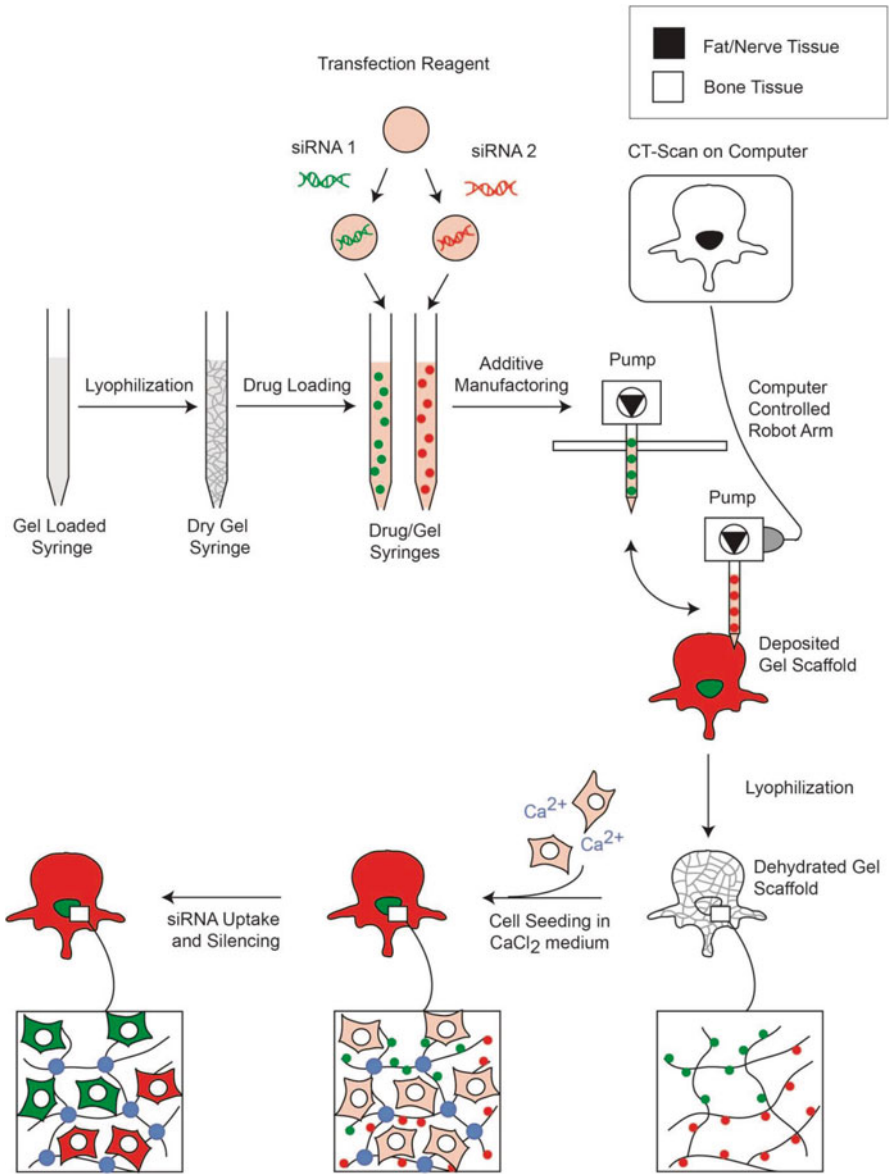


Fig. 13.9 Additive manufacturing of a complex implant guided by patient scanning data and using two hydrogels loaded with different siRNA particles

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Chapter 14

Guided Cellular Responses by Surface Cues for Nanomedicine Applications

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Abstract Using surface cues to guide and ultimately control cellular responses is of paramount importance in numerous biomedical applications. Since cells react on feature sizes both on the micro and nanometer scale, these length scales are crucial parameters when designing new materials for applications in medicine e.g. for tissue engineering and drug delivery. Thus, variation of the features at the nanometer length scale is an integral part of nanomedicine research and development. In this chapter the interaction between biological systems and artificial materials will be addressed in general with focus on simplified model systems where only one or a few parameters are varied at two-dimensional (2D) surfaces. At first biomolecular adsorption/immobilization on surfaces will be addressed followed by a discussion of approaches to synthesize functionalized surfaces and the influence of such surfaces on cellular response. Some of the key parameters, which will be discussed in more detail, are topography, chemistry, and elastic modulus of the substrate. Even though there is a vast amount of published data it will become clear that there is still not a detailed understanding of the influence of these parameters on biosystems at the cellular level. Obviously, the degree of complexity increases when several of the surface cues are combined. The challenges in understanding the cellular responses in detail in real systems with the simultaneous variation of multiple parameters leads to the introduction of the field of high throughput screening of biomaterials.

Keywords 2D surfaces • Surface cues • Artificial niche • Cell behavior • Nano/micro fabrication

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14.1 Introduction

The search for routes to replace or repair malfunctioning tissue or organs in the human body has led to multiple breakthroughs within the field of medicine. Systems such as orthopedic implants, heart valves, intraocular lenses, stents and pacemakers are routinely used to sustain body function and to improve quality of life of patients worldwide. To enable continuous improvement of existing systems and to enable further advances within the field of biomedical devices, a detailed understanding of the interaction at the nanometer scale between cells and their surroundings is needed. Understanding this interplay will allow researchers to tailor improved biomaterials to be used for e.g. guided stem cell differentiation and tissue engineering purposes. The research and development focused on novel materials with functionality depending on nanometer scale features is an integral part of the field of nanomedicine.

Fundamental cellular functions and behavior are governed by diverse and complex external stimuli from the surrounding microenvironment, known as the niche. The niche consists of extracellular matrix (ECM), soluble factors as well as other cells that are present and these provide a variety of physical and chemical factors. A multitude of these external stimuli essentially orchestrates the fate of cells by initiating a cascade of biochemical signals by interacting and consequently influencing the fundamental cellular functions including motility, proliferation, differentiation and apoptosis. A focal point of research has been aimed at deconstructing the complex relationship between external signals and cellular responses to increase our knowledge in understanding the cellular regulatory mechanisms. Since mammalian cell niches are challenging to control experimentally, the properties of niche regulatory components have been extensively explored *in vitro* using a *reductionistic approach* and a toolbox of artificial (bio) materials and fabrication strategies for creating the artificial cues. Here, the surface and interfacial properties of the materials become of paramount importance for studying the effect of physical, chemical and biomolecular ‘cues’ on cellular function and fate, since it is the surface that first comes in contact with the cells and regulatory processes are initiated at the interface between the material and the cells.

This chapter describes the surface and interfacial aspects of artificial materials for guiding cellular behavior. Current investigations are, to a large extent, based on trial and error but continuous research within this area holds promise to gradually expand the nanomedicine tool-box and ultimately enable rational design of surfaces for specific purposes. The focus is two-dimensional (2D) nanoscale structures thus excluding the field of porous materials/scaffolds (3D). The effect of various artificially constructed external material surface cues including chemical, biomolecular and physical properties as well as a combination of multiple cues that are influential to the cell behavior are highlighted. It is important to note that the cellular responses obtained from experiments using 2D and 3D systems do not always correlate with one another and recent growing evidences indicate that the 3D experimental platforms depicts more of a native cellular niche. Interested readers should refer to

excellent reviews [1–7] which discuss the 2D and 3D culture systems and the impact on cell behavior. Nevertheless, 2D experimental platforms are highly advantageous as a simplified approach in determining the individual role of cues in a well-defined manner on cell behavior via a repertoire of material fabrication processes and commonly employed engineering strategies for generating 2D systems. Recent emergence of high throughput screening (HTS) platforms is also highlighted; as such platforms have now become a proven tool for rapidly and systematically probing the effect of multiple nanoscale niche components.

14.2 The Role of Surfaces in Biorecognition and Cell Responses

14.2.1 Biomolecules at Interfaces

Biorecognition by the cells interacting with an artificial material depends on the type, orientation and conformation of the biomolecules presented on the surface. The state of how biomolecules reside on the surface is influenced by the properties of the surface [8]. This successive interfacial event occurs as, for example, when a medical device is implanted into the body (Fig. 14.1). Among a variety of biomolecules, adsorption of proteins to surfaces in particular has been extensively studied since the cell adhesion to the surface and the subsequent cellular response depends on the availability of specific cell binding peptide sequences presented by the proteins. At constant temperature and pressure, protein adsorption to surfaces occurs by lowering the Gibbs energy, G of the system with the contributions from enthalpic (H) and entropic (S) effects; $\Delta_{\text{ads}}G = \Delta_{\text{ads}}H - T\Delta_{\text{ads}}S$. Numerous enthalpic and entropic effects including van der Waals and electrostatic forces, hydrogen bonding, hydrophobic interactions and surface packing restrictions via steric and excluded volume effects determine protein adsorption [9, 10]. The intricacy of the protein adsorption becomes further evident as some of these effects can be simultaneously altered by the external parameters of temperature, pH and ionic strength: [11, 12] Diffusion of proteins towards the surface increases with increasing temperature and entropic gain is expected to occur as the water molecules and ions are released upon protein adsorption. Changes in the pH or ionic strength alters the electrostatic state of the proteins [13]. At the isoelectric point and/or at high ionic strength, protein-protein repulsion is minimized leading to higher protein adsorption as a result of higher surface packing density, as well as causing the proteins to aggregate.

Protein adsorption on surfaces occurs in several consecutive steps and the concentration of proteins in solution differs from those on the surface with time, depending on the properties of the surface, types of protein involved and their environment [8, 14]. This means that the surface exposed to a single or a few protein types may be very different from those surfaces exposed to multiple types of proteins. In the mixed protein system (Fig. 14.2a), certain types of proteins initially

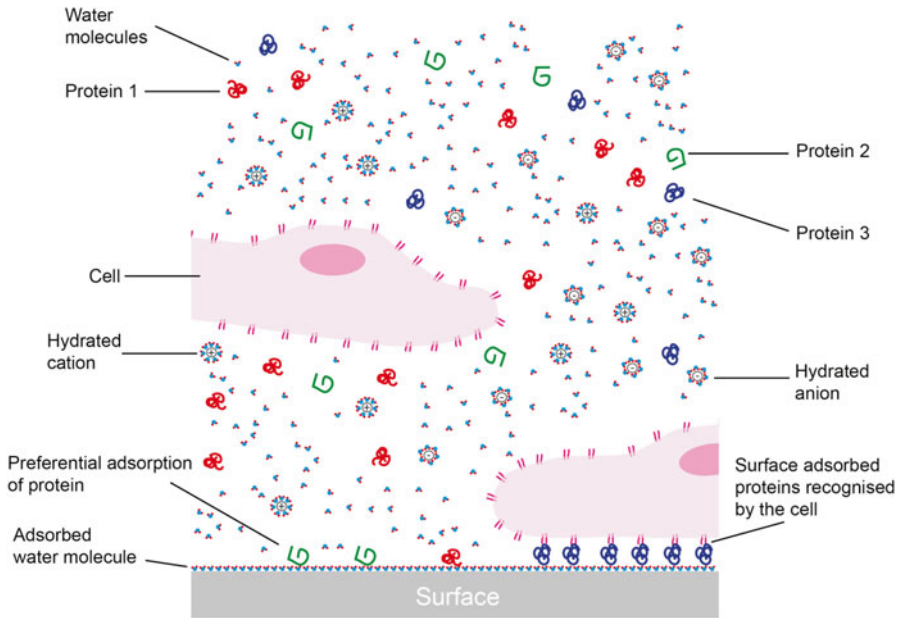


Fig. 14.1 Schematic diagram depicting the interface between the (implant) material surface and the biological milieu *in vivo* (adapted from Kasemo [8]). Successive surface adsorption events occur as water molecules first reach the surface, followed by proteins and finally the cells. The surface-cell interaction depends on the properties of the surface

adsorb onto the surface almost immediately after exposure. Depending on the state and type of the adsorbed proteins, displacement and reorganization by other types of protein may occur at the interface over longer adsorption time periods of minutes to days, a phenomenon known as the Vroman effect [14]. Once adsorbed, each protein on the surface could adopt orientational and conformational changes, depending on the biological and/or local environment (Fig. 14.2b and c). The final structure of the adsorbed protein layer also depends on the rate at which the proteins adsorbed onto the surface and the kinetics of protein adsorption is largely governed by the bulk concentration and protein size. The rate of adsorption is higher with higher bulk concentration and the neighboring proteins may sterically prevent conformational changes, whereas at a lower bulk concentration, the rate of adsorption is lower, leading to longer time for the given proteins to undergo conformational changes on the surface [14]. The size of the protein also contributes to the kinetics and arrangement of proteins on the surface, where a higher number of smaller proteins adsorb onto the surface compared to larger proteins because of higher rates of diffusion, but larger proteins typically possess higher binding affinity towards the surface due to having a larger potential contact area [12]. Preferential orientation of adsorbed proteins on the surface originates from its complex structure, with different regions of proteins presenting different properties depending on the local amino

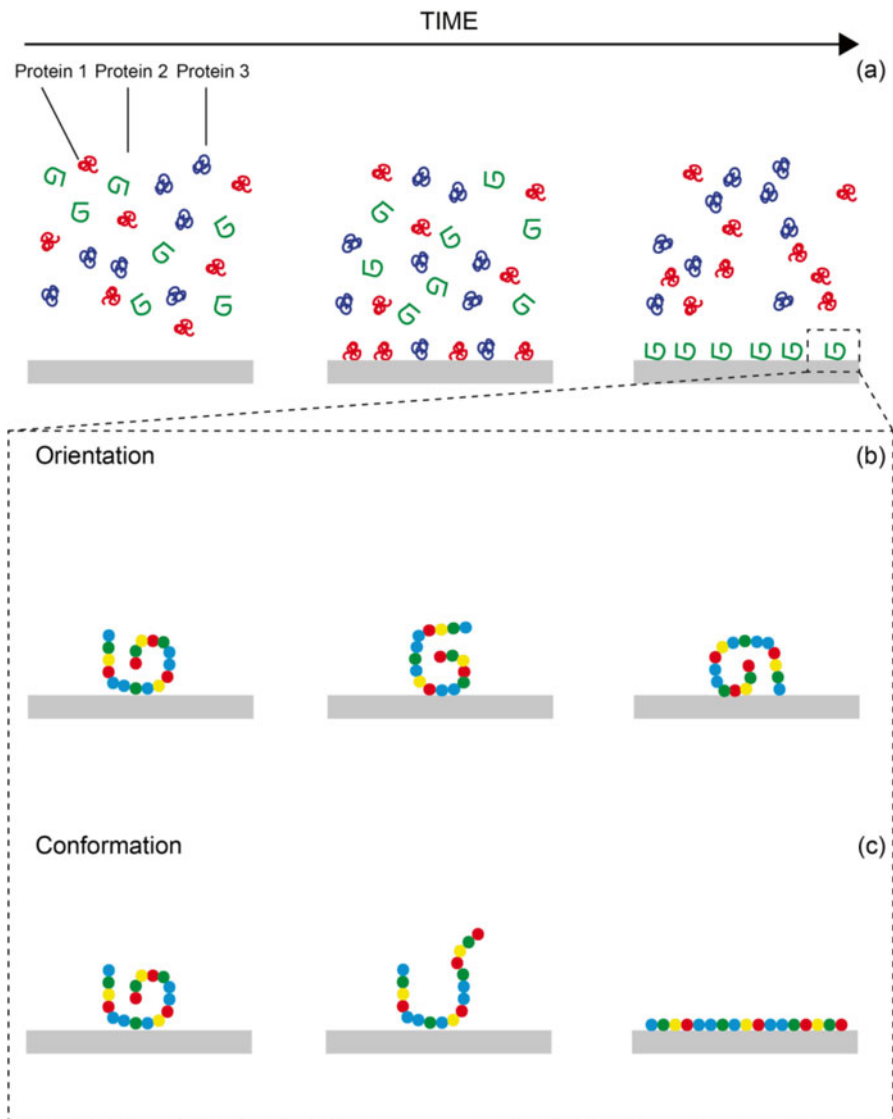


Fig. 14.2 Protein adsorption on a surface with time (adapted from Castner and Ratner [14]). In a mixed protein solution, displacement of the initially adsorbed protein by another type can occur over a prolonged period of time (a). Once a protein is adsorbed on a surface, orientation (b) and conformation (c) change may take place

acid compositions. If the adsorbed protein is structurally stable and the adsorption is driven by electrostatic interaction between the protein and the surface, the orientation of the protein could change further if a sufficient surface protein density is reached. High surface density leads to close proximity between the neighboring adsorbed proteins exposing the electrostatically repulsing domains, leading to

reorientation towards a less repulsive arrangement (Fig. 14.2c). Conformational change (denaturation) of proteins at adsorption is epitomized by the structural ability of the protein and the result of the difference in the free energy minimum between the proteins in solution to the proteins adsorbed on the surface [11, 12, 14]. Small and rigid proteins, classically referred to as ‘hard proteins’ such as lysozyme and β -lactoglobulin have less tendency to undergo structural change upon surface adsorption compared to larger and less tightly structured ‘soft proteins’ such as albumin and immunoglobulins [15]. Protein denaturation may lead to loss of biological function and hinder subsequent recognition by the cells, ultimately leading to e.g. an inflammatory response.

14.2.2 Cell Adhesion on Surfaces

In vivo most cells (except non-adherent cells such as blood and tumor cells) require adhesion to a surface of extracellular matrix (ECM) as well as surrounding cells in order to maintain and regulate the functionality and development of multicellular organisms. Cells interact with two types of surfaces in vivo; cell-cell and cell-ECM interactions. Cell-cell adhesion is a specific process where cells bind selectively to another type of cells mediated by cell adhesion molecules, including selectins, integrins, immunoglobulin super family and cadherins [16, 17]. The cell-ECM adhesion on the other hand, is primarily mediated by integrins and much of the research effort in the recent years has been focused on unraveling the composition and molecular architecture of integrin adhesions as well as its adhesion related signaling events [18–20]. Integrins are heterodimeric transmembrane proteins composed of α and β subunits (most common being $\alpha_5\beta_1$ and $\alpha_v\beta_3$) that specifically recognize different short amino acid chain (e.g. the often mentioned Arg-Gly-Asp sequence, often termed RGD) motifs present in many of the ECM proteins. Each set of subunits possess a distinct capacity for bi-directional transduction (outside-in and inside-out). Characteristic features in cell-matrix adhesions include focal complexes, focal adhesions, hemidesmosomes and podosomes and these differ in terms of location, size, morphology, constituents and are induced by Rho family GTPases. Formation of focal adhesion has been the most studied and characterized type of adhesions. An initial step in the formation of focal adhesion involves physical interaction between the ligand and integrin as its conformation changes into an active upright state, opening the intracellular domains of the integrin for the subsequent attachment to cytoplasmic proteins including talin, vinculin and focal adhesion kinase (FAK) that connects to the actin cytoskeleton [21]. This further leads to the assembly of focal adhesion by recruiting additional components leading to further clustering of integrin complexes promoted by the binding of vinculin to talin and reinforcement of the integrin-cytoskeleton bonds. Molecular complexity in the formation of focal adhesion is evident, with integrin-mediated adhesion alone consisting of ~160 distinct components including actin regulators, adaptor proteins and signaling molecules with ~700 links to each other [19].

14.3 Fabrication Strategies for the Generation of Artificial Surface Cues

14.3.1 Chemical and Biomolecular Patterning

In the context of conducting fundamental cell studies, chemical and biomolecular surface patterning involves precise positioning and presentation of chemical functional groups and biomolecules onto a substrate over micrometer to nanometer lengths scale. Generally the most straightforward approach is to fabricate a pattern consisting of areas containing a single chemical/biomolecular moiety which the cells can interact with, while the surrounding regions are non-fouling/bioresistant. Poly- and oligo-ethylene glycol (PEG [22, 23] and OEG [24, 25], respectively) containing materials are often used for creating such bioresistant regions, though other types of materials such as polysaccharides [26] and zwitterionic polymers [27, 28] have been found to provide robust bioresistance. Inspired by the microelectronics fabrication methods used for preparing e.g. microprocessors [29], MEMS [30], and NEMS [31], the early surface patterning approaches have been to employ ‘top-down’ instruments such as electron beam (e-beam) lithography [32] and photolithography [33]. In contrast, ‘bottom-up’ methods such as colloidal lithography (CL [34]), block copolymer lithography (BCL [35]) and block copolymer micelle lithography (BCML [36]) have been developed for the fabrication of micro and nano-scale surface patterns in an autologous manner typically via self-assembly processes. A number of specific advantages and drawbacks exist in both fabrication strategies; for example ‘top-down’ approaches possess flexibility over feature shape, pattern size and positioning. However, serial and multi-step fabrication as well as access to a clean room is often required which can be costly. On the other hand, ‘bottom-up’ approaches can be straightforwardly conducted ‘on a bench’ at a low cost and over large areas but suffer from a lack of versatility in the geometrical features that can be patterned as well as the frequent presence of pattern defects and order at a longer length-scale. In almost all cases, self-assembled monolayers (SAMs) such as thiols [37, 38], silanes [37, 39], phosphates [37, 40] and polyelectrolytes [41] are used in combination with lithographic techniques either; (1) directly, (2) indirectly or (3) via material guided self-assembly. The direct patterning of SAMs typically involves using SAMs as an ‘ink’ to spatially position SAMs onto a substrate and, subsequently, backfill the ‘clean’ non-printed areas with bio-resisting SAMs. Examples include micro- and nano-contact printing (μ CP [42] and nCP [43]), dip and polymer pen nanolithography (DPN [44, 45] and PPN [46]) and ink jet printing [47]. The indirect patterning of SAMs involves temporary pre-patterning of surfaces with polymer resists to create a surface pattern of exposed/unexposed substrate surface. The surface is subsequently immersed in the solution with components for generating SAMs which will self-assemble onto the substrate-exposed areas and, finally, the polymers are subsequently removed and the surface passivated. The pre-patterns can be fabricated at sub-micro and nanometer scale using photolithography [41, 48], e-beam lithography [49] and nanoimprint lithography (NIL [50]) using

polymer resists. In comparison to the above approaches, the material-guided self-assembly takes advantage of the material-specific self-assembling nature of SAMs (e.g. thiols on gold and silanes on oxide surfaces), where surfaces are first patterned with different materials and sequentially incubated in different types of SAMs. Patterning of different materials can be carried out by, for example, photolithography [51] and CL [52] using physical vapor deposition (PVD) and/or chemical etching to make e.g. a stable chemical contrast or height variation.

For biomolecular surface patterning, biomolecules can be immobilized either directly (i.e. physically adsorbed) onto a surface or via specific conjugation. Physical adsorption of biomolecules is commonly carried out via electrostatic interactions between biomolecules and substrates bearing opposite charge, or via hydrophobic interaction by rendering the substrate hydrophobic using SAMs, depending on the substrate wettability. Specific conjugation via covalent (e.g. N-hydroxyl succinimide or NHS coupling between primary amines of biomolecules to a carboxylic acid moiety) or affinity-ligand based interactions (e.g. biotin-streptavidin) permits presentation of biomolecules in its native state without denaturation or displacement, while orientation can be controlled by targeting a specific amino acid sequence on the protein for conjugation.

Although patterning and presentation of more than one type of chemical moieties can be conducted on a surface with relative ease, the patterning of multiple biomolecules may become a challenging affair, unless direct immobilization methods are employed. For generating multiple protein patterns via indirect or material guided self-assembly routes, a chemical moiety that can be couple specifically to one type of biomolecules must be patterned while minimizing/avoiding non-specific bioadsorption from another type of biomolecule which is to be patterned on the surface. This implies that the general covalent coupling method such as NHS cannot be straightforwardly used to pattern multiple proteins, thus different affinity coupling moieties with a non-fouling background is required [53–55]. For further details in chemical and biomolecular patterning, see further Refs. 55–60.

14.3.2 Topographical Patterning

Surface topography, or in other words the structural landscape of a surface, is a physical cue that has already found applications within the field of biomaterials. It is for instance well established that rough surfaces with features on the micro and nanometer scale are able to facilitate accelerated bone ingrowth for orthopedic and dental implants [61] as compared to smooth controls. Such surfaces are typically prepared by blasting the surface with an appropriate blasting media yielding a random structure composed of valleys and peaks. While random structures are of interest application wise, the trend with respect to more fundamental studies is moving towards employing a higher degree of structural coherence when investigating cellular responses. Such systems are frequently composed of ordered, semi-disordered or disordered structures with well-defined feature sizes in the micro and/or

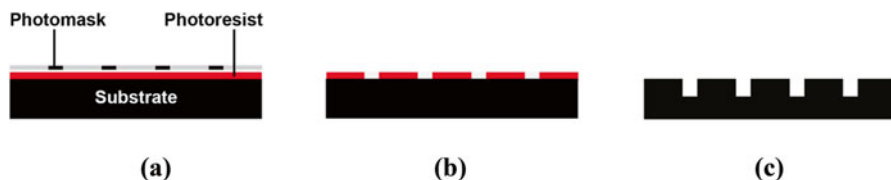


Fig. 14.3 Schematic overview of the photolithography process using a negative resist. **(a)** The substrate is coated with a thin layer of photoresist(s) (*red*) and a pattern from a pre-designed photomask is then transferred to the resist layer by irradiation with ultra violet (UV) light through the photomask. Molecules of the exposed regions of the resist react to form cross-linking bonds, which stabilize the material. **(b)** Following UV exposure, the unreacted resist is removed and the underlying substrate is exposed. The remaining resist protects the underlying substrate and it is now possible to selectively modify the substrate using different methods such as e.g. physical vapor deposition or reactive ion etching (RIE). **(c)** Final surface topography prepared using RIE (Color online)

nanometer size range [62, 63]. A wide range of methods can be utilized when synthesizing such cell culture substrates, most of which are based on the “top-down” approach as previously mentioned. One of the most frequently used techniques for producing well defined topographical features in the micrometer range is photolithography. The basic principle behind this method is illustrated in Fig. 14.3 and relies on using a photoactive polymer, known as a resist, to coat a substrate material, e.g. silicon.

The resist at the surface is subsequently selectively exposed to electromagnetic radiation through a photomask which screens part of the resist. Two main types of resist exist known as “negative” and “positive” which indicate if the resist is cross-linked or destabilize, respectively, by the exposure. Typically radiation in the UV or deep UV range is used as the short wavelengths allow for a higher ultimate resolution (diffraction limited) and, by that, higher feature resolution. Upon radiation the pattern transferred from the photo mask to the resist is developed, thus, exposing the underlying substrate at the intended areas. Subsequently, the underlying substrate can be modified and this process is typically based on either a reactive ion etching process, through which substrate material is removed, or on a physical vapor deposition process enabling addition of a material onto the substrate surface. While several modifications to standard photolithography has been developed, which allows for a very high lateral feature resolution of ~ 40 nm [64, 65], these are extremely costly and mainly employed in the semiconductor industry and most commercially available methods have a maximum resolution of approximately 250 nm. The advantage of photolithography is the high throughput nature of the process when the mask has been produced. Even though the production is relatively straight forward, access to a clean room is required. The mask does, however, represent a significant expense and if lateral changes to the structures are needed, a new mask must be obtained. If features in the lower nanometer size range are needed, the process of electron beam (e-beam) lithography can be employed. This method also utilizes a resist layer, but rather than using a mask to transfer a pattern to the substrate surface, it uses an electron beam to “write” the structures directly into the resist. This is

known as a direct write method and while it is capable of producing features with a size of a few nanometers, it is also a time consuming process [64] since it is serial. The topographical structures produced by photo and e-beam lithography can be used directly for e.g. cell culture experiments. However, the topographical structures can also be utilized as masters for replica molding. This method belongs to the family of soft lithography and the basic principle is to replicate surface topographical features using a polymeric material. Several different polymeric materials can be used for this process but the material most frequently employed for cell culture studies is based on polydimethylsiloxane (PDMS). Typically the liquid polymer material is cast onto the surface structures and set to cure. Subsequently, the polymer material can be separated from the master, resulting in a negative replica of the surface structures. As the master structure can be reused, replica molding is a very cost effective route for obtaining a large number of samples. With respect to the structural integrity of the transferred pattern, the elastic modulus of the material used for the process greatly affects the lower limit for the minimum feature resolution. Utilizing a chemically modified PDMS composition, topographical replication of carbon nanotubes with a vertical dimension of ~ 2 nm has been achieved [66]. This special material is referred to as “hard” PDMS and has an elastic modulus of ~ 9 MPa [67].

14.3.3 Modulation of Mechanical Surface Properties

When designing substrate surfaces with varying mechanical properties, e.g. viscoelasticity, it is important to keep in mind that the simultaneous introduction of any chemical changes also has the potential to affect the cellular response. With respect to elucidating the fundamental mechanisms involved in cellular mechano-sensing, it is therefore important to employ model systems that present varying mechanical properties with a minimal degree of change to other properties such as e.g. surface chemistry, wetting properties and topography. In the literature there are two main approaches that have found applications for research purposes. These are; (1) varying the degree of cross-linking of polymeric materials and (2) geometric design of the surface topography of elastic materials. With respect to the approach of varying the chemistry of the substrates, the two systems that are most frequently employed are based on polyacrylamide hydrogels or polydimethylsiloxane (PDMS) elastomers.

Polyacrylamide hydrogels are prepared by copolymerization of the two components; acrylamide and *N,N'*-Methylenebisacrylamide (bis-acrylamide). During this reaction the acrylamide monomer will form polyacrylamide chains which are cross-linked to each other by the bis-acrylamide cross-linker component, thereby forming an interlocking polymer network [68]. Depending on the amount of bis-acrylamide used in the preparation the number of cross-linking events can be varied which in turn determines the elastic properties of the final hydrogel [69]. Increasing the concentration of the bis-acrylamide component will increase the elastic modulus of the

material. The viscoelastic properties of the polyacrylamide system can also be designed by varying the ratio between the acrylamide monomer and the bis-acrylamide cross-linker. Using this, it is possible to vary the loss modulus (a materials ability to retain a specific strain as a function of time) while keeping the storage modulus (elastic properties) of the material constant between the different substrate formulations [70]. With respect to the PDMS system the classical method for modulating the mechanical properties is also to vary the amount of cross-linker used in the preparation, similarly to the acrylamide system. This will result in varying degrees of cross-linking and produces materials with varying elastic modulus. The use of alternating curing periods or curing temperature have also been employed for the production of cell culture substrates [71]. This approach is, however, linked to the risk of time dependent alterations to the mechanical properties as the curing reaction for many PDMS formulations proceeds at room temperature. With respect to PDMS based systems, Fu et al., have described an alternative for varying the effective mechanical properties without affecting the chemical composition of the material. In their study, a geometrical approach, where closely spaced pillars were prepared by replica molding, is described. By keeping the chemical composition constant but varying the height of these pillars, substrates were generated showing varying pillar deflections in response to cellular traction forces [72]. Similarly to the challenges associated with varying chemical compositions, varying pillar deflections between the different substrates will ultimately change the local topographical appearance of the substrate surrounding the individual cells. This, in turn, might also hold the potential to affect the cellular response.

14.4 Surface Cues for Controlling Cell Behavior

Enormous complexity of the cell-matrix interaction is evidently present in a typical *in vivo* nanomedicine application; it is far from a trivial task in deconstructing numerous direct and indirect signaling interactions that can activate or inhibit targeting molecules which ultimately determines the cellular fate. One might, as Geiger et al. [19] expressed, attempt two approaches for unraveling this multifarious adhesion signaling events: (1) investigate individual cellular adhesion mechanism in detail, integrate and respond to the signal given by the cells, or (2) initiate specific cellular response by altering the properties of the cell interacting surfaces. With substantial progress made in the areas of materials science and nanotechnology in the recent decades, reconstruction and mimicry of *in vivo* niches down to the nanometer length scale has become possible for measuring cell-surface interactions. Furthermore, discovery of numerous synthetic materials have allowed investigating beyond the *in vivo* mimicry for identifying and seeking new cellular behavior on novel materials that are absent *in vivo*. Single and often multiple cues can be presented on a surface with varying physical, chemical and biomolecular properties as described in the previous sections. Physical properties may include; surface roughness/topography, viscoelasticity as well as shape, size, and spacing of topographical

features. Chemical and biomolecular properties include; composition (i.e. basic chemical moieties, bare material composition) and material-specific surface chemistry (i.e. metals, polymers and their associated surface functionality), amino acids, proteins, carbohydrates and lipids. Experimental conditions and the corresponding cell response to the interfacial parameters should be prudently designed and accounted for. For example, topographical effects on cellular behavior could be interpreted in terms of wettability (i.e. surface energy) or the roughness of the surface investigated. In addition, cell culturing conditions such as presence/absence of serum and the concentration, type of growth media, cell seeding density, durations of study and type of cells/passage number may have a dramatic effect on the cell behavior. The following sections will provide critical examples of studies conducted in the past and recent and their importance in determining relationships between artificial instructive cues and cellular behavior.

14.4.1 Chemical and Biomolecular Cues for Guided Cellular Behavior

14.4.1.1 Chemical Cues

The early studies into the effect of surface chemical properties on cellular behavior have been conducted directly on material surfaces such as polymers to determine the relationship between the polymer surface properties and cell behavior. For example, van Wachem et al. studied adhesion and proliferation of human endothelial cells on a range of polymers with varying wettability and surface charge and observed polymers with moderate wettability showed highest number of cells adhered [73, 74]. The concentration of functional groups within a polymer can be readily controlled by copolymerization [74, 75] and the surface chemical groups can be quantitatively determined by surface analytical techniques such as X-ray photoelectron spectroscopy (XPS) and Fourier-transform infrared spectroscopy (FTIR). While polymer surfaces provide an ideal platform for biomaterial research and plays an important role in elucidating material-specific interaction with cells, it is difficult to readily obtain precise control over other surface parameters (e.g. topography) that may also contribute to cellular behavior depending on the type of polymers used. To determine the sole effect of chemical groups on cellular behavior, self-assembled monolayers (SAMs) of thiols or silanes have been typically employed on flat surfaces. For studies using a single type of functionalized alkane thiols, general trends in the cell number has been observed regardless of different cell types, where, for example the growth of bovine aortic endothelial cells (BAEC) [76] increased in the order of $\text{OH} < \text{COOCH}_3 < \text{CH}_3 < \text{COOH}$, glioma cell line C6 [77] generally increased in the order of $\text{SH} < \text{CH}_3 \leq \text{OH} < \text{NH}_2 < \text{COOH}$, and MCF-7 cancer cells [78] increased in the order of $\text{SH} \leq \text{OH} \leq \text{CH}_3 < \text{NH}_2 \leq \text{COOH}$. These studies have been conducted with the presence of serum, where some studies have

highlighted the correlation between chemical functional groups and the level of elutability of adsorbed albumin and adsorption of cell adhesive proteins such as fibronectin [76, 79, 80]. Similar trends in the cell behavior such as cell proliferation and spreading on various functional groups have also been observed using functional silanes for a range of cell types including fibroblasts [81–83] and stem cells [84, 85]. In particular, positively charged NH_2 surfaces led to highest viable human mesenchymal stem cells (hMSCs) adhesion and increase in the osteogenic differentiation pathway, while negatively charged COOH surfaces showed rounded morphology with chondrocytic differentiation [84]. As for neural stem cells (NSCs), SO_3H surfaces showed highest cell contact area and favored oligodendrocyte differentiation, while COOH , NH_2 , SH and CH_3 guided NSCs to differentiate into neurons, astrocytes and oligodendrocytes [85].

Limited studies into the sole direct effect of micro-patterned chemical moieties on cell behavior have been reported to date. Basic studies into the baby hamster kidney (BHK) cell adhesion and proliferation were carried out on hydrophilic/hydrophobic silane patterns on a quartz surface using photolithography showed higher cell number and proliferation on the patterned hydrophilic regions [86]. Similar cell adhesion results were obtained on osteoblasts [87] using alkyl thiols with different terminal functional groups patterned via photolithography. Preferential cell growth were observed with hierarchical preference of $\text{CH}_3 < \text{OH} < \text{COOH}$. However, the study involved pre-adsorption of fibronectin onto these moieties and the results indicated that the focal contact formation by the cells varied in the same order and depend on the adsorbed protein density and cell-cell interactions.

The direct effect of nanoscale chemical patterns on cells has been studied more extensively. In particular, several cell adhesive short peptide motifs such as RGD [88] have been patterned in nanoscale to study cell adhesion, spreading, focal adhesion assembly, migration and polarization using block copolymer micelle lithography (BCML) and MC3T3 osteoblasts [89–92]. These studies have demonstrated that cyclic RGDfK peptide-functionalized gold nanoparticles that can only interact with a single integrin per particle spaced less than 73 nm showed cell adhesion and spreading, whereas spacings larger than 73 nm or with a random peptide sequence have shown limited adhesion and spreading. This critical finding of the ligand spacing threshold of ~ 70 nm also applied to ordered vs disordered pattern, where nanoscale disordering with global average ligand spacing of larger than 70 nm inhibited cell adhesion [92]. Controlled differentiation of human mesenchymal stem cells (hMSCs) have been demonstrated on nanoarrays of COOH , NH_2 , CH_3 and OH with feature size of ~ 70 nm having defined spacings ranging from 140 to 1000 nm [93]. Compared to standard tissue culture polystyrene (TCPS), CH_3 nanopatterned arrays enhanced the expression of CD29, CD73, CD90, CD105 and CD166 and maintained an enhanced expression of phenotypic markers after 28 days, highlighting that simple chemical nanoarrays can promote and maintain MSC phenotypes *in vitro* without exogenous biological factors and heavily supplemented cell media.

14.4.1.2 Biomolecular Cues

The effect of biomolecular cues on cellular behavior has been researched intensely over the last several decades and continues to be an important area in the field of nanomedicine. Analogous to the studies conducted on simple chemical moieties/cues as described in the previous section, the early studies into the effect of biomolecules on cellular behavior were carried out on flat surfaces. The studies in the 1970s and 1980s in particular involved making basic observations of adhesion, motility and growth of different types of cells on ECM components physically adsorbed on surfaces. With a repertoire of surface patterning strategies becoming readily accessible in the 1990s, the research focus shifted towards investigating the micro and nano-scale effect of surface patterned biomolecules. One of the notable studies into the effect of geometric shapes patterned with adhesive ECM protein was demonstrated by Chen et al. [94, 95], where the behavior of endothelial cells were switched from growth to apoptosis by changing the size of fibronectin adhesive islands. Micro meter scale ECM geometrical limits on cell spreading and adhesion were studied further by Lehnert et al. [96] using mouse B16F1 melanoma cells, buffalo rat liver cells and NIH-3T3 fibroblasts. They discovered that all of the studied cell types regulated the number of focal adhesions formed to the range of surface coverage and the spacing of the fibronectin pattern, with maximum distance of 25 μm between the adhesive surfaces with fibronectin coating for cells to achieve half-maximal spreading. Dot area of as low as 0.1 μm^2 was capable of inducing focal adhesion formation but did not support spreading when the inter-dot spacing was increased more than 5 μm . The effect of pattern shape was also investigated on stem cells. McBeath et al. [97] employed similar approach from the previous study [94] to investigate the lineage commitment of human mesenchymal stem cells (hMSCs). The study have importantly found that the cell-adhesive pattern that allowed cells to spread on the surface, steered hMSCs into the osteogenic lineage, but the pattern that restricted the spreading led hMSCs into the adipogenic lineage and the cell shape, cytoskeletal tension and RhoA signaling were found to be integral to the lineage commitment of the hMSCs.

Tseng et al. [98] investigated how ECM proteins affect the spatial organization of intercellular junctions by constraining the location of epithelial cell-fibronectin adhesion, by patterning fibronectin in various shapes. It was found that intercellular junctions were displaced due to a large tensional force when they were close to fibronectin, whereas the junctions remained stable with low tensional force in the area without fibronectin, highlighting the importance of ECM in morphogenesis. Micro-patterning has also been used to study cell motility. Bailly et al. [99] investigated the regulation of lamellipodial protrusion during chemotaxis by using 10 μm line patterns of vitronectin surrounded by non-cell adhesive regions. The cells only adhered on the vitronectin lines but upon exposure to epidermal growth factor (EGF), the cells were able to protrude into the non-cell adhesive regions. However, a stable protrusion was only achieved by establishing a firm contact to the substrate. Kumar et al. [100] and Kushiro et al. [101] demonstrated controlled directional migration of fibroblast, endothelial and epithelial cells by generating asymmetric

'tear drop' shaped adhesive islands of fibronectin. Although the same 'tear drop' micro-patterns were used in both studies, differing directional bias were observed due to the disparate migration properties of the cell types and the difference in the environmental cues presented to the cells. The importance of cell-cell interactions in cell motility were further highlighted by micro-patterning a combination of cell-ECM (collagen IV) and cell-cell (E-cadherin-Fc) adhesion cues to determine the motile behavior of epithelial cells on alternating micro-patterned lines [102]. While collagen IV was found to be essential for cell migration and E-cadherin-Fc was not, when both biomolecules were present, the direction of cell migration as a result of reduced lamellipodial activity was promoted by E-cadherin-Fc in a concentration-dependent manner without affecting the migration rate, indicating the importance of crosstalk between cell-cell and cell-ECM adhesion complexes upon migration. Beyond motility, microcontact printing was used to investigate the effect of spatial distribution of ECM on cell shape anisotropy and cell polarity in the context of cell division. The study concluded that ECM controls the location of intracellular actin dynamics at the membrane which in turn affect the orientation of the cell divisional axis [103].

The impact of nanoscale effect of biomolecules on cell behavior has also been studied in recent years in attempt to gain further understanding of the underlying mechanisms of cell function and fate. Colloidal lithography has been used to identify the nanoscale geometric effect of cell-ECM and cell-cell interaction processes by quantifying fundamental cellular processes including cell adhesion, spreading, focal adhesion formation and maturation and adherence junction formation [104–106]. In particular, the focal adhesion formation and development was found to vary depending on the ECM protein type, where a single focal adhesion was able to bridge across the gap between patches of vitronectin with sizes of 200 nm but not for fibronectin patches, until the patch size reached 500 nm. This may be due to the differences in the force which cells can exert on these proteins, with fibronectin being more prone to unfolding and mechanically more pliable than vitronectin [106]. Such approach was extended to study the effect of nanoscale geometry of the adhesive patches on epidermal stem cell differentiation, where it was found that such nano-scale patterns dictate the stem cell fate decision via the control of cell shape and AP-1 transcription activity, an important transcription factor controlling the expression of epidermal terminal transcription factors [107]. Giam et al. [108] also studied the effect of fibronectin feature size on human mesenchymal stem cell (hMSC) differentiation with and without the presence of osteogenic inducing media. By monitoring the FAK phosphorylation level, it was found that 300 nm fibronectin patterned features without osteogenic inducing media was more effective at inducing osteogenesis than the unpatterned surface with osteogenic inducing media, indicating the importance of programmed nanoscale feature sizes of ECM proteins for controlling the fate of hMSCs. Recent studies have further highlighted the importance of nanoscale ligand geometry in immune system responses [109, 110]. For example, using the block copolymer micelle lithography (BCML) approach, Delcassian et al. [109] generated CD3 and CD16 antibody patterns to monitor the effect of ligand spacing on the activation levels of T cells and Natural Killer (NK)

cells. The activations were assessed in terms of membrane-localized phosphotyrosine for T cells and the size of contact area for the NK cells, and the study showed decreased cell response with increasing ligand spacing in both cell types, with the threshold activation at the ligand spacing being 69 nm and 104 nm for T cells and NK cells, respectively. All together, these studies conclusively epitomize the importance of nanoscale biomolecular patterned cues for controlling cellular behavior in many biological settings.

14.4.2 Topographical Cues for Guided Cellular Behavior

It has been known for many years that topographical structures influence cellular response. Early on, already in the 1940s, Weiss demonstrated that cells are guided by directed structures and the term “contact guidance” was established [111]. Fabrication of topographical structures in the micrometer range for cell culture followed where the influence of topographical patterning on several cell types was examined, pioneered by the group of Curtis et al. [112, 113]. In 1964 it was shown that the cells align along the direction of a few micrometer wide grooves. Since then, a vast number of articles and reviews have addressed the influence of topography on cellular response. Following the expanding range of possibilities to synthesize and characterize structures with features at the nanometer scale (nanostructures), the focus has shifted to exploring this length-scale in the recent years [114, 115]. One focus area has been application of such nanostructures in nanomedicine.

One of the key issues has been to distinguish between the influence of topography on the adsorbing bilayer (e.g. denaturation and orientation of proteins) as already discussed in this chapter from the direct influence of topography on the attached cells. Multiple studies have shown that the amount [116] and the orientation/denaturation of the proteins adsorbed [117, 118] depend critically on nanoscale features. The change in conformation may change the presentation of binding sites of the protein either exposing or hiding such sites [119] thereby potentially also changing the cellular response. The field is further complicated by the fact that the actual roughness of a surface relevant to a given biomolecule is dependent on the size of the molecule making lateral size-scale dependent roughness measurements desirable [120]. Thus, it is still expected that features at the micrometer length-scale is less dependent on the detailed protein adsorption as already concluded 40 years ago [113]. With respect to nanostructured artificial surfaces, detailed knowledge about the mechanisms guiding the cellular response, including attachment, spreading and proliferation, is still to be obtained. An additional challenge concerning the characterization of the cell-substrate system is the choice of a specific protocol, including cell type(s). Different cell types have shown to give different response making comparison between the published results even more challenging. An early example was shown by Turner et al. [121], where preferential attachment of a primary cell type and a cell line was observed either on nanostructured silicon ‘grass’ or flat control, respectively. Another apparent conflict has been observed in studies

on surfaces consisting of TiO₂ nanotubes of varying diameter. In 2007 Park et al. published results showing that osteogenic differentiation of rat bone marrow derived MSCs was promoted on small diameter nanotubes (15 nm), while differentiation was inhibited on larger diameter tubes (100 nm) [122]. In contrast, Oh et al. showed that culturing human MSCs on TiO₂ nanotubes with a diameter of 30 nm promoted proliferation, while 100 nm nanotubes selectively differentiated the MSCs along the osteogenic lineage [123]. Several reasons for this dispute have been discussed including the origin (species) of the cells [124, 125].

Lately, much focus has been on stem cell behavior on artificial materials for e.g. stem cell therapy and ultimately production of artificial tissue/organs [63, 126, 127]. For these applications MSCs is an often used adult stem cell source. Topographies on the micrometer scale direct mesenchymal stem cells [128], and it has also been demonstrated that nano-topography as the sole factor is an important cue to induce a specific cellular response, e.g. osteogenic differentiation or proliferation. A range of artificial surface materials has been employed including TiO₂ [123] and polydimethylsiloxane (PDMS) [129]. In general cell types do react to nanotopographies. For example, Teixeira et al. [130] examined human corneal epithelial cells on nano-grooved silicon oxide surfaces. It was observed that the cells aligned along grooves with 70 nm wide ridges and a 400 nm pitch, while the cells were rounded on the flat control.

However, not only the features of the nanoscale topographies play an important role. Also the detailed *arrangement* of nano-features is of importance. It has been shown that, using nanoscale disorder (perturbing a 50 nm offset laterally) of a perfect array of 100 nm deep and 120 nm diameter pits with average pitch spacing of 300 nm, it is possible to stimulate hMSCs towards osteogenic differentiation in vitro [62]. Later it was further demonstrated by the same group that the same nanotopographical features with reduced value of offset and thereby a more “perfect” square lattice resulted in a switch of the hMSC behavior from osteogenic differentiation to a state more suitable for expansion of the hMSCs, still retaining the characteristics of the cell type (hMSC markers and multipotency) [131].

Apart from the structures with long-range order as discussed previously in this section, stochastic nano-scale structures have an impact on cellular behavior as well. For example, silica nanoparticles of different size have been used to create surfaces with varying nano-roughness. Such surfaces have been shown to affect fibroblastic morphology, cell adhesion and spreading for periods of up to 7 weeks [132]. For human endothelial cells, a significant reduction in the adhesion and proliferation was observed as compared to flat control silica surfaces [117].

The complex behavior of cells on topographically structured surfaces has led to a “learning by nature” approach. This route has been used for creating biological materials with functional integration [133] e.g. for creating superhydrophobic and oleophobic surfaces. New directions in research e.g. synthesizing nanoscale topographical features mimicking virus surface topography has been proven successful as well. In a recent study it has been demonstrated that cellular uptake efficacy depends on the nanoscale roughness meaning that the performance of delivery vehicles can be improved by topography [134].

As described above, the influence of topographical cues on cellular behavior is a complex affair involving both biomolecular adsorption and direct mechano-transduction. Furthermore, various cell types respond differently even to the same surface cues and small differences in experimental protocols may lead to very different conclusions. Therefore, more systematic testing is needed and several approaches to implement “high-throughput screening” in biomaterials have been reported the last decade. This is discussed further in Sect. 14.4.5 below.

14.4.3 Mechanical Cues for Guided Cellular Behavior

The types of tissue comprising the human body present a wide range of mechanical properties from the soft brain tissue, with an elastic modulus of approximately 0.2 kPa, to the hard mineralized bone tissue with an elastic modulus in the GPa range [135, 136]. The mechanical properties of bone, in particular, vary greatly with respect to specific anatomical sites, degree of mineralization and also among individuals [136]. Numerous studies have indicated the existence of a correlation between cellular behavior and the mechanical properties of the cell culture substrate. However, the major challenge preventing researchers from obtaining direct evidence of such correlation, was the biochemical complexity of the systems used for the investigations. This complexity did not allow for generating substrates with varying mechanical properties while retaining an identical ligand composition. The study by Pelham and Wang, 1997 was the first that described a culture system allowing for elucidating the direct effect of the mechanical properties. This system was based on polyacrylamide hydrogels which were mechanically modulated by varying the amount of bis-acrylamide cross-linker added to the material during preparation. The acrylamide substrates were subsequently modified by immobilizing type I collagen on the surface by chemical cross-linking. Using this biochemical reductionistic approach, the effect of substrate stiffness on motility and cellular morphology of kidney cells and fibroblasts was investigated. The results for both cell types showed a reduction in the motility of cells cultured on the soft substrates and that these cells were less spread and showed a more ruffled cell morphology, compared to the cells on the more rigid substrates [69]. In a later study, the effect of a mechanical gradient was examined, also using an acrylamide substrate. This study was conducted using fibroblasts and it was found that cells preferentially migrated towards the more rigid area of the substrate. Moreover, the cell spreading area increased as the cell encountered the rigid area of the substrate. Similarly it was found that local deformation of the substrate, either by increasing or decreasing the local tension of the substrate, would either cause the cell to change its direction of migration towards the deformation site in response to increased tension, or away from the site, in the case of decreased substrate tension [137].

Mechanical properties have also been found to affect the differentiation of adult stem cells. In a study by Engler et al., 2006 it was shown that mesenchymal stem cells preferentially differentiated towards the phenotypic lineages that matched the

elastic modulus of the cell culture substrate. This study utilized a system similar to the one developed by Pelham and Wang, 1997 in order to generate cell culture substrates with elastic modulus in the range of brain tissue (0.1–1 kPa (soft)), muscle (8–17 kPa (intermediate)) and collagenous bone (25–40 kPa (stiff)). In this study, the cells cultured on different substrates were found to exhibit morphological characteristics of their *in vivo* lineage counterparts. Thus, soft substrates induced cell morphologies with branched protrusions, intermediate induced spindle like cell morphologies, and stiff substrates induced highly spread cells with polygonal characteristics. These characteristics were already appearing after 4 hours of culture and became increasingly pronounced in the period spanning the culture period. Microarray profiling of cellular mRNA transcripts revealed that the cellular expression of phenotypic markers was also affected by the mechanical properties of the substrate and that this expression followed the morphological appearance of the cells [135].

Engler et al. further investigated the stability of the differentiated cellular phenotypes by treating cells that had been pre-cultured on the mechanical modulated substrates with chemical differentiation factors intended to drive the cells towards alternate phenotypes. These investigations revealed the guiding potency of the flexible substrates, as the cells retained their acquired phenotypic characteristics and the differentiated state was found to be highly stable [135].

Later, a study by Cameron et al., 2011 showed that the cellular response was not only determined by the elastic modulus, but was also influenced by the viscoelastic properties of the culture substrate. In this study polyacrylamide substrates was developed having similar storage moduli but with loss modulus properties spanning from 1 to 130 Pa. This setup allowed for investigating the effect of the materials ability to dissipate the energy through viscous flow, via cellular traction forces, on the behavior of mesenchymal stem cells. Thus, cells cultured on substrates with a high loss modulus continuously experience that the traction forces, exerted by their focal adhesions, would decrease as time progressed. The findings of this study showed that substrates with a high loss modulus (high degree of energy loss) increased cell area and proliferation while the size and maturity of focal adhesions were found to be reduced. These findings indicate that the initial attachment of the cell, to the high loss modulus substrates, allows the cell to exert traction forces to the substrate but that the loss modulus dissipates the energy, thus, forcing the cell to spread out to regain traction. As this process is continuous, focal adhesion development is inhibited which in turn yields immature complexes. The study also found that cells cultured on the high loss modulus substrates showed increased expression of smooth muscle cell markers compared to the cells on the lower loss modulus substrates. Moreover, the use of adipogenic and osteogenic differentiation media, to push the cells towards the specific lineages, was most effective for cells cultured on the high loss modulus substrates. Chemical inhibition of the cells, ability to exert traction forces to the different substrates induced responses from the cells cultured on the low loss modulus substrate that were similar to the response found for cells on the high loss modulus substrates, indicating the importance of cytoskeletal tension forces in relation to guiding cellular responses [70].

Contrary to the chemically modified substrates described above, Fu et al., 2011 has demonstrated the feasibility of generating cell culture substrates where effective elastic modulus was varied purely by geometrical factors. This was achieved by generating PDMS substrates comprising closely spaced pillars of varying heights. In order to ensure that the cells would only adhere to the top of the pillars, these were coated with fibronectin via a micro contact printing approach. In the study, the differentiation of hMSCs was investigated using bi-potent differentiation medium for simultaneous induction of adipogenic and osteogenic differentiation. It was found that the cells cultured on the short pillars, showing higher rigidity, preferentially differentiated towards the osteogenic lineage. On the other hand, preferential differentiation towards the adipogenic lineage was found for the taller, more flexible pillars [72].

The different studies investigating the effect of mechanical properties on cellular behavior has shown the guiding potency of this physical cue, both with respect to regulating cellular migration, proliferation and differentiation. These studies have paved the road for future investigations, thus enabling the scientific community to further elucidate the mechanisms involved in, and the implications of, the phenomenon of cellular mechano-sensing.

14.4.4 Combined and Asymmetrical Cues for Guided Cellular Behavior

While approaches investigating the effect of single guidance parameters on cells in vitro are important for understanding the fundamental processes resulting in specific cellular responses, the synergy of cues presented to cells in vivo necessitates studies designed to elucidate the combined effects of multiple cues. While replicating the exact conditions of most in vivo situations implies that 3D cultures are utilized, such systems are beyond the scope of the current text. There are, however, several studies that have begun to shed light at the interplay between various cues, using two dimensional culture approaches.

The study by Rowlands et al. (2008) has investigated the combined effect of substrate modulus and tissue specific adhesion ECM protein ligands on the differentiation of mesenchymal stem cells (MSCs) [138]. The findings of this study showed that the combination of a flat substrate with elastic modulus in the range of collagenous bone (>34 kPa), in combination with collagen I, favored bone formation relative to the same substrate modified with fibronectin. On the other hand, the use of fibronectin as ligand for substrates with an elastic modulus in the range of muscle tissue (8–17 kPa) favored myogenic differentiation, relative to similar collagen I modified substrates [138]. The study by Lee et al. (2013) investigated the effect of different adhesion ligands and geometrical restraints presented on substrates having an elastic modulus of approximately 0.6 kPa. Cellular differentiation was assessed with respect to the adipogenic and neurogenic lineages. Initially, the

study investigated the individual effects of coating substrates with fibronectin, laminin or collagen, either in small circular patches, limiting cell spreading, or as continuous distributions, allowing the cells to spread freely. Generally, cells confined to the circular patches showed a higher tendency to differentiate along the adipogenic lineage compared to cells that were allowed to spread freely. Moreover, substrates modified with fibronectin were observed to favor adipogenic differentiation while collagen modified substrates generally favored neurogenesis. Taking the study one step further, the group investigated the influence of varying the shape and size of the ligand pattern on which the cells were allowed to attach and spread. More specifically, round, elongated and four-branched star structures were examined. The study found that the star shaped adhesion areas, favored neurogenesis compared to round areas, having the same nominal surface area [139].

The migration of fibroblasts in response to different configurations of mechanically modulated hydrogel gradients, functionalized to yield an opposing collagen I gradient has also been investigated. The substrates used for this study were designed so that the areas showing a high elastic modulus had a low collagen concentration and vice versa. The study found that the fibroblasts preferentially migrated towards increasing ligand concentrations and thus were less prone to be affected by the modulus of the underlying substrate [140].

While the various findings of the studies investigating combined an asymmetrical cues, first of all, highlights the complex interplay of various cues in relation to guiding cells towards a specific lineage, more importantly, it illustrates the complexity of designing and optimizing the biomaterials of the future. Carefully tailoring the interplay between parameters such as mechanical, chemical and structural properties can greatly increase the performance of a biomaterial for specific applications. The complexity of the interplay of the different cues has led to the development of high throughput screening approaches as discussed in the next section.

14.4.5 High Throughput Screening

Understanding the biomaterial-cell interaction is of utmost importance for the development of novel biomedical surfaces with tailored properties for specific applications in e.g. nanomedicine. In the preceding sections different cues from surfaces contributing to the overall guiding of cellular response has been discussed; both chemical/biomolecular, topographical, and mechanical properties has been shown to influence cellular behavior. Already with this span of surface parameters, it is extremely challenging to deduce guidelines. However, the cellular output obtained from these cues are more complex, as also such parameters as cell shape [141], cell colony size [142], and even the combined synergistic effect of the factors above [138] are of importance. Realizing that even looking for a needle in a haystack is simple compared to understanding the interplay between all the factors, additional routes are necessary to speed up the development of biomedical products which can benefit the individual patients.

The biomedical material research approaches have typically either been testing “off-the-shelf” materials in *in vivo* models or using a reductionistic approach focusing on a single parameter in *in vitro* experiments. Both routes have proven valuable and necessary parts of the research field. What would add further value would be a more direct route to obtain design guidelines for materials properties to specifically tailor cellular (and tissue) responses.

In the last decade, a new field in biomaterials has developed, sometimes termed “Materiomics” [143]. An important aspect of the field of Materiomics is high-throughput screening (HTS). Here, the detailed understanding of the cellular mechanisms is less important than the actual cellular response. One of the first demonstrations of the HTS approach was published by Anderson et al. [144]. In this study the influence of chemical composition of polymer mixtures on human embryonic stem cell (hESC) differentiation was addressed: 576 (24×24) combinations of different acrylate, diacrylate, dimethacrylate and triacrylate monomers were deposited in an array in triplicates on a standard glass slide with a non-fouling background. After crosslinking the monomers, the hESCs were seeded on the arrays and it was shown that specific polymer compositions which promote cytokeratin-positive cells could be identified. Since the detailed chemical composition of each individual spot was known, it was possible to produce hit arrays. A follow up study using the same platform addressed human MSC–biomaterial composite interactions [145]. Later studies included a more thorough surface characterizations including water contact-angle, surface topography, surface chemistry and indentation elastic modulus [146].

Screening of topographical structures at the micro and nanometer scale has been addressed by several groups. In an early demonstration, a combinatorial array was created by standard photolithography comprising 169 (13×13) individual microstructures with lateral feature sizes from 1 μm to 6 μm. The approach proved to give topographical guidelines for enhancing osteogenic expression and mineralization using murine pre-osteoblasts [147]. Using the same array, it was also possible to optimize structures promoting mineralization by human dental pulp stem cells [128], human fibroblastic proliferation [63] and enhance non-viral transfection efficiency in primary human fibroblasts [148]. Another approach was developed by de Boer et al., where the topographical structures were designed by mathematical algorithms to obtain nonbiased, random surface features. The resulting array was reproduced on a 2×2 cm² area, called a “TopoChip”, which consisted of poly(lactic acid) with 2176 different topographies [149, 150].

Platforms involving combinatorial mixtures of extracellular matrix (ECM) proteins and soluble growth factors have been developed as well using robotic spotting technology. Such platform has been applied to study the effects of 32 different combinations of five extracellular matrix molecules (collagen I, collagen III, collagen IV, laminin and fibronectin) on cellular expansion/differentiation [151, 152].

It is evident that a 2D surface is far from the typical 3D environment experienced by cells *in vivo*. Thus it would be desirable to be able to do advanced HTS of 3D microenvironments with varying compositions. An attractive route is to use hydrogels mixed with cells. Several groups have tested cellular response inside various

miniaturized cell-laden gels in a combinatorial manner, e.g. for osteogenic differentiation of human mesenchymal stem cells [153]. Recently, a method to create a 3D microarray platform using robotic nanolitre liquid-dispensing technology to create more than 1000 unique microenvironments has been published. In this study the combined effects of matrix elasticity, proteolytic degradability of the surrounding hydrogel and three distinct classes of signaling proteins (extracellular matrix proteins, cell–cell interaction proteins and soluble factors) has been addressed [154].

These newly developed methods to screen and identify surfaces or micro-niche environments with specific cell guiding properties may enable development of novel technologies and products for nanomedicine applications. One area with a huge potential is the expansion of multi- and pluripotent cells on a clinically relevant scale for novel cell-based therapies and tissue engineering.

14.5 Summary and Outlook

In this chapter the different types of cues from surfaces of artificial materials on cellular behavior has been discussed. It has been demonstrated that controlling a number of material parameters is necessary to control the cell-material system e.g. the protein adsorption, surface chemistry, surface topography and mechanical properties and combinations thereof. All these material parameters are of importance to control when designing materials for biological applications at the nanoscale e.g. for nanomedicine applications. The reader might feel that it is an enormous challenge and maybe even an obstacle to address this issue. Furthermore, the whole engineering aspect of transferring the obtained 2D results to real in vivo 3D applications is not straight forward. In some ways this is true. However, one might also view the parameters and the possible synergistic effects as a hitherto almost untapped source of possibilities to control cellular response to, and maybe ultimately even tissue interaction with, artificial materials. Furthermore, even though 2D surfaces are simple models of a real 3D system, all surfaces e.g. at and inside 3D scaffolds are at a certain level 2D. Thus it is expected that the knowledge gained by the examination of parameters in 2D systems may provide guidelines for the construction of 3D systems as well. With the advance of well-defined cell substrates and particularly high-throughput screening of single and multiple parameters the chances are that novel combinations of cues will be identified to optimize cellular/tissue responses and thus be of importance in the field of nanomedicine.

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