

REVIEW ARTICLE

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Extracellular vesicles: from bench to bedside



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Abstract

Extracellular vesicles (EVs) are cell-derived membrane-bound vesicles with heterogeneous contents, including genetic materials, proteins, lipids and small metabolites. The classic EVs are exosomes, which originate from endosomal systems, and microvesicles, which are shed from the plasma membrane. Newly discovered organelle migrasome, once released from cells, adds another player to the EV realm. EVs are present in biological fluids and are important in multiple physiological and pathological processes, including immune regulation and cancer metastasis. Knowledge of EV biology is essential to promote the clinical application of EVs as potential candidates for non-invasive liquid biopsy and drug delivery vehicles. This is a fast-expanding field, but more attention should be paid to the fundamental biology of EVs in order to keep up with the explosive growth of translational needs.

Keywords: Migrasome, Exosome, Extracellular vesicles, Biomarkers, Clinical translation

1 Introduction

Communication from one cell to another, or between a cell and its microenvironment, is crucial in both physiological and pathological conditions. Such crosstalk is achieved by direct cell-cell contact and canonical secretion as well as by membrane-bound extracellular vesicles (EVs). Almost all tested cells are capable of secreting various types of EVs, containing biologically active cargos such as DNA, RNA, protein and metabolites, that vary in response to the microenvironment. Although released externally, these vesicles resemble their cell origin to some extent, and reflect the real-time state of the parent cell. EVs end up in most bodily fluids, including blood, urine, saliva and breast milk. This makes EVs perfect candidates for non-invasive liquid biopsy. By virtue of their cellular origin, EVs have high biocompatibility and low immunogenicity, which means they have potential as drug delivery vehicles and therapeutic reagents. Rapid advances in our understanding of the fundamental

biology of EVs clearly demonstrate the potential of EVs as cancer biomarkers and drug delivery vehicles, while technical advances accelerate progress towards clinical application.

This review aims to discuss the biological properties of the two major EV classes, exosomes and microparticles, as well as newly discovered organelles called migrasomes which are specifically released from migrating cells. We will highlight the utility of EVs for the development of disease diagnostics and therapeutics, which is rooted in basic EV biology and promoted by advanced integrated multidisciplinary technologies.

2 EV biology

2.1 EV classes, biogenesis, and cargos

The secretion of extracellular vesicles was initially described as a means of eliminating unneeded compounds from the cell (Trams et al. 1981). In recent years, EVs have been recognized as membrane-protected vesicles containing bioactive agents which can execute certain functions such as signal communication. Although the classification of EVs is continuously evolving, based on our current knowledge of their biogenesis, they can be broadly divided into two main categories, exosomes and microvesicles. The newly discovered organelles called

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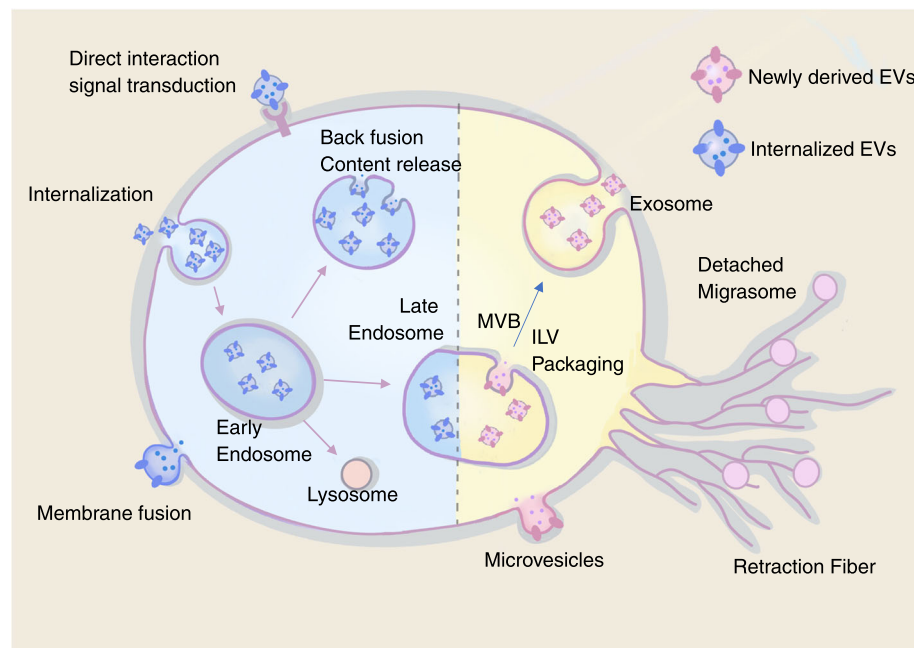


Fig. 1 Features of EVs. EV biogenesis and signal transmission. The right side shows a schematic representation of exosome, microvesicle and migrasome biogenesis. The left side shows signal transmission which can be achieved by direct activation of surface receptors for outside-in signaling, fusion with target cells or complete engulfment of EVs. After uptake by cells, EVs enter early endosomes which subsequently fuse with other organelles in the endocytic compartment. Steps of EV biogenesis are denoted by blue arrows. Steps of EV uptake and cargo release are denoted by purple arrows. "Internalized EVs" can be self-generated (cell-autonomous) or generated by other cells (cell-nonautonomous)

migrasomes, once released extracellularly, become a type of EV (Fig. 1). Our understanding of these vesicles is largely based on evidence provided by transmission and immunoelectron microscopy, fluorescence imaging, genetic approaches and biochemical means. Each EV has unique characteristics, such as size, biogenesis, and biomarkers (Table 1). Exosomes have received more attention than other types of EVs. In recent years, however, research interests have shown some tendency to focus on larger EVs rather than exosomes since their distinct functions and wide range of cargos have gradually been revealed (Lee et al. 2011; Cocucci et al. 2009; Jiao et al. 2021).

Exosomes, which have diameters of 30–150 nm, were initially identified as membrane vesicles released by reticulocytes, an observation subsequently extended to a wide variety of cell types (Johnstone et al. 1987). Exosomes originate from endocytic pathways. The inward budding of endosomal membranes forms intraluminal vesicles (ILVs) in the lumen of a multivesicular body (MVB). ILVs are released as exosomes by fusion of the MVB membrane with the plasma membrane (Harding et al. 1984; Pan et al. 1985). The whole process involves particular sorting machineries, which ensure cargo specificity in exosomes. The discovery of the ESCRT (endosomal sorting complexes required for transport) machinery

Table 1 Key features of EV populations

	Exosomes	Microvesicles	Detached Migrasomes
Size	30-150 nm	100-1000 nm	500 nm-3 μm
Biogenesis	Exocytosis of MVBs originating in endocytic pathways	Budding of the plasma membrane	Generation on the tips or intersections of retraction fibers during cell migration
Isolation	Ultracentrifugation (100,000–200,000 g)/ Sucrose gradient ultracentrifugation (They et al. 2006)	Differential centrifugation (18,000–20,000 g) (Yuana et al. 2011)	Differential centrifugation (20,000 g)/ Iodixanol gradient ultracentrifugation (Chen et al. 2018b)
Detection	TEM, Western blotting, Mass spectrometry, Flow cytometry (bead-coupled)	Western blotting, Flow cytometry (bead-coupled) (Sellam et al. 2009; Gyorgy et al. 2011)	TEM, Western blotting, Fluorescence microscopy (Zhao et al. 2019; Chen et al. 2018b)
Markers	CD63, CD81, CD9, TSG101	Annexin V, tissue factor (Piccin et al. 2007)	Integrins, NDST1, PIGK, EOGT, CPQ (Zhao et al. 2019)

in ILV formation and cargo sorting was the first breakthrough (Hurley, 2008; Tamai et al. 2010; Colombo et al. 2013). There are four different ESCRTs, ESCRT 0, I, II, and III, which act in a stepwise manner (Hurley, 2008; Henne et al. 2011). ESCRT 0 recognizes ubiquitinated proteins, which is a signal for segregation of cargos into ILVs, on the outside of the endosomal membrane (Tamai et al. 2010; Raiborg & Stenmark, 2009). ESCRT I and II are subsequently recruited to initiate and drive the budding of ILV membranes (Shields et al. 2009; Razi & Futter, 2006; Katzmann et al. 2001). In turn, ESCRT III is recruited for the scission of the ILVs into the MVB lumen (Lin et al. 2005). This process is finalized by removal of the ubiquitin tag from the cargo proteins via recruitment of a deubiquitinating enzyme (McCullough et al. 2006; Agromayor & Martin-Serrano, 2006; Ma et al. 2007), and disassembly of ESCRT-III by the ATPase Vps4 (Stuchell-Brereton et al. 2007; Azmi et al. 2006). An ESCRT-independent pathway also exists, in which ceramide mediates membrane deformation in ILV and MVB formation (Trajkovic et al. 2008). Tetraspanin family proteins regulate the cargo sorting for exosomes (Theos et al. 2006; van Niel et al. 2011). These tetraspanin proteins form clusters and dynamic membrane macrodomains that mediate budding (Charrin et al. 2014). Additional mechanisms, for example co-sorting of chaperones or proteins with high affinity for certain kinds of lipids, also contribute to sorting. Apart from proteins, RNAs are important cargos of exosomes (Geminard et al. 2004). Sorting of RNAs is achieved by sorting of RNA-binding proteins with specific motifs, which bind and sort RNAs depending on their sequence (Villarroya-Beltri et al. 2013). The sorting machinery and its regulation is undoubtedly important and our knowledge is far from complete. The exosome proteomic database, available from the “ExoCarta” website, shows that exosome contents are highly heterogeneous between different cell types and body fluids. The nature and abundance of exosome cargos are influenced by the physiological or pathological state of the donor cell. How these stimuli regulate the cargo sorting machinery in order to achieve the highly heterogeneous features of exosomes is largely unknown. Endosome dynamics and sorting machineries should be considered when investigating exosomes as biomarkers or when manipulating exosomes.

Microvesicles, initially known as “platelet dust”, were first identified as subcellular material originating from platelets and have long been studied mainly for their role in blood coagulation (Wolf, 1967; Sims et al. 1988; Satta et al. 1994). Microvesicles range from 500 nm to several microns in diameter. They are formed by outward budding of vesicles from the plasma membrane (Tricarico et al. 2017). Generation of microvesicles requires

membrane lipid rearrangement and plasma membrane deformation, in which cytoskeleton elements are widely involved. Sorting of cargos into microvesicles is achieved through plasma membrane anchoring (Al-Nedawi et al. 2008). Microvesicles have received much less attention than exosomes and the detailed mechanisms of their biogenesis remain to be elucidated.

Migrasomes, first discovered in 2015, are micron-scale vesicles which contain numerous small internal vesicles. During cell migration, migrasomes grow on the tips or intersections of retraction fibers, which mark the path of cell migration on the extracellular matrix (ECM) (Ma et al. 2015). Migrasomes remain active communications with cell body before being detached from retraction fibers and released as extracellular vesicles. Migrasome formation requires cell migration. Thus, the first study to document the physiological function of migrasomes was in zebrafish embryonic development, which involves massive cell migration (Jiang et al. 2019). Migrasomes, enriched in a combination of ligands including chemokines, morphogens and growth factors, deliver signaling cues and function in organ morphogenesis and positioning (Jiang et al. 2019). Detached migrasomes have also been identified in multiple body fluids, expanding the range of their potential functional scenarios (Zhao et al. 2019). Unlike other extracellular vesicles, migrasomes maintain contact with the parent cells through retraction fibers for quite a long period of time before their eventual extracellular release. Large-scale plasma membrane deformation is observed in migrasome formation. Tetraspanin family proteins, especially TSPAN4, together with cholesterol, form micron-scale membrane macrodomains to accomplish the membrane shaping for retraction fiber extension and migrasome biogenesis (Zhang et al. 2020; Huang et al. 2019). Integrins play a role in providing the force that adheres migrasomes to the ECM (Wu et al. 2017). Thus, TSPAN4 and integrins are markers of migrasomes. Migrasomes are a medium for intercellular lateral transfer of RNAs and proteins. Interestingly, distinct from exosomes, migrasomes contain membrane-bound organelles and large amounts of mRNA derived from the parent cell (Zhu et al. 2021). Recently, Yu’s group showed that damaged mitochondria in migrating cells are transported into migrasomes and subsequently disposed of. This process, known as mitocytosis, requires the damaged mitochondria to be positioned at the cell periphery for sorting into migrasomes. Mitocytosis not only maintains mitochondrial quality for cell viability, but is also likely to be involved in sending out mitochondrial stress information (Jiao et al. 2021). At present, it is largely unknown whether other membrane-bound organelles and mRNAs are sorted into migrasomes. This is being actively explored.

2.2 EV destinations

Once released into the microenvironment, EVs reach the recipient cells and trigger functional responses or promote phenotypic changes. Signal transmission can be achieved by direct activation of surface receptors for outside-in signaling, fusion with target cells or complete engulfment of EVs, all of which require vesicle recognition and docking (Fig. 1). The specificity is determined by recognition between ligands enriched at the surface of EVs and receptors on the plasma membrane of the recipient cells, which depend on the origin and subpopulation of EVs and on the identity of the recipient cells. In terms of specific enrichment of surface molecules, plasma membrane-derived EVs might be different from endocytic pathway-derived EVs. So far, several ligand-receptor pairs have been reported. Protein-glycan interactions include lectins and glycan moieties (Hao et al. 2007; Saunderson et al. 2014; Barres et al. 2010; Shimoda et al. 2017), and heparan sulfate proteoglycans and fibronectin (Christianson et al. 2013). Lipid-protein recognition can occur between phosphatidylserine (PS) that is flipped to the outside of the membrane and T cell immunoglobulin mucin (TIM) family proteins (Sims et al. 2017). Adhesion molecules include tetraspanins, integrins, ICAM-1, ECM components, etc (Hao et al. 2007; Yuan et al. 2017). Specifically, the interaction of integrins with ECM proteins, mostly fibronectin and laminin, has been shown to have important roles in ensuring that exosomes interact with the right recipient (Purushothaman et al. 2016). In *in vivo* situations, integrin heterodimers drive EVs towards specific target organs, possibly through the cell-associated ECM.

EVs can directly activate cell surface receptors via protein and bioactive lipid ligands (Polgar et al. 2005). EVs can also deliver contents through membrane fusion with the plasma membrane. In addition, EVs can be internalized, which requires release of their contents inside the target cells to actually elicit a downstream response. The destiny of internalized EVs may be lysosome degradation to provide metabolites. For EVs with a signaling mission, their contents should retain their original function within the recipient cell. This functional competence is supported by mounting evidence showing targeted conversion of recipient cells. Interestingly, EVs can release their contents into the cytosol of the parent cell through back fusion with the MVB membrane (Bissig & Gruenberg, 2014) (Fig. 1).

2.3 EVs in immune modulation

The role of EVs in diseases, the onset and progression of which require cell-cell signal transduction, has been widely documented. The study of exosomes in immune

responses and tumors has progressed at a rapid pace compared with other functions in embryonic development, neurodegeneration, metabolic diseases and cardiovascular diseases. Regardless of the system or disease being studied, the same basic cell biology questions should be answered, such as uncovering the identities of the donor and recipient cells, the candidate molecules, the sorting machinery and the functional response of recipient cells. For example, regulation of the immune response by exosomes is accomplished through direct antigen presentation by MHC molecules on the exosome surface. Exosomes derived from B lymphocytes induce antigen-specific MHC class II-restricted T cell responses, suggesting a role for exosomes in antigen presentation *in vivo* (Raposo et al. 1996). Exosomes derived from ovalbumin (OVA)-treated dendritic cells elicited OVA-specific CD8⁺ T cell activation (Wahlund et al. 2017). This sets up a foundation for studying exosomes as cancer vaccines. Modulation of immune responses by exosomes also involves presentation of immunoregulatory surface molecules on exosomes. Presentation of immunoregulatory molecules such as PD-L1 (programmed cell death ligand 1) on the exosome surface contributes to suppression of T cells and promotion of tumor growth (Chen et al. 2018a; Poggio et al. 2019). Exosomes may also regulate the immune response by influencing gene expression and signaling pathways in recipient cells, principally by the transfer of microRNAs (miRNAs) and proteins, which is well documented in other reviews. Exosomes can either eliminate or promote innate or adaptive immune responses under different pathological conditions (Robbins & Morelli, 2014).

2.4 EVs in cancer

Exosomes have been associated with neoplasia, tumor growth and metastasis, the hallmark features of cancer. Neoplastic reprogramming and tumor growth can be promoted by exosome cargos including miRNAs, as well as HRas and Kras mRNAs (Abd Elmageed et al. 2014). Another study showed that cancer cell exosomes induced random mutations and initiated malignant cell transformation, and the transformed cells could form tumors *in vivo* (Stefanius et al. 2019). Metastatic ability can be transferred through exosomes to promote the epithelial-to-mesenchymal transition (EMT) of poorly metastatic cells (Le et al. 2014).

Exosomes are detected in the tumor microenvironment, which is composed of cancer cells, stromal cells and stromal elements such as ECM components. Modulation of the immune response by tumor-derived exosomes involves a prometastatic inflammatory response (Le et al. 2014), which influences dendritic cell maturation and immunosuppression. The reciprocal exchange

of exosomes between the stroma and cancer cells has been reported to enhance fitness for tumor growth. This exosomal exchange results in metabolic changes, PTEN suppression or autocrine Wnt-PCP signaling through transfer of metabolites, miRNAs or proteins (Zhang et al. 2015; Luga et al. 2012). Exosomes have been implicated in the angiogenic and ECM remodelling of the tumor microenvironment. In this context, suppression of endothelial tight junction proteins, reduced integrity of blood vessel endothelial cells, and degradation of the ECM through matrix metalloproteases (MMPs) are involved in promoting tumor growth and dissemination (Zhou et al. 2014; Kucharczyk et al. 2013; Genschmer et al. 2019; Yokoi et al. 2017).

Apart from creating their own in situ microenvironment for tumorigenesis, tumors induce formation of microenvironments in distant organs to support the survival and outgrowth of tumor cells even before colonization. These distant microenvironments are named pre-metastatic niches (PMNs) (Kaplan et al. 2005). When mice were pre-injected with exosomes from highly metastatic melanoma cells, they developed many more metastatic tumors than mice injected with exosomes from weakly metastatic melanoma cells. This proves that EVs carry important factors for PMN formation (Peinado et al. 2012). Follow-up studies showed that EVs induce a series of events during PMN formation to create a supportive microenvironment for metastasis, including vascular leakiness, attracting and educating bone marrow-derived cells (BMDCs), and reprogramming of local resident cells to induce proinflammatory responses and metabolic fitness (Costa-Silva et al. 2015). Moreover, the integrin pairs on exosomes predict the site of tumor metastasis (Hoshino et al. 2015). More interestingly, at the very early stage of tumorigenesis, tumor cells have the potential to metastasize by dormancy in pre-formed PMNs before undergoing activation and growth. This explains why some tumors produce metastases long after the primary tumor is removed (Peinado et al. 2017). Although there is still some debate about the role of tumor-derived exosomes in shaping PMNs, there is hope that this feature of exosomes could be used to predict whether a patient's cancer will metastasize.

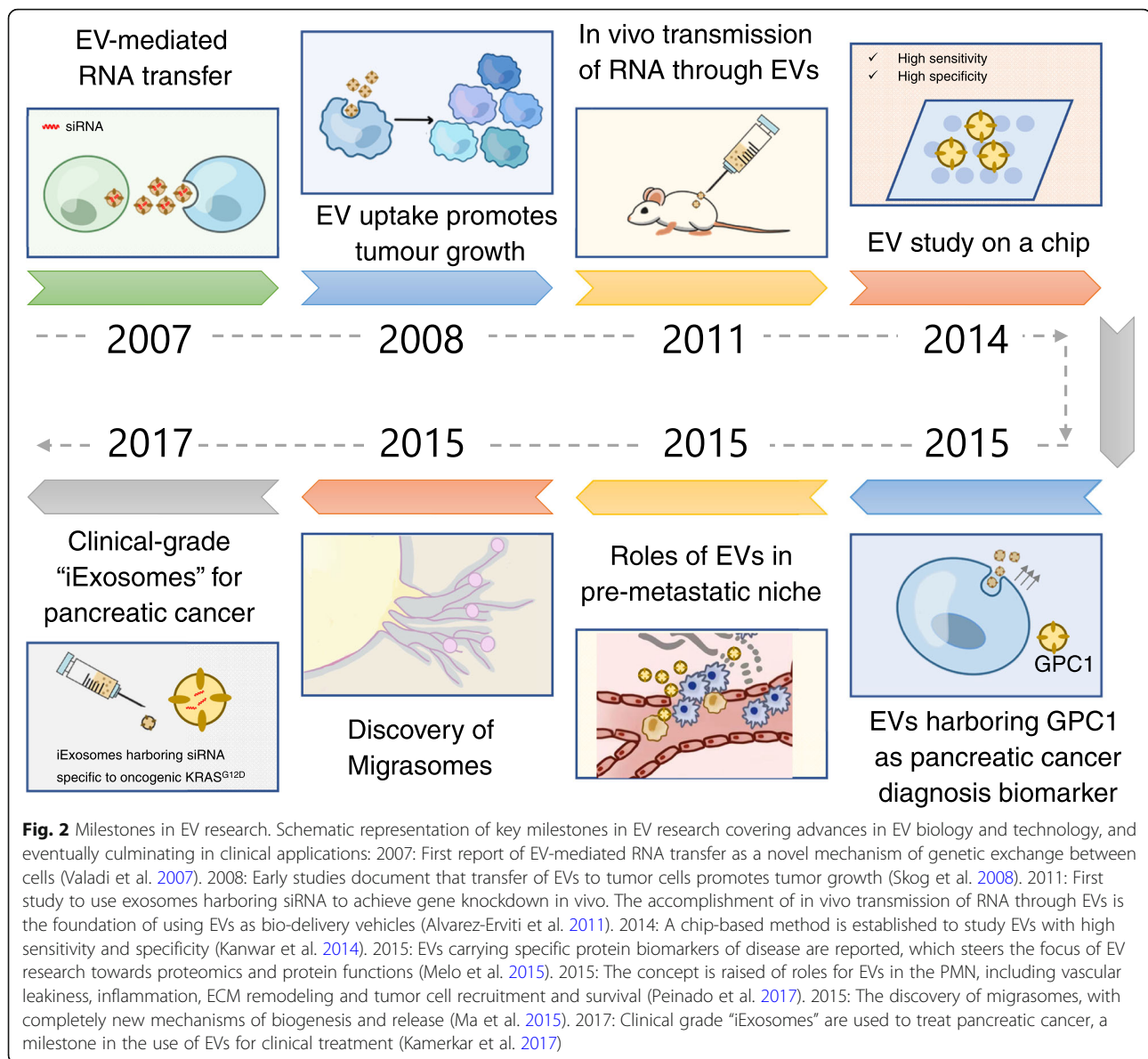
2.5 Techniques for studying EV biology

Progress in the EV field requires understanding of the underlying fundamental cell biology mechanisms. Advanced biochemistry approaches, imaging methods and omics techniques have been very important in advancing the EV field. High-throughput, large-scale knockdown or knockout screening has greatly accelerated the discovery of the protein machinery network. Super-resolution omics makes identification and quantification of EV contents increasingly accurate. Breakthroughs in live-imaging

methods have improved the temporal and spatial resolution when analyzing cultured cells or animal models. These advances will surely aid in the understanding of the mechanisms and function of EVs (Wu et al. 2021).

3 Research milestones in the development of exosomes for clinical translation

After they were first discovered in 1980, EVs were largely dismissed for a couple of decades as cell debris or “waste carriers” of the cell. A series of landmark events both in cell biology and technology development support the potential of EVs as cancer biomarkers (Fig. 2). Jan Lötvald's group was the first to report that exosomes could serve as a novel mechanism of genetic exchange between cells, through the transfer of functional RNA molecules, including mRNA and miRNA (Valadi et al. 2007). Johan Skog et al. reported that exosomes specifically promote tumor growth through the transport of mRNA, miRNA and angiogenic proteins. These tumor-derived exosomes have the potential to provide diagnostic information through a blood test (Skog et al. 2008). This was followed by the work of Lydia Alvarez-Erviti et al. who were the first to deliver siRNA with targeted exosomes in vivo (Alvarez-Erviti et al. 2011). Another important study introduced proteins into the diagnostic story. Glypican-1 (GPC1), a cell surface proteoglycan, was identified to be specifically enriched in exosomes harvested from the blood of patients with pancreatic cancer. The level of GPC1 distinguished benign pancreatic disease from pancreatic cancer with high sensitivity and specificity (Melo et al. 2015). The role of exosomes in pre-metastatic niche formation, and the organ-seeking nature of exosomes through integrin zip coding, may make it possible to predict whether and where metastasis will occur (Costa-Silva et al. 2015; Hoshino et al. 2015). Recently, the discovery of migrasomes, which are formed and released via completely new mechanisms, has steered the EV field in broader directions (Jiao et al. 2021; Ma et al. 2015). The integration of microfluidics and various sensors has made it possible to isolate and detect EVs on a chip, with significantly improved accuracy and sensitivity (Kamwar et al. 2014). Simultaneously, exosomes have been explored for delivery of a miRNA or siRNA payload to facilitate anticancer treatment. Clinical-grade mesenchymal stem cell (MSC)-derived exosomes harbouring KrasG12D siRNA (iExosomes) have been used to treat pancreatic cancer in multiple animal models (Kamerkar et al. 2017). It is interesting to see from these milestones that the interplay between basic scientific discoveries and technological developments has led to huge advances in the EV field. Though there are still many debated ideas and technical limitations in the newly emerging areas of the EV field, it is clear that EVs are implicated in many



facets of disease development and progression, and therefore they are ideal candidates as biomarkers and/or therapeutic tools.

4 EVs as diagnostic biomarkers

4.1 EVs as the target of liquid biopsy

Nowadays, the analysis of cancer characteristics is central to patient management and treatment decisions. Liquid biopsy provides the opportunity of detecting, analyzing and monitoring cancer in various body fluids such as blood or urine instead of a fragment of cancer tissue. EVs have many advantages as targets of non-invasive liquid biopsy. The biggest advantage of EV analysis over other blood-based liquid biopsy targets such as circulating tumor cells (CTCs) is the access to a larger

population of biomarkers in EVs. EVs harvested from a patient's blood contain enough materials to analyze and use as a diagnostic indicator. The heterogeneity of EV cargos in various disease conditions provides an opportunity to establish EV-related disease specificity. Tumors contain a heterogeneous mix of cancer cells which generate different cancer cell clones within tumor tissues. Thus, tissue biopsy might not be able to provide an accurate landscape of the entire tumor. EVs from patients' blood offer comprehensive information about tumor heterogeneity. In addition to genetic information, analysis of EVs can also be used to monitor proteins and metabolites associated with the tumor state. Longitudinal sampling to monitor disease progression is informative and much easier to carry out, since repeated

sampling of body fluids is more acceptable than repeated biopsies.

4.2 Technologies used for analysis of EV biomarkers

Techniques for isolating and characterizing EVs are the key to connecting the mounting evidence for EV cargos in different diseases with clinical applications using EVs as diagnostic biomarkers. Conventional methods to isolate EVs from plasma or other samples include ultracentrifugation, density-gradient centrifugation, filtration, polymer-based precipitation, size exclusion chromatography (SEC) and immunoaffinity purification (Doyle & Wang, 2019; Coumans et al. 2017). To increase the purity of EVs, immunoaffinity purification can be performed using tissue-specific exosome markers to enable extraction of subgroups of EVs. This allows tracing of EV origins and avoids contamination from other EV sources. Such a strategy has been actively adapted for purification of brain-derived exosome due to the presence of brain tissue-specific exosome markers (Shi et al. 2014; Yu et al. 2020; Deng & Miller, 2019). Other strategies include 2-step purification with ultracentrifugation followed by immunoaffinity purification to further increase the purity (Shurtleff et al. 2016). In summary, the performance of different exosome isolation methods varies significantly due to contamination, poor yield, intensive labor, low throughput or high equipment demands, and there are various trade-offs between parameters. There are several different characterization techniques. Analysis of morphological features relies on electron microscopy (EM), fluorescence imaging and nanoparticle tracking analysis (NTA). Targeted detection of either RNAs or proteins relies on PCR- or antibody-based detection. Most recently, unbiased multi-omics approaches have been applied to investigate the molecular composition of EVs. These approaches combine next-generation sequencing, metabolomics and mass spectrometry-based proteomics. Advanced super-resolution detection methods demand high quality samples. These methods have limited tolerance of contamination and protein/chemical labelling, and they require small sample volumes and fairly high enrichment. Low throughputs are another concern. Thus, there is an urgent need to develop advanced isolation technologies compatible with these increasingly refined analysis tools. The introduction of microfluidics has facilitated chip-based isolation procedures, a big step forward which will significantly improve the accuracy of exosome isolation from biological samples (Vaidyanathan et al. 2018). Microfluidic isolation methods are typically rapid and efficient and require small starting volumes. They allow for the development of innovative separation mechanisms based on the acoustic, electrophoretic, and electromagnetic properties of the exosomal vesicles. Most recently, in 2021,

Liu's group designed a device to impose periodic negative pressure oscillations (NPOs) on a nanoporous membrane to achieve separation of label-free exosomes from small particles (free proteins or nucleic acids) (Chen et al. 2021). The isolation methods are compatible with downstream omics studies which provide deep and unbiased characterization of EV cargos. This is a valuable resource for identifying biomarkers.

4.3 Challenges and opportunities in the clinical translation of EVs as biomarkers

Though much of the clinical data suggests important links between EVs and diseases, these studies are at an early stage and the results are only correlative. Fundamental questions about the biological mechanisms underlying exosome structure and function remain unanswered, although landmark discoveries have been made. Further investigations are required to fully resolve the functional capabilities of these vesicles.

The function of EV RNA cargos has received more attention than protein cargos, due to the fact that proteomics techniques lag behind sequencing techniques in terms of both depth and throughput. However, protein cargos have equally important functions and should receive greater focus. Recently, there have been multiple attempts using proteomics approaches to revisit the cargos of exosomes from both cell culture and clinical samples (Shi et al. 2014; Hoshino et al. 2020; Jeppesen et al. 2019; Kugeratski et al. 2021). These studies for the first time involved samples from large clinical cohorts to reveal the high level of heterogeneity in clinical samples and to shed some light on EV proteins which are consistently and significantly enriched under disease conditions. In addition, the proteins present on the surface of EVs provide invaluable information about the physiological states of the parental cells of EVs. A multicomponent, combinatorial approach using a combination of EV markers will provide the most accurate indication of disease. At present it is challenging to isolate desired EV subpopulations with high purity from the mixture of EVs in body fluids. There is a clear need to establish better exosomal markers for this purpose. Therefore, the EV surface proteome should be put on the agenda.

Large-scale unbiased discovery of exosome biomarkers requires high quality clinical cohorts and standardized isolation and characterization protocols. Currently, variation in isolation strategies and analytical techniques has made multi-centre integrated analysis difficult, especially for proteomic analysis. This adds another drawback to these already low-throughput techniques.

Regarding clinical application scenarios, most of the isolation approaches are still not compatible with clinical analysis due to issues of scalability, standardization and validation. Furthermore, several of the approaches are

time-consuming and require large sample amounts and extensive pre-treatment steps.

5 EVs as therapeutic agents and bio-delivery vehicles

5.1 EVs as therapeutic agents

Targeting extracellular vesicles to exploit their innate therapeutic potential or using them for drug delivery are important emerging strategies for therapy. Due to their bioactive cargos, EVs on their own have therapeutic potential. In myocardial infarction or other models of injury, it is confirmed that exosomes are responsible for the beneficial outcomes of transplanted MSCs rather than the MSCs themselves (Bruno et al. 2009; Lai et al. 2010). This pioneering discovery boosted research into the application of EVs derived from MSCs in driving tissue regeneration. The success of therapy based on MSC-derived exosomes is reflected by the growing number of ongoing clinical trials for various diseases (Zipkin, 2020).

5.2 EVs as bio-delivery vehicles

Exosomes derived from B cells present antigens and stimulate T cells *in vivo*. Dendritic cells (DCs) secrete exosomes expressing MHC-I/II molecules and T cell costimulatory molecules, which suppress tumor growth depending on T cell function (Raposo et al. 1996; Zitvogel et al. 1998). Although it is not exactly clear whether the antigen presentation by exosomes is direct or indirect *in vivo*, these milestone discoveries promote the investigation of exosomes as immune modulators (Fig. 3). The resulting immune activation or suppression can be exploited in different disease scenarios like cancer or autoimmune diseases.

A study in 2007 was the first to show that exosomes facilitate lateral RNA transfer between cells (Valadi et al. 2007). Later, another study successfully delivered siRNA to mouse brain by injection of exosomes (Alvarez-Erviti et al. 2011). This was the first time a desired cargo was loaded into self-generated exosomes to achieve delivery

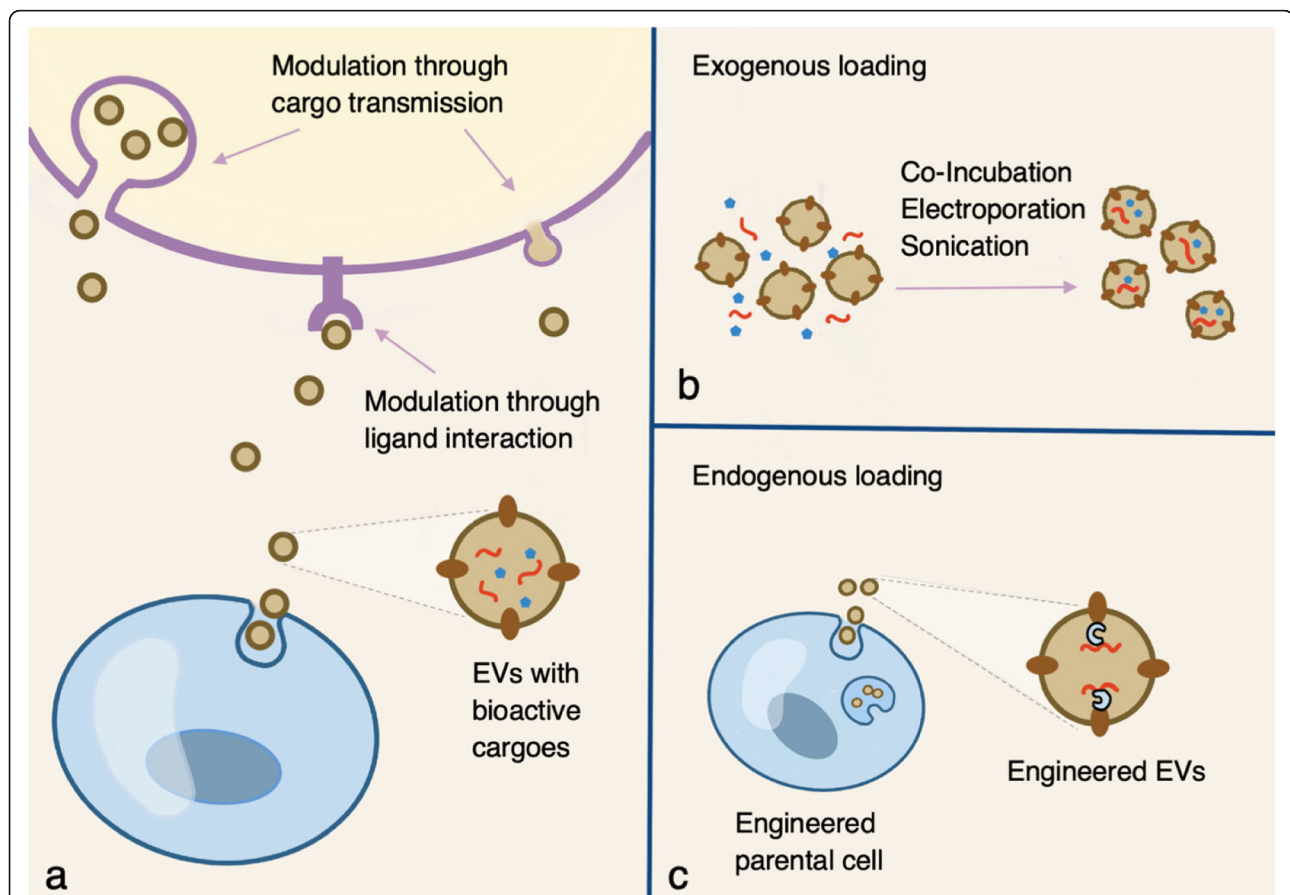


Fig. 3 EVs as therapeutic agents. Exploitation of EVs as therapeutic agents. **a.** EV-based therapeutics can be achieved by direct activation of surface ligands or by transfer of growth factors, soluble proteins, bioactive lipids and genetic materials. **b., c.** Extracellular vesicles can be engineered to contain certain contents or to present ligands on the surface by exogenous (**b.**) or endogenous (**c.**) loading. **b.** Post-purification EVs are exogenously loaded with the purified cargo of interest. The resulting EVs retain their original cargo. **c.** EV-producing cells are modified to overexpress fusion proteins in which a targeted cargo is joined with a canonical EV membrane protein

in vivo. The inherent ability of EVs to cross biological barriers, even the blood-brain barrier, makes them more attractive than other delivery mediators (Zhuang et al. 2011). These pioneering studies established the foundation of exosomes as bio-delivery vesicles and promoted investigations into clinical-grade exosomes derived from stem cells (Fig. 3).

There are two main strategies to exploit extracellular vesicles for cargo delivery, endogenous loading or exogenous loading (Fig. 3). Endogenous loading can be achieved by engineering the parental cells so that they express a fusion protein consisting of a targeted cargo protein joined to a canonical EV membrane protein. For exogenous loading, EVs are first isolated and then loaded with a purified RNA of interest through co-incubation, sonication or electroporation. The resulting EVs retain their original cargo (Zickler & El Andaloussi, 2020). It should be noted that other intracellular components will also be actively or passively loaded into the engineered EVs, which may influence the therapeutic efficacy. Importantly, engineering of parental cells and post-purification loading can be used together, thereby taking advantage of both specific biodistribution and increased anti-tumor efficacy. Other than that, EV surface proteins can be modified by optical tags and radioactive isotope labelling for monitoring purposes (Salunkhe, 2020).

5.3 Challenges

Although multiple discoveries have paved the way to potential clinical applications of EVs, the clinical translation of EVs is still at a very early stage and major challenges need to be overcome. For example, MSC-derived exosomes have been shown to both inhibit and promote tumor growth (Zhu et al. 2012; Bruno et al. 2013). Such discrepancies are probably a consequence of uncertainty about which EV cargos have bioactive effects, variability in the cell culture conditions which produce EVs, unstandardized purification protocols, and a lack of EV quality control markers. Therefore, suitable cell sources for clinical-grade EV production are urgently needed to increase yield, purity and safety. In addition, standardized protocols for EV purification should be strictly established. Last but not least, knowledge of the fundamental biology of EVs should keep up with the explosive growth of translational needs.

6 Conclusions and perspectives

Much progress has been made in recent years in understanding the basic biology of extracellular vesicles. However, despite the enormous therapeutic potential of EVs, this field still needs a systematic understanding of the basic mechanisms involved in EV biogenesis, cargo sorting and release. Over the past decades, scientists have

characterized the involvement of endocytic pathways in EV generation. The subsequent explorations of EV function in immune regulation or tumor metastasis and EV engineering are mostly based on endocytic pathways. However, the potential of EVs should not be limited only to one certain pathway. The discovery of migrasomes has steered the field towards broader applications. Migrasomes remain strong connection with cell body as cellular organelles before being released extracellularly as a type of EV. The unique biogenesis mechanisms of migrasomes allows cells to release a wider range of molecules, which brings new opportunities to exploit EVs as biomarkers, therapeutic agents and bio-delivery vehicles. The discovery of cellular organelles in migrasomes, which function in organelle quality control and reflect cellular stress conditions, provides another exciting example of EV functions beyond our current knowledge of exosomes.

Advanced technologies like imaging methods or omics methods should be developed for deep analysis of small amounts of EVs or even individual EVs. The in-depth identification, accuracy and reproducibility of quantification achieved by advanced omics techniques will guarantee new discoveries about the basic mechanisms of EV biogenesis and function, and will promote the clinical translation of EVs.

The EV field is highly interdisciplinary. Cell biologists, physicians and engineers should work together on the basic functions of extracellular vesicles and on their translation from the bench to the bedside.

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