

Sustainable Agriculture Reviews 54

Vinod Kumar Yata
Ashok Kumar Mohanty
Eric Lichtfouse *Editors*

Sustainable Agriculture Reviews 54

Animal Biotechnology for Livestock
Production 1

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Vinod Kumar Yata 
Animal Biotechnology Centre
National Dairy Research Institute
Karnal, Haryana, India

Ashok Kumar Mohanty
Animal Biotechnology Centre
National Dairy Research Institute
Karnal, Haryana, India

Eric Lichtfouse 
Aix-Marseille University
CNRS, IRD, INRAE
Coll France, CEREGE
Aix-en-Provence, France

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Preface

Many developed countries are actually facing a ban of animal food products, promoted mainly by urban and vegetarian activists who have never experienced living in farms and who do not acknowledge that their modern way of life is largely the result of hard work of their elders with the help of farm animals. However, since the start of society, animal production has been an essential agricultural sector worldwide providing food, labor, aesthetics and social values, and even today many farmers would not survive without animals. This book entitled ‘Animal Biotechnology for Livestock Production 1’ is our first volume providing advanced knowledge on biotechnological methods to improve the livestock production, with focus on animal reproduction, health, diagnosis and nutrition. Chapter 1 presents on artificial insemination in cattle, with focus on physiology aspects of the estrous cycle, estrus synchronization program, ovulation synchronization program for timed artificial insemination, strategies for improving fertility and use of sexed semen in artificial insemination. Chapter 2 reviews biotechnological applications for production of dromedary camels, with details on camel herd reproduction, reproduction control and artificial insemination. Sperm dilution, thawing, conservation, and insemination techniques are also discussed. Recent biotechnological applications for livestock production are summarized in Chap. 3, with emphasis on somatic cell nuclear transfer, artificial insemination, embryo transfer, embryonic stem cell technology and marker assisted selection.



Cattle production in France. Copyright 2021 Eric Lichtfouse

Chapter 4 reviews applications of stem cells in livestock, with emphasis on mesenchymal stem cells. Immunomodulatory, antimicrobial activity, migration and reparative functions of stem cells are detailed. Chapter 5 presents techniques for profiling proteins and metabolites associated with feed efficiency in dairy cattle. Recent findings on key metabolites and proteins of metabolic pathways are also disclosed. Chapter 6 focuses on processing, packaging, and safety of dairy products. Applications of biotechnologies in food diagnosis are also explained. Chapter 7 reviews ‘on-farm point-of-care’ diagnostic technologies in animals. This chapter covers various point-of-care and on-farm diagnostic technologies for monitoring animal health and disease with focus on molecular, electrochemical-biosensors diagnostics. Chapter 8 presents biotechnological applications in the poultry industry. This chapter covers the concepts and developments of biotechnologies for poultry production, breeding, feed and nutrition. This chapter also discusses applications in poultry vaccines, biologics, disease diagnosis and food processing.

We express our thanks to all authors who have contributed high quality chapters. Our special thanks are due to the Indian Council of Agricultural Research (ICAR), the Government of India and the Director of the ICAR National Dairy Research Institute (NDRI), Karnal, India for providing the institutional support. We would like to acknowledge Dr. Sudarshan Kumar, Scientist, ICAR-NDRI, Karnal, India for his help in choosing contributors and reviewers. We would like extend our thanks to the staff of Springer Nature, for their generous assistance, constant support, and patience in initializing and publication of this book. We acknowledge our thanks to

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Karnal, India

Vinod Kumar Yata

Karnal, India

Ashok Kumar Mohanty

Aix-en-Provence, France

Eric Lichtfouse

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About the Editors



Vinod Kumar Yata is an interdisciplinary researcher working at National Dairy Research Institute, Karnal, India. Previously, he worked as an Assistant professor at the Department of Biotechnology, Dr. B R Ambedkar National Institute of Technology Jalandhar, Punjab, India. He received his Ph.D. in Biotechnology from Indian Institute of Technology Guwahati. He specializes in interdisciplinary research which includes Nanotechnology, Microfluidics, Animal Biotechnology, Cancer biology and Bioinformatics. He has developed a microfluidic device for the separation of live and motile spermatozoa from cattle semen samples. He opened up a new avenue to prodrug enzyme therapy by introducing the nanocarriers for the delivery of non-mammalian prodrug activating enzymes. He elucidated the structural features and binding interactions of several bio molecules by *in silico* methods. He has published three books as an editor and one book as an author with Springer Nature publisher. He has published several research papers in peer reviewed international journals and presented papers in several international conferences.



Ashok Kumar Mohanty is a principal scientist at Animal Biotechnology Centre, ICAR-National Dairy Research Institute, Karnal, India. His group is involved in various basic and applied research related to animal production systems. His research group has made pioneering contributions in the field of Animal Biotechnology, with emphasis on Gene cloning, Expression and Functional characterization of Animal Proteins, Proteomics in Animal Production, Cell and Molecular Biology and Structural Biology of Proteins. His group has developed a Buffalo Mammary Epithelial cell line for the first time which can be used a model system to understand lactation biology in animal as well as human. His team has also developed a pregnancy diagnostic kit for the early detection of pregnancy cattle and buffalo. His group is also extensively involved in developing low cost technology for semen sexing in cattle. He has organized a number of national and international workshops and international conferences. He is a recipient of several awards which includes, DBT Overseas Associateship by Ministry of Science & Technology, Govt. of India., Jawaharlal Nehru Award (gold medal) by Indian Council of Agricultural Research (ICAR), New Delhi for outstanding postgraduate research in the field of Animal Biotechnology and Young Scientist Award sponsored by International union of Crystallography (IUCr) and Dept. of Science and Technology (DST), Govt. of India for attending IUCr congress at Geneva, Switzerland. He is the Fellow of National Academy of Dairy Sciences, India and executive member of proteomics society of India and associate fellow of National Academy of Veterinary Science, India. He has supervised more than 50 graduate, PhD students and post docs. He published more than 200 peer reviewed research and review papers. He has also authored 8 book chapters in the areas of animal and food biotechnology published by in national and international publishers.



Eric Lichtfouse is a professor of environmental sciences and scientific writing at Aix Marseille University and Xi'an Jiaotong University. He has invented carbon-13 dating, a molecular-level method allowing to study the dynamics of organic compounds in temporal pools of complex environmental media. He has discovered temporal pools of individual substances in complex media such as soils. He is Chief Editor and founder of the journal *Environmental Chemistry Letters*, and the book series *Sustainable Agriculture Reviews* and *Environmental Chemistry for a Sustainable World*. He is the author of the book *Scientific Writing for Impact Factor Journals*, which includes an innovative writing tool: the micro-article. He has awards in analytical chemistry and scientific editing.

Contributors

Alzahraa M. Abdelatty Department of Nutrition and Clinical Nutrition, Faculty of Veterinary Medicine, Cairo University, Giza, Egypt

Mohamed Abdelmegeid Department of Animal Medicine, College of Veterinary Medicine, Kafr-Elsheikh University, Kafrelsheikh, Egypt

Abdulrahman S. Alharthi Department of Animal Production, College of Food and Agriculture Sciences, King Saud University, Riyadh, Saudi Arabia

Kiran Ambatipudi Department of Biosciences and Bioengineering, Indian Institute of Technology Roorkee, Roorkee, Uttarakhand, India

Mohammed H. Bakr Animal Production Department, Faculty of Agriculture, Cairo University, Giza, Egypt

Vinay Bhaskar Animal Biotechnology Centre, National Dairy Research Institute, Karnal, Haryana, India

Dixita Chettri Department of Microbiology, Sikkim University, Gangtok, Sikkim, India

Mona M. M. Y. Elghandour Facultad de Medicina Veterinaria y Zootecnia, Universidad Autonoma del Estado de Mexico, Toluca, Mexico

Ahmed A. Elolimy Department of Animal Production, National Research Centre, Giza, Egypt

Djallel Eddine Gherissi Institute of Agronomic and Veterinary Sciences, University of Souk-Ahras, Souk Ahras, Algeria

Laboratory of Animal Productions, Biotechnologies and Health, University of Souk-Ahras, Souk Ahras, Algeria

Nesrein M. Hashem Animal and Fish Production Department, Faculty of Agriculture, Alexandria University, Alexandria, Egypt

Ayushi Kapoor Department of Biosciences and Bioengineering, Indian Institute of Technology Roorkee, Roorkee, Uttarakhand, India

Satish Kumar Animal Biotechnology Centre, National Dairy Research Institute, Karnal, Haryana, India

Ramzi Lamraoui Institute of Agronomic and Veterinary Sciences, University of Souk-Ahras, Souk Ahras, Algeria

Department of Biology of Living Organisms, Faculty of Natural and Life Sciences, University of Batna, Batna, Algeria

Juan J. Loor Department of Animal Sciences, University of Illinois, Urbana, IL, USA

Elis Lorenzetti Laboratory of Animal Virology (LABVIRAL), State University of Londrina-UEL, Londrina, PR, Brazil

Post Graduate Program in Animal Health and Production, Universidade Pitágoras Unopar, Araçatuba, PR, Brazil

S. M. Lutful Kabir Department of Microbiology and Hygiene, Bangladesh Agricultural University, Mymensingh, Bangladesh

Dhruba Malakar Animal Biotechnology Centre, National Dairy Research Institute, Karnal, Haryana, India

Miguel Mellado Department of Animal Nutrition, Autonomous Agrarian University Antonio Narro, Saltillo, Mexico

Fábio Morotti Laboratory of Animal Reproduction and Biotechnology, State University of Londrina-UEL, Londrina, PR, Brazil

P. Ravi Kanth Reddy Veterinary Dispensary, Taticherla, Andhra Pradesh, India

Abdelfattah Z. M. Salem Facultad de Medicina Veterinaria y Zootecnia, Universidad Autonoma del Estado de Mexico, Toluca, Mexico

Marcelo Marcondes Seneda Laboratory of Animal Reproduction and Biotechnology, State University of Londrina-UEL, Londrina, PR, Brazil

S. K. Shaheenur Islam Department of Microbiology and Hygiene, Bangladesh Agricultural University, Mymensingh, Bangladesh

Department of Livestock Services, Krishi khamar Sarak, Dhaka, Bangladesh

Bhaskar Sharma School of Life and Environmental Sciences, Faculty of Science, Engineering, and Built Environment, Deakin University, Geelong, VIC, Australia

Department of Biotechnology, TERI School of Advanced Studies, Delhi, India

Anil Kumar Verma Department of Microbiology, Sikkim University, Gangtok, Sikkim, India

Aparna Verma Department of Biosciences and Bioengineering, Indian Institute of Technology Roorkee, Roorkee, Uttarakhand, India

Monica Yadav Department of Biosciences and Bioengineering, Indian Institute of Technology Roorkee, Roorkee, Uttarakhand, India

Mohamed Zeineldin Department of Animal Medicine, College of Veterinary Medicine, Benha University, Benha, Egypt

Chapter 1

Artificial Insemination Program in Cattle



Fábio Morotti, Elis Lorenzetti, and Marcelo Marcondes Seneda

Abstract Artificial insemination is a key technique to increase reproductive efficiency and the genetic improvement in cattle. The first report of artificial insemination has been dated to more than 220 years ago; after that, many studies have been published, and this biotechnology is widely used in domestic animals. Despite the advances of embryo transfer programs *in vivo* and *in vitro* in cattle, artificial insemination still represents the main form of assisted reproduction in many herds, being one of the lowest cost strategies to improve genetic merit in the farms. Artificial insemination has many advantages compared to natural breeding, such as health benefits, e.g. control of infectious diseases; genetic improvement, e.g. use of genetically improved bulls; reproductive management improvements; and more stringent control of the zootechnical and economic features of the herd, e.g. standardized management and animals.

In addition, we highlight the main advances related to the use of artificial insemination in cattle, such as the development of other reproductive biotechniques, such as timed artificial insemination and resynchronization of estrus or ovulation. In addition, the sexed semen has evolved satisfactorily with promising results both in insemination and in embryo production. An important advance in artificial insemination is related to the hormonal control of the estrous cycle that allows the elaboration of highly efficient protocols for the synchronization of estrus and ovulation. The hormonal protocols have contributed to the dissemination of the artificial insemination by allowing the insemination of many females in a short period of time. Based on many studies evaluating timed artificial insemination protocols and

F. Morotti (✉) · M. M. Seneda
Laboratory of Animal Reproduction and Biotechnology, State University of Londrina-UEL,
Londrina, PR, Brazil
e-mail: fabiomorotti@uel.br

E. Lorenzetti
Laboratory of Animal Virology (LABVIRAL), State University of Londrina-UEL,
Londrina, PR, Brazil

Post Graduate Program in Animal Health and Production, Universidade Pitágoras Unopar,
Arapongas, PR, Brazil

early and accurate pregnancy diagnosis techniques, it was also possible to establish estrus and ovulation resynchronization strategies, which have greatly contributed to economic profitability. Finally, here we review many strategies associated with manipulation of the dominant follicle, estrus behavioral and antral follicle count have been elaborated and they represent an important progress to improve the use of timed artificial insemination in cattle. Therefore, this chapter aims to present the main advances of artificial insemination and timed artificial insemination, highlighting the technical parameters, advantages, and influencing factors, and to discuss practical and current strategies for the improvement of the herd via artificial insemination programs.

Keywords Cattle · Estrous cycle · Synchronization · Artificial insemination · Timed artificial insemination · Resynchronization · Sexed semen · Pregnancy · Antral follicle · Fertility

Abbreviations

TAI	Timed Artificial Insemination
TCR	Temporary calf removal
AFC	Antral follicle count
AI	Artificial Insemination
BCS	Body condition score
CL	Corpus luteum
EB	Estradiol benzoate
EC	Estradiol cypionate
eCG	equine chorionic gonadotropin
EV	Estradiol valerate
FSH	Follicle stimulating hormone
GnRH	Gonadotropin-releasing hormone
hCG	human chorionic gonadotropin
IVF	<i>In vitro</i> fertilization
LH	Luteinizing hormone
P4	Progesterone
PGF2 α	Prostaglandins
RSS	Reverse-sorted semen
RTS	Reproductive tract scoring system
US	Ultrasound

1.1 Introduction

For definition artificial insemination (AI) is a procedure characterized by manually semen deposition in the female reproductive tract. This practice is considered the oldest reproductive biotechnology and is one of the more efficient strategies for genetically improving livestock and increasing their reproductive efficiency. Therefore, AI is still considered the most widely used biotechnology in the world due to its genetic, health and economic benefits for beef and dairy cattle.

The productive and reproductive performance of the herd are considered the main factors responsible for affecting the livestock economic scenario. However, reproductive traits have low heritability and can be highly affected by environmental, nutritional, and health factors. Thus, adequate management of all livestock sectors is required to increase productive and reproductive efficiency and achieve economic autonomy. Therefore, to use reproductive strategies with maximum efficiency, it is necessary to control influencing factors as well as ensure that the overall management is suitable for livestock production.

Conventional AI programs that using estrus detection and insemination procedure have been used for a long time and still represents the main form of reproduction in many herds, mainly for dairy cattle (Smith et al. 2018). However, low service rates due to estrus detection failures represents one of the main obstacles to improving reproductive efficiency for both beef and dairy cattle. This problem is most evident in *Bos indicus* herds that have a short duration of estrus and high percentage of estrus behavior that occurs at night in relation *Bos taurus* females (Pineiro et al. 1998; Galina et al. 1996). In addition to the failures in the estrus detection and the particularities of estrus behavior, a high proportion of cows managed in tropical pastures face nutritional problems that intensify the postpartum anestrus, hindering the return of cyclicity and compromising the efficient use of conventional AI (Baruselli et al. 2004a, 2018).

On the other hand, timed artificial insemination (TAI) have emerged as an alternative to conventional AI programs, in addition to representing the most effective strategy for employment and dissemination of AI and its benefits, mainly for large-scale programs, because the TAI aims for efficient control of estrus and ovulation of all females destined for breeding. After the onset of programs for the synchronization of ovulation, the estrus detection was dispensed, and TAI became widely used because it allows ovulation control and insemination of large numbers of females in a predetermined amount of time. Many advantages have been added to the AI technique associated with synchronization of ovulation, such as elimination of estrus detection, improvement of livestock management, planning of services and financial resources, and improvement of pregnancy rates depending on categories and reproductive characteristics of the herd (Colazo and Mapletoft 2014; Lamb et al. 2010; Baruselli et al. 2004a). TAI protocols synchronize follicular recruitment and growth, regression of the corpus luteum (CL), and ovulation of the dominant follicle, improving reproductive performance because all females are inseminated regardless of their estrus behavior (Lamb and Mercadante 2016). Therefore, TAI

programs have been used as a reproductive tool that better enables the expansion of AI in dairy and beef cattle.

After the first protocol of ovulation synchronization, a wide variety of protocols have been developed focusing on adjustments in female category to minimize management and cost and to obtain more reproductive efficiency. Among various hormonal combinations for TAI in cattle, the association of progesterone/progestin (P4) or gonadotropins with estrogen and prostaglandins (PGF2 α) is noteworthy (Baruselli and Teixeira 2018; Baruselli et al. 2004a). Other reproductive strategies such as estrus or ovulation resynchronization have been successfully used to reduce the number of females that fail pregnancy after the first AI procedure. In addition, resynchronization has been used to increase pregnancy rates in the short time of the breeding season and reduces the use of bulls for natural breeding (Marques et al. 2015; Campos et al. 2013). This chapter aims to present the main advances of AI and TAI, highlighting mainly the technical parameters, advantages, and influencing factors and to discuss practical and current strategies for improving reproductive performance in cattle.

1.2 Artificial Insemination Program: Benefits and Advantages

Genetic improvement is considered one of the main benefits of using AI (Fig. 1.1). In cattle, the genetic improvement is achieved by the choice of proven bulls with desirable characteristics for enhancement in milk or meat production. In addition, AI also provides genetic gain in a short amount of time and low cost compared to natural breeding because semen storage is an economically viable alternative

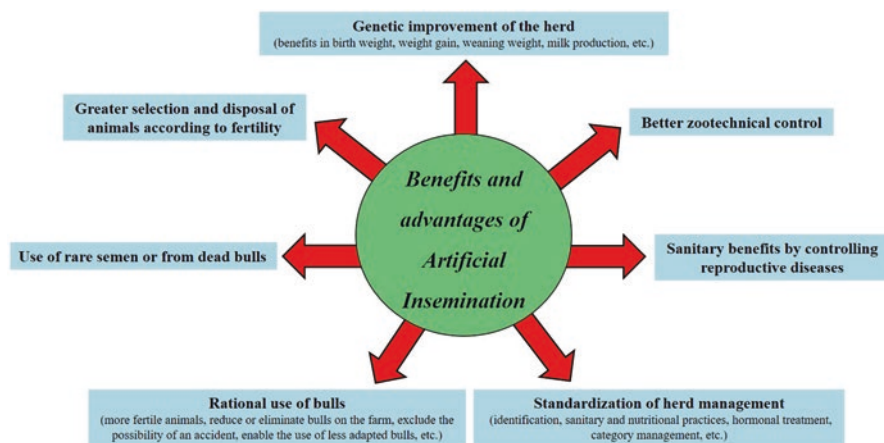


Fig. 1.1 Main benefits and advantages associated with the use of artificial insemination programs in cattle

compared to purchase and keeping a bull with high zootechnical potential in the herd (Baruselli et al. 2018).

Another benefit associated with the use of AI programs is the intensification and standardization of reproductive management of the herd for better zootechnical control of animals. Particularly, of this is important in tropical countries, because the majority of beef and dairy production comes from herds kept almost exclusively in grazing conditions. This strategy represents a low-cost production system; however, some management practices need some adjustments to achieve success after using reproductive biotechnologies. Thus, optimized management practices and better zootechnical control allows for more efficient production in addition to the selection of the most fertile animals and for discarding females with infertility (Marques et al. 2018).

Many infectious agents such as bacteria, viruses, protozoa, and fungi affect reproductive performance in cattle and can potentially be transmitted through natural breeding (Anderson 2007; Yoo 2010). Reproductive diseases such as Brucellosis, Leptospirosis, Campylobacteriosis, Trichomoniasis, Bovine viral diarrhea and Infectious bovine rhinotracheitis (Infectious pustular vulvovaginitis) and Trichomoniasis can provide severe consequences to reproductive performance and economic gain due to the increased occurrence of early and late embryonic loss and fetal death (Alfieri et al. 2019). In this context, another benefit of AI is herd sanity because of the elimination of natural breeding and the use of bulls that can carry contagious infectious diseases (Eaglesome and Garcia 1997). Thus, along with genetic improvement, AI aims to use semen commercialized by qualified companies, which select sires based on sanitary program analysis to control reproductive diseases.

The implementation of AI programs also allows for the reduction or elimination of bulls on farm (Marques et al. 2018), releasing pastures for other animal categories. Furthermore, replacing the natural breeding program to AI program can avoid the possibility of bull attacks (accidents) on workers involved in dairy activities, which is common. In addition, the insemination facilitates the selection of bulls with low expected progeny differences in birth weight, reducing the occurrence of dystocia due to the birth of smaller calves (Mee 2008).

In countries with a tropical climate, with high temperatures, the use of some bull breeds represents a great challenge for the breeder's good performance in the natural breeding. The same situation can be observed in countries with cold climates and that intend to make use of bulls of Zebu breeds, which are often more adapted and regions with hot climates (Menegassi and Barcellos 2015). In this context, the use of AI also allows for the use of bulls less adapted to some climatic conditions, i.e., it is common to observe a reduction in libido of some breeds of bulls when placed in high temperature environments. Therefore, AI enables the use of *Bos taurus* bulls for crossbreeding, which are less adapted to high temperatures and require more management. Furthermore, AI also favors heterosis by crossing *Bos taurus* and *Bos indicus* cattle (Buckley et al. 2014), which represents an interesting strategy due to the improvement in zootechnical performance, quality of progeny carcasses (Gama et al. 2013), parasitic resistance, adaptability, and good acceptance by consumers.

Finally, despite AI having many advantages for the herd, there are some limitations, such as the need for specialized labor, failure in applying biotechnology, and nutrition and sanitarian factors (Demetrio et al. 2007; Orihuela 2000). Understanding these limitations is necessary for implementation of strategies to enable the use of AI programs.

1.3 Physiology Aspects of the Estrous Cycle in Cattle

The estrous cycle reflects a cyclical pattern of ovarian activity that makes the female receptive to the male, favoring mating and further gestation (Forde et al. 2011). Domestic cattle (*Bos indicus* and *Bos taurus*) are considered annual polyestric animals and exhibit estrus behavior on average every 21 days, varying from 18 to 24 days depending on the number of waves of follicular growth (Figueiredo et al. 1997; Roche 1996; Ginther et al. 1989b). However, this cyclical behavior is only possible from puberty, when the female reached appropriate weight and age for each breed.

1.3.1 Puberty

The pubertal process is characterized by an activation of physiological events in the hypothalamic-adenohypophyseal-ovarian axis which culminate in reproductive maturity. In this process, the females undergo by a sequence of events that begins with the expression of estrus, followed by the occurrence of ovulation and the beginning of a luteal phase (Seneda et al. 2019; Atkins et al. 2013; Rawlings et al. 2003). The onset of puberty is variable according to age, weight, and subspecies of heifer, for example taurine animals have greater sexual precocity than zebuine animals. In *Bos taurus*, puberty can occur at approximately 6 to 12 months of age or when the heifer reaches 40 to 50% of the adult weight (Moran et al. 1989). However, in *Bos indicus*, puberty is achieved at approximately 15 to 18 months of age or close to 60 to 70% of the adult weight (Nogueira 2004). In addition, factors such as season of the year, growth rate, nutritional intake, social cues and treatment with exogenous progestins can change the age to the puberty.

In prepubertal females, the low concentration of estradiol produced in the gonads due a negative feedback mechanism prevents the hypothalamic preovulatory center from secreting adequate pulses of Gonadotropin-releasing hormone (GnRH; Amstalden et al. 2011). The absence of GnRH also inhibits the release of Follicle stimulating hormone (FSH) and Luteinizing hormone (LH) from the anterior pituitary, thus causing an absence of follicular waves and ovulation (Senger 1997). On the other hand, when the hypothalamus starts the GnRH neuron function (influenced mainly by nutritional control), there is a decrease in the negative feedback of estradiol, which leads to an increase in the frequency of release of the GnRH pulses

(Amstalden et al. 2011; Kinder et al. 1995). In this context, the follicle growth occurs due to increased levels of intrafollicular estradiol and release of a GnRH discharge through of the center of the hypothalamus (Jainudeen and Hafez 2013). Subsequently, a GnRH discharge results in an increase in the tonic release of LH pulses that is the main endocrine factor that regulates the onset of puberty in heifers. Increased frequency of LH release increases the development of ovarian follicles that produce estradiol sufficient to induce behavioral estrus and a preovulatory peak of gonadotrophins (Foster et al. 2006; Kinder et al. 1995).

Commonly the first behavioral estrus is not preceded by significant concentrations of P4. Thus, the first luteal structure that forms disappears and puberty is reached only after the occurrence of the first behavioral estrus. From that, there is the ovulation and the development of a CL with a typical lifespan. Finally, in puberty, all components of the hypothalamic-adenohypophyseal-ovarian axis will be properly normalized so that reproductive activity occurs cyclically and regularly, and sexual maturity is achieved. The prepubertal heifers present a specific pattern of endocrine and ovarian follicular dynamics. The first ovulation usually occurs at 56 weeks of age, being usual non regular cycles at the beginning of ovarian activity. The short duration of ovulatory cycle was associated with low P4 concentrations and small CL (Evans et al. 1994).

In this context, it is important to emphasize that the occurrence of puberty is considered one starting point to the productive life of a female (Brumatti et al. 2011). Then, early attainment of puberty predicts anticipation of this process (Ferraz et al. 2018). However, studies have shown that the optimal age at first birth is 24 months (Haworth et al. 2008). Below this age range, it is improbable that heifers are of enough body size to express their full genetic potential, either for the production of milk throughout life or even to deliver and maintain a healthy calf. In addition, breeding costs are increased for heifers calving later than those heifers calving early (Boulton et al. 2017). Heifers calving later have lower chances of completing the first lactation (Brickell and Wathes 2011) and relatively low fertility with longer delivery intervals (Wathes et al. 2014). In this way, producers that breed late females may be wasting resources with less likely to give birth in an ideal age or to survive beyond a first lactation in order to recover the costs of breeding (Boulton et al. 2017). In this context, studies have been developed with the purpose of generating alternatives to promote puberty as early as possible in heifers, without disrupting future productive and reproductive performance of these females (Seneda et al. 2019). Genetic selection by animals with greater sexual precocity and the use of hormonal protocols to induce puberty represent good alternatives for the selection of earlier and more productive matrices in the livestock (Gonzalez et al. 2020).

1.3.2 Neuroendocrine Control of the Estrous Cycle

The estrous cycle, a series of reproductive events and ovarian functions in cattle, such as recruitment, follicular growth, ovulation, luteinizing and luteolysis, are regulated by a perfect hormonal interaction among hormones from the

hypothalamus (GnRH), anterior pituitary (FSH and LH), ovary (P4, estradiol, inhibins and oxytocin), and uterus (prostaglandin F2 α – PGF2 α). In addition, non-hormonal factors, such as insulin-like growth factors, also have an important function in ovarian follicular activity. The hormones act through a positive and negative feedback system for controlling estrous cycle, allowing reproductive events by specific control of ovary and uterus activity (Forde et al. 2011; Crowe 2008; Roche 1996).

GnRH is one of the main hormones involved in the estrous cycle control by stimulating the production and secretion of gonadotropins (Fig. 1.2). This hormone acting on the anterior pituitary or adenohypophysis to stimulate FSH and LH production and release. GnRH is produced in the nucleus of hypothalamus neurons and transported by axons to the median eminence, which is where it is stored (Yin and Gore 2010). Subsequently, GnRH is released and transported to the hypophysis by the portal hypothalamic-hypophyseal system, where GnRH binds to cell surface receptors and signals the release of FSH and LH (Vizcarra et al. 1999; Moenter et al. 1992; Keri et al. 1985). FSH is stored in secretory granules located in the cytoplasm for shorter periods, whereas LH is stored for longer periods during the estrous cycle (Farnworth 1995).

The GnRH functions depend of stage of the estrous cycle. At the beginning of the cycle, GnRH stimulates FSH secretion to follicular recruitment and initial growth of small follicles. At the end of the cycle GnRH stimulates LH secretion to induce final

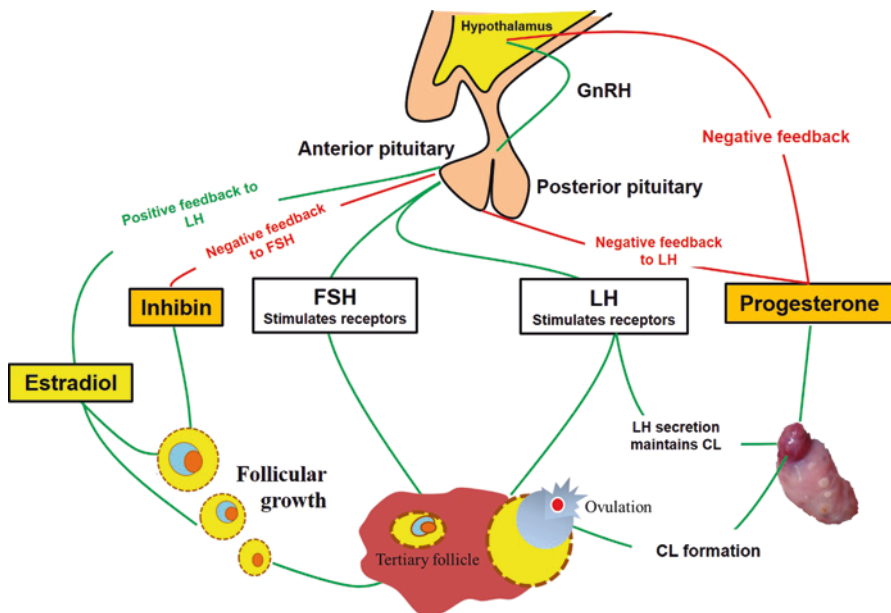


Fig. 1.2 Schematic representation of the hormonal interaction of the hypothalamic-pituitary-gonadal axis to stimulate and/or inhibit ovarian follicular activity in cattle. Gonadotropin releasing hormone (GnRH), Luteinizing hormone (LH), Follicle-stimulating hormone (FSH), and corpus luteum (CL)

growth and ovulation of the dominant follicle (Crowe and Mullen 2013). The CL regression during the follicular phase of the estrous cycle results in basal P4 concentrations. So, there is a concomitant increase in estradiol concentration due to the development of a dominant follicle, which induces increasing release of GnRH and allows estrus behavior during which females are sexually receptive to mounts (Frandsen et al. 2009). However, ovulation of the dominant follicle only occurs with basal P4 plasma concentration and LH pulse frequencies occurring every 40–70 min for 2–3 days (Roche 1996). The dominant follicle ovulates approximately 10 to 14 h after the end of the estrus behavior. After breakage of the preovulatory follicle, metaestrous is initiated and by 3 to 4 days is characterized by the formation of CL from the hemorrhagic corpus (Forde et al. 2011).

Immediately after ovulation, CL formation occurs from luteinization of granulosa and internal theca cells. This ovarian structure is responsible for P4 production, which is required for the diestrous phase or maintaining a pregnancy (Schams and Berisha 2004; Butcher et al. 1992). During the diestrous phase, P4 concentrations remain elevated, and recruited follicles continue to develop due to FSH being released from the pituitary. However, growing follicles are inhibited due to high levels of P4 during this phase, which does not allow appropriate frequencies and/or amplitudes of LH pulses due to negative feedback with this hormone. Therefore, the dominant follicle does not ovulate and undergoes atresia, gradually decreasing its diameter, during diestrous (Crowe 2008; Manikkam and Rajamahendran 1997; Savio et al. 1993; Taylor and Rajamahendran 1991). On the other hand, during the proestrous phase, P4 concentrations decrease due to CL regression in response to PGF2 α uterine secretion; then, this hormonal condition (low P4 and high estradiol) allows for LH pulses and consequently ovulation (Fig. 1.3) (Crowe and Mullen 2013; Forde et al. 2011; Goff 2004).

1.3.3 Estrous Cycle Phases and Estrus Behavior

The estrous cycle in cattle includes a normal length of 18 to 24 days with two distinct phases: (1) the luteal or progesteronic phase and (2) follicular or estrogenic phase. The luteal phase lasts 14 to 18 days and is the period that follows ovulation, being characterized by formation and CL presence (also known as metestrus and diestrus phases). The follicular phase extends from CL regression (luteolysis) until ovulation and lasts 4 to 6 days. This phase is also known as proestrous and estrus, being a required event for ovulation of dominant follicle. In this phase, the dominant follicle undergoes final maturation, and the ovulatory follicle releases the oocyte to fertilization at the uterine tube (Forde et al. 2011; Adams et al. 2008; Rawlings et al. 2003; Sunderland et al. 1994).

The signs of estrous behavior are manifested during the follicular phase and characterized by primary and secondary signs. Unrest, increased physical activity (vocalization and hyperactivity), swelling and reddening of the vulva, friendly contact with herd mates, smelling and licking the genitalia, clear mucus secretion, and

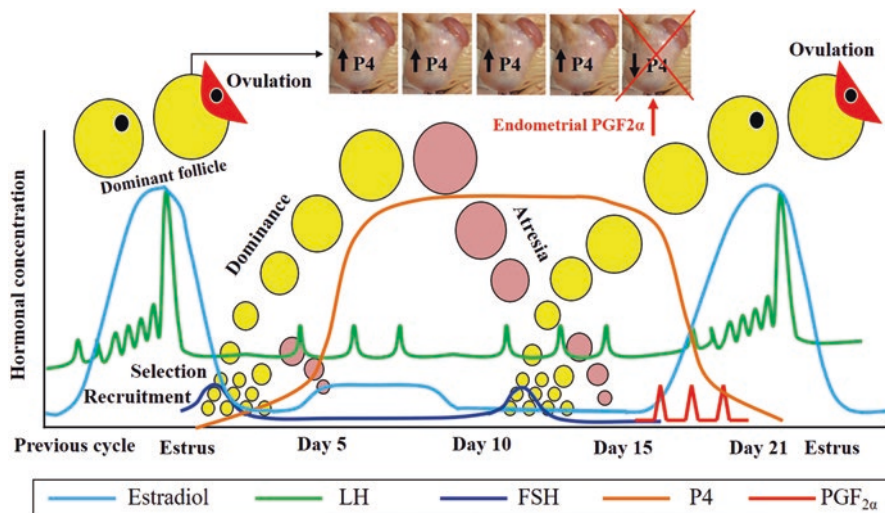


Fig. 1.3 Schematic description of the growth pattern of ovarian follicles during follicular dynamics [recruitment, selection, dominance, and atresia (luteal phase) or ovulation (follicular phase)] and secretion pattern of estradiol, Luteinizing hormone (LH), Follicle-stimulating hormone (FSH), progesterone (P4), and prostaglandin F_{2α} (PGF_{2α}) during the estrous cycle in cattle

attempts to mount constitute secondary signs. However, being receptive to mounting is the most reliable signal of estrus in cattle, which is considered the primary sign (Løvendahl and Chagunda 2010; Galina et al. 1982). Frequently, secondary signs precede (proestrous) estrus and may continue after females accept the mount. Therefore, the identification of secondary signals also assists in the process of identifying the estrus itself.

The estrogen is the main hormone responsible for estrus behavior, which is produced by the dominant follicle and signals the hypothalamus to induce the secondary signs and behavior of receptivity, an event necessary for successful mating. However, an increase in the estradiol concentration and a reduction of P4 concentration is required for manifestation of estrus behavior. This hormonal condition occurs at the end of the luteal phase, being induced by luteolysis between 15 and 17 days of the estrous cycle (Allrich 1994; Vailes et al. 1992). Thereafter, the P4 plasma level is reduced below 1 ng/mL, and this event is responsible for estrus return in nonpregnant females.

Estrus behavior duration and intensity can be affected by several factors, such as, age, production level, environmental factors, nutritional factors, health status, size of the sexually active group, presence of a bull, climatic factors (strong rain and wind), management of animals in pastures and stalls, and subspecies (*Bos taurus* versus *Bos indicus*) (Crowe 2008; Galina et al. 1996). In beef cattle, an average duration of 8 to 10 h has been reported (Crowe 2008; Bó et al. 2003). In dairy cattle, the average estrus duration is 8 to 16 h in cows and 12 to 14 h in heifers, but this duration tends to decrease as milk production increases (Forde et al. 2011; Crowe

2008; Allrich 1994). This negative relation between milk production and duration of estrus has been reported in high-producing dairy cows (above 40 kg of milk per day) because there is a decreasing circulating estradiol level due to increased metabolism of this hormone (Wiltbank et al. 2006).

Differences in the characteristics of estrus between *Bos taurus* and *Bos indicus* females has been well reported (Pinheiro et al. 1998; Galina et al. 1996). In general, zebu cattle present a short duration of estrus and high rates of estrus expression at night, which makes it difficult to identify the estrus for conventional insemination program. Under Brazilian management conditions, the use of a radiotelemetry device in heifers has demonstrated a shorter average estrus duration for *Bos indicus* (12.9 h in Nelore) compared to that of *Bos taurus* (16.3 h in Angus), but no difference was found between *Bos indicus* x *Bos taurus* crossbred (12.4 h in Nelore x Angus) and *Bos indicus* (Mizuta 2003). In addition, it was reported that 53.8% of the estrus expression in *Bos indicus* (Nelore cows) begins at night (between 6 p.m. and 6 a.m.) with 30% of its beginning and finishing occurring at night (Pinheiro et al. 1998).

1.3.4 Events of the Ovarian Follicular Dynamics

Follicular dynamics is one of the most important events of ovarian physiology, being widely studied in *Bos taurus* and *Bos indicus* females (Cuervo-Arango et al. 2011; Mackey et al. 2000; Figueiredo et al. 1997; Ginther et al. 1989a). The follicular dynamics involve growth, development, and maturation of ovarian follicles that are essential processes for reproductive efficiency in farm animals. These dynamics events are totally dependent on the perfect functioning of the hypothalamus-pituitary-gonadal axis that acts through positive and negative feedback mechanisms to stimulate development of ovarian follicles by gonadotropins secretion (FSH and LH).

During the estrous cycle, 2 or 3 waves of follicular growth may be present, but only the last wave is considered an ovulatory wave (Figueiredo et al. 1997; Ginther et al. 1989b). Every wave consists of the recruitment of a group of small follicles (4 to 6 mm in diameter) by FSH stimulation. These follicles grow and future dominant follicle is selected, which suffers a more efficient action of FSH in relation to the other follicles of the wave. So, there is follicular divergence and a dominant follicle emerges, which undergoes atresia or ovulation depending on the circulating levels of P4 that may or may not allow the estrogen trigger on LH pulsatility (Atkins et al. 2013; Forde et al. 2011). Follicular waves are initially established during the prepubertal period; however, the dominant follicle undergoes atresia and does not ovulate. Therefore, only after puberty the ovarian follicular dynamics become a cyclic event characterized by estrus and ovulation (Atkins et al. 2013; Bergfeld et al. 1994).

In general, ovarian follicular development through a non-gonadotropin-dependent phase and a gonadotropin-dependent phase (Webb et al. 2004). During fetal life, for example, follicle development occurs for 3–4 months and the two

phases are present. During the gonadotropin-dependent stage, follicular development occurs in waves (2–3 waves of growth per estrous cycle), being that each follicular wave comprises emergence, selection, dominance, and atresia or ovulation (Fig. 1.3) (Ginther et al. 1989a). Frequently, dairy cows have two waves and heifers, or beef cattle have three waves (Forde et al. 2011). In *Bos indicus* (Nelore), there is a predominance of two waves for cows (83.3%) and three waves for heifers (64.7%) (Figueiredo et al. 1997).

The emergence of the wave is the first event gonadotropin dependent, which is characterized by recruitment of a group of follicles (1 to 50 follicles ≥ 5 mm) that starts growth in response to increased FSH concentration (Sunderland et al. 1994). Follicular growth and cell proliferation in this period occur primarily by FSH stimulation, being characterized by the presence of FSH receptors located in the granulosa cells of follicles at day 3 of the follicle wave (Ginther et al. 2002; Richards et al. 1998). Following follicular development, the next important event is follicular selection, a mechanism that reduces the number of recruited follicles to the ovulatory quota of each species, usually one follicle in bovine species (Sunderland et al. 1994). In this phase, there is a transient increase in FSH concentration, which induces an increase in the aromatase enzyme activity of granulosa cells responsible for conversion of androgen to estrogen (Hillier 1994). This process increases estrogen and inhibin concentrations of healthy follicles, assisting the selection process and gonadotropin secretion (Ginther et al. 2002).

Once the follicular selection process has been established, the dominant follicle diameter increases, and divergence is started when the largest follicle reaches an average diameter of 5.7 to 6.1 mm in *Bos indicus* (Sartorelli et al. 2005) and 8.0 to 8.5 mm in *Bos taurus* (Ginther et al. 1999). In sequence, the follicular dominance is started by growth of the dominant follicle of approximately 1 to 2 mm per day (Figueiredo et al. 1997; Knopf et al. 1989). Then, its diameter is enlarged due to an increase in the follicular fluid. This fluid contains high estrogen concentration, in addition to high levels of inhibin that is associated with negative feedback in the pituitary, reducing FSH levels to basal concentrations (Sunderland et al. 1994). However, due to the presence of LH receptors in the granulosa and theca cells, the dominant follicle becomes sensitive to LH, and even at a low FSH concentration, it continues to grow and accumulate estrogen in the follicular fluid (Xu et al. 1995).

With the growth of the dominant follicle, now being more stimulated by the action of LH than FSH, the destination of the dominant follicle is dependent on frequency and amplitude of the LH pulse (Forde et al. 2011). During the early luteal phase, LH pulsatility has a low amplitude (delta LH, 0.3 to 1.8 ng) and high frequency (20 to 30 pulses/24 h), and during the middle of the luteal phase, LH pulses have high amplitude (delta LH, 1.2 to 7.0 ng) and low frequency (6 to 8 pulses/24 h) (Rahe et al. 1980). This amplitude and frequency are still insufficient to promote final maturation and ovulation of the dominant follicle. Then, during the luteal phase of the estrous cycle, the dominant follicle undergoes atresia and all production of estrogen and inhibin decreases gradually, removing the negative feedback on the FSH secretion in the hypothalamus/pituitary. This event promotes new FSH secretion and the emergence of a second follicular wave (Atkins et al. 2013; Forde et al. 2011).

High estrogen production is an essential characteristic of the dominant follicle that largely depends on LH pulse frequency. In this phase, LH binding to its receptors on theca cells provides the conversion of cholesterol to testosterone through a series of catalytic reactions. Then, testosterone diffuses from theca cells to granulosa cells and is converted into estrogen by the aromatase enzyme (Hillier 1994; Sunderland et al. 1994). Estrogen has a local effect on follicular development and a systemic effect on the hypothalamus and hypophysis. During the follicular phase, when the P4 concentration is low, the estrogen produced by the preovulatory follicle induces an increase in the GnRH level in the hypothalamus, which promotes an increase in the amplitude and frequency of LH pulses that stimulates the final maturation and ovulation of the preovulatory follicle (Crowe and Mullen 2013; Sunderland et al. 1994). The maximum diameter of the preovulatory follicle is variable, being 10 to 14 mm in *Bos indicus* (Sales et al. 2012a; Figueiredo et al. 1997) and 14–20 mm in *Bos taurus* (Ginther et al. 1989a).

In addition, hormonal action intraovarian factors are also important to estrous cycle regulation, acting indirectly by altering estradiol synthesis or directly through negative feedback that controls the hypothalamus and pituitary gland (Forde et al. 2011). Although the acquisition of LH receptors by the granulosa cell layer is considered the main mechanism that promotes the follicular selection process (Lucy 2007), the bioavailability of the insulin-like growth factor (Rivera and Fortune 2003) and the presence of other growth factors also contributes to the development, proliferation, and steroidogenic capacity of the dominant follicle (Knight and Glister 2006).

1.4 Estrus Synchronization Program

Estrus synchronization is a hormonal strategy performed in a group of females with the aim of inducing all animals in the same phase of the estrous cycle. Using this hormonal strategy, management for estrus identification and insemination are facilitated. Two possible methods for estrus synchronization are available: (I) interrupting the activity of the luteal phase or (II) extending the CL length. A luteolytic agent such as PGF2 α or its synthetic analogs (Table 1.1), frequently are used to interruption of the luteal phase and then induce estrus. On the other hand, the duration of the luteal phase can be modified using a progestin treatment, which is similar to the endocrine activity of CL and prevents estrus behavior during treatment. In this case estrus synchronization occurs after removing the P4 source.

1.4.1 Use of Prostaglandins and Synthetic Analogs

Since the 1970s, a wide variety of studies have evaluated the effects of PGF2 α and its synthetic analogs in domestic species (Odde 1990; Rajamahendran et al. 1976). A treatment with PGF2 α in cattle decreases the P4 plasma concentration in 24 h

Table 1.1 Active principle, dose, and commercial name of main prostaglandins and synthetic analogs

Active principle	Dose (mg)	Commercial name
Cloprostenol	0.526	Bio-Cio ^a
Cloprostenol	0.530	Ciosin ^b
Cloprostenol	0.500	Cioprostim ^c
Cloprostenol	0.526	Cioton ^d
Cloprostenol	0.526	Clocio ^e
Cloprostenol	0.526	SincroCio ^f
Cloprostenol	0.500	Sincrosin ^b
Cloprostenol	26.30	Induscio ^g
D-cloprostenol	0.150	Croniben ^h
D-cloprostenol	0.150	Prolise ⁱ
D-cloprostenol	0.150	VeteGlan ^j
Dinoprostromethamine	25.00	Lutalyse ^k

Source: ^aBioVet, ^bMSD Saúde Animal, ^cBoehringer Ingelheim, ^dJA Saúde Animal, ^eBimeda, ^fOurofino Saúde Animal, ^gGlobalGen Vet Science, ^hBiogénesis Bagó, ⁱUnião Química Farmacêutica Nacional SA, ^jHertape Calier Saúde Animal SA, ^kZoetis Saúde Animal

after injection in 80 to 100% of the animals. However, morphological CL regression is gradual and can be observed by ultrasound approximately 24 to 48 h after treatment. Despite PGF2 α inducing efficient luteolysis, estrus behavior can be observed for a long time due to the stage of development of dominant follicle at the time of application of PGF2 α . If a dominant follicle is present at the time of application, the estrus behavior occurs in a short period, approximately 24 to 72 h after luteolytic agent application. Otherwise, only the dominant follicle of the next follicular wave may be considered, in this case there will be a longer interval until the occurrence of estrous behavior, up to 144 h (Kastelic et al. 1990). Therefore, the synchronization program with PGF2 α requires careful practice for estrus observation in treated animals.

Despite the high efficiency of prostaglandin causing luteolysis and inducing estrus, it is worth noting that its use only has an effect in the presence of a responsive CL. Therefore, for more efficient use of luteolytic agents, it is necessary to identify CL by transrectal palpation or ultrasound. A CL up to the 5th day of life after the ovulation is not responsive to luteolysis because the CL is still in formation. In the same way that a CL after the 15th day of the cycle also does not respond to the application because it may have already undergone natural luteolysis (Fernandes et al. 2006; Meidan and Levy 2002; Levy et al. 2000).

On the other hand, when CL is not evaluated, estrus synchronization with luteolytic agent can be performed by two applications of PGF2 α with intervals of 11–14 days. The second application induces luteolysis in animals that do not respond to PGF2 α at the first treatment, resulting in synchronization rates of 70 to 80% of females (Binelli et al. 2014; Machado et al. 2007; Baishya et al. 1980). Following this strategy, animals that manifest estrus after each application of PGF2 α

can be inseminated, or for greater concentration in the number of females in estrus it is recommended to inseminate all animals only after the second application.

Another option for application of PGF2 α in AI programs is the detection of estrus for 5 to 7 consecutive days followed by AI, and in females that did not express estrus, CL identification and PGF2 α application is performed for further estrus observation and insemination. The estrus observation can continue during the next 5 to 7 days, for a total of 12–14 days of work. These three protocols for the synchronization with PGF2 α are illustrated in Fig. 1.4.

Currently the main synthetic analogs of PGF2 α (Table 1.4) commercially available are dinoprosttromethamine, cloprostenol, and D-cloprostenol. These analogs are considered potent luteolytic agents used routinely in estrous cycle synchronization programs. Intramuscular administration of conventional doses (dinoprosttromethamine 25 mg, cloprostenol 500 μ g, D-cloprostenol 150–300 μ g) is the most used but also have tested other routes for reduction of doses. Half a dose by vulvar submucosa injection resulted in similar synchronization rates; however, application in the vulvar submucosa is a route less practiced, especially for a nonexperienced worker (Baryczka et al. 2018; Valldecabres-Torres et al. 2012; Chacur et al. 2010; Colazo et al. 2002).

Although estradiol esters (estradiol-17 β , benzoate, cypionate and estradiol valerate) are capable of causing luteolysis, the luteolytic effect depends on the ester and

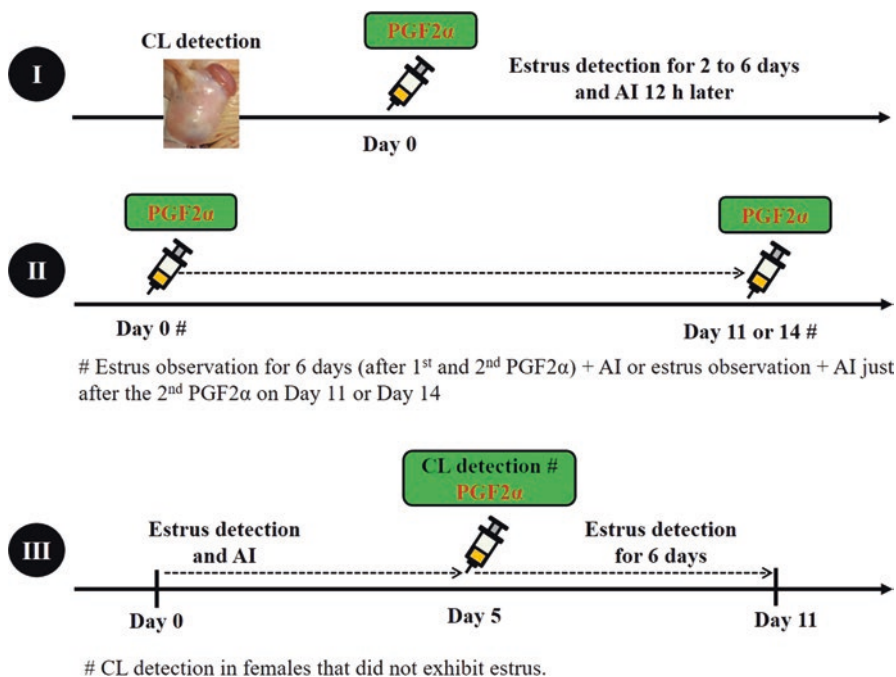


Fig. 1.4 Schematic representation of three treatments options for estrous synchronization with PGF2 α and its synthetic analogs in cattle. CL corpus luteum, AI artificial insemination

dose used (Baruselli et al. 2004b). Of all available esters, estradiol valerate has the greatest potential for luteolysis; however, prostaglandins and their analogs are still more used to induce estrus in cyclic cows.

Commonly females in estrus are inseminated 12 h later, and conception rates are similar to those obtained in natural estrus. However, to facilitate the management of inseminations, females identified in estrus by the morning are inseminated in the late afternoon. Those already detected to be in estrus by the afternoon are inseminated in the morning of the next day. In addition to facilitating the management of the process, this strategy also results in better synchronization of oocyte and sperm viability.

1.4.2 Progesterone and Progestin

Since the 1950s, P4 and its synthetic analogs (progestins) have been used for estrous synchronization programs. The first studies involving P4 were carried out with formulations administered by an oral or injectable route for 14 to 21 days, producing an effective estrus synchronization rate but variable fertility (Hansel et al. 1966). These initial results were unsatisfactory, probably due to their use over a long time and the suboptimal concentration of P4, leading to the formation of persistent follicles containing low-quality oocytes (Fortune and Rivera 1999). In the last decade, the authors of this chapter have conducted a series of studies with injectable P4 formulations, showing promising results for ovarian follicular dynamics (Morotti et al. 2013a, b) and improved fertility of cows submitted to TAI program (Campos et al. 2016a, b). However, there is still no commercially available injectable formulation for use in estrous cycle synchronization programs. Natural P4 has been used in intravaginal devices, and synthetic P4 has been used in subcutaneous auricular implants (norgestomet) or oral use (melengestrol acetate). Despite the different forms of P4 presentation (Table 1.2), an intravaginal device is the most common for synchronization programs.

For the purpose of estrus synchronization, formulations containing P4 can be used to prolong the lutein phase of the estrous cycle or to simulate the effect of a CL, so the synchronization of the animals occurs when removing the source of P4. Although this strategy provides an effective synchronization of the females, in cyclic animals the result is more satisfactory when the application of a luteolytic agent is combined with the removal of the P4 source (Baruselli et al. 2004b). In addition to estrous cycle synchronization, P4 has also been used for cyclicity induction in prepubertal heifers (Gonzalez et al. 2020). This subject will be discussed further in another section of this chapter.

Table 1.2 Main forms of progesterone or progestin presentation (active principle) and their commercial names, administration routes, and concentrations

Active principle	Commercial name	Administration route	Concentration
Norgestomet	Crestar ^a	Auricular	3 mg
Melengestrol acetate	MGA, Premix ^b	Oral	0.5 mg/day
Progesterone	CIDR-B ^b	Intravaginal	1.9 g
Progesterone	DIB ^b	Intravaginal	0.5–1.0 g
Progesterone	Cronipres ^c	Intravaginal	0.5–1.0 g
Progesterone	PRIMER ^d	Intravaginal	0.5–1.0 g
Progesterone	Sincrogest ^e	Intravaginal	1.0 g
Progesterone	SincrogestInjetável	Injectable	150 mg/mL
Progesterone	Progecio ^d	Injectable	70 mg/mL
Progesterone	Repro One ^f	Intravaginal	0.5 g
Progesterone	Repro Neo ^f	Intravaginal	1.0 g
Progesterone	Repro Sync ^f	Intravaginal	2.0 g
Progesterone	Biprogest ^g	Intravaginal	1.25 g
Progesterone	Progestar ^h	Intravaginal	0.96 g
Progesterone	Betaprogin ^h	Injectable	25 mg/mL
Progesterone	Prociclar ⁱ	Intravaginal	0.75 g
Progesterone	Fertilcare 600 ^a	Intravaginal	0.6 g
Progesterone	Fertilcare 1200 ^a	Intravaginal	1.2 g

Source: ^aMSD Saúde Animal, ^bZoetis Saúde Animal, ^cBiogénesis Bagó, ^dUnião Química Farmacêutica Nacional SA, ^eOurofino Saúde Animal, ^fGlobalGen Vet Science, ^gBimeda, ^hBoehringer Ingelheim, ⁱCeva Saúde Animal

1.5 Ovulation Synchronization Program for Timed Artificial Insemination

For the elaboration of an efficient ovulation synchronization program, the hormonal protocol must combine three basic principles: (I) synchronization of the emergence of the follicular growth wave, (II) synchronic control of the progesteronic phase, and (III) synchronized ovulation induction.

Based on these principles, protocols for ovulation synchronization aim to control the growth of the follicular wave to regulate the luteal phase and induce ovulation in a fixed amount of time, avoiding the necessity of estrus detection for AI. For this purpose, some hormones such as P4, progestin, PGF2 α , estradiol esters (estradiol-17 β , benzoate, cypionate and estradiol valerate), equine chorionic gonadotropin (eCG), human chorionic gonadotropin (hCG), and GnRH are combined to provide in estrus and ovulation synchronization.

1.5.1 Synchronization of Follicular Wave Emergence

Progesterone combined with estrogen and/or GnRH is an efficient hormonal combination to synchronization in the emergence of a new wave of follicular growth. Treatment with P4 and estradiol esters on a random day of the estrous cycle (day 0) has been effective for promoting follicular atresia and the emergence of a new follicular wave at a fixed time in cattle (Bó and Baruselli 2014). Exogenous estradiol induces atresia of small follicles by decreasing FSH release, whereas P4 is used to suppress the release of LH and inhibit the activity of large follicles, which are dependent on this gonadotropin. Therefore, application of estrogen and P4 at the same time promotes atresia of all of the follicles (large, medium, and small) present in the ovary and a new wave emergence 2 to 4 days later (Baruselli et al. 2004b; Bó et al. 1995, 2003).

In cyclic *Bos taurus* cows and heifers, the emergence of a new follicular wave occurred in 3 to 4 days after treatment with progestogen and 5 mg of estradiol-17 β or progestin and 1 to 2.5 mg of estradiol benzoate (EB) (Martinez et al. 2005; Bó et al. 1995). Apparently, *Bos indicus* and *Bos taurus* females have similar intervals for the synchronization of the follicular wave. For example, in Brangus heifers treated with an intravaginal P4 device and 2 mg of EB the follicular wave emergence did not differ from wave emergence in Angus heifers (3.3 ± 0.6 versus 4.3 ± 0.2 days) (Bó et al. 2003). A similar time of emergence was also observed between Nelore and crossbred $\frac{1}{2}$ Nelore \times $\frac{1}{2}$ Angus heifers (3.3 ± 0.6 versus 3.5 ± 0.1 days) (Carvalho et al. 2008). Heifers and Nelore cows receiving 2.0 mg of EB and insertion of an auricular implant of norgestomet presented intervals of 2.5 ± 0.2 days for a new wave of emergence (Sá Filho et al. 2011). However, doses of 2.5 and 5.0 mg of estradiol valerate (EV) resulted in intervals of wave emergence of 4.2 ± 0.3 and 6.1 ± 0.6 days in heifers and 3.1 ± 0.4 and 4.0 ± 0.5 days in cows, respectively (Bó and Baruselli 2014).

Another efficient alternative for the synchronization of follicular wave emergence is to associate P4 with GnRH analogs. A study carried with dairy cows (*Bos taurus*) in lactation evaluated the association of 2 mg of EB and CIDR versus 100 μ g of GnRH and CIDR and revealed follicular wave emergence at 4.8 ± 0.4 and 2.0 ± 0.2 days, respectively (Kim and Kim 2007). In high-yielding dairy cattle the emergence of a wave of follicular growth is a major challenge. In this context, studies show that the application of estrogen and GnRH analogs in association with the insertion of a P4 device appears to be more effective in synchronizing the emergence of the follicular growth wave than the isolated use of P4 and estrogen or P4 and GnRH. In addition, twice the dose of GnRH has been more effective in synchronizing the emergence of the wave in these animals (Silva 2020; Wiltbank and Pursley 2014). Despite the efficiency of both combinations, estrogen has a lower cost compared to that of GnRH.

1.5.2 Control of the Progesteronic phase

The progesteronic phase can be controlled efficiently by the use of a P4 device, by the use of luteolytic agents, and by the association of these, which is more effective for synchronization. The use of luteolytic agents is highly efficient for inducing estrus synchronization in cyclic females as previously discussed. However, the use of these agents in association with P4 devices has been more effective than their isolated uses. In this context, PGF2 α or its synthetic analogs was proposed for increasing the fertility of cows submitted to synchronization with CIDR-B for 8–9 days associated with EB at the time of P4 device insertion (Macmillan and Peterson 1993a, b). Efficient control of the progesteronic phase has been reported with the removal of the P4 device between D6 and D9 (Bó and Baruselli 2014; Sá Filho et al. 2010a, b, 2011) and PGF2 α application can be performed at any of the three times: (i) at the time of P4 removal, (ii) 24 to 48 h before P4 removal, or (iii) on day 0 at P4 device insertion (Bó et al. 2002; Macmillan and Peterson 1993a, b). Therefore, administration of luteolytic agents such as PGF2 α before or at the end of P4 treatment is required to ensure luteolysis and adequate control of the progesteronic phase (Kastelic et al. 1999).

The time for application of the luteolytic agent has been widely studied in *Bos indicus* cattle. Initially, luteolysis was observed in 83.3% Nelore cows treated with 5 mg of EV between 2 and 7 days after treatment (Pinheiro et al. 1998). Therefore, due to the high percentage of induction of luteolysis using EV at the mentioned dose, it is not necessary to use PGF2 α in protocols using progestin and EV at the beginning of hormonal treatment (Kastelic et al. 1999). The effects of the induction of luteolysis with dinoprostromethamine were investigated simultaneously or 48 h before P4 device removal. In this study, a higher P4 concentration (4.58 ± 0.21 ng/mL) was observed in cows treated simultaneously to the P4 device removal compared to those treated 48 h earlier (3.05 ± 0.21 ng/mL) (Peres et al. 2009). In addition, cows with early luteolysis improved the ovulation rate (77.0% versus 85.4%) and increased the pregnancy rate by approximately 16%.

In summary, to obtain greater efficiency in TAI protocols, the addition of PGF2 α 2 to 3 days before or at the end of P4 treatment is required to induce luteolysis in cyclic females. In a normal estrous cycle, this strategy increases the proestrous phase (low P4 and high estradiol), better preparing the uterus for pregnancy or allowing subsequent ovulation. However, it is possible that this anticipated luteolysis is not necessary when using more potent PGF2 α analogs such as cloprostenol or D-cloprostenol.

1.5.3 Synchronized Induction of Ovulation

The use of ovulation inducers in association with P4 treatment has been proposed to reduce the dispersion of ovulation and improve the ovulation rate. In addition to the wave emergence, the use of estradiol esters [EB or estradiol cypionate (EC)] at the end of P4 treatment induces ovulation between 72 and 84 h after P4 removal in 75% of cows (Morotti et al. 2013a, b; Bó et al. 1995). Several hormones are used to induce ovulation in cattle, such as estradiol esters (estradiol-17 β , EB, and EC), GnRH analogs, hCG, and LH. The main ovulatory mechanism associated with the use of ovulation inducers is related to the induction of an endogenous LH surge (Baruselli et al. 2004b). Estradiol esters are considered indirect inducers, and GnRH and LH are considered direct inducers. According to cost benefit analysis, the most used hormone for induction of ovulation in *Bos indicus* cattle is EB 24 h after P4 removal (1 mg) or EC along with device removal (0.5 mg in heifers to 1 mg in cows) (Torres et al. 2014). However, GnRH analogs can also be used for this purpose from the time of insemination up to 16 h before.

Frequently, within the TAI protocols the insemination is performed 48 h after P4 removal or approximately 8 to 12 h before the expected time for ovulation (Baruselli et al. 2004a, b; Hanlon et al. 1996). However, to reduce management practices, administration of estradiol esters was proposed to induce ovulation simultaneously or 24 h after removal of the P4 device. Similar results to those with EB for induction of ovulation were obtained with EC administration (Sales et al. 2012a). Similar intervals between CIDR removal and LH surge (54.6 ± 3.4 versus 59.3 ± 3.5 h) and between CIDR removal and ovulation (81.6 ± 5.0 versus 86.4 ± 4.8 h) were observed in heifers treated with 0.5 mg of EC 0 or 24 h after intravaginal device removal (Sales et al. 2012b). Application of EB or EC to induce ovulation after P4 removal resulted in similar ($P > 0.05$) follicular responses (ovulatory follicle diameter: 13.1 versus 13.9 mm), the interval from P4 device removal to ovulation (70.2 versus 68.5 h) and ovulation rate (77.8% versus 82.8%). In addition, the pregnancy per AI was similar ($P > 0.22$) between cows treated with EB (57.5%; 277/482) and EC (61.8%; 291/471) (Sales et al. 2012b). In another study, an interval of 45 h was observed between injection of 0.5 mg of EC simultaneously to P4 device removal and LH surge (Sales et al. 2015).

Other ovulation inducers, such as synthetic GnRH analogs, hCG, or LH, can be used. However, the use of LH is restricted to inducing ovulation in protocols for ovarian superovulation due to its high cost. The same limiting factor, cost, also has to be considered for hCG and GnRH, although GnRH has been widely used alone or in association with EC or EB in females that exhibit low or no estrus expression during TAI protocols (Rodrigues et al. 2019). Generally, indirect ovulation inducers are preconized in conventional TAI protocols, being that the ovulation is normally obtained by cheaper alternatives by the use of estrogens. On the other hand, direct inductors are used either in specific situations, either for TAI or more often for superovulation protocols.

1.6 Strategies for Improving Fertility at TAI

TAI programs represent one of the most important advances in reproductive biotechniques in cattle. The main advantages of this reproductive tool are undoubtedly associated with ease of management, since the TAI protocol eliminated the practice of estrus identification and its low efficiency, contributed to a greater intensification in the use of AI, which made the technique better known and widespread in the world. In addition, TAI has increased the service rate to 100% of the synchronized animals, which significantly contributes to increasing the reproductive efficiency of the herd and the economic performance of the farm (Baruselli et al. 2018).

The goal of the TAI program is to inseminate all treated animals at the same time and to obtain pregnancy rates of approximately 40 to 60%. The best reproductive rates are achieved with well-managed herds, with high quality nutrition, good health, and semen quality and a well-trained team. Although good results can be obtained with TAI, a reduction of approximately 15 to 20% in the conception rate of *Bos indicus* cattle with a high rate of anestrous has been reported (Baruselli et al. 2004a; Fernandes et al. 2001).

Postpartum cows may undergo a prolonged anestrous period during which they do not show behavioral signs of estrus, which is decisive when reproduction is dependent on bull mating. In this context, anestrous postpartum represents one of the biggest challenges for beef breeding, mainly due to endocrine imbalance regarding adequate LH secretion during this period, which is considered the main factor for anestrous in *Bos indicus* cattle (Baruselli et al. 2004a, 2011). Additionally, many herds managed on tropical pastures may suffer nutritional restrictions due to the seasonality of forages, which may aggravate postpartum anestrous conditions. Therefore, due to characteristics of estrus expression, poor efficiency of estrus detection, and a high percentage of cows in postpartum anestrous (Stevenson et al. 2015; Lamb et al. 2010; Crowe 2008; Yavas and Walton 2000b), AI programs based on estrus detection present with low efficiency, mainly for tropical and subtropical herds (Baruselli et al. 2004a; Bó et al. 2003). On the other hand, TAI program represents one of the most efficient alternatives. However, even receiving a hormonal protocol for TAI, fertility may be affected if hormonal adjustments or management practices are not performed properly.

Administration of eCG and temporary calf removal (TCR) are strategies performed to provide appropriate gonadotropin support for growth and ovulation induction of the dominant follicle in females submitted to TAI (Barreiros et al. 2014; Campos et al. 2013; Sá Filho et al. 2014; Yavas and Walton 2000a, b). Practices such as this have been associated with TAI protocols and improved postpartum fertility because gonadotropin support provides an increase in follicular activity. However, the use of this gonadotropin support has been highly indicated mainly in animals with intense anestrous postpartum or that have a low body condition score (BCS) (Baruselli et al. 2004b).

eCG is a glycoprotein hormone produced by the corium of a pregnant mare with wide application in TAI protocols. This glycoprotein has a high molecular weight

and has an excellent gonadotrophic support for follicular growth due its mixed bio-activity (at approximately 2/3 FSH and 1/3 LH) (Soumano et al. 1996; Murphy and Martinuk 1991). It can be used at the end of the hormonal protocol to provide adequate gonadotropic aid for dominant follicle growth, mainly in acyclic cows and heifers (Barreiros et al. 2014). A schematic representation of the pharmacological action and physiological variations of TAI protocols is shown in Fig. 1.5.

Many studies evaluated the effect of eCG administration or TCR use on the conception rate for TAI in cattle (Baruselli et al. 2004a, 2018; Prata et al. 2018; Barreiros et al. 2014; Campos et al. 2013; Sá Filho et al. 2014; Yavas and Walton 2000a, b). Initially, similar conception rate was observed after TAI between cyclic lactating Nelore cows that were submitted to TCR (50.5%) compared to control group (53.5%). However, there was a 22% increase in the conception rates of *Bos indicus*

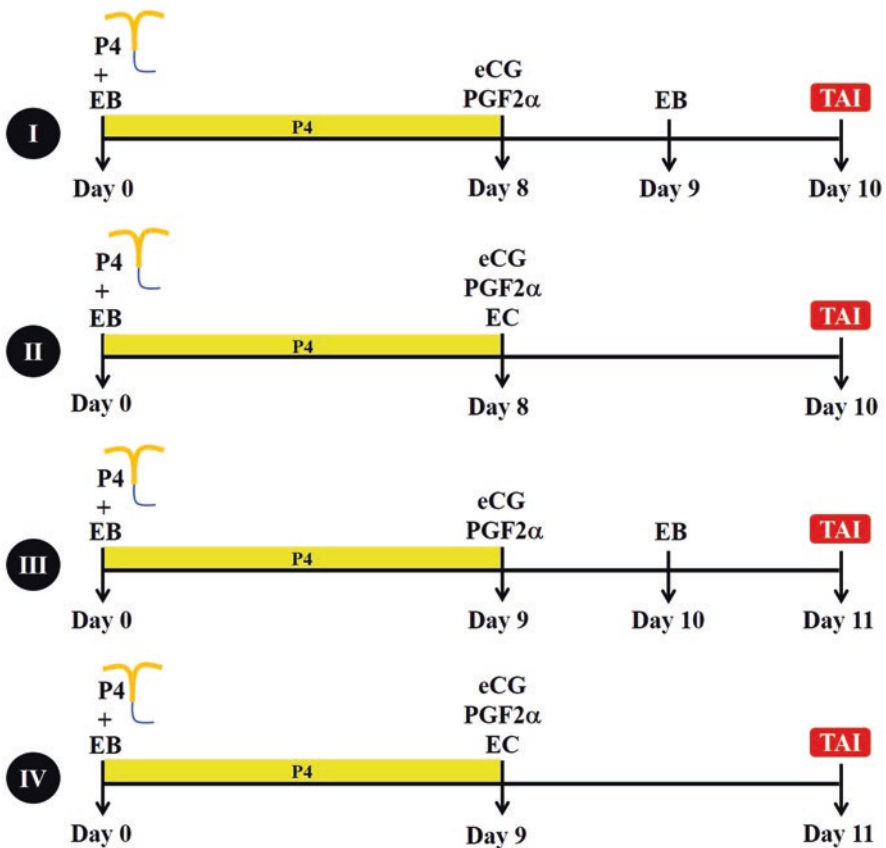


Fig. 1.5 Diagrammatic representation of the protocol for the synchronization of ovulation and timed artificial insemination (TAI) in cattle. Treatment with progesterone (P4) for 8 days [ovulation induced by estradiol benzoate (EB, I) and estradiol cypionate (EC, II)] or 9 days [ovulation induced by EB (III) and EC (IV)]. D, day; P4, progesterone (1.0–1.9 g); EB (D0), 2 mg; EB (D9), 1 mg; PGF2 α , 12.5 mg of dinoprost or 125–250 μ g of cloprostenol; EC, 1 mg; eCG, 300–400 IU equine chorionic gonadotropin

cows with high anestrus rate after the same treatment (Ereno et al. 2007). In addition, after TAI programs, females who do not become pregnant have hormone treatment-induced cyclicity and may result in pregnancy rates of approximately 60 to 65% in the first 45 days of the breeding season (Baruselli et al. 2004a). Therefore, eCG administration for acyclic cows has been recommended to provide a pregnancy rate of around 50%, which is similar to that observed in cyclic cows (Baruselli et al. 2004a; Kastelic et al. 1999).

The use of eCG has been proposed in cows with a high postpartum period (intense anestrus) and BCS < 3 to obtain conception rates around 50% (Baruselli et al. 2004a). However, an interaction between BCS and the postpartum period can also influence conception rates to TAI. Therefore, administration of eCG in cows among 30 to 60 days postpartum is recommended regardless of the BCS (2 to 3.5). Additionally, management using TCR for 56 h was effective for increasing the conception rate of cows submitted to TAI with BCS < 2.5 compared to cows that were of control group (48.2% versus 28.2%; $P < 0.05$) (Peña 2007).

Some studies were conducted to evaluate the effect of eCG and TCR after P4 device removal on the growth of the dominant follicle in *Bos indicus* cattle. Studies have shown increased dominant follicle diameter and increased ovulation rate (85%) in Nelore cows treated with GnRH associated with 48 h of TCR compared to cows that were injected with GnRH without TCR (51%) (Meneghetti and Vasconcelos 2001). A common effect of eCG and TCR on TAI protocols is associated with increased ovulation rates due to a greater synchronization of the preovulatory LH surge (Sá Filho et al. 2006; Cavalieri et al. 1997; Williams et al. 1996). The effects on dominant follicle growth are likely partly related to the percentage of cyclicity and the postpartum period at the beginning of the TAI protocol. It was reported that follicles with a high maximum diameter promote a linear increase in both the follicular diameter and conception rates in *Bos indicus* and *Bos taurus* cattle (Baruselli et al. 2018; Sá Filho et al. 2010a; Borsato et al. 2004).

The luteotrophic effect of eCG was already shown by an increase in the P4 concentration (8.6 ± 0.4 versus 6.4 ± 0.4 ng/mL; $P < 0.05$) 12 days after TAI in Brangus cows (Baruselli et al. 2004a). In addition, the TCR practice also results in a higher concentration of P4 after TAI and provide a higher pregnancy rate especially in acyclic females. Thus, the importance of these gonadotrophic stimuli is remarkable because the interaction between the P4 concentration and early embryonic development are key events during the maternal recognition process of gestation (Mann and Lamming 2001).

1.6.1 TAI in Heifers: Reaching Sexual Maturity and Induction of Puberty

The age of puberty is one of the most important parameters to determine the occurrence of the first calving both in beef and dairy cattle. Although puberty is often associated with endocrine mechanisms, gene-environment interactions can also

affect reproductive physiology in cattle. In heifers, follicular growth in the prepubertal period is characterized by the FSH secretion, occurrence of the follicular wave emergence and dominance follicular, but without the occurrence of ovulation (Wiltbank et al. 2002). Until puberty, endocrine mechanisms promote negative feedback on the hypothalamus, which decreases GnRH and LH pulsation, blocking the final growth of the dominant follicle and ovulation (Day et al. 1987). In this context, an interruption of inhibitory factors on the hypothalamus occurs only when the animal has adequate weight and body development, which allows for full ovarian activity (Seneda et al. 2019). Therefore, it is observed that nutrition can greatly affect the occurrence of puberty.

In general, zebuine and crossbred heifers experience delayed puberty approximately 15 to 27 months old and body weight around 280 to 300 kg (Teodoro et al. 1993). Taurine heifers start puberty approximately with 10 to 15 months and weight from 220 to 250 kg (Patterson et al. 1992). In the last weeks of the prepubertal period and soon after puberty, there is a modification in the size of the female genital system, possibly by increasing the concentration of P4 and estradiol, signaling the acquisition of sexual maturity. Then, puberty is the beginning, but not the fullness, of the activity of the reproductive female trait (Honaramooz et al. 2004). It is important to note that sexual maturity occurs with the full development of the reproductive system, as evidenced by the increase in the uterine size and tone, besides regular ovarian activity.

A combined analysis of the uterus and ovaries should always be considered before using early hormonal treatment programs for the synchronization of estrus and ovulation in heifers. Heifers with regular ovarian activity, but presenting uterine dimensions characteristic of the prepubertal period, generally have low pregnancy rates after AI. A specific pattern of uterine diameter to start reproduction has been studied, but comparative analysis among animals before and after puberty can provide standardization for both the diameter and the tone of the uterus.

A five-point reproductive tract scoring system (RTS) was developed to estimate the pubertal status and reproductive potential of beef heifers by transrectal palpation of the uterine horns, ovaries, and ovarian structures (Table 1.3) (Rosenkrans and Hardin 2003; Andersen et al. 1991; LeFever and Odde 1987) system is practical and highly accurate, and females with score I, II, or III are considered prepubertal and

Table 1.3 Reproductive tract scoring system (RTS) used to evaluation of sexual maturity in heifers

Score	Ovarian and uterine characteristics
RTS I	Immature reproductive tract, without uterine tone, and without palpable ovarian structures
RTS II	Uterine horn diameter with 20 to 25 mm, without uterine tone, and follicles ≤ 8 mm
RTS III	Mild uterine tone and follicles 8 to 10 mm in size
RTS IV	Uterine horn diameter of 30 mm, presence of uterine tone, and follicles ≥ 10 mm
RTS V	Presence of a palpable corpus luteum

Adapted from Andersen et al. (1991)

those with score IV or V are considered pubertal and respond better to reproductive practices.

Prior to the breeding season a puberty-inducing hormone treatment can be used for heifers that have not yet reached sexual maturity. P4 and estradiol-based hormonal protocols have been proposed to induce cyclicity in prepubertal heifers (Rasby et al. 1998). Although some mechanisms are not fully known, it seems that P4 sensitizes the hypothalamus, reducing estradiol receptors and inhibiting GnRH and LH pulsatility (Andersen et al. 1991). In addition to possible effects on the hypothalamus, 17 β -estradiol and P4 treatment increases the expression of receptors for GnRH in the pituitary and improves LH synthesis and secretion (Looper et al. 2003). Therefore, P4 treatment for 10 to 14 days mimics a functional CL and consequently induces puberty, possibly by reducing estradiol receptors in the hypothalamus, which after P4 removal increases LH secretion and promotes ovulation. For greater efficiency, at the end of treatment with P4, 1 mg of EB is applied to increase the ovulatory potential of the dominant follicle (Gonzalez et al. 2020; Rodrigues et al. 2013).

The induction of cyclicity with the P4 device (previously used for 24 days) was tested in prepubertal Nelore heifers and resulted in 85.6% of cyclicity in P4-treated heifers and 81.1% in heifers injected with 1 mg of EB at the time of device removal (Sá Filho et al. 2006). Cyclicity induction in *Bos indicus* heifers has been tested with different sources of P4. Inductions using a P4 device (fourth use) versus P4 injectable (150 mg, i.m.) were efficient for cyclicity induction (81.5% versus 86.3%; respectively; Fig. 1.6) (Gonzalez et al. 2020). A similar cyclicity rate was obtained

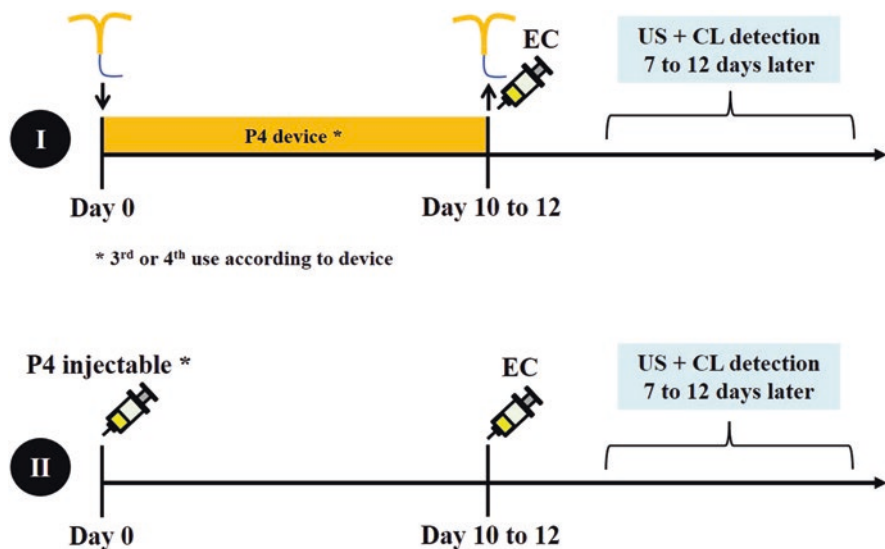


Fig. 1.6 Diagrammatic representation of cyclicity induction protocols in prepubertal heifers. Treatment with progesterone (P4) for 10–12 days [P4 intravaginal device at 3rd or 4th use (I) or P4 injectable (II)]. EC, estradiol cypionate (0.5 mg); * P4 injectable, 150 mg; US, ultrasound evaluation for identification of corpus luteum (CL)

in females treated with P4 and EB (64%) compared to those treated with P4 and EC (67.2%). However, after TAI, greater pregnancy/AI and pregnancy/treated were achieved with EC in relation to the EB (Sá Filho et al. 2015). It is worth highlighting that the authors observed maintenance of P4 concentrations above 1 ng/mL for 7 days in prepubertal Nelore heifers in females treated with a new or reused intra-vaginal device for 24 days.

1.6.1.1 TAI in *Bos indicus* Heifers

TAI programs in heifers have resulted in high reproductive rates similar to those found in cows. However, *Bos indicus* heifers treated with a first-use P4 device (CIDR-B) and/or containing high P4 concentration can reduce the diameter of the preovulatory follicle and interfere with the conception rate for TAI (Dias et al. 2009; Carvalho et al. 2008). This occurs because a high P4 concentration can significantly suppress gonadotropin secretion, specifically LH, impairing the follicular growth rate (Dias et al. 2009; Kinder et al. 1996). Therefore, synchronization of ovulation in heifers is commonly performed with a lower circulating P4 concentration, employing a monodose P4 device, using a reused device (second or third use), by PGF2 α anticipation (day 0 or 7), and/or employing eCG gonadotropic stimulation at device removal (Dias et al. 2009; Carvalho et al. 2008).

Although it provides a reduction in endogenous P4 concentration in cows, the reuse of intravaginal P4 devices in heifers increases the efficiency of TAI, reduces protocol costs, and results in similar conception rates to those of cows (Sales et al. 2011b; Peres et al. 2009). Another alternative for *Bos indicus* heifers is the use of norgestomet which does not compromise the follicular growth or pregnancy rate of TAI (Sá Filho et al. 2013, 2015). Figure 1.7 shows the principal hormone protocols for the synchronization of ovulation and TAI protocols for *Bos indicus* heifers.

1.7 Use of Sexed Semen for AI and TAI

The choice of the calf's sex in the livestock system is considered a determining factor in beef and dairy cattle. For example, a male calf has low or no zootechnical value in dairy farms compared to a female calf. However, a male calf has great importance for beef cattle farms due to its high potential production. In this context, sexed semen has been used in AI programs to increase efficiency in production systems and genetic improvement programs for better targeting sex-related characteristics (Baruselli et al. 2007; Weigel 2004).

Sexed semen has been highlighted in cattle breeding due to the benefits generated by its use (Seidel Jr et al. 1997, 1999). Among various advantages, sexed sperm can provide genetic improvements for milk and meat production, favoring expressions related to sex, decreased incidence of dystocia by the birth of female calves weighing around 2 kg less than males, better efficiency in the production of breeders

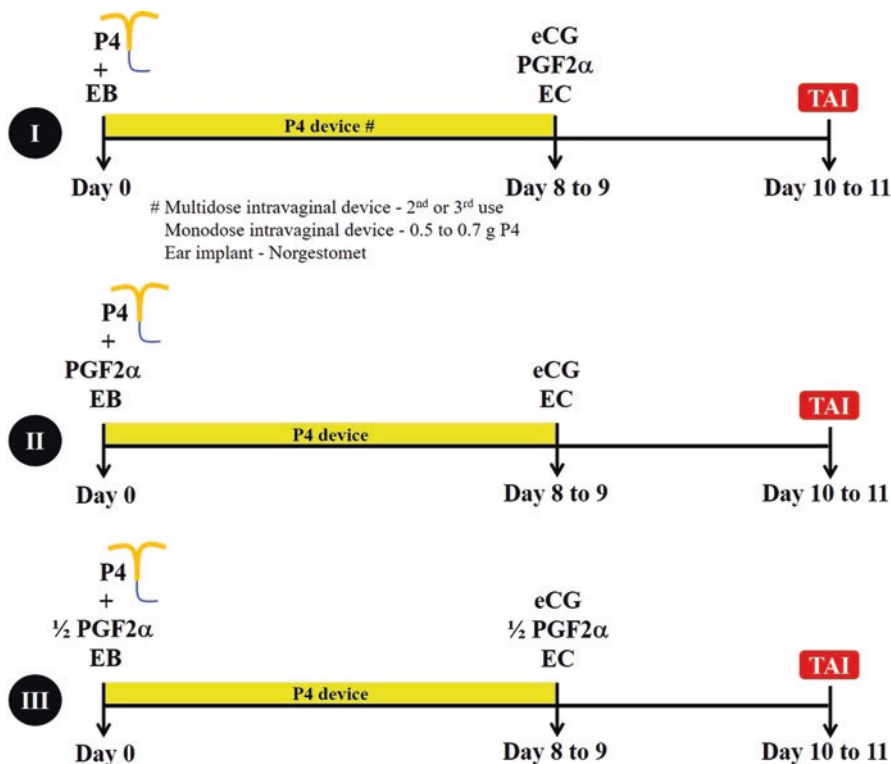


Fig. 1.7 Diagrammatic representation of protocols for the synchronization of ovulation and timed artificial insemination (TAI) in heifers. (I) Treatment with progesterone device (multidose intravaginal, 2nd or 3rd use; monodose intravaginal, 0.5–0.7 g; Ear implant, norgestomet) for 8–9 days. (II) Treatment with progesterone for 8–9 days with PGF2α injection at the beginning of the protocol. (III) Treatment with progesterone for 8–9 days with 1/2 PGF2α injected on day 0 and 1/2 PGF2α together with P4 device removal. P4, 0.5 and 0.7 g of progesterone; EB, 2 mg of estradiol benzoate; PGF2α, 12.5 mg of dinoprost or 125–250µg of cloprostenol; eCG, 300–400 IU of equine chorionic gonadotropin; EC, 1 mg of estradiol cypionate

for replacement (heifers or bulls), and increased number of calves when it is desired to increase herd size for production (Seidel Jr. 2007; Cerchiaro et al. 2007). In addition, advantages from the production of desired sex embryos (superovulation and *in vitro* fertilization) can be obtained, and the best performance using sexed semen is obtained with *in vitro* embryo production (Morotti et al. 2014; Pellegrino et al. 2016; Pontes et al. 2010).

Currently, flow cytometry is the most efficient technology for sexing sperm, and this approach is based on DNA differences between X and Y (Seidel Jr. 2007; Seidel Jr. and Garner 2002). In bulls, X sperm contains around 4% more DNA in their chromosome than Y sperm (Garner 2006), and such a difference allows sperm classification with an efficiency close to 95% for determining the desired sex (Morotti et al. 2014; Pontes et al. 2010). In general, the sexing process involves a preparation

and adding the DNA-binding dye Hoechst 33342 in the ejaculated (Garner et al. 2013). In the sequence the mix of the ejaculate and DNA dye is analyzed in the flow cytometer that can make over 30,000 consecutive evaluations of individual sperm per second, sorting the ejaculate into three parts: X sperm, Y sperm, and waste (indiscriminate and dead sperm) (Seidel Jr. 2014). Therefore, even with high-speed spermatid sorting, producing sexed sperm straws with a conventional concentration (20 to 50 million sperm) is not viable or economical. For this reason, sexed semen was initially commercialized in straws with approximately 2 million sperm (Schenk et al. 2009; Weigel 2004).

In this context, many studies have proven that there is considerable variation in fertility among bulls (both sexed and non-sexed sperm) and that the use of low fertility bulls and low concentration dose may result in a marked reduction in fertility (DeJarnette et al. 2008, 2011; Andersson et al. 2004). Approximately 10 to 20% of bulls have low fertility with a conventional semen dose; therefore, in this case, a reduction in fertility of more than 10 to 20% is expected after using a dose of two million sperm (Schenk et al. 2009; Seidel Jr. and Schenk 2008a, b). However, there is evidence of a relationship between sexed and non-sexed semen. For example, bull fertility using low doses of non-sexed sperm is usually a good indicator of fertility using a low sexed sperm dose (Seidel Jr. 2014).

A comparison among conception rates in Holstein heifers and cows using sexed semen from different bulls and different insemination doses (2.1, 3.5, and 5.0 million sperm) recorded the best pregnancy rates in nulliparous heifers using the 5 million sperm dose (59.5% pregnancy rate) compared to those of the 2.1 and 3.5 million sperm doses (46.4% and 52.2%, respectively) (DeJarnette et al. 2008). Bull or sperm dose did not affect the conception rate among cows (27.0, 29.1, and 30.3% for the 2.1, 3.5, and 5.0 million sperm doses, respectively). This study indicated that using an increased sperm dose of some bulls could improve the conception rate in nulliparous heifers, whereas neither the bull nor the sexed sperm dose affected the conception rate in cows. Therefore, a high insemination dose is only advantageous for some bulls, and it is recommended to identify ahead of time the maximum bull fertility based on semen quality factors and perform tests with low insemination dose.

In addition, a lower sperm dose, sexed semen has lower fertility also due to a lot of damage caused during the sexing process (diluent, dye, classification pressure, freezing process, etc.). Using good management practice, pregnancy rates in cattle with sexed sperm are approximately 80% of those rates from a conventional semen dose, which represents a reduction of 10 to 20% in pregnancy rates compared to those of the conventional non-sexed semen dose (Butler et al. 2014; Seidel Jr. 2014). There is variation in the conception rates reported in the literature; however, there is a consensus regarding the conception rate of heifers after estrous detection. It is estimated that AI with sexed semen ranges from approximately 70 to 90% of those with conventional semen, depending mainly on the farm management conditions. Conception rates of 40 to 68% were reported for Holstein heifers after insemination with sexed semen and between 67 and 82% using conventional semen. Despite the lower fertility reported for sexed semen, adjustments of time of AI regarding the onset of estrous can improve conception rates (Seidel Jr. et al. 1999).

For AI programs using sexed semen frequently females are inseminated based on estrous detection after natural estrus or synchronization with PGF2 α using two doses 14 days apart. Thus, insemination can be performed about 12 to 24 h after estrous detection (Seidel Jr. and Schenk 2008a, b) or about 80 to 82 h after the second administration of PGF2 α (Kurykin et al. 2007). Using heat detectors (radiotelemetry, Heat Watch[®]) in Jersey heifers, AI with sexed semen was performed at different times considering the onset of estrous (12–16 h, 16–20 h, 20–24 h, and 24–30 h). In this study, higher conception rates were reported for females inseminated 16–24 h after the onset of estrous (average 53.7%) compared those inseminated after 12–16 h (37.7%); however, similar conception rates were observed compared to those of females inseminated 24–30 h after the onset of estrous (45.5%) (Sá Filho et al. 2010a, b). Based on estrous observation, an intravaginal P4 device can also be used from days 0 to 8 in association with PGF2 α injection 24 h before device removal (Underwood et al. 2010a). This strategy facilitates the management of synchronization and estrus identification.

Other studies have also reported that using a high number of sexed sperm per insemination after estrous detection (2.1–4.2 million) in two stages with a 12 h interval, GnRH administration at the time of estrous detection (Sá Filho et al. 2010a, b), and the deposition site in the uterus (Kurykin et al. 2007) did not affect the conception rates in dairy heifers. On the other hand, conception rate is affected by number of services to which that female is submitted, first (55.3%a), second (46.1%a), or third services (34.8%b) (Sá Filho et al. 2010a, b), suggesting that fertility following AI with sexed semen tends to decrease with the number of services (DeJarnette et al. 2009). In this context, sexed semen is more appropriate for heifers (Butler et al. 2014; Seidel Jr. 2014), and its use is recommended in the first service postpuberty, followed by the use of conventional semen in the next AI. In addition, this strategy enables maximum fertility using sexed semen; it is possible to reduce dystocia (female calves cause fewer calving problems) and increase the proportion of females calves in heifers and primiparous around 65% (Weigel 2004).

Initially, low conception rates were reported for beef and dairy cattle submitted to TAI with sexed semen, but studies have contributed to promote satisfactory strategies. Heifers inseminated with sexed sperm between 55 and 56 h after CIDR removal and PGF2 α injection had a 34% pregnancy rate, whereas those submitted to TAI between 67 and 68 h after CIDR removal presented a 49% pregnancy rate (Seidel Jr. and Schenk 2008a, b). Lactating dairy cows submitted to OvSynch protocol (GnRH, PGF2 α , and GnRH) had similar pregnancy rates to TAI using a 10 million sexed sperm dose (43.9%), 2 million sexed sperm dose (40.5%), or 10 million non-sexed control sperm dose (55.6%) (Schenk et al. 2009).

Strategies have been developed to improve the reproductive performance of the herd after using sexed semen. The effect of time for AI (54 h after implant removal/16–18 h before ovulation versus 60 h after implant removal/10–12 h before ovulation) and semen [conventional (40 million sperm) versus sexed (2 million sperm)] were studied in 389 beef cows (*Bos indicus*) 30–60 days postpartum. In this study, conception rates were similar between TAI using sexed semen at 54 h (48.4%) and 60 h (55.1%) after norgestomet implant removal and among conventional

(58.9%), sexed X (52.0%), and sexed Y semen (49.0%). However, 6 hours delay in TAI increased by 9% the conception rate of animals inseminated with sexed semen [54 h (37.4%) versus 60 h (46.4%)]. These data suggest that the most appropriate time to perform TAI with sexed semen is 60 h after P4 device removal (10–12 h before ovulation) (Souza et al. 2008). The same experimental design was tested in dairy cattle (*Bos taurus*) using ovulation synchronization with CIDR-B insertion, 2 mg of EB and PGF2 α on day 0; CIDR-B removal and PGF2 α on day 8; 1 mg of EB on day 9; and TAI on day 10 (54 or 60 h after CIDR-B removal). Similar results were obtained, demonstrating that a 6 h delay in TAI with sexed semen also increased conception rates in dairy cattle (Sales et al. 2011a).

These increased conception rates observed when using sexed semen after 54 or 60 h of device removal or 10–12 h before ovulation is justifiable because sexed sperm require less time for capacitation due to the flow cytometry process (Lu and Seidel Jr. 2004). In addition, ultrasonography evaluation to determine the diameter of the dominant follicle at the time of TAI with sexed semen has improved pregnancy rates. Using sexed semen for TAI, an effect of the type of semen [conventional (45.4%) versus sexed (54.2%)] and of the diameter of the dominant follicle at TAI [≥ 11 mm (57.9%) versus <11 mm (44.1%)] was observed on the pregnancy rates (Sá Filho et al. 2012).

Currently, sexing sperm has achieved important advances and process has been performed based on next-generation technologies (SexedULTRATM - Sexing Technologies, Navasota, TX), which provides sex-sorted semen that is commercially available for dairy and beef cattle. This sex-sorted semen is presented at a concentration of 4×10^6 spermatozoa per straw. In addition, this semen includes adjustments to the composition of the medium that include the prestaining seminal treatment, modifications in the staining medium itself, and freezing extenders, which contribute to greater balance and pH maintenance for prolonged times (Vishwanath and Moreno 2018; Thomas et al. 2017; de Graaf et al. 2014).

Sex-sorting from next-generation technologies was compared with the different methodologies of sex-sorted semen in Nelore cattle submitted to the TAI protocol. For all cows, the TAI protocol was similar as demonstrated in Fig. 1.8. In this study, the pregnancy rate for TAI ($P < 0.0001$) is promising for the current sexing methodology used (Table 1.4) (Baruselli et al. 2017).

Another study conducted in Brazil (Marques et al. 2018) analyzed the conception rates after TAI with sex-ultra semen in suckled Nelore cows ($n = 281$). In addition to testing different doses (4, 6, and 8×10^6 sperm from a single Nelore bull), this study evaluated the rational use of semen according to estrus behavior (sex-sorted to cows in estrus and non-sorted to cows with no estrus expression) after P4 removal (Fig. 1.8). The conception rate was similar among the cows inseminated with 4 (59.5%, 47/79), 6 (58.4%, 45/77), and 8×10^6 sperm (51.9%, 40/77). Cows in estrus showed 56.7% (123/233) and those not in estrus showed 33.3% (16/48) conception rates. Interestingly, this study highlights a strategy with high efficiency for the use of sex-sorted semen in commercial field conditions. It emphasizes the positive results achieved (pregnancy rate higher than 50%), which can be attributed

Table 1.4 Pregnancy rate TAI in Nelore cows inseminated with conventional semen or different methodologies of sex-sorted semen

Nelore bulls	Methodology	Pregnancy rate % (n/N)
Conventional semen (20 × 10 ⁶ sptz)	Frozen semen without sexing	52.0 ^a (112/199)
Sex-sorted (2.1 × 10 ⁶ sptz)	Previous sexing methodology	28.2 ^c (58/206)
Sex-ultra (2.1 × 10 ⁶ sptz)	Current sexing methodology	37.6 ^{b,c} (72/191)
Sex-ultra (4 × 10 ⁶ sptz)	Current sexing methodology with enhanced concentration	43.0 ^b (86/200)

a–b: indicates statistical difference among different methodologies of sex-sorted semen
Adapted from Baruselli et al. (2017)

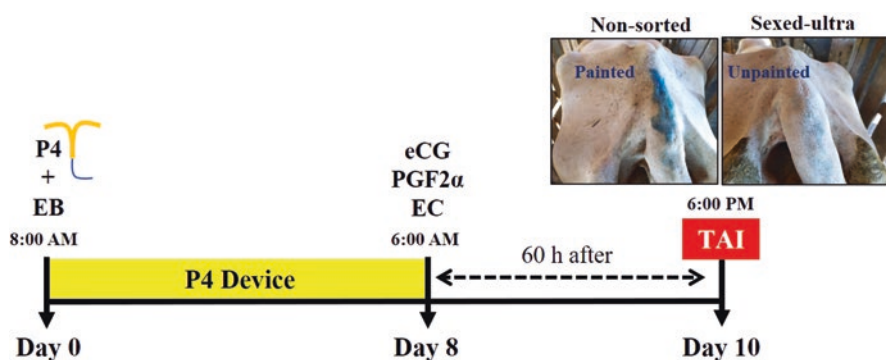


Fig. 1.8 Diagrammatic representation of the TAI protocol performed in Nelore cows inseminated with sexed-ultra (cows unpainted at the base of the tail/estrus expression) or conventional semen/non-sorted (cows painted at the base of the tail/no estrus expression). Treatment based on P4-estradiol and insemination performed 60 h after removal of the P4 device. P4, progesterone (0.6 g); EB, estradiol benzoate (2 mg); PGF2 α , cloprostenol (0.526 mg); eCG, equine chorionic gonadotropin (300 IU); EC, estradiol cypionate (1 mg). (Adaptations from Marques et al. (2018))

to the new methodology of sexed semen and to the fact that this semen was only used in cows that expressed estrus, as indicated by unpainted tails.

Another important consideration about sexed semen is related to the time when sexing is performed based on freezing. Conventionally, sexed semen preparation begins with semen collection from bulls followed by sperm sex sorting (X and Y chromosomes) and freezing. Currently also is possible to obtaining sexed sperm from previously frozen doses (reverse-sorted semen, RSS), which represents a breakthrough in livestock management due to the possibility of its use in association with other biotechnologies. In addition, there is no need for the bull to be close to the sexing laboratory and allows for the sexing of freezing semen from bulls that have died. In this context, this technique can also be used as an alternative in AI (Underwood et al. 2010b) or IVF programs (Morotti et al. 2014) to preselect the sex of the offspring produced.

1.8 Resynchronization Program

Resynchronization is a reproductive strategy that aims to concentrate the breeding period from females that failed to become pregnant after receiving the AI procedure. Using resynchronization, non-pregnant females are quickly diagnosed and inseminated in the shortest possible time, reducing the service period and making the breeding season shorter. This strategy is very interesting because it is associated with a higher pregnancy rate from AI, reduction in the number of bulls to natural breeding after TAI or AI program, in addition to several benefits associated with the management of calves (Baruselli et al. 2018; Marques et al. 2015).

Insemination is a worldwide established biotechnology that provides economic returns by improving offspring production rates. However, this return is closely linked to strategies that can be associated with using this technique. Among the main factors that affect the productive performance of calves until weaning, there is the interaction between the month of birth and geographic location, such as variation in the calf weight at weaning. For example, in cattle from tropical countries Nelore calves born from August to October had high weaning weights (Bocchi and Albuquerque 2004), possibly due to a higher concentration of calving in the late winter and early spring after conception during months of high food availability. Therefore, a short breeding season aims to improve progeny performance. Furthermore, calves whose conceptions occurred in the first half of the breeding season had increased production rates, such as earlier slaughter, earlier reproductive life, and their mothers had a higher pregnancy rate at the end of the breeding season (Carneiro et al. 2012).

1.8.1 Estrus Resynchronization

A wide variety of hormonal treatments can be used for synchronization of estrus and ovulation in cattle, and this strategy allows for inseminating all treated animals, resulting in conception rates of approximately 50% and a service rate of 100% (Baruselli et al. 2004a, b, 2018). Using an ovulation synchronization protocol, synchronous estrus return is expected in those females who failed to conceive after the first TAI. Commonly, this estrus behavior is concentrated between 18 and 23 days later, and reproductive practices can be strategically combined at this time (Baruselli et al. 2004a, b; Cavalieri et al. 2004). In this estrus return, conventional AI can be used, or bulls can be rationally allocated together with cows for natural breeding from 10 to 15 days after TAI (Campos et al. 2013). This management results in high conception rates at the end of the breeding season (Torres-Júnior et al. 2009). Generally, visual estrous detection in *Bos indicus* cows results in low service rates (25%) due to a short estrous duration (<12 hours) and high incidence of estrous occurring at night (30%) in Nelore cows (Pinheiro et al. 1998; Galina et al. 1996). Furthermore, postpartum anestrous usually has a high incidence in *Bos indicus*

cattle, especially if kept in regions with seasonal forages (Ayres et al. 2008). Therefore, difficulty in estrous detection in nonpregnant cows is a challenge for improving the number of pregnancies following AI because fewer than half of nonpregnant cows have become inseminated again (Campos et al. 2013).

The TAI program followed by estrous detection and subsequent AI usually results in 60 to 65% pregnancy rate in the first 45 days of the breeding season compared to 30% after natural mating and 20% after conventional AI. Failure in inseminating nonpregnant cows in a short period results in prolonged intervals among services and in delayed pregnancies (Baruselli et al. 2018; Marques et al. 2015; Campos et al. 2013; Galvao et al. 2007; Bartolome et al. 2005; El-Zarkouny, 2004). In this context, ovulation resynchronization is a strategy to increase the number of pregnancies in the first half of the breeding season. Therefore, the aim of ovulation resynchronization is to eliminate the estrous observation, increase the number of calves born by AI, and reduce the number of bulls needed.

1.8.2 Main Strategies for Ovulation Resynchronization

In general, hormonal treatments for ovulation resynchronization in cattle are similar to those previously described for the TAI protocol. One of the most common treatments for resynchronization involves the use of an intravaginal P4 device and EB, combined with application of PGF2 α , EC and/or GnRH (Sá Filho et al. 2014; Campos et al. 2013; Galvao et al. 2007; El-Zarkouny, 2004). This hormonal combination allows for synchronization of the emergence of a new wave of follicular growth in nonpregnant females. The ovulation resynchronization can be started after confirming that the female is not pregnant (traditional resynch/30 days after 1st TAI) (Marques et al. 2015) or before even knowing the female's gestational status (early resynch/22–23 days after 1st TAI and super-early resynch/14 days after 1st TAI) (Baruselli et al. 2018; Sá Filho et al. 2014; Campos et al. 2013).

Although TAI has shown satisfactory results, reductions of 15 to 20% in the conception rate are reported for cattle with high anestrus rate (Ayres et al. 2008; Baruselli et al. 2004a, b). In this context, administration of eCG in anestrus suckled cows has been recommended to provide pregnancy rates of approximately 50%, which is similar to those observed in cyclic cows (Barreiros et al. 2014; Kastelic et al. 1999). Therefore, the use of gonadotropic support (equine chorionic gonadotropin) or temporary calf removal (48–72 h) has been recommended at this time to final growth of the dominant follicle in resynchronization protocols of postpartum cattle (Campos et al. 2013).

Resynchronization 30 days (traditional resynch) was one of the first resynchronization strategies used, which consists in performed a pregnancy diagnosis from 28 to 32 days after the first TAI (Fig. 1.9). Therefore, only nonpregnant females are submitted to the second hormonal protocol, resulting in a 40-day interval between TAI (Marques et al. 2015; Bartolome et al. 2005). This resynchronization results in

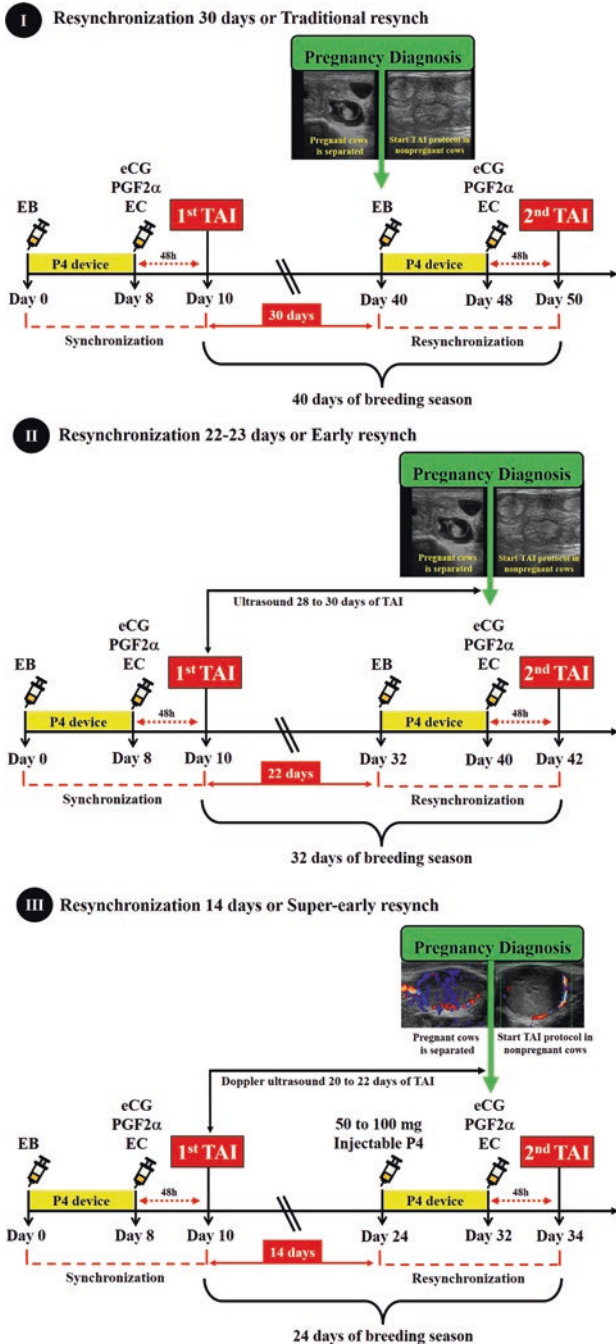


Fig. 1.9 Schematic representation of the protocol used for timed artificial insemination (TAI) and resynchronization of ovulation in nonpregnant cattle. I) Resynchronization 30 days/traditional resynch; II) Resynchronization 22–23 days/early resynch; III) Resynchronization 14 days/super-early resynch. *P4* progesterone (0.5–1.9 g), *EB* estradiol benzoate (2 mg), *EC* estradiol cypionate (0.5 mg for heifers and 1 mg for cows), *PGF2 α* prostaglandin (125–500 μ g), *eCG* equine chorionic gonadotropin (300–400 IU)

a conception rate of 50 to 60% for each TAI, and the final cumulative pregnancy rate can be over 80%.

Another possibility for ovulation resynchronization (early resynch/22–23 days after 1st TAI) is to start a second hormonal protocol 22–23 days after TAI without a previous pregnancy diagnosis. Therefore, all females initially treated for 1st TAI protocol receive a P4 intravaginal device and EB injection between 22 to 23 days regardless of gestational status (Sá Filho et al. 2014; Campos et al. 2013; Chebel et al. 2003). Upon P4 device removal (8–9 days from insertion), the pregnancy ultrasound evaluation is performed, and nonpregnant females are injected with PGF2 α and an ovulation inducer, such as EC (Fig. 1.9).

The resynchronization program 14 days after 1st TAI (super-early resynch) anticipates insemination at 8 days in relation to resynch 22–23 and 16 days in relation to resynch 30, making the breeding season extremely short and a 14-day interval between TAIs. This resynch is initiated 14 days after the first insemination, being that all females receive a combination of treatments with intravaginal P4 (device) and 50–100 mg of injectable P4, IM. Then, on day 22, the diagnosis of pregnancy is made by assessing CL vascularization with color Doppler. Cows with low or absent vascularization are considered nonpregnant and continue resynchronization treatment with device removal, PGF2 α , EC, and eCG, receiving TAI 48 h later (day 24). Cows with moderate CL or strong vascularity are considered pregnant and do not receive any hormonal treatment except reassessment with B-mode ultrasound (Baruselli et al. 2018).

The use of color Doppler ultrasonography to characterize the CL vascularization is essential for classifying female's gestational status (Hassan et al. 2018). After the luteolysis process (approximately 14 to 17 days of the estrous cycle), the pregnant female maintains high vascularization in the CL, and therefore, the Doppler evaluation allows an indirect assessment of gestational status from the CL vascularization score. This technique allows for a much earlier pregnancy diagnosis than B-mode ultrasound identification. Nevertheless, females with vascularized CL need to be reevaluated with B-mode ultrasound 8 to 10 days later to visualize the gestational vesicle and detect eventual false positives (females classified as pregnant by the CL vascularization score but without a gestational vesicle) (Baruselli et al. 2017; Pugliesi et al. 2017; Siqueira et al. 2013).

Reproductive programs that allow insemination every 21 days would be a more ideal because they would have a 100% (21/21) service rate. In this context, resynch at 14 days is the strategy that most closely approximates this ideal service rate, being 87.5% (24/21) to resynch 14, 66% (32/21) to resynch 22–23, and 52.5% (40/21) to resynch 30 (Baruselli et al. 2018). However, the implementation of super-early resynch is more laborious and, for success, requires an ultrasound with Doppler function in addition to high professional experience.

1.8.3 Success Rate Using Resynchronization

Satisfactory results are achieved with resynchronization programs in cattle, which results with pregnancy rates of approximately 50% after each TAI. These strategies result in pregnancy rates of 80 to 90% during days 24 to 40 of the breeding season (according to resynch), contributing to greater genetic gain of the herd (more pregnant from AI) besides reducing the number of bulls on the farm. In some situations, natural breeding (use of bulls) may be associated with resynchronization for 10–15 days after the 2nd or 3rd TAI until the end of the breeding season (Baruselli et al. 2018; Marques et al. 2015).

On the other hand, there is variation in conception according to the number of calving. Primiparous and secundiparous cows showed a reduction in the conception rate from the 1st to 2nd TAI (20%) compared to heifers and multiparous cows. This reduction is possibly associated with nutritional limitations and not the category’s own fertility. Conception rates among different categories of Nelore females were different (Fig. 1.10) in a study evaluating reproductive performance (Marques et al. 2015). At the end of the breeding season, heifers had a higher conception rate (85%) than those of primiparous (76%) and multiparous cows (78%). In this study, all animals received a source P4 (intravaginal device for the cows or ear implants for heifers) in combination with 2 mg of EB on day 0. All females received 250µg of cloprostenol, 300 IU of eCG, EC (1 mg to cows and 0.5 mg to heifers), and P4 removal on the eighth day. TAI was performed 48 hours after P4 removal. Thirty days after the 1st TAI, all females were evaluated by ultrasonography, and the nonpregnant bovines were resynchronized with the same hormonal treatment. The pregnancy rate was assessed by ultrasound 30 days after TAI.

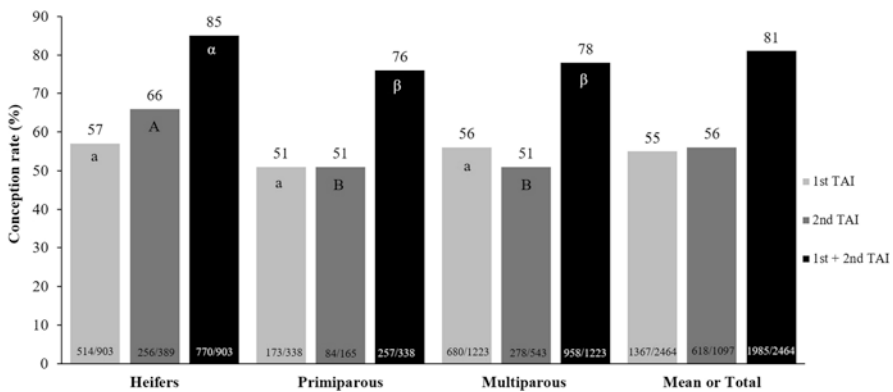


Fig. 1.10 Conception rates among different categories of Nelore females after the timed artificial insemination (1st TAI) and ovulation resynchronization (2nd TAI). (Adapted from Marques et al. (2015))

a-b: indicates statistical difference for categories in 1st TAI; A-B: indicates statistical difference for categories in 2nd TAI; and α - β : indicates statistical difference for categories in 1st and 2nd TAI

Early resynch anticipates the 2nd TAI in 8–9 days, but both pregnant and non-pregnant cows receive a P4 device and 1–2 mg of EB between 19 and 23 days after the first TAI. In addition, to promote follicular atresia and synchronize follicular growth, estradiol esters can induce luteolysis (Vieira et al. 2014; Pinheiro et al. 1998). However, studies with *Bostaurus* (El-Zarkouny 2004; Chebel et al. 2003) and *Bos indicus* (Sá Filho et al. 2014; Campos et al. 2013) females reported no interruption of pregnancies obtained in the 1st TAI using GnRH or EB in the early resynchronization. There was no reduction in the conception rates in Nelore cows treated (54%) or untreated (47.8%) with 2 mg of EB 23 days after TAI (Campos et al. 2013). In addition, pregnancy losses were similar in Nelore females treated with 1.0 mg of EB (4.1%) compared to untreated females (2.0%) and maintained with bulls for natural mating (Sá Filho et al. 2014). Conception rates are similar to those obtained with resynch 30 days, as shown previously, resulting in approximately 50% in the 2nd TAI and a total of 75% of conception in 32–33 days of service (Sá Filho et al. 2014; Campos et al. 2013). In this resynch, variations in conception rates were also observed according to number of calving. Nonsuckled cows, heifers, and postpartum cows have a decrease of 20% in conception rates. Furthermore, primiparous cows showed a similar conception rate (40%) in two subsequent (Marques et al. 2015; Crepaldi et al. 2014; Sá Filho et al. 2014). In cows that did not become pregnant after the 1st TAI, a 2nd TAI (resynchronization of ovulation) may be performed 32 days later, and a 3rd TAI may be repeated, resulting in a total conception rate of 90% after three services in 64 days of a breeding season (Baruselli et al. 2018).

A similar pregnancy rate using resynch 22 versus resynch 14 was observed in the 1st synchronization (48% versus 53%) and resynchronization (56% versus 51%), respectively. Cumulative pregnancy rates after 32 and 24 days of the breeding season were also similar for resynch 22 (77%, 97/126) and resynch 14 (75%, 89/118). However, resynch 14 improved the service rate every 21 days with 66% for resynch 22–23 and 87.5% for resynch 14 (Baruselli et al. 2017).

In general, reduced conception rates are observed in the 2nd and 3rd TAI. In addition to the number of calving, it is necessary to consider other factors, such as BCS and health status of the herd. Reduction in the period of service for 62 to 80 days using three TAI provides a short calving interval and a considerable improvement in reproductive efficiency (with an average calving interval of less than 12 months). Pregnancy rates between 75 and 90% were reported for cattle submitted to resynchronization, reducing the numbers of bulls needed for natural mating or even not requiring them at all.

1.9 Dominant Follicle Manipulation During the TAI Protocol

The size of the dominant follicle at the time of TAI has been extensively investigated due to its influence on the reproductive behavior and performance of animals submitted to synchronization of ovulation. Although the TAI protocol aims for ovulation synchronization and does not require estrus detection, many studies have

investigated the relationship of this behavior, diameter of dominant follicle and the fertility of animals undergoing TAI programs (Moraes et al. 2019; Morotti et al. 2013a, b, 2018a; Pfeifer et al. 2012, 2015; Sales et al. 2012a, b; Sá Filho et al. 2010a, b, 2011). In this context, dominant follicle size at the time of insemination has been widely studied in which the largest follicle diameter at TAI is positively associated with higher estrus expression, higher ovulation rate, larger size of CL, higher concentration of P4, and higher pregnancy probability (Moraes et al. 2019; Morotti et al. 2018a; Pfeifer et al. 2015).

1.9.1 Estrus Expression and Fertility in TAI

Considering that the follicular diameters may differ at the time of TAI and that this may determine a higher or less ovulatory potential according to follicle size (Gimenes et al. 2008), block TAI has been suggested as a possibility to improve fertility of females undergoing a TAI program. Block TAI can be applied as a strategy to increase the pregnancy rate in Nelore cows (Pfeifer et al. 2015). On day 10 of the TAI protocol, the blocks are divided according to the size of the dominant follicle; in this way, TAI is performed at different times in each group (Table 1.5). Performing TAI in a block of Nelore cows increased the pregnancy rate by 16.7% (65.5%; 129/203) compared to the group subjected to conventional TAI on day 10 (48.8%; 102/209; Pfeifer et al. 2015).

For better understanding of the relationship of dominant follicle size at TAI, estrus expression, and fertility, an estrus score can be performed based on the painting intensity remaining at the base of the tail. Normally, at the removal of the P4 device, all animals have the sacral and tail base region painted with marker stick, and at the time of TAI, the animals are classified into different scores (Nogueira et al. 2019). According to the proportion of paint removed, animals can be classified into (Fig. 1.11): score I, no or low paint removal (no estrus expression), score II, partial paint removal (up to 75%; low estrus expression), and score III, complete or > 75% paint removal (high estrus expression). This practical strategy allows for usefully identifying cows with greater estrus expression and consequently

Table 1.5 Pregnancy rate in cows inseminated either in blocks or conventionally according to the diameter of the dominant follicle at timed artificial insemination (TAI)

Group	Follicular diameter [#]	TAI timing	Pregnancy rate* % (n/N)
Control	Conventional TAI	48 h after P4 removal	48.8 (102/209)
Block TAI	> 15 mm	0 h	63.5 (129/203)
	13.0–14.9 mm	6 h later	
	10.1–12.9 mm	24 h later	
	≤ 10 mm	30 h later	

[#]Diameter of dominant follicle 48 h after P4 device removal (day 10)

* P-value <0.01

Adapted from Pfeifer et al. (2015)

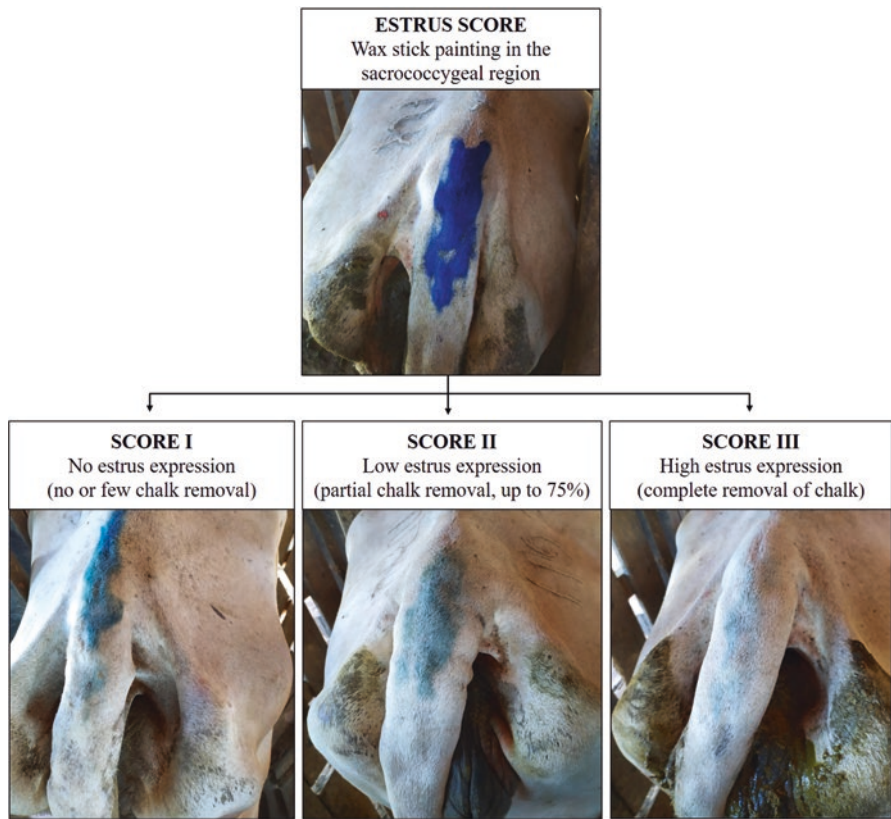


Fig. 1.11 Scheme of the estrus expression intensity score during timed artificial insemination (TAI) using wax stick painting in the sacrococcygeal region. Commonly, paint management is performed at the time of removal of the intravaginal progesterone device and the estrus score classification performed immediately before insemination. Cows with high estrus expression are conventionally inseminated and low estrus expression or no estrus expression receive insemination together with GnRH analogue application to potentialize the occurrence of ovulation

improving the pregnancy rates in TAI (score I, 40.0%; score II, 49.7%; and score III, 60.9%), allowing the cows with low score to be targeted for additional treatments aimed at improving pregnancy.

Even using TAI, the classification of the estrus score is interesting because scores I and II present lower reproductive performance. Based on this it was suggested application of 100µg of gonadorelin (GnRH group; n = 470) against 1 mL of saline (control group; n = 467; Rodrigues et al. 2019) to animals with I and II scores. Cows with score III (estrus group; n = 1347) received no additional treatment. The pregnancy rate was greater in the estrus group (57.09%; 769/1347) than in the control (36.18%, 169/467) and GnRH groups (45.95%, 216/470). However, GnRH injection increased pregnancy by approximately 10% in relation to the control. Therefore, using P4-estradiol-based TAI protocols, estrus expression can be efficiently monitored with

painting on the sacrococcygeal region. Thus, GnRH application in cows with a low estrus score is a simple strategy that can increase the pregnancy rate in cattle.

1.9.2 Antral Follicle Count and Fertility in TAI

The antral follicle count (AFC) is a highly variable reproductive characteristic among the females of a herd, but with very high repeatability throughout the productive female life (Lima et al. 2020; Moraes et al. 2019; Jimenez-Krassel et al. 2017; Burns et al. 2005). For this reason, the AFC has become an interesting reproductive tool since it is closely related to the efficiency of reproductive biotechniques, in addition to the great possibility of affecting female fertility (Garcia et al. 2020; Morotti et al. 2018a, b). In general, AFC can be determined in a simple way through the ultrasound exam to quantify the number of follicles ≥ 3 mm in diameter and usually is performed on a random day of the estrous cycle of the cows. Therefore, considering the AFC variability, cows can be classified as low, intermediate, or high AFC according to the number of antral follicles present in the ovary during ultrasound evaluation (Silva-Santos et al. 2014; Burns et al. 2005).

The evaluation of the AFC is inserted in a very current context of the selection of females and that seems to be better established for the production of embryos (Garcia et al. 2020; Seneda et al. 2019; Santos et al. 2016; Silva-Santos et al. 2014). For example, for both *in vivo* production and *in vitro* production, the selection of donors with high AFC is positively associated with better reproductive performance in these biotechniques (Zangirolamo et al. 2018; Santos et al. 2016). However, relation between AFC and efficiency of the TAI program is not yet fully understood (Morotti et al. 2018a, b). In taurine cattle there are studies that show that AFC does not exert any influence, or it has been observed that high AFC is responsible for better reproductive performance (Evans et al. 2012; Ireland et al. 2010). However, the authors of this chapter have conducted a series of studies on this subject in zebuine cattle which reveal that low AFC appears to have better reproductive performance when subjected to TAI.

In this context, evaluating the ovarian follicular dynamics and fertility of Nelore cows submitted to TAI programs, Morotti et al. (2018a) revealed higher follicular diameters (Fig. 1.12) and pregnancy rates for females with low AFC compared to those with high counts (Table 1.6). In addition to these findings, other studies have also revealed an interaction of AFC with body condition score in *Bos indicus* (Moraes et al. 2019) and higher reproductive longevity and greater reproductive performance in *Bos taurus* with low AFC (Jimenez-Krassel et al. 2017).

In a study on the relationship between AFC and reproductive performance, and the pattern of expression of important genes for various cellular functions in *Bos indicus* cattle (Lima et al. 2020) showed promising data with positive influence of low AFC on fertility. In this study we found that very low AFC in Nelore cows resulted in a large dominant follicle diameter, a tendency to have higher P4 concentration and greater pregnancy rate in TAI program. In addition, Nelore heifers with

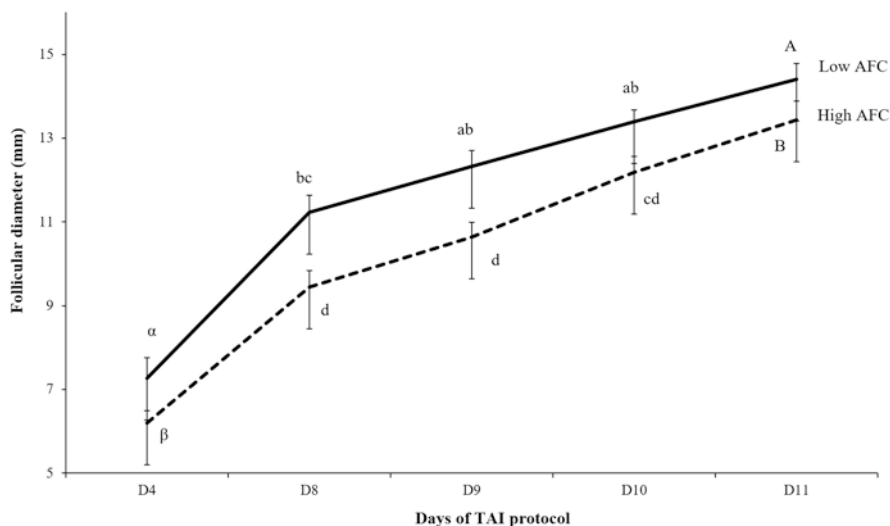


Fig. 1.12 Diameter of dominant follicle (mm) during ovarian follicular dynamics in Nelore cows with consistently high (≥ 45 follicles; dashed line) or low (≤ 15 follicles; continuous line) antral follicle count synchronized with a timed artificial insemination (TAI) protocol. Values denoted using different Greek letters (α - β ; punctual evaluation on day 4), lowercase letters (a-d; evaluations with 24-h intervals) or capital letters (A-B; evaluations with 12-h intervals) were different ($P < 0.05$). (Adapted from Morotti et al. (2018a))

Table 1.6 Mean (M) and standard deviation (SD) of the antral follicle count (AFC) and conception rates from Nelore cattle with consistently high, intermediate, or low AFCs following a TAI protocol in two different studies

Studies	AFC groups	AFC	Conception rate
		Mean \pm SD	% (n)
Morotti et al. (2018a)	Low (≤ 15 follicles)	11.3 \pm 2.8 ^c	61.7 ^a (150/243)
	Intermediate (20–40 follicles)	29.4 \pm 6.0 ^b	52.9 ^b (210/397)
	High (≥ 45 follicles)	52.8 \pm 7.7 ^a	49.5 ^b (96/194)
	P-value	0.001	0.027
Moraes et al. (2019)	Low (≤ 10 follicles)	7.7 \pm 2.6 ^c	57.7 ^a (176/305)
	Intermediate (11–29 follicles)	18.0 \pm 3.4 ^b	49.7 ^b (155/312)
	High (≥ 30 follicles)	38.0 \pm 12.4 ^a	47.9 ^b (57/119)
	P-value	<0.0001	0.008

Values with different superscripted letters (a, b) were different ($P \leq 0.05$) between the AFCs and conception rates

low AFC exhibited oocytes and cumulus cells with a better expression patterns of genes linked to intercellular communication, meiotic control, epigenetic modulation, adaptation and cellular stress response and follicular growth. However, studies on the ovarian follicular population are current, it is a subject that has been widely investigated, and many aspects are not yet fully understood.

1.10 Considerations

Currently, reproductive biotechniques have achieved great technological advances that have contributed greatly to increasing the development of livestock. Artificial insemination certainly represents one of the most popular assisted reproductive techniques that contributes significantly to genetic improvements, being easy to apply and has been considered one of the greatest potentials for expansion in cattle. In this context, many strategies have been developed to stimulate the use of artificial insemination programs. Alternatives to pharmacological control of the estrous cycle, the use of timed artificial insemination, insemination with sexed semen, ovulation resynchronization, and dominant follicle manipulation, are highly effective strategies that are currently indicated as reproductive practices in beef and dairy cattle. Finally, maintaining these advances and utilizing the strategies discussed herein are great challenges but are necessary to increase the productive and reproductive efficiency of livestock.

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Chapter 2

Reproduction Management and Artificial Insemination in Dromedary Camel



Djallel Eddine Gherissi and Ramzi Lamraoui

Abstract The dromedary camel is a unique multipurpose species under extreme arid conditions. It is used for production, leisure, transport and agricultural work. Its profile as a production animal has given it major importance in ensuring the sustainability of marginalized communities in the poorest regions of the world, but also because of the nutritional and therapeutic properties of its milk and meat. Nowadays, the exploitation of this animal resource known an orientation towards intensification under the pretext of the expanding demand on camel products. The productivity of camel herds is conditioned by its reproductive potential. Often, camel herd performances are limited by low fertility rate, high inter-calving intervals, late puberty leading to low longevity, low milk production and unavailability of young calves for herd renew and fattening farms. To reach an optimal level of camel herd's numerical productivity and therefore accelerate the genetic progress, it is imperative to reduce the duration of the unproductive periods, namely the waiting period and the reproduction period. This involves through the choice of a suitable farming system, improving breeding practices, improving data recording and reproductive monitoring, application of strategic reproduction control and new reproduction techniques.

Here we review strategies of camel herd reproduction management and means to improve camel herd reproductive performances using sexual activity control in males and females and artificial insemination practice. Controlling ovarian cycles of the female camel consists of controlling follicular growth and timing of ovulation. Therefore, this technique can be used to induce and synchronize the ovarian function during a favorable reproduction period or even during the non reproductive

D. E. Gherissi (✉)

Institute of Agronomic and Veterinary Sciences, University of Souk-Ahras,
Souk Ahras, Algeria

Laboratory of Animal Productions, Biotechnologies and Health, University of Souk-Ahras,
Souk Ahras, Algeria

R. Lamraoui

Institute of Agronomic and Veterinary Sciences, University of Souk-Ahras,
Souk Ahras, Algeria

Department of Biology of Living Organisms, Faculty of Natural and Life Sciences,
University of Batna, Batna, Algeria

season. Zootechnical means do not really make possible to synchronize the ovarian cycle in camels. They are used as complementary tools to the pharmacological methods. The different hormonal based protocols are: melatonin, progesterone, prostaglandin F₂ α and GnRH. Artificial insemination in dromedary camel is a topic of contemporary research that tends to develop standard sperm collection protocols, monitoring of sperm quality, short-term preservation, cryopreservation and semen thaw with considerable interest to increase the pregnancy rate and therefore to market this technology in camels for large-scale use. Various genetic, sanitary and economic advantages could be insured by this biotechnology and overcome many problems in regard with camel fertility efficiency such as short breeding season, long gestation period, traditional reproductive management and widespread of genital and venereal infectious diseases.

Keywords Artificial insemination · Oestrus synchronization · Dromedary camel · Camel bull · Breeding · Reproduction performance · Sperm · Follicular cycle · Female camel · Calving interval

Abbreviations

°C	<i>degrees Celsius</i>
AF	<i>annual herd fertility</i>
AI	<i>Artificial insemination</i>
AV	<i>artificial vagina</i>
BLUP	<i>Best Linear Unbiased Prediction</i>
CASA	<i>Computer Assisted Sperm Analysis</i>
CFM	<i>Calving to first mating interval</i>
CI	<i>Calving interval</i>
CIDR	<i>Controlled Internal Drug Releasing</i>
COI	<i>Calving-oestrus interval</i>
DC	<i>dromedary camel</i>
eCG	<i>equine chorionic gonadotropin</i>
EDTA	<i>Ethylenediaminetetraacetic acid</i>
FAO	<i>Food and Agriculture Organization</i>
GnRH	<i>gonadotropin releasing hormone</i>
GPG	<i>GnRH-PGF₂ α-GnRH</i>
h	<i>hour</i>
hCG	<i>human chorionic gonadotropin</i>
IIED	<i>International Institute for Environment and Development</i>
ILRI	<i>International Livestock Research Institute</i>
IMV	<i>Institut de médecine vétérinaire</i>
INRA	<i>Institut national de la recherche agronomique</i>
LH	<i>luteinizing hormone</i>
M1, M2	<i>Mating1, Mating 2...</i>

mA	<i>ampere</i>
min	<i>minutes</i>
ml	<i>millilitre</i>
NGF	<i>nerve growth factor</i>
OD	<i>Open days</i>
OIF	<i>ovulation inductor factor</i>
PGF2 α	<i>Prostaglandin F2α</i>
PHT	<i>Progeny History Testing</i>
PRID	<i>Progesterone Relasing Intravaginal Device</i>
RP	<i>reproduction period</i>
SeNPs	<i>Selenium nanoparticles</i>
SSO	<i>small and smooth ovaries</i>
V	<i>volt</i>
WP	<i>waiting period</i>
ZnONPs	<i>Zinc oxide nanoparticles</i>

2.1 Introduction

The dromedary camel (DC: *Camelus dromedarius*) is the best livestock animal well adapted to arid ecological conditions. It fights against desertification by using a high floristic biodiversity with high digestibility and better food efficiency. In addition, its low gregarious outpatient pasture and long intestinal transit contribute for a better germination of the seeds after excretion. Camels are multipurpose animals well adapted to harsh environment. They are used for nutritional, dietetic, economic, social and cultural purposes. They offer milk and meat production, hair and are used for racing, transportation and tourism (Al Abri and Faye 2019). The nutritional value of camel meat has been shown to be much higher than beef, lamb and goat because it contains less fat with a comparable index of essential amino acids (Raiymbek et al. 2015). Camel meat is rich in essential amino acids such as leucine and lysine, essential fatty acids such as omega 3, minerals, vitamins and bioactive compounds such as carnosine, anserin, glutathione, etc. (Kadim Isam et al. 2018). Camel milk is 3–10 times richer than cow's milk in vitamin C. (Faye 1997; Konuspayeva et al. 2009; Al Abri and Faye 2019). Its hypo-allergic property is due to its low content on β -casein and β -lactoglobulin (Konuspayeva et al. 2009). It's whey protein fractions are higher (El-Agamy 2006). So, dromedary camel contributes to the subsistence of transhumant and nomadic Saharan communities, often with restricted access to various protein resources of animal origin. Thus, it constitutes for them an element of food security in the face of climate change, a long-term capital ensuring the well-being of the family and an element of social prestige. Finally, the dromedary participates in sports and leisure activities and represents a cultural heritage in many countries and contributes in socio-economic activities and

diversification of many marginalized area of arid areas worldwide (Al Abri and Faye 2019).

Poor reproductive performance in camel species is mainly due to a low fertility rate and increased inter-calving interval. The low numerical productivity of camel herds is often discussed among cameleers with adversarial opinions in regard to its aspects and its risk factors. However, the veterinarian should provide advice based only on specific information with reference to herd reproductive history. The serious difficulty often arises is about the reproduction evaluation in DC herds is often confronted to the following constraints: a pastoral breeding system in continuous displacement, traditional breeding practices and lack of recording information related to reproduction, absence of state veterinary services interventions on reproduction associated to difficulties to reach remote breeding areas for regular monitoring by private veterinarians, lack of appropriate training for health and management of camel herds. The diagnosis and treatment in this animal are generally approached in the same way as for other farm animals.

The reproduction control in camel species implements technical procedures or herd management strategies that make possible optimizing reproductive performances, particularly during anoestrus period (seasonal, post-partum) taking into the count the breeding systems of the species. It aims to reduce the rate of infertility, choice of calving period, calving synchronization, and decrease of unproductive periods (advancement/synchronization of puberty, decrease of anoestrus length, mating and conception during the non rutting season). Efficient methods to reproduction control in the camel are required as the interest grows in the potential application of artificial insemination and embryo transfer as tools for improving genetic traits, such as milk, meat and wool production and racing ability (Al Eknah 2000).

Compared to other farm animals, little research on camel reproduction control and artificial insemination has been found in the camel side. Oestrous synchronisation solves oestrous detection problems and makes artificial insemination more effective (Helmy 1991; Minoia et al. 1992). In female camels, ovulation may happen through intrauterine semen deposition without the need for coitus (Chen et al. 1985; Musa et al. 1990). This makes AI a reliable technology for continuous genetic advancement using males of high genetic potential and inseminating female groups that have undergone synchronization of follicular development. However, the large-scale use of this biotechnology faces many difficulties, such as semen selection, male camel sexual activity, heterogeneity of recovered semen quality (small volume, low sperm concentration and high viscosity) (Skidmore et al. 2013), limited knowledge of semen storage, optimum insemination time and sperm dose (Al-Bulushi et al. 2018).

2.2 Reproduction and Genetic Improvement of Dromedary Camels

The DC (*Camelus dromedarius*) is known for its rather long reproductive cycle characterized by a long prepubertal period, long gestation period, long interval between generations and a low fertility rate (Gherissi et al. 2020a). Likewise, the incidence of genital pathologies, abortions and neonatal mortality are high (Gherissi et al. 2019, 2020a, b). All these factors together lead to a low numerical productivity of camel herds, particularly in extensive breeding system (Faye 2018; Brigitte 2005).

Genetic improvement according to the classical method involves the animal identification, assessment of their performance, selection of breeders and use of breeders in order to achieve genetic progress. In fact, genetic improvement can only take place if performance and pedigree are recorded. Selection would be effective when it is carried out following estimation of breeding value using progeny (BLUP: Best Linear Unbiased Prediction) widely practiced in various animal species.

$$BV_p = 2 \left(\frac{n}{n+k} \right) [P(\text{progeny}) - CA]$$

$$k = \frac{4-h^2}{h^2} \qquad r^2 = \frac{n}{n+k}$$

BV_p: breeding value of the parents, *r*²: reliability, *n*: number of offspring (informants), *P* (progeny): Mean value of the performance of the progeny, *CA*: Comparison average which characterizes the environment.

According to the previous formula, the precision of the selection depends closely on the quality and quantity of the individual controls of the available informants and the herd size used to calculate comparison average of the performance. Thereby, the low reproduction levels within camel herds can generate serious difficulties in order to give to appreciate with great reliability the genetic potential of the parents who will serve as reproducers (Emami Mibody et al. 2016). Contrariwise, good herd reproduction statue make easier to appreciate genetic value and animals with the highest predicted value can be selected as parents with minimum error.

On the other hand, genetic progress level depends on the variance of the genetic value in the population, the intensity of selection, the reliability of the estimated breeding value and the generation interval. In this regard, Al-Sobayil et al. (2006) and Almutairi et al. (2010) showed that camels species have a high genetic variability reflected in the heritabilities of various traits (body weight and growth rates were moderate to high; *h*² = 0.24–0.40, birth weight *h*² = 0.37, Daily gain ranged *h*² = 0.25 and 0.49, milk yield at 305 days *h*² = 0.24 and test day yields *h*² = 0.22). This reflects a potential for ample genetic gain if systematic selection is to be implemented (Al Abri and Faye 2019; Bahbahani et al. 2019).

$$GP \text{ per year} = \frac{S_A \cdot i \cdot r_{\widehat{VE,VE}}}{t}$$

GP: genetic progress, S_A : variance of the genetic value in the population, i : the intensity of selection, $r_{\widehat{VE,VE}}$: reliability of the estimated breeding value t : generation interval.

The selection intensity was indicated by the proportion of animals required for the next generation to be parents. Reproductive efficiency and management of camel herds therefore have an important influence on the rate of parents necessary for the next generation to be produced and, therefore, on the rate of genetic progress. The generation interval in this species is long (approximately 8 years) and is highly dependent on genetic potential, environmental conditions, husbandry systems and breeding methods. A higher reproduction rate for a given population size means a lower number of breeding animals and, therefore, a higher selection intensity. A greater number of offspring per breeder also promotes a more accurate estimate of genetic values (Emami Mibody et al. 2016). Another benefit of increased reproductive rates is the faster dissemination of superior genetic material. In this context, reproduction control, in particular using reproductive biotechnologies, can have a direct effect on increasing the rate of genetic progress of camel herds.

Artificial insemination (AI) leads to high reduction in the use of breeding males. In dromedaries, male and female camels with high genetic values could constitute a nucleus of selection of less than 1% of the entire population. AI increases the available information on siblings and therefore increases the genetic value accuracy estimated in their parents (Van Arendonk and Bijma 2003). Furthermore, AI allows increasing the selection intensity and precision of males based on the progeny test especially for traits with low heritability (such as functional traits). The semen exchange between different herds allows for the establishment of genetic links between them and reduces the risk of venereal diseases transmission between herds. Sperm freezing gives organizations and AI centers ability to create camel gene banks as back-up stores of camel genetic diversity. In addition, cryopreservation would facilitate the exchange and transport of semen and the international exchange of camel's genetic material.

AI requires availability of technical skills in AI centers and camel farms with effective means of communication between these two stakeholders. However, in many camel breeding countries, financial means are lacking and the structuring of camel breeding is inadequate for setting up successful AI operations. In addition, most camel herders are reluctant to camel breeding intensification and prefer mobile and pastoral systems which make difficult to follow the follicular cycle and implement timed AI at large scale. Therefore, AI is probably not predictable for livestock in extensive grazing systems.

2.3 Reproductive Cycle of Female Camels

The female camel is strict monotocus and seasonal polyoestral breeder (Akral and Khanna 1995). During the non breeding period the sexual behavior disappears but the follicular waves persist at low growth levels (Gherissi et al. 2018) similar to what happen in ewe (Dogan et al. 2020) but different to mare who undergoes a deep ovarian rest in the non breeding season (Boeta et al. 2006). Ovulation is induced by GnRH-like factor or Ovulation inductor factor (OIF) contained in camel bull seminal plasma which is identified as nerve growth factor (bNGF) (Sanjay et al. 2012; Bogle et al. 2012) also with luteotropic effect on the corpus luteum (Silva et al. 2014). This molecule induces pre-ovulatory luteinizing hormone (LH) surge 2–3 h after mating (Marie and Anouassi 1986) followed by ovulation (32–40 h after copulation) and subsequent formation of corpus luteum (El-Allali et al. 2017). Without mating at the breeding period (mature follicle of 11–19 mm); female's exhibit repeated cycles of follicular growth and regression without a cyclic corpus luteum. Three main phases have been described for follicular wave in female camel (Fig. 2.1): growth, maturity and regression. After emergence of small follicles

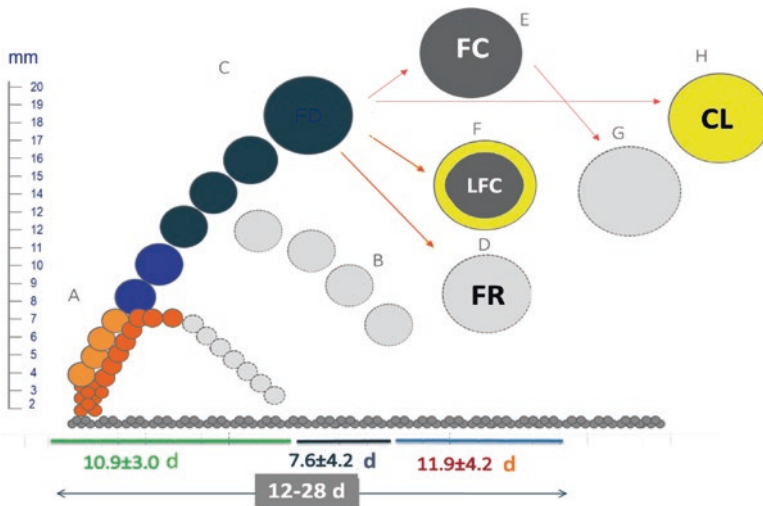


Fig. 2.1 The main follicular cycle phases in female camel (*FC* follicular cyst, *LFC* luteinized follicle cyst, *FR* dominant follicle regression, *CL* corpus luteum)

Following this first recruitment phase, several follicles (3–6) continue to grow (5–17 mm) until the emergence of one or two large dominant follicle which becomes mature; including 5–10 mm medium size follicles and 11–17 mm pre-ovulatory follicles. The dominant follicle undergoes a stagnation stage for 5–7 days. In 50% of unmated females, regression occurs until disappearance in the ovarian stroma. In other unmated females (50%) the largest follicle continues to grow attaining a large size (30–65 mm); this cyst-like follicle cannot ovulate spontaneously however, it regresses thereby permitting the expression of the next follicular growth waves in an average of 18 days later. In mated females, following ovulation, the mature follicle is usually transformed into a functional corpus luteum. (Adapted from Zarrouk et al. 2003)

(<4 mm in diameter) at the periphery of the ovaries during the recruitment phase, several follicles (3–6) continue to progress as medium size follicles (5–10 mm) until the dominance of one or two large follicles (11–19 mm) that become(s) mature pre-ovulatory follicle(s). The growth phase (<4 mm–10 mm) extend over 10.9 ± 3.0 days and dominance phase lasts over 7.6 ± 4.2 days (Skidmore et al. 1994).

Follicular regression occurs in 50% of unmated females until total desperation of these follicles in the ovarian stroma. The largest follicle tends to develop in other uncoupled females (50%) to reach an oversized follicle (>25 mm), which constitutes a follicle cyst that decreases ovulatory response (Tibary 2018). Presence of 2 or more co-dominant follicles in camelids is not rare and can occur in up to 40% of the follicular wave (Manjunatha et al. 2015; Campbell et al. 2015). A combination of stimuli, including the chemical factor in the seminal plasma, neurohormonal responses to the mechanical stimuli of coitus and the male effect, could result in the ovulatory response in the camel (Al Eknah 2000).

The anovulatory hemorrhagic follicles cysts seems to be the most frequent among ovarian cysts in female camels (Gherissi et al. 2019; Tibary 2018). The pathophysiology of anovulatory hemorrhagic follicles is poorly understood and their presence seems to not disturb follicular wave patterns (Tibary 2018). The follicular cyst or the corpus luteum of the non pregnant females regresses 18 days and 11 days later, respectively (Fig. 2.1), allowing the next waves of follicular growth to begin (Zarrouk et al. 2003; Marie and Anouassi 1987). The total length and the seasonal frequency of follicular wave in the female camel was found to vary considerably (17–28 days and maintain or reduced frequency of follicular waves) according to the geographic location, nutrition statute, start and end of the breeding season, body condition, photoperiod (Abdoon 2001; Vyas et al. 2004; El-Allali et al. 2005; Sghiri and Driancourt 1999; Gherissi et al. 2018, 2020b). In mated females, the ovulated follicle is usually transformed into a functional corpus luteum and its maintenance is conditioned by the suppression of PGF 2α release before day 10 after ovulation (Skidmore et al. 1998).

2.4 Camel Herd Reproductive Performance Evaluation

2.4.1 Data Collection for Reproduction Analysis

Obtaining individual reproductive history is integral to an accurate evaluation of reproductive herd performances and to explain eventual difficulties that arise there. Regarding the reproductive particularities under pastoral camel breeding conditions, a method based on data collection from breeder's offspring history called "Progeny History Testing" (PHT), has been suitably adapted to camel reproduction investigations. This technique was developed firstly in Ethiopia by a team from the Ministry of Agriculture, and then widely used by the International Livestock Research Institute (ILRI). PHT method was approved by the International Institute

for Environment and Development and the World Food and Agriculture Organization (IIED and FAO) in their data collection guides for different farming systems.

Under semi intensive management system, identification of all the animals in the herd using the most relevant means to camel species is essential. The breeding management in general and reproduction in particular must be done in compliance with certain preliminary conditions relating to the animal and its environment leading to monitoring and data collection by the breeder himself and stakeholders, in particular by taking observations, clinical examinations and preventive or curative treatments. The observations recording in semi-intensive camel farms is a concept that is absent in almost all camel breeders. This preliminary practice for the evaluation of the reproductive status of farms is fundamental and must be encouraged by using clear or even cryptic language. A mutually agreed system of abbreviation ratings can be very useful. Such information is manifold; they are particularly linked to calving (day, number, nature, type, interventions, etc.), sexual cycle (postpartum cycle number, number of cycles used, etc.) mating (days, duration of sexual receptivity, mating time, mating after induced or natural oestrus), pregnancy diagnosis (day, nature, result), body condition score, pathologies (absence of oestrus, ovarian and uterine pathologies, infections, embryonic mortality, mastitis, etc.). All observations must always refer to the identity of the animal. It will be accompanied by the date or even the time of the observation as well as the nature of the observation. The use of recording, scoring and organizing observations for processing is highly recommended. The most modest would be a simple farm management camel breeding calendar. Computerized systems adapted to camel breeding could emerge, which would effectively improve the management of several other aspects related to reproduction such as milk production, nutrition, etc.

2.4.2 Reproduction Indicators of Camel Herds

2.4.2.1 Calving Interval (CI)

The CI is the most important criteria to look at when talking about reproduction of the camel. The lengthening of this parameter leads to increase unproductive period (even drying period). It also decreases the herd numerical productivity by decreasing offspring production per year. Finally, it increases reform rate and herd replacement cost. Therefore, camel herders have every interest in the fact that their female camels give births often with the shortest intervals throughout their whole life.

In terms of herd performance; the interest is to measure the average of this parameter over the entire present females in order to identify the multiple individual and collective reasons for its lengthening/shortening. It is possible to shorten the CI of females by acting on waiting period (WP: time between calving and first mating) and reproductive period (RP: time between the first mating and the conception) (Fig. 2.2). WP can be shortened by rigorous peri-partum management with particular attention to late pregnancy and parturition (preparation, obstetric assistance and

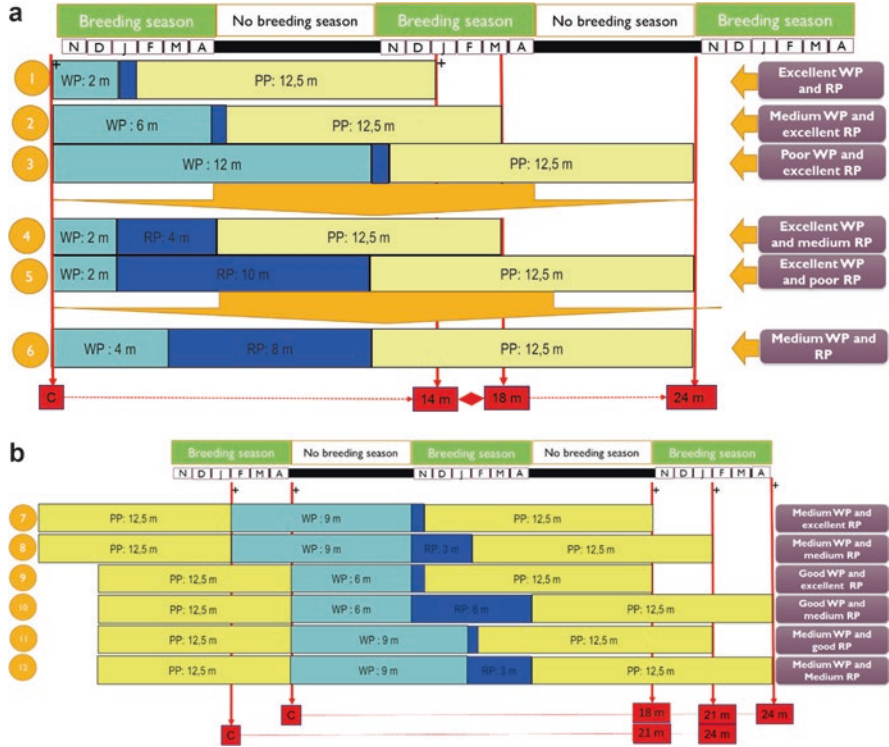


Fig. 2.2 Potential factors responsible for calving interval (CI): 18 months as objective and 24 months as threshold when calving occur at the beginning (a) or the middle/end (b) of the breeding season. WP: Waiting Period, RP: Reproduction Period, PP: Pregnancy Period. Calving at the beginning of the breeding season results on 2 to 12 months period of open days (OD) and females can achieve CI equal or less than 24 months. Excellent WP and RP lead to OD of about 2 months and CI of 14 months (box 1). Medium WP or RP lead to OD of about 6 months and CI of 18 months (boxes 2 and 4). Females with poor WP or RP will have an OD of 12 months and CI of 24 months (boxes 3 and 5). This last situation (CI of 24 months) could occur when moderate reproductive difficulties affect both WP and RP (Medium WP and RP, box 6). Animals with difficult reproductive status show association of anoestrus and infertility that lead to poor WP and RP and lead to OD and CI exceeding 12 months and 24 months (out of threshold), respectively. However, when the ovarian resumption and/or uterine involution periods overlaps with the seasonal anoestrus, the CI would be approximately 18–24 months. This is especially true when calving occurs on the middle or at the end of the breeding season. In this case, medium WP with excellent to good RP would give CI of 21–24 months (boxes 7, 8, 11 and 12). Similarly, the calving interval would be 21–24 months if there is a good WP with excellent to medium RP (boxes 9 and 10). In case of prolonged anoestrus associated with subfertility, the CI would exceed 24 months (out of threshold). (Adapted from Gherissi et al. 2020a)

balanced nutrition) and using good management practices (inputs and offspring’s weaning). Calving at the beginning of the breeding season coincides with the wet period and food availability, which promotes a good body condition and early resumption of ovarian activity with mating and pregnancy during of the same

season (Fig. 2.2). Regular checkups of postpartum animals to diagnose and treat related health problems (including mastitis, uterine infections, intercurrent diseases) and stillbirths are also recommended during WP. The RP management must be implemented by a regular balanced supply diet during the late pregnancy and postpartum period, camel bull flushing and sexual behavior and fertility control (male/female ratio, reform due to ‘infertility ...), as well as the specific diagnosis and treatment of genital diseases (metritis, bursitis ...) (Tibary et al. 2005; Ali et al. 2010).

2.4.2.2 Calving-Oestrus Interval (COI)

Healthy female camels show first postpartum oestrus between 14 and 42 days under pastoral livestock systems (Elias et al. 1984). When female camels were mated between 5 and 6 weeks post partum, adequate conception rates were obtained (Derar et al. 2014). The COI could, however, be extended to the next breeding season when calving occurs in the middle or at the end of the breeding season. Taking into account these factors, the COI target was set at 2 to 6 months and the threshold at 9 months (Gherissi et al. 2020a; Table 2.1). Open days (OD).

High rates (70–80%) of cyclic females were reported at 30 days postpartum in well-fed dromedaries, therefore the conception rates at 45 days postpartum were about the average scores (Tibary and Anouassi 1997a; Derar et al. 2014), but ovarian cyclicity can resume after 8–10 months when the diet is inappropriate for ovarian functioning (Skidmore et al. 1994). Therefore; when female camels conceive within the same calving season OD would be less than 6 months. However, if the dam is mated during the next breeding season OD period could last, instead form 7 to 12 months. Accordingly, the objective for this parameter was set at 180 days and the threshold at >360 days (Gherissi et al. 2020a, Table 2.1).

Table 2.1 Standard objectives and thresholds of reproductive parameters of female camels

Variable	Unit	Objective	Threshold
Age at first oestrus	Months	24–36	>42
Age at first mating	Months	24–36	>48
Birth to conception interval	Months	36–48	>54
Age first calving	Months	48–54	>60
Calving to oestrus interval	Months	2–6	>9
Open days	Days	≤180	>360
Calving interval	Months	≤18	>24
Annual fertility	%	>85	<75
Mating-pregnancy diagnosis	Days	21	30
Pregnancy length	Months	12.5	13.5
Culling age	Years	15–17	> 17
Calving number	Integer	5	7 > n > 3
Culled females per year	%	<5	>30

Adapted from Gherissi et al. (2020a)

2.4.2.3 The Annual Herd Fertility AF (Pregnancy Rate)

It reflects the ratio within a year between the number of pregnant female camels and the number of mated females. The fertility rate for camel herds looking to improve camel numerical productivity should range between 70% and 95% (Faye 2018). The agreed objective and threshold for this parameter are >85% and <75%, respectively (Gherissi et al. 2020a).

2.4.2.4 Female Camel's Calving Interval Objective and Threshold

The CI depends on waiting period (postpartum to first mating), reproduction period (first mating to conception) and the female's fertility. Calving to first mating interval (CFM) in pastoral conditions occurs within 14 to 42 days (Elias et al. 1984). Derar et al. (2014) showed that camels could be mated between the fifth and sixth week postpartum with satisfactory open days and conception rates. The time to first oestrus (associated to first mating) after calving varies from 1 to 38 months with a mean value of 10.3 ± 5.77 months (Gherissi et al. 2020a). High levels of CFM would be due to poor nutritional conditions leading to late resumption of follicular activity that takes up 8–10 months (Gherissi et al. 2018). The pregnancy duration in female camel is about 12.5–13 months (Gherissi et al. 2017).

The prompt ovarian resumption and occurrence of first mating earlier in the same calving season allow considering a CI objective of 14–18 months and an alarm threshold of 24 months (Fig. 2.2a). If calving occurs in the middle or late breeding season, the CI target would be 18–21 months and the alarm threshold would be 24 months (Fig. 2.2b). In fact, the objective CI is set at ≤ 18 months and the alarm threshold is set at >24 months (Table 2.1).

2.5 Reproduction Control in Dromedary Camel

Reproductive control tools can reduce unproductive periods by extending breeding season length and stimulating follicular growth and inducing ovulation during the non-breeding season. They facilitate allowing rational herd management to improve animal's resilience to harsh livestock conditions under arid environments (calves care, fodder availability and postpartum control) and to adapt herd production according to market needs (milk, fattening and calves). In addition, they allow accelerating genetic progress, firstly by facilitating the widespread dissemination of AI and secondarily by developing ovarian stimulation protocols and embryonic transplantation.

Synchronisation of ovarian activity principles applied in cows are used to synchronise ovarian activity in female camels, but methods are adjusted to physiological and behavioral characteristics of camelids, resulting in varying degrees of performance. The reproduction control in female dromedary camels is possible

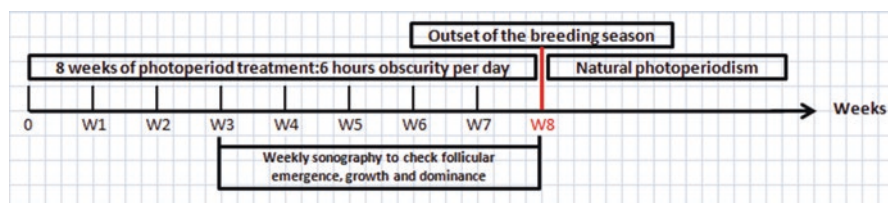


Fig. 2.3 Photoperiodic treatment of female camels 2 months before the beginning of the breeding season

using zootechnical means such as photoperiodic control (Vyas et al. 2008; Swelum et al. 2018a), food supplementation (Hammadi et al. 2001), male effect (Hafez and Hafez 2001) or hormonal exogenous supply such as melatonin (Dholpuria et al. 2012; Swelum et al. 2018b; Ainani et al. 2018; El-Allali et al. 2018) or progesterone (Monaco et al. 2013; Swelum and Alowaimer 2015; Swelum et al. 2018c), GnRH and PGF (Quzy et al. 2013; Al-Sobayil 2003, 2008; Manjunatha et al. 2018). The different applied protocols would be followed by blind natural mating or AI in fixed time using fresh or frozen semen.

2.5.1 Photoperiodic Control

In female camels, the photoperiodic control consists in applying a blindfolded eyes using black mask 6 h per day beginning 2 months before the start of the breeding season (November) (Vyas et al. 2008). Darkness perception by female camels using photoperiodic control leads to melatonin synthesis and release which advances the date of the sexual season. This remarkably induced a good follicular growth activity from the third week of treatment and the development of the ovulatory follicles between fifth and seventh week of treatment (Fig. 2.3; Vyas et al. 2008). Likewise, the ovulation, mating and pregnancy rates are quite considerable (Vyas et al. 2008).

In camel bulls, significant increase in testicular measurements and volume, libido, melatonin and testosterone concentrations were achieved by blindfold application during the non-breeding season to induce a shortened daily photoperiod of approximately 2.55 h relative to the natural day cycle (Swelum et al. 2018a).

2.5.2 Male Effect

Just one study was found in literature, documenting the pheromonal “male effect” causing female camels to cycle earlier in the breeding season (Hafez and Hafez 2001). In fact, the use of the male effect in camel species is rarely considered. Behavioral and endocrine processes of this practice to control reproduction in

camels are by far to be compared with other species such as sheep and goats. These two species are spontaneous ovulation while the camel is characterized by induced ovulation through mating.

2.5.3 Flush Feeding

The daily feed supplementation using 4 kg of commercial concentrated food during the late pregnancy (about 2 months before calving) and at the start of the postpartum period (5 kg/day for 3 months) helps to improve the female camels overweight at the time of calving and allows a rapid and significantly higher rate of ovarian activity resumption (Hammadi et al. 2001), short waiting period (PA) (around 40 days), high rate (approximately 70%) of females mated for the first time at 60 days after parturition (Hammadi et al. 2001) and high conception rate (Mostafa et al. 2016a, b). In addition, this practice helps to better growth of camels up to the age of 90 days (Hammadi et al. 2001).

2.5.4 Ultrasound-Guided Aspiration of the Follicle

The elimination of the dominant follicle using ultrasound-guided fine needle aspiration makes possible to end the current follicular wave and emergence of new follicular pool in an average time of 2.3 ± 0.5 days. The dominant follicle deviation occurs after 8.8 ± 1.1 days (Skidmore et al. 2009).

2.5.5 Exogenous Melatonin Based Protocols

The effectiveness of treatment based on exogenous melatonin consists of mimicking short days either at the beginning or at the end of the breeding season. For female dromedary camels; two melatonin treatment methods are possible. The first option is to administer it 2 months ahead the natural breeding season to support the follicular waves emerging and ultimate growth of the dominant follicle(s) at the start of the breeding season (Dholpuria et al. 2012). For example, in Algerian local condition it carries out at early September and continues until the end of October over 8-week period (Fig. 2.4). The second option is to start treatment at the end of the breeding season to allow an ovarian follicular activity extension for the rest of the year (non breeding season) (El-Allali et al. 2018). This treatment begins at early April and ends in late May (Fig. 2.5). At the end of these two proposed treatments, monitoring of ovarian activity is recommended to ensure the quality of ovarian cycles.

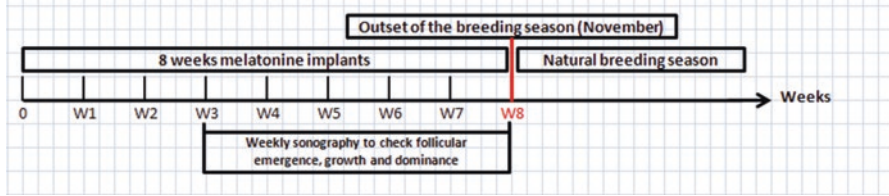


Fig. 2.4 Exogenous melatonin treatment of female camels 2 months before the beginning of the breeding season

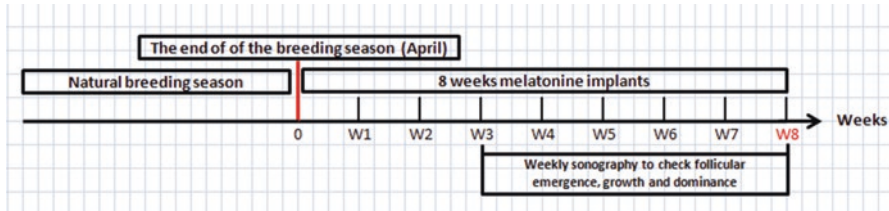


Fig. 2.5 Exogenous melatonin treatment of female camels 2 months after the end of the breeding season

To administer exogenous melatonin for camels, Dholpuria et al. (2012) recommend to use commercial implants containing 18 mg of melatonin at a dosage of 1 implant per 28 kg bodyweight (approximately 20 implants per female camel), which to be inserted subcutaneously by a 12-gauge applicator, preferably sterilized beforehand (Fig. 2.6).

For camel bull; the treatment during the non breeding season with melatonin implants (18-mg per 28 kg of live bodyweight) or approximately 28–32 implants per bull two times at an interval of 35 days helps to improve significantly their reproductive performance (Plasma testosterone, testicular measurements, sexual behavior and spermogram) (Swelum et al. 2018b).

2.5.6 Exogenous Progesterone Based Protocols

Daily progesterone injection (100–150 mg) and various devices secreting progesterone (CIDR 1.38 g: Controlled Internal Drug Releasing Device, new PRID Δ 1.55 g: Progesterone Relasing Intravaginal Device) are experimented to synchronize follicular waves in female camels (Cooper et al. 1990, 1992; Monaco et al. 2013; Swelum and Alowaimer 2015; Hussein et al. 2015; Swelum et al. 2018c; Tibary 2018; Abo El-Maaty et al. 2019). Cleaning of perineum and flushing of vagina with iodopovidone solution before device insertion is very important since it clinically improves vaginal environment at the end of the treatment (Monaco et al. 2013). Female camels receive CIDR for 7 days (Abo El-Maaty et al. 2019), 10 days

Fig. 2.6 The technique of application of subcutaneous melatonin implants in camel. (From Dholpuria et al. 2012)

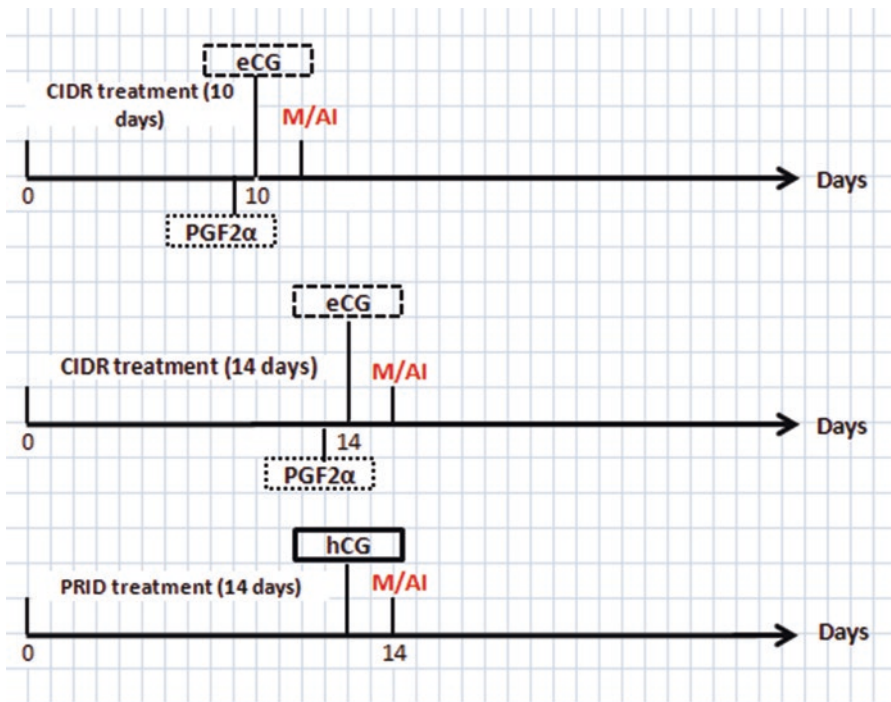


Fig. 2.7 Exogenous progesterone treatments during intra-vaginal devices (CIDR, PRID) with eCG or hCG and timed artificial insemination

(Monaco et al. 2013, Fig. 2.7), 14 days (Swelum and Alowaimer 2015; Al-Fatlawi and Al-Hamedawi 2017; Fig. 2.7) or 17 days (Al-Sobayil 2008), while new PRID Δ is held 14 days for Swelum et al. (2018c; Fig. 2.7). The progesterone plasmatic concentration increases earlier after device insertion (2 days) but its plasmatic peak is observed after 12 days (Swelum and Alowaimer 2015).

There are conflicting studies on the effectiveness of PRIDs and CIDRs (Monaco et al. 2013; Tibary 2018). The authors conclude that the treatment of exogenous progesterone could fully prevent sexual receptivity and suppress the development of large follicles without suppressing follicular wave's growth. These devices have sometimes been related with increasing spontaneous ovulation (Tibary 2018).

There are conflicting reports on the efficacy of PRIDs and CIDRs (Monaco et al. 2013; Tibary 2018). The authors opinions converge that exogenous progesterone treatment could completely prevent sexual receptivity and suppresses the growth of large follicles but could not suppress the follicular wave. Sometimes, these devices have been associated with increased spontaneous ovulation (Tibary 2018). Injection of PGF2 α (500 μ g) one day before CIDR withdraw (Monaco et al. 2013), eCG (2000 IU) at the day of CIDR withdraw (Al-Sobayil 2008) or hCG (3000 IU) at the last day of PRID treatment (Al-Sobayil 2008) were also practiced in female camels (Fig. 2.7). The daily injection of progesterone (100 mg, IM) for 10–16 days provided promising results in multiple ovulated female camels (McKinnon et al. 1994; Tibary 2018). More comprehensive studies are necessary to better understand the effectiveness of different associated treatments to exogenous progesterone administration on follicular wave synchronization in female camel. A recent study found that long-acting injections of progesterone can be used in camels and can be more effective than daily injections (Chhaibi et al. 2016). Synchronizing follicular waves using progesterone devices reminds a controversial issue between the authors particularly on their effects at different breeding seasons (Tibary 2018). Their effectiveness at the beginning and in the middle of the breeding season has been reported vain due to its action in reducing the follicular diameter but not follicular number (Monaco et al. 2013). During the breeding season Swelum and Alowaimer (2015) reported a high percentage of camels presenting ovulatory follicles 2–4 days after CIDR withdrawal showing therefore its significant interest in follicular wave synchronization. In another more study; Swelum et al. (2018c) confirmed the interest of new PRID Δ in the synchronization of follicular waves during the breeding season with a high rate of females which carry ovulatory follicles 2 days after the end of treatment. The ovulation rate, the non-return rate are significantly increased, the open days period reduced mainly and the fertility rate is higher when the blind natural mating or timed AI are practiced 48 h after PRID withdraw (Al-Sobayil 2008; Al-Fatlawi and Al-Hamedawi 2017; Swelum et al. 2018c).

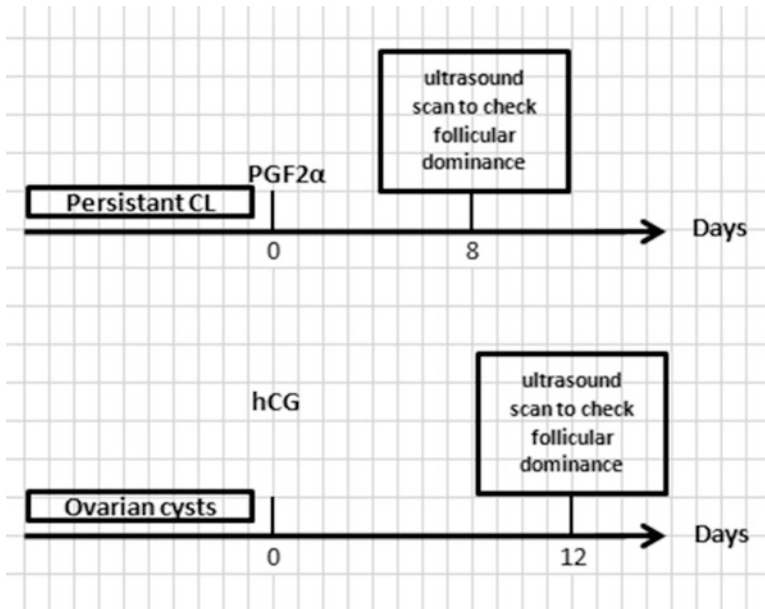


Fig. 2.8 Prostaglandine (PGF2 α) and human chorionic gonadotrhpin (hCG) treatments of persistent corpus luteum and ovarian cysts, respectively

2.5.7 Prostaglandin and Human Chorionic Based Protocols

PGF2alpha (500 μ g) administration for female camels with a persistent CL and hCG (4500 IU) for those with cystic ovaries were reported to have high efficacy on the number of females with ovaries showing ovulatory follicles 8 and 12 days post-treatment, respectively, but the pregnancy rate for both protocols remained low (Quzy et al. 2013; Fig. 2.8).

2.5.8 GnRH-Based Protocols

“*Select synch*” treatment 2 months ahead the natural breeding season with GnRH (250 μ g) on day 0 and PGF2 α (5 ml) on day 7 (Skidmore et al. 2009) or day 10 (Mostafa et al. 2016a, b) did not improve the quality of follicular wave at the end of the treatment (Fig. 2.9). GnRH alone may be effective in synchronizing follicular waves to mitigate the harmful effects of prostaglandin on the antioxidant potential of camels and their response to breeding (Abo El-Maaty et al. 2019). Follicular wave emergence and follicular deviation occur about 3 days (range 2.5 and 3.5 days) and 5.5 days (range 4 and 11 days) respectively, after GnRH administration

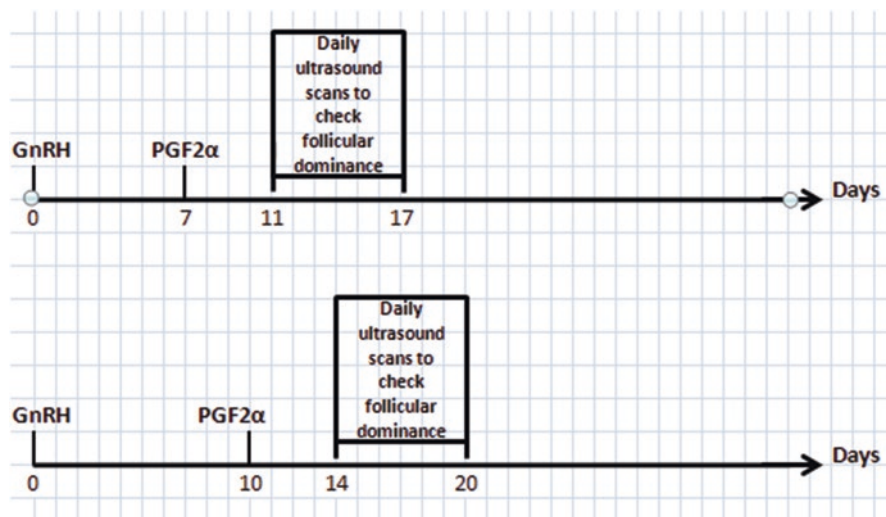


Fig. 2.9 Select synch (GnRH-PG) protocol for oestrus induction in female dromedary camels

(Manjunatha et al. 2015; Skidmore et al. 2009) and 4.5 days and 10.7 days for GnRH-PGF2 α treatment.

The Ovsynch (GnRH-PGF2 α -GnRH: 0–7–9) and eCG (2000 IU) treatments for female camels with small and smooth ovaries (SSO) 2 months ahead the natural breeding season induced high effective rates of animals with matures follicles at day 8 and day 12 post treatment, respectively (Quzy et al. 2013; Fig. 2.10). The GPG (0–7–14) treatment showed high proportion ($\approx 73\%$) of female camels had a total of 22 dominant follicles (1.3–1.9 cm) of which 21 ovulated after GnRH injection day 14 (Skidmore et al. 2009; Fig. 2.10).

Ovsynch (0-7-10- 22:M1) protocol with timed natural mating followed by pregnancy diagnosis 21 days later and ovarian ultrasonographic monitoring of non pregnant females for second timed mating was examined by Manjunatha et al. (2018) with high rate of follicular wave synchronization and consistent pregnancy rate (Fig. 2.11).

Ovsynch -Resynch (GPG-M1-GP-M2; Fig. 2.12) treatments were experimented recently by Manjunatha et al. (2018) in dromedary camels during the breeding season. Firstly these authors used Ovsynch protocol (GnRH-PGF2 α -GnRH: 0-7-10) followed by timed mating 12 days later (day 22). Then, systematic GnRH injection was received by all female camels at day 36 regardless to their pregnancy statue; followed by pregnancy diagnosis at day 43. PGF2 α (day 43) was administrated to non pregnant females which were resubmitted to second timed mating at day 48. This protocol showed high proportion of ovulated female camels after Ovsynch protocol. The GnRH in Resynch protocol resynchronized effectively the follicular development for second mating of non pregnant females with high early and late pregnancy rates and low rate of pregnancy loss (Manjunatha et al. 2018).

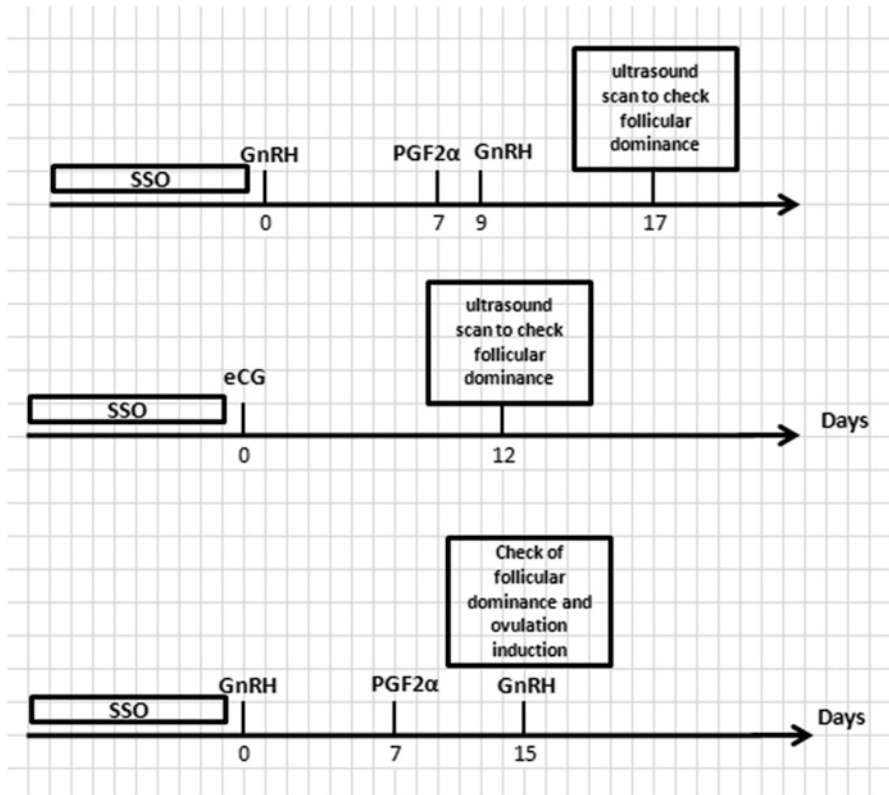


Fig. 2.10 Ovsynch (GnRH-PG-GnRH) and eCG (equine Chorionic Gonadotropins) protocols for oestrus induction in female dromedary camels with small and smooth ovaries (SSO) 2 months ahead the natural breeding season

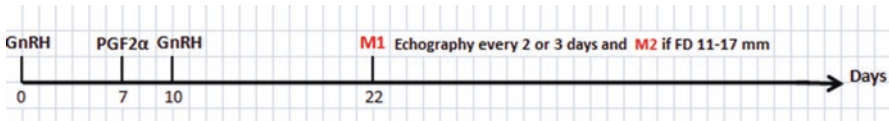


Fig. 2.11 Ovsynch (GnRH-PG-GnRH) protocol followed by blind natural mating and ultrasonographic monitoring of non pregnant females for second timed mating

The repetitive follicular wave resynchronization (GPG-M1-GP-M2-GP-M3; Fig. 2.12) after each mating was also explored by Manjunatha et al. (2018). They reported high synchronization rate at first mating and also high resynchronization rate at second and third timed natural mating. The pregnancy rate and the pregnancy loss rate after Ovsynch-first mating and Resynch protocols-second and third mating showed very interesting results (Manjunatha et al. 2018). An encouraging pregnancy cumulative rate is favorable to adopt this protocol in camels under farm and field conditions (Manjunatha et al. 2018).

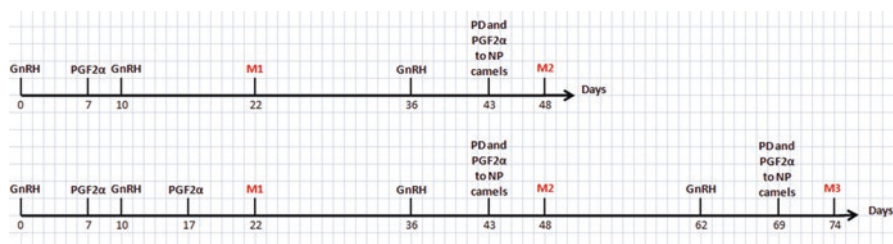


Fig. 2.12 Ovsynch -Resynch (GnH-PGF2α-GnRH-M1-GnRH-PGF2α-M2) protocol for oestrus synchronization of female dromedary camel

The listed reproductive control methods could help to improve the CI in camel herds and therefore decrease the unproductive periods, overcome the constraint of the short breeding season, facilitate herd management practices (vaccination of mothers, maintenance and growth of camels, weaning, postpartum control ...) and thus improve the birth rate and the numerical productivity, milk and meat production and in the long term decrease the generation interval for accelerating genetic progress.

2.6 Artificial Insemination

2.6.1 Definition

Artificial insemination (AI) is the act of depositing sperm by means of an instrument, at the most opportune time and in the most suitable place in the female genital tract. Since the 1960s, the use of AI as a breeding technique in camelidae has been reported with first camelid offspring from Bactrian camel obtained by this technique in 1961 (Elliot 1961). However, this technique has gained popularity over the last 25 years as interest is increasing to improve camel genetic characteristics such as milk, meat and wool quality, as well as racing ability. AI in DC faces several problems, including complications related to semen collection, bull sexual activity, semen quality (low volume, low sperm concentration and high viscosity), limited awareness of optimal insemination time and sperm dosage per insemination session, and lack of a standard storage technique (Elliot 1961; Tibary and Anouassi 1997c; Al-Bulushi et al. 2019b).

2.6.2 Importance of Artificial Insemination in Camels

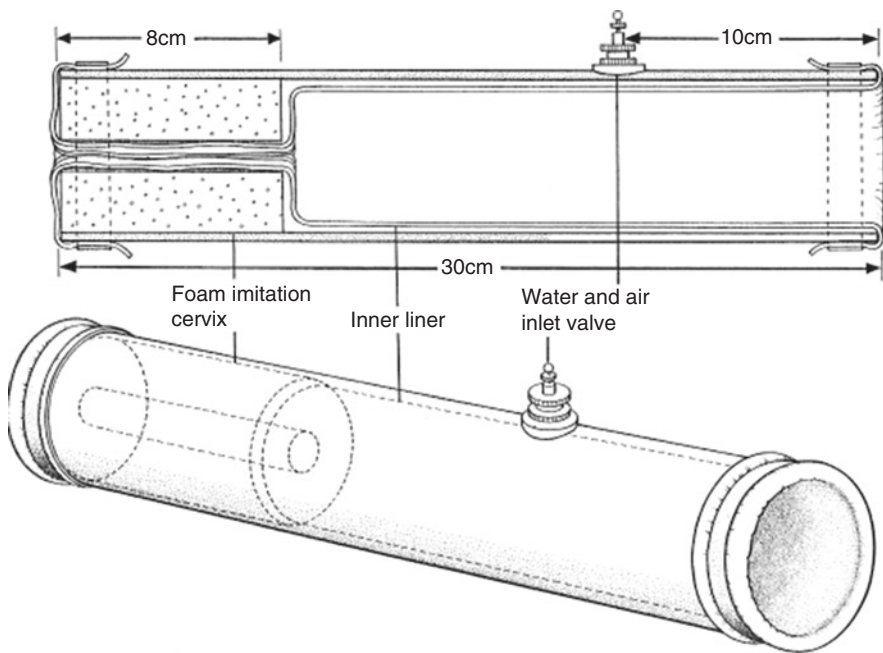
The method offers many advantages: on the one hand, it facilitates the management of reproduction in the camel herds and increases the reproductive capacity of the males (ratio male/females) and therefore to obtain the most offsprings which helps rapid genetic improvement. On the other hand, it is a preventive measure against sexually transmitted diseases. Moreover, the use of AI would prevent the need to transport male or female animals for mating, thus reducing the charges and danger of moving valuable animals. The short and long term sperm conservation allows the international transportation of semen and conservation of the genetic material of males with high genetic level. It helps eliminate behavioral problems so AI would minimize the risks of injury. Finally, it facilitates the application of cross breeding programs between different camelid's species.

Table 2.2 Advantages and disadvantages of sperm collection using artificial vagina

Advantages	Disadvantages
High sperm concentration	Decreased sperm mobility with prolonged contact of sperm with the inner membrane of the AV
High frequency of sperm collection during periods of sexual activity	Requires a trained camel bull
Not very aggressive technique, preserves the well-being and the natural sexual instinct	Requires high technicality of the operator
Allows observation of sexual behavior to assess male reproductive ability	Injuries risk of the animal and the operator
Can be replaced by condom device or held between the female's hind legs	Risk of refusal mating and especially when using dummy or with AV poorly suited
Can be used in the presence of female teaser in estrus or a camel dummy	High cost associated with the use of teaser females
Collection can be facilitated by creating an underground room or small laboratory underneath the dummy	
Method adapted to the insemination center: high collection rate, high sperm quality, less stressful manipulation of animals with high genetic potential, collection hygiene	
More adequate for short term and long term sperm preservation	
Higher pregnancy rate with fresh raw sperm	

Table 2.3 Advantages and disadvantages of sperm collection using an electroejaculator

Advantages	Disadvantages
Sperm collection during non breeding season	Low sperm concentration
More safe for the animal and operator	Requires pharmacological tranquilization of the camel bull
Lower cost	Aggressive technique that does not take into account the sexual instinct and comfort of the male camel
Collection of males with certain conditions that do not allow normal mating (locomotor diseases)	Low frequency and cadence of sperm collection
	More difficult sperm conservation
	Lower pregnancy rate

**Fig. 2.13** Diagram of an artificial vagina used for DC. (Reprinted with permission of editor: Aurich Christine; from Bravo et al. 2000)

2.6.3 Semen Collection

The semen collection is the first step in AI and/or its examination. The primary objective of a semen collection is to obtain a large number of clean and optimum quality ejaculates (Monaco and Lacalandra 2019). The collection method has mighty effects on the physical and possibly biochemical properties of the ejaculated material (Tables 2.2 and 2.3) (Tibary et al. 2014). The artificial vagina (AV) and

Fig. 2.14 Semen collection of male DCbull using modified bull-artificial vagina and female camel teaser with operator who direct penis by hand manipulation of sheath towards the opening of the artificial vagina. **(a)** Modified artificial vagina for collection of semen, **(b)** Coil on short artificial vagina to simulate cervical rings and **(c)** Collection of semen on a receptive female mount. (Photos reprinted with permission of the editor: Ahmed Bamouh, from Tibary and Anouassi 2018)



electroejaculation are the accepted methods for semen collection in camels (Bravo et al. 2000). The AV constitutes the means widely accepted by several authors who have worked on the dromedary using a female camel as a teaser or dummy. It's simple and practical device which has two parts. An outer cylinder made of rigid material, most often hard and thick rubber (thermal insulation) or plastic with an opening closed by a stopper. The inner latex or artificial rubber is introduced into the outer cylinder and its ends folded down and held in place by an elastic band. The cavity thus is formed by the outer cylinder and the inner liner. One end of the artificial vagina is lubricated: it will be used to introduce the penis; on the other is fixed a rubber cone at the end of which is fitted a graduated glass or better plastic tube to collect the sperm (Figs. 2.13 and 2.14).

Electroejaculation is rarely applied in the camel species (Tibary et al. 2014; Tibary and Anouassi 2018). This procedure requires the tranquillization or even the

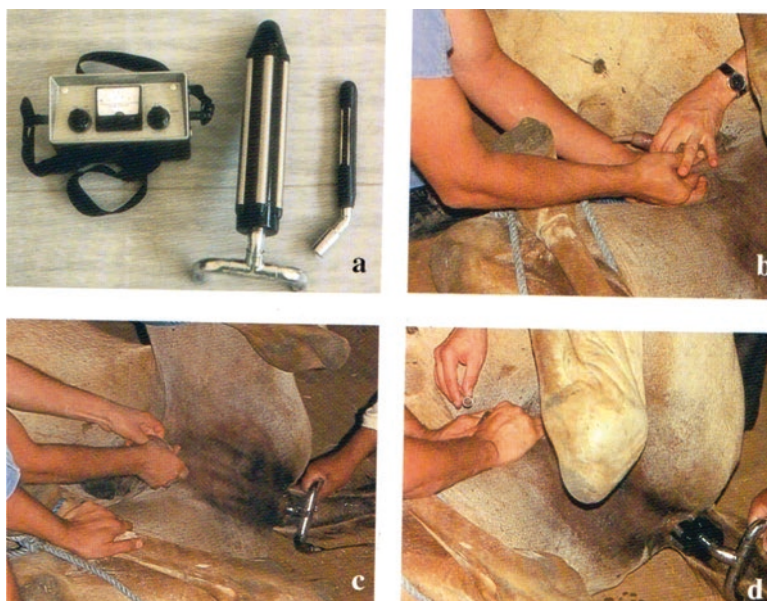


Fig. 2.15 Semen collection of male dromedary camel bull using an electroejaculator. (a) Electroejaculator, (b) exteriorization of part of the penis outside the sheath, (c) placement of the electroejaculator after animal preparation, (d) semen collection step. (Photos reprinted with permission of the authors: Ahmed Tibary and Abdelhak Anouassi, from Tibary and Anouassi 1997b)

sedation of the animal, which would present a health risks (Fig. 2.15). The degradation of the physico-chemical quality and the concentration of the sperm collected by electroejaculation make it difficult its conservation and freezing (Tingari et al. 1986). So, electroejaculation (12V, 180 mA) can be applied with exceptional cases with camel bulls presenting highly vigorous and aggressively behavior, with low libido or with certain physical problems and for collection of semen during the non breeding season (Skidmore et al. 2013; El-Hussanein 2003). In this chapter, we don't consider this technique because its use does not meet the requirements of animal welfare.

Old methods of collecting semen such as collecting sperm directly from the vagina and collecting sperm by massaging the genital pool glands have also been reported in dromedaries (Tibary and Anouassi 1997b).

The modified bull-AV was commonly used for camel bull with good results (Bravo et al. 2000; Tibary and Anouassi 2018; Fig. 2.13). Some precautions must be taken into consideration in order to adapt the AV and avoid causing injuries to male bulls leading to the refusal of the animal for subsequent samples. The short modified AV allows also keeping away sperm from contact with the rubber liner of the AV and avoiding soiling the sample that adversely affect sperm quality. Therefore, it is recommended to:

- Shortening AV to allow the semen to pass directly into the collection flask. The internal part would have a diameter of 5 cm and a length of 30 cm (Bravo et al. 2000). The internal sleeve ends with a rubber receiving cone on which the collecting tube will be fixed. The whole will be secured with two rubber straps.
- To prevent contact with the rubber material, an additional removable plastic inner liner can be added, but the authors have found that males do not tolerate this well (Skidmore et al. 2013).
- For a perfect stimulation of ejaculation in dromedaries, the modified bull-AV should be equipped by foam imitation cervix of about 8 cm in length at its distal part. This provide a narrowing simulating cervical to receive the highly mobile urethral process of the camel penis to stimulate ejaculation during the extended copulatory (Bravo et al. 2000; Monaco et al. 2016; Monaco et al. 2018; Skidmore 2019). The filling water (55–60 °C) must be sufficient and it is necessary to blow air between the inner liner and the outer rigid wall to achieve an inner temperature of 38–40 °C and a pressure equal to that of the female vagina (Skidmore et al. 2013).
- The filling water (55–60 °C) must be in sufficient quantity and blowing air between the inner liner and outer rigid wall are important to obtain an internal temperature of 38–40 °C and a pressure equivalent to that of the female vagina (Skidmore et al. 2013).
- Keeping the vagina in an oven at 45 °C and fill it only in the minutes preceding the sample. Overpressure is not recommended because it may not leave enough space for the penis during ejaculation and cause the internal wall to burst. Correct filling is when thr upright opening of AV simulates a vaginal slit.

The semen collection should be done as much as possible in a quiet place to avoid any stressful situation for the animal and on non-pulverulent soil for the comfort of the operator and to avoid contamination of the sample as camels mate in a couched position. The use of a female teaser is essential. The impact of time of female teaser presentation and number of mating attempts before going to effective semen collection seem to have important effect on collected sperm volume and concentration (Al-Bulushi et al. 2018). These factors were reported to have a significant effect by increasing male excitation and improving sperm quality in other species (Hanzen 2015).

When the animal is in mating position, the penis will be directed by means of the operator hand applied to the sheath towards the opening of the artificial vagina, which directed slightly outwards while avoiding any excessive deviation of the penis. The sensory arousal of moist heat drives the intromission reflex. Before ejaculation is done, the male will make multiple thrusts, interspersed by periods of rest. The ejaculate typically happens in fractions and whole process can be prolonged between 5 and 50 min (Bravo et al. 2000). Thus, it is very difficult to know if ejaculation has occurred; collection continues until the male stands up. The effective ejaculation (intromission into the artificial vagina until the end of ejaculation) takes place after a variable duration ranging from 5 min to 20 min (Anouassi et al. 1992; Hassan et al. 1995) without yet a correlation between the duration of collection and

the quality of the sperm. Soon after, the male stands up. Since the process of semen collection might be quite long, addition of about 1–2 ml of extender into the collection vessel before collection may be beneficial (Bravo et al. 2000). Immediately after the male stands up, the vagina is turned over so as to collect the sperm in the collection tube. The latter will, if necessary, be protected from any thermal shock by wadding or another isothermal envelope.

On the other hand, sexual peculiarities of the camel species (Tibary and Anouassi 1997a, b, c; Bravo et al. 2000; Monaco et al. 2018); such as the copulatory behavior i.e. mating in sternal recumbency, lengthy ejaculation throughout copulation (from 5 min to 50 min), aggressive sexual behavior, the highly viscous nature of the semen and intrauterine semen deposition constitute major constraints to sperm collection and have made it mandatory to adapt sperm collection techniques with various degrees of success. We particularly mention:

- Using some specific housing and welfare conditions to prevent injuries, stress and stereotypical behaviours (oral, locomotor, self-ejaculation) associated with poor
- reproductive performance (Monaco and Lacalandra 2019).
- Enhancing endocrine pathways and ejaculation quality of the male sexual by acting on the herd structure in which is maintained the camel bull, photoperiodic treatments and/or pharmacological (exogenous melatonin) protocols (Monaco et al. 2018; Monaco and Lacalandra 2019).
- Using female camel teaser and placed in sternal recumbency position with the operator in an underground room beneath the collection area which allows full observation of mating behavior and ejaculatory pattern (Hemeida et al. 2001; Fig. 2.16).
- Using artificial vagina devices implanted under the female teaser (Tibary and Anouassi 1997a, b, c; Al-EknaH et al. 2001; Fig. 2.17) or mounted on a dummy (El-Hussanein 2017; Fig. 2.18).
- Replace teaser female by a dummy designed in the similar position for natural mating, inside which is inserted an AV. The dummy is fixed on the floor of the collection yard and as a ceiling for a small laboratory to exchange AV from one



Fig. 2.16 Semen collection of male dromedary camel bull using artificial vagina oriented and maintained by an operator in an underground room which allows him full observation of mating behavior and ejaculatory pattern. (Photos from Hemeida et al. 2001)

Fig. 2.17 Semen collection of male dromedary camel bull using artificial vagina held between the female's hind legs. (Photo reprinted with permission of the authors: Ahmed Tibary and Abdelhak Anouassi, from Tibary and Anouassi 1997b)



Fig. 2.18 Semen collection of male dromedary camel bull using artificial vagina mounted in a camel dummy. (Photo reprinted with permission of the editor: Daryoush Babazadeh, from El-Hussanein 2017)



male to the other and to carry out rapid evaluation and partial extension of the collected semen (El-Hussanein 2003, 2017; Ziapour et al. 2014; Fig. 2.19).

- Place condoms within the vagina of female teasers which showed acceptable result regarding ejaculated sperm volume and its characteristics (Tibary and Anouassi 2018; Fig. 2.20).
- Training camel bull to semen collection using dummy device and condoms. In this regard, serious behavioral limitations were reported if males are not used exclusively for AI (Tibary and Anouassi 2018; Monaco and Lacalandra 2019).
- To prevent semen contamination, a rope circled around the thorax could be used to attach the tail and the prepuce could be washed and dried before beginning the semen collection session (Monaco and Lacalandra 2019).

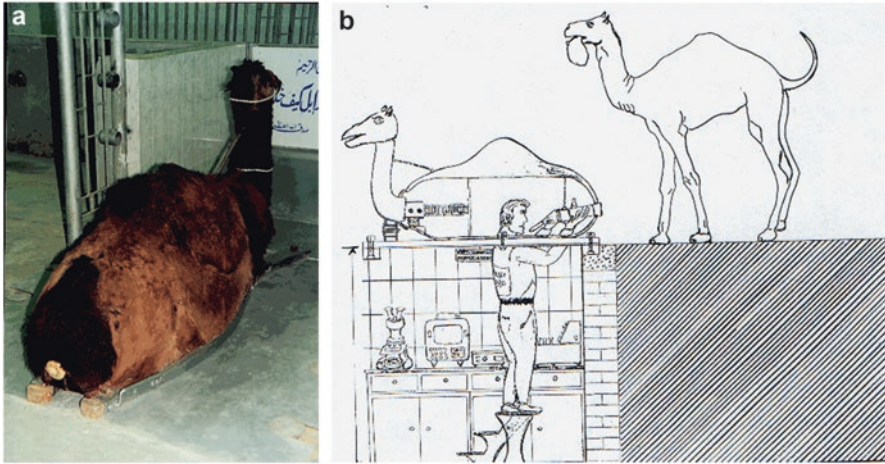


Fig. 2.19 Semen collection of male dromedary camel bull using a dummy and mounted artificial vagina with operator stays in a small laboratory underneath the dummy. **(a)** A hind view for the dummy to indicate its external feature and the mounted AV, **(b)** A diagrammatic drawing for the dummy and the small lab underneath. (Photo and diagram reprinted with permission of IVIS editor from El-Hussanein 2003)

2.6.4 Initial Sperm Examination

The available studies on the ejaculate initial quality of camel sperm showed high variability of the camel sperm characteristics.

2.6.4.1 Macroscopic Examinations

Volume

The sperm volume of the dromedary varies considerably depending on the technique and means of collection as well as the male's libido and the time of stimulation before collection. Using electro-ejaculator the volume varies from 1 ml to 10 ml (Tibary and Anouassi 1997b) even this volume varies from 2 ml to 12.6 ml when semen collection is carried out using AV (Wani et al. 2008; Tibary and Anouassi 1997b). According to Monaco et al. (2018); more than four ejaculates per collection session of about 45 min is possible resulting in total ejaculation volume of 12–16 ml. Even though allowing time for the male to recover after each ejaculation, the concentration of sperm was slightly lower after the third ejaculate.



Fig. 2.20 Semen collection of male dromedary camel bull using condom device mounted within the vagina of receptive females and secured with a harness. (a) Condom device (left) inserted in the vaginal of an estrous female (right), (b) Female with secured intravaginal condom in the breeding pen (c) Male teasing the female with intravaginal condom (d and e) Male breeding within the intravaginal condom. (Photos reprinted with permission of the editor: Ahmed Bamouh, from Tibary and Anouassi 2018)

Appearance and Consistency

Semen of camels is of gray to milky white color (Bravo et al. 2000). Sand contamination can change semen color, for this reason the proposed camel dummy with inserted internal AV prevent semen contamination (El-Hussanein 2017; Skidmore 2019). Camel semen is in gel form and the proportion of seminal fluid varies leading to high variability of viscosity and sperm concentration (Deen et al. 2003). Depending on individual males, semen collection process, stimulation duration and the volume of the gel fraction in the ejaculate, this proportion of the seminal fluid is highly variable. (Tibary and Anouassi 2018; Skidmore et al. 2013).

The main constraint related to camel sperm viscosity is due to the distribution evenly throughout ejaculation of the liquid fraction which induces a difficulty of its separation (Skidmore et al. 2013). It was suggested that the high semen viscosity is resulted of higher seminal plasma levels of glycosaminoglycans or mucin 5B (Kershaw-Young et al. 2013).

It is important that the sperm of the dromedary undergoes a total liquefaction before its microscopic examination and dilution. Tibary and Anouassi (1997b) reported that sperm with little seminal fluid liquefies faster than sperm that contains large volume of seminal fluid. In addition, some studies have shown that liquefaction time of the camel sperm at 25–38 °C is 8–20 min (Tibary and Anouassi 1997b),

15–30 min at 30–37 °C (Al-Qarawi et al. 2002) and 90 min at 37 °C (Wani et al. 2008). However, during these time ranges, change in sperm viscosity also constitutes a partial liquefaction that is subject to a high variance between males and between ejaculates from the same male, ejaculate manipulation technique during incubation, type of extender and quantity of mucin in the ejaculate. (Tibary and Anouassi 2018; Al-Bulushi et al. 2018). The complete liquefaction of dromedary ejaculates took 8 h (Tibary and Anouassi 1997b). In recent study, this time was estimated to 23.9 ± 1.5 h (Mal et al. 2016). In practice, this time would certainly result in loss of quality of semen (Tibary and Anouassi 2018). Fast liquefaction (after 1.5 h at 37 °C) of camel diluted 1: 1 using Tris – lactose egg yolk, Tris – tes egg yolk,, sucrose egg yolk, citrate egg yolk and Tris – fructose egg yolk extender- compared to semen without an added extender (Wani et al. 2008). Attempts at rapid liquefaction of dromedary sperm using mucolytic and proteolytic compounds such as plasmin, trypsin, chemotrypsin, amylase, collagenase and ficin have given variable results (Tibary and Anouassi 1997b; Bravo et al. 2000; Ghoneim et al. 2010; El-Bahrawy et al. 2015; Monaco et al. 2016; Keshavarz et al. 2016). Results using collagenase (0.5–1%) or papain (1.7 Unit/ml) seems to provide the best semen liquefaction in DC(20–40 min) (Kershaw-Young and Maxwell 2012; Tibary and Anouassi 2018; Monaco et al. 2016). These enzymatic treatments improve post-thaw motility and velocity patterns of spermatozoa but also can be deleterious to spermatozoa (Rateb et al. 2019; Skidmore 2019). Mechanical processes were also investigated to improve the duration of semen liquefaction using prolonged method of dilution (1: 1–1: 5) (Wani et al. 2008; Malo et al. 2017) and application of ultrasound (Rateb 2016; Skidmore 2019). The camel semen treatment by amylase improves viscosity score and mobility. The association of this enzymatic treatment with centrifugation for seminal plasma removal decreases semen viscosity and motility but increases the rate of spermatozoa abnormalities (El-Bahrawy 2017a, b).

2.6.4.2 Microscopic Examination of Semen

After first dilution (1:1) using semen extender and complete semen liquefaction the microscopic evaluation of camel sperm can be performed by measuring concentration, total motility, individual motility, viability and morphological aspects (Table 2.4). After that the second dilution can be applied to final ratio (1: 3).

Table 2.4 Mean characteristics of camel semen collected by the artificial vagina (Bravo et al. 2000)

Volume (ml)	Concentration	Motility (%)	Stained dead sperm (%)	Sperm showing morphological abnormalities (%)	Sperm showing acrosome abnormalities (%)
7.5	325	51	18	28	8.5

Concentration

The concentration expresses the number of spermatozoa per mm^3 (or per ml). It can be determined directly by counting spermatozoa using a hematic cell or indirectly by visual comparison of the sperm with standard solutions, by electronic counting or by nephelometry (spectrophotometer or colorimeter). This last method can however be indirectly increased following the presence of debris (sand) in the sperm of the dromedary and thus a variation of the opacity.

Direct counting is done using a hemacytometer. This type of count assumes the prior dilution of the sperm. The dilution rate depends on the apparent concentration of the sperm and the desired final concentration. There are different types of hemacytometer which are characterized in particular by their surface (S) and the depth of their counting chamber (P): example Thoma (S: 1 mm^2 , P: 0.1 mm). The hemacytometer consists of a glass slide, hollowed out of a small cuvette whose bottom is lined with a grid. Thoma's cell has a grid of 16 large squares each comprising 16 small squares. The area of the large squares is equal to 1 mm^2 . The counting chamber is 0.1 mm high.

After depositing a drop of diluted sperm and covering it with a coverslip, the number of spermatozoa is determined at magnification 400 on a surface corresponding to 4 large squares. Conventionally, only the heads of spermatozoa located inside the two parallel lines delimiting each large square or whose head is on the left and upper lines delimiting a large square are taken into account. The concentration is calculated as follows: $\text{Concentration} = N \times 4 \times 10 \times D$ (N is the number of sperm counted in 4 large squares, 4 since the hemacytometer has 16 large squares with a total area equal to 1 mm^2 , 10 since the height of the enumeration chamber is equal to 0.1 mm and D is the degree of the initial dilution).

The sperm concentration and the total number per ejaculate of male DC varies among studies ranging from 80 to $1300 \times 10^6/\text{ml}$ (Tibary and Anouassi 1997b; Wani et al. 2008; Al-Bulushi et al. 2018) and from 240 to 2576×10^6 (Tibary and Anouassi 1997b; Morton et al. 2011; Tibary et al. 2014; Tibary and Anouassi 2018). Camel semen concentration is qualified as oligozoospermia when it shows less than 40 millions spermatozoa/ml (Waheed et al. 2014). Specific individual variations, frequency of collection and technique/time needed for semen collection can lead to significant variation of sperm numbers in the dromedary ejaculate. (Tibary and Anouassi 2018; Al-Bulushi et al. 2018). It is also interesting that authors have mentioned that the absence of sperm in many ejaculates is due to incomplete ejaculation (Hassan et al. 1995). Weekly semen collection during the peak breeding season for 3 weeks or twice weekly semen collection over 8 weeks did not affect the semen quality (Al-Bulushi et al. 2018). The pregnancy rate using fresh camel sperm with low concentration (75×10^6 sperm/ml) showed significantly low rate compared to sperms with concentration of 150 and 300×10^6 sperm/ml (et al. 2019b).

Massal Motility

Sperm mobility is evaluated according to two criteria: mass and individual mobility. The mass mobility is evaluated on diluted sperm by placing a drop on a preheated slide and observes it under a phase contrast microscope in low light intensity at 100× magnification. It refers to the intensity of the waves caused by the movement of sperm. A score of 0 to 5 is assigned to the observed sample: 0: no movement of the sperm, 1: slight noticeable movement, no waves, 2: few waves, 3: many waves, 4: fast and intense waves and 5: very fast waves.

The nomenclature of asthenozoospermia was given to camel sperm with decreased spermatozoa motility (Waheed et al. 2014). In Al-Bulushi et al. (2019b) study it was shown that insemination using raw camel sperm with 84–86% total motility result on high and comparable pregnancy rate (81%) to that obtained by natural mating (83%).

Individual Mobility

Individual mobility is assessed on a drop of diluted and sufficiently liquefied sperm placed on a preheated slide under ×400 magnification. The dilution is done in a diluter (“extender”) previously heated to a ratio of 1 to 20 (0.25 ml of sperm and 4.75 ml of diluter). These media should be ideally prepared before the examination to avoid any change in pH, detrimental to the motility of the spermatozoa. Three to five fields close to the center of the drop will thus be examined and the average calculated. The individual motility is the proportion (%) of sperm that pass through the field of the microscope relatively quickly with rotational movements of the head (flexing or tracer sperm). Some sperm spinning in small circles or moving backwards is due to the abaxial implantation of their tail. Others move in a curvilinear fashion or more slowly. They are not considered mobile. Computer Assisted Sperm Analysis (CASA) type image analyzers make it possible to quantify more precisely (the nature and speed of movements).

A very good quality sperm must have at least 50% of motile spermatozoa in the camel (Bravo et al. 2000). Numerous reports from the literature have not revealed sperm mobility problems of dromedary sperm, however, we find that the proportion of motile sperm varies from 0 to 80% (Tibary and Anouassi 1997b; Deen et al. 2003) and that is significantly influenced by several factors such as means of collection, technique and conditions of collection, sperm viscosity, nature of the diluter, the synthetic material of the inner lining of the AV and the length of time that the sperm spends inside AV (Tibary and Anouassi 1997b; Deen et al. 2003). Addition of caffeine could improve the individual motility of the camel sperm (Deen et al. 2003).

The examination of individual motility is interesting because it indirectly provides information on the integrity of the sperm membrane and its morphological integrity. Thus, a high percentage of motile sperm combined with a high percentage of dead sperm suggests improper handling of sperm rather than abnormal sperm.

Likewise, low motility is often correlated with a high percentage of abnormal forms or dead sperm.

Morphological Examination

Morphological examination requires staining of the sperm by placing a drop of semen on the end of a slide and spreading it in a thin layer with another blade tilted at 45°. The preparation is dried in air for few minutes. The smear is then fixed by immersion in a solution of methyl alcohol or in a solution of 5% formalin. Free air fixation is also possible such as passing the smear quickly over an alcohol lamp. Any thermal shock will nevertheless be avoided during these preparations. The preliminary dilution of the sperm (2 drops in 0.5 ml of diluent) will facilitate the morphological examination. Some stains are intended to better show the morphology of the spermatozoon (total staining), the others called vital make possible to differentiate between dead and living spermatozoa. Among the total stains, some are said to be simple (India ink, methylene blue, toluidine blue, gentian violet, fuschin, etc.): they provide uniform coloration of the spermatozoa while the second called double (Giemsa, Williams) and better show the structural differences at the level of the head, acrosome or intermediate piece. The principle of vital staining is to use a stain which only crosses the membranes of dead cells (eosin, rose bengal, cresyl green) and a background stain which facilitates reading (methylene blue, nigrosine). Eosin-nigrosin staining is conventionally used with a sperm-stain mixture of between 1/10 and 1/20.

The examination will ideally be carried out with an immersion objective at $\times 1000$ magnification. One hundred spermatozoa will be counted or even more if the number of sperm is greater. Dead sperm will be stained red due to eosin having entered owing to changes in membrane permeability.

Sperm morphological abnormalities can be said to be primary (1) if they originate during the spermatogenesis phase (testis) or secondary (2) if they occur during their maturation phase (epididymis). The majority of sperm lesions are said to be primary. Some can be both primary and secondary such as the presence of droplets, tailless heads. Sperm abnormalities can also be classified as major or minor depending on whether or not they have a negative effect on fertility. Finally, they can affect the various parts of the spermatozoon individually or simultaneously.

The head may show abnormalities in the shape, size, duplication, position or structure of the acrosome. The anomalies of the intermediate piece concerned cytoplasmic rest and angulation. The tail can be too short, too long, absent, duplicated, coiled at the end of the flagellum, or even under the head, and even around it or angulated (angulation exceeding 90°).

A rate of 10% of abnormal spermatozoa has been reported in camel sperm (Al-Bulushi et al. 2018) without significant effect of the frequency of collection on this parameter. The teratozoospermia of camel sperm was given to ejaculates with more than 40% abnormal spermatozoa (Waheed et al. 2014). The proportion of

protoplasmic droplet spermatozoa was 1.02 ± 0.2 on average, while 2.7 ± 0.6 and $9.7 \pm 2.9\%$ had mid-piece and tail defects, respectively (Wani et al. 2008).

2.6.5 Sperm Dilution and Conservation

2.6.5.1 Fresh Semen Dilution

The purpose of camel sperm dilution is to increase the total mass of the sperm fraction to split the ejaculate into fertilizing doses while adding substances that ensure sperm survival during storage. The latter can last a few hours or from a few months to several years.

In the dromedary, the separation of the liquid fraction from the sperm is difficult due to the distribution evenly throughout ejaculation of the liquid fraction. In addition, the sperm consists of a very variable volume of glandular secretions and that the testicular secretions constitute the essential of the semen.

The dilution medium requires the following characteristics: osmotic pressure which must be isotonic with the sperm and be able to maintain it during the storage period, contains colloidal substances (egg yolk, lipoproteins, lecithins) to ensure spermatozoa protection, contains buffer substance which makes possible to maintain a pH favorable to spermatozoa but its presence is not too important given the low concentration of spermatozoa in the sperm of the dromedary leading to a slight decrease in pH, contains nutrients to support metabolism, vitality and longevity of spermatozoa and contains antimicrobial agents.

The well diluted sperm offer best condition to spermatozoa and allow them to fulfill their 4 functions prior to fertilization: metabolic activity producing energy, mobility to progress in the female genital tract, protective enzymes on the acrosome to facilitate penetration into the oocyte and the presence of proteins on the plasma membrane to ensure their optimal survival in the female genital tract and their attachment to the pellucid of the oocyte.

To use camel sperm few minutes to a few hours after collection, it must be diluted at a ratio of 1:1 to 1:3 in a suitable diluter previously warmed. The dilution depends on the concentration of the ejaculate and has objective to achieve a standard concentration of 100×10^6 sperms/ml (Anouassi et al. 1992). This value can be revised downwards or upwards depending on the quality of collected sperm and insemination method (Bravo et al. 2000; Skidmore and Billah 2006a). Diluted sperm can be stored at room temperature (37°C) until insemination (Tibary and Anouassi 1997b).

2.6.5.2 Short Term Preservation

If the sperm is stored for up to 24 h, it is mandatory to gently decrease the temperature of the mixture sperm-diluter to 4°C by placing the tube in a water bath at room temperature and then refrigeration to reach 5°C after about an hour (Tibary and

Anouassi 1997b). Equitainer of equine semen can be used for the semen transport during 18–36 h. The researches on camelid semen extenders remind scarce. Most extenders used for camel sperm are adapted from bull and stallion commercial sperm extenders as Glucose-EDTA, Dimitropolous 11, INRA-96, Sodium-citrate egg yolk, Skim milk, Kenny's equine extender, lactose egg yolk (Tibary and Anouassi 1997b, 2018) and have showed a low impact on sperm motility and viability. A commercial extender has been available for camel semen since the early 90s (Camel Buffer Green[®], IMV, L'aigle, France). Recently, Al-Bulushi et al. (2019a) recorded that INRA is less suitable than Green Buffer to preserve semen at refrigeration for 24/48 h. It is mandatory to remember that for short-term preservation of camel sperm, it is essential to leave the time necessary for the complete liquefaction of the sperm before adding the diluter in order to obtain a homogeneous mixture (Tibary and Anouassi 1997b) and possibly reduce the effects of dilution on sperm mobility. It will be remembered that there are still great variations in the characteristics of the sperm of dromedaries subjected to short-term storage.

2.6.5.3 Long Term Preservation

If semen is to be used beyond 24 h after collection, freezing should be considered. There is no consensus on successful strategies for maintaining the fertilizing lifetime of frozen-thawed camel spermatozoa. The large scale application of AI in dromedary camels is impeded by this constraint. Freezing requires use of cryoprotectant agents. In view of the potential deleterious effects of cryoprotective agents on the sperm, they must be used at an optimal dilution which allows their benefits to be obtained and prevent osmotic and thermal shock from occurring both during dilution, cooling, freezing and thawing processing (Watson 2000).

Semen from DC was frozen successfully in a variety of extenders: Green Buffer[®] Egg yolk-glycerol, INRA and Egg yolk (Crichton et al. 2015), Tris-egg yolk-glycerol (Deen and Sahani 2006). Two steps are required for the cryopreservation of dromedary spermatozoa: dilution with cooling extender (80 ml Lactose 11%, 20 ml Egg Yolk) and dilution with freezing extender (95.5 ml cooling extender, 06.0 ml Glycerol and 1.5 ml Orvus paste –Equex). Immediately after collection, the cooling extender is added to the semen. The freezing extender is characterized by the fact that it contains a 7% of glycerol and an Orvus paste (emulsifying agent) that plays a role in plasma membrane stabilization of the sperm. (Tibary and Anouassi 2018). The first diluter is maintained at 32 °C and the second diluter at 4 °C. The Tris based extender with 7% glycerol and sucrose based extender (73.0 ml Sucrose 12%) showed superiority to preserve post-thaw spermatozoa quality (mobility and acrosome integrity) and fertility of camel bull (Zhao et al. 1994; Tibary and Anouassi 2018; Akbar et al. 2018). Likewise cholesterol-loaded cyclodextrin enhanced cryo-survival, post thaw sperm motility (0 h and 3 h post thaw) at room temperature and post-thaw sperm progressive status (Crichton et al. 2015, 2016).

Packaging sperm in pellets is rarely practiced in dromedary camels. Pellets have the disadvantage of not being able to be identified correctly. More conventionally,

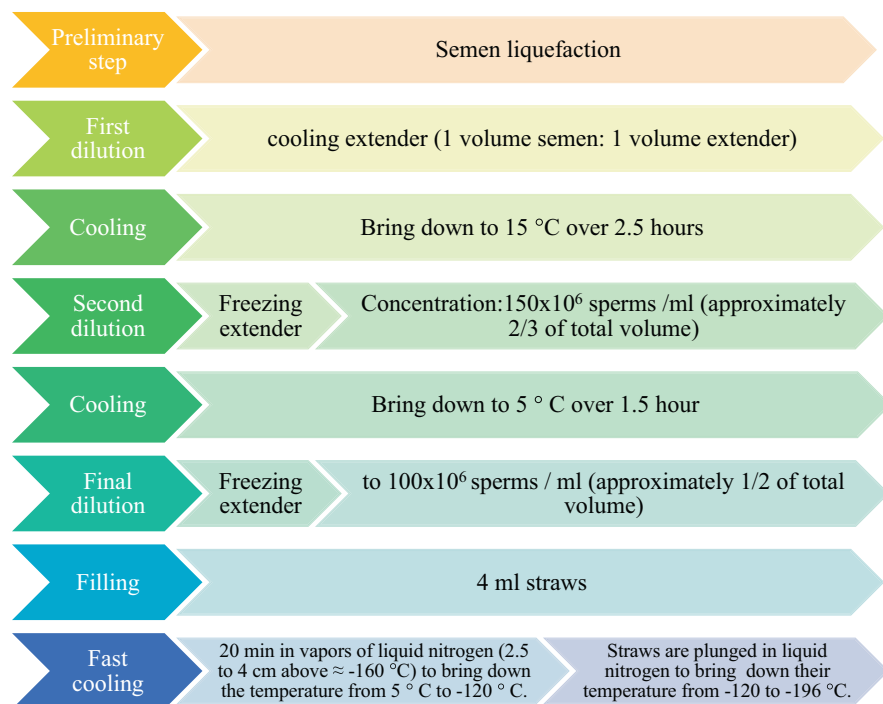


Fig. 2.21 Steps of camel sperm freezing in 4 ml straws

the sperm is packaged in polypropylene or polyvinyl straws with volumes between 0.25 and 4 ml. The required sperm dose has typically been a 4 ml straw that will contain 100×10^6 sperms/ml (Tibary and Anouassi 2018).

After semen liquefaction, freezing process depends mainly on the volume of the straws. When using large straws (4 ml) we proceed as follows (Fig. 2.21):

When we use semen package of 0.25 or 0.5 ml straws, the freezing process starts after total liquefaction according to the following steps (Tibary 2001; Deen et al. 2003; Niasari-Naslaji et al. 2006, 2007; El-Bahrawy 2010; Crichton et al. 2015):

- Dilution and cooling to 5 °C over 1 h.
- Semen package and equilibration at 5 °C for 2 h.
- Fast cooling by straws exposure to liquid nitrogen (4 cm above liquid nitrogen level) for 10 min
- Transfer of the cooled straws directly into liquid nitrogen.

Recent study performed at Camel Reproduction Centre, (Dubai, United Arab Emirates) by Malo et al. (2020), showed that camel sperm quality (total motility, viability and acrosome status) did not change significantly after 24 h cooling at 4 °C using two extenders (INRA 96 and Green Buffer) with or without cryoprotectant (3% glycerol or 3% ethylene glycol). Same authors reported also that shorter equilibration times (0–60 min) did not affect results of post-thaw sperms although

Glycerol 3% and ethylene glycol provided the best motility recovery rates compared to methyl formamide, dimethyl sulfoxide (Malo et al. 2017, 2020). In Addition, the post thawing results using Green Buffer showed highest rates of total and progressive motility and viability (Malo et al. 2020). For sperm viability, INRA with Ethylene Glycol 3% showed the lower rates compared to other extenders-cryoprotectant mixture (Green Buffer with glycerol 3% or Ethylene Glycol 3% and INRA 96with glycerol 3%) or buffers without any cryoprotectants (Green Buffer and INRA 96). Al-Bulushi et al. (2019a, b) recorded that better semen quality was observed using Triladyl (containing glycerol as cryoprotectant). In addition, the addition of catalase to extenders can act during prolonged sperm handling to prevent oxidative damage and thus promote chilled and cryopreserved sperm to retain better motility characteristics (Medan et al. 2008; Malo et al. 2019, 2020).

2.6.6 Sperm Thawing

Different thawing speeds were recommended depending on the packaging used for camel sperm.

The content of pellets is thawed out by dropping them into heated receptacles or incorporate it to thawed extender (Tibary and Anouassi 2018). Semen frozen in ampules is thawed out by placing them for 30 s to 1 min in a water bath set at 45–55 °C (Bravo et al. 2000). Straws (0.25 or 0.5 ml) can be thawed by placing them in water at 40 °C for 8 s or 30–60 s if the water is at 35–37 °C. The 4 ml straws are thawed at 50 °C for 40 s by continuous agitation in the water bath (Tibary and Anouassi 2018; Bravo et al. 2000). Cryovials are thawed out for 2 min by immersion in a water bath at 40 °C (Deen et al. 2003).

In recent study, Malo et al. (2020) reported that sperm extended in Green Buffer maintained better post-thaw vitality than sperm in INRA 96. The analyze of the effect the interaction of extenders with cryoprotectants on sperm vitality showed that INRA96-ethylen glycol 3% exerted most adverse effect on sperm vitality compared to INRA96-Glycerol 3%, Green Buffer-ethylene glycol 3%, Green Buffer-Glycerol 3% and both extenders INRA96 and Green Buffer without cryoprotectants. In other hand, all aforementioned sperm treatments did not showed any post-thaw effect on acrosome integrity (Malo et al. 2020).

Liquefaction of camel semen using Tris lactose supplemented with mucolytic agent (amylase) followed by centrifugation of seminal plasma removal in the presence of antioxidant agent (caffeine) improved significantly post-thaw motility and recovery rate suggesting a particular interest of this refined protocol for camel semen cryopreservation (El-Bahrawy 2017a, b). Similarly, semen extender supplemented with trace elements zinc and selenium nanoparticles (ZnONPs and SeNPs) improved significantly camel sperm vitality, progressive motility, ultrastructural morphology, sperm membrane integrity and decreased apoptosis when frozen and thawed (Shahin et al. 2020). These results meet the previous established conclusion about advantageous effects of nanotechnology in sperm function by increasing

antioxidant effect and bioavailability and reducing the undesired liberation of toxic concentrations (Falchi et al. 2018; Shahin et al. 2020).

Damage that occurs during freezing-thawing processes primarily affects cell membranes (plasma and mitochondrial) and, in more serious cases, the nucleus (Ahmed et al. 2017). Evaluating the percentage of motile spermatozoa at various intervals during incubation is a classic method of evaluating the method of freezing/thawing semen. The development of computerized mobility analysis systems helped to reduce the subjectivity part of this assessment. (Tibary and Anouassi 2018) reported other evaluation methods proposed for camel species such as hypoosmotic swelling test (incubation in fructose or sucrose solution of 50–100 mOsm at 37 °C for 45 min), special staining techniques (Isothiocyanate-conjugated peanut agglutinin) (Morton et al. 2007, 2008), Chlortetracycline staining for spontaneous capacitation (Crichton et al. 2015). The correlation between all these methods and the pregnancy rate remains to be demonstrated after a wide application of AI in camel herds.

2.6.7 Insemination Technique

Compared to other animals such as saw and ewe, camel insemination is simpler since the cervix is shorter and straighter and the uterus is less coiled, so it is easier to pass a catheter through the cervix and to direct it up the uterine horn per rectum (Skidmore et al. 2013). The ovarian function of the female camels must be monitored by series of ultrasonography exams to follow the growth of the dominant follicle which will be ovulated before AI. Ovulation induction is commonly performed out with a single injection of GnRH/hCG given intravenously when there is a 1.3–1.7 cm antral follicle in the ovaries (Skidmore et al. 1996; Medan et al. 2008). AI must be carried out 24 h after the injection of GnRH/hCG. The equipment consists of an insemination gun of 40–45 cm length and 5–6 mm diameter with an outer body and an inner mandrel. It is completed by an external plastic sheath attached to the insemination gun by means of a small washer. The fresh or thawed semen straw is wiped to remove all traces of water (water = spermicide) and the identity of the camel bull is immediately verified. It is then sectioned approximately 1 cm from its closed end and introduced into the insemination gun previously heated by friction to avoid any thermal shock. A plastic sheath (disposable) ensures the sanitary protection and the tightness of the device. It is possible to fit a flexible catheter within the rigid outer catheter preheated to 38 °C to avoid thermal shock for deep uterine insemination (Fig. 2.22) (Tibary and Anouassi 1997c).

The insemination technique is that of cervical catheterization with immobilization of the latter rectally (Fig. 2.22). The right or left hand introduced into the rectum, grasps the cervix and the other hand introduces the catheter into the vulva (previously cleaned), pushing it forward and following the ceiling of the vagina (angle of 45°) to avoid the urinary meatus. Vaginal folds are avoided by pushing the cervix held with the right or left hand forward (this clears the folds).



Fig. 2.22 Artificial insemination technique using special catheter for deep uterine deposit of camel semen. (a) Sterile deep horn insemination gun (b) Female prepared for deep horn insemination (c) Insertion of the deep horn insemination gun and transrectal guidance of the flexible catheter (d) Deep horn AI gun after insemination. Note the flexible catheter within the rigid outer catheter. (Photos reprinted with permission of the editor: Ahmed Bamouh, from Tibary and Anouassi 2018)

Locating the orifice of the cervix into which the catheter tip is to enter requires skill and must be done without any brutality. The hand that moves the cervix must, by the way a glove is put on, manipulate the cervix so that it comes over the catheter while avoiding the cervical folds until it reaches the cervico-uterine portion. Although it is easy to pass the catheter through the open cervix of the female during oestrus, under no circumstances should the catheter be forcefully pushed into the cervix (risk of injury).

It was therefore suggested that the semen should be deposited at the tip of the ipsilateral uterine horn to the ovary showing the mature follicle, thus closer to the uterine tubule junction rather than the uterine body (Skidmore 2019; Al-Bulushi et al. 2019b). The index finger of the hand inserted into the rectum controls, through the rectal wall, the correct position which allows the semen to be deposited (push

Table 2.5 Pregnancy rate of female camel according to different methodologies of artificial insemination

Author	Ovulation induction	Insemination time	Type of semen	Collection method	Females inseminated	Place of deposition	Pregnancy rate (%)
Deen et al. (2003)	hCG	24 h post hCG	WF	300 × 10 ⁶	13	UH	40
	hCG	48 h post hCG	EC	300 × 10 ⁶	10	UH	0
	hCG	24 post hCG	FT	300 × 10 ⁶	10	UH	7
Medan et al. (2008)	hCG	48 h post hCG	WF	100 × 10 ⁶	13	UB	46
		48 h post hCG	EC (free-catalase)	100 × 10 ⁶	9	UB	22
		48 h post hCG	EC (with catalase)	100 × 10 ⁶	8	UB	37
Anouassi et al. (1992)	Vasectomized male		WF	100 × 10 ⁶		UB	80
McKinnon et al. (1994)	hCG		WF	AV			32
Skidmore and Billah (2006a)	GnRH		E	150 × 10 ⁶	15	UB	53
	GnRH		E	150 × 10 ⁶	14	UH	43
	GnRH		E	80 × 10 ⁶	14	UB	7
	GnRH		E	80 × 10 ⁶	15	UH	40
	GnRH		E	40 × 10 ⁶	14	UB	0
	GnRH		E	40 × 10 ⁶	14	UH	7
Al-Bulushi et al. (2019b)	GnRH	Immediately after ovulation treatment	WF (3–8 ml)	474 × 10 ⁶	11	UB	81
	GnRH	Immediately after ovulation treatment	WF (1.5–3.5 ml)	442 × 10 ⁶	10	UB	60
	GnRH	Immediately after ovulation treatment	WF (1 ml)	454 × 10 ⁶	12	UB	33

(continued)

Table 2.5 (continued)

Author	Ovulation induction	Insemination time	Type of semen	Collection method	Females inseminated	Place of deposition	Pregnancy rate (%)
Al-Bulushi et al. (2019b)	GnRH	Immediately after ovulation treatment	E	300 × 10 ⁶	17	UB	35
	GnRH	Immediately after ovulation treatment	E	301 × 10 ⁶	18	UH	72
	GnRH	Immediately after ovulation treatment	E	302 × 10 ⁶	19	UH	68
Al-Bulushi et al. (2019b)	GnRH	24 h post GnRH	EC	303 × 10 ⁶	20	UH	70
	GnRH	30 h post GnRH	EC	304 × 10 ⁶	17	UH	23
	GnRH	24 h post GnRH	E (3.5 ml)	300 × 10 ⁶	20	UH	70
	GnRH	24 h post GnRH	E (1.7 ml)	150 × 10 ⁶	23	UH	65
	GnRH	24 h post GnRH	E (0.9 ml)	75 × 10 ⁶	22	UH	40
	GnRH	24 h post GnRH	EC (Triladyl)	300 × 10 ⁶	18	UH	11
Al-Bulushi et al. (2019b)	GnRH	24 h post GnRH	EC (Optixcell)	300 × 10 ⁶	17	UH	5
	GnRH	24 h post GnRH	EC (Triladyl)	500 × 10 ⁶	19	UH	21
	GnRH	24 h post GnRH	EC (Optixcell)	500 × 10 ⁶	16	UH	12
	GnRH	24 h post GnRH	FT	300 × 10 ⁶	23	UH	0
Morton et al. (2010)	GnRH	24 h post GnRH	FT	500 × 10 ⁶	20	UH	0
	GnRH	24-48 h post GnRH	FT	150 × 10 ⁶	11	UB	27
Akbar et al. (2018)	GnRH	30 h post GnRH	FT	100 × 10 ⁶	21	UB	42 ^a

UH uterine horn, UB uterine body, ^apregnancy at 90 days post insemination, WF whole fresh, E extended, EC extended cooled, FT frozen thawed, AV artificial vagina

exerted on the plunger) at just this optimal level. The utero-tubal zone acts as a reservoir for spermatozoa. These are released regularly and continuously in order to ensure fertilization in the upper third of the oviduct without there being polyspermia (Skidmore 2019).

2.6.8 Factors Affecting Fertility After AI

AI with raw undiluted camel ejaculates resulted in wide variability of pregnancy rate: 40–81.8% pregnancy rate (Table 2.5; Anouassi et al. 1992; Deen et al. 2003; Skidmore and Billah 2006a, b; Medan et al. 2008; Morton et al. 2010; Al-Bulushi et al. 2019b). These results are somewhat comparable to results of timed natural mating programs: 67–71.4% (Manjunatha et al. 2015, 2018; Al-Bulushi et al. 2019b). Low volume (1 ml) insemination with undiluted ejaculate resulted in lower pregnancy compared to whole and split undiluted ejaculate. (Al-Bulushi et al. 2019b). However, some studies showed lower or null results of pregnancy rate using chilled and non diluted semen for AI (El-Hussanein 2003; Medan et al. 2008; Morton et al. 2013).

The pregnancy rate after AI using chilled camel semen diluted by different extenders show large variability of pregnancy rate (Table 2.5): 0% using skimmed milk (Skidmore 2000); 5.9–12.5% using Optixcell (Al-Bulushi et al. 2019b); 11.1–21.1% using Triladyl (Al-Bulushi et al. 2019b); 17–34% using INRA 96 (Morton et al. 2010, 2013; Malo et al. 2020); 50% using lactose plus egg yolk (Anouassi et al. 1992), 53% using Laciphos (Skidmore 2000); 10–72.7% using Green buffer plus egg yolk (Al-Bulushi et al. 2019a, b; Skidmore and Billah 2006a; Morton et al. 2010, 2011, 2013; Malo et al. 2020). The wide variability of the pregnancy rate after AI by liquid chilled semen would be difficult to explain especially when considering that *in vitro* semen quality appears to be high.

The post-thaw frozen semen showed promising consistency, but this is not evidenced by the pregnancy outcomes to date. Single early pregnancy in DC has (1/13 animals inseminated) resulted from AI with cryopreserved spermatozoa (Deen et al. 2003). The use of glycerol as cryoprotectant with Green Buffer extender during chilling did not interfere with cryosurvival, but it may be toxic to the fertility of fresh chilled sperm leading to null pregnancy rate (Malo et al. 2020). After that, Morton et al. (2010) reported a pregnancy rate of about 27% in single ovulated female camels using frozen-thawed semen diluted in Green Buffer and fertilization rates of about 11.1% in multiple ovulating camels. The first successful encouraging results for large scale utilization of frozen semen insemination of dromedary camels was recently reported by Akbar et al. (2018) after a series of experiments on semen processing, cryopreservation and AI giving live birth of camel calves at Camel Breeding Center, Dubai, United Arab Emirates date 2018 (Akbar et al. 2018). In this experiment; 21 inseminated females by frozen -thawed semen with tris based extender and 7% glycerol gave the following results: On the 30th day of insemination, 15 (71.42%) females were found to be pregnant. Between the 18th and 30th

days, 6 she-camels faced embryonic mortality (Akbar et al. 2018) and pregnancy was confirmed in the remaining females at 60 and 90 days of insemination (09). At mid gestation, two females aborted and the remaining seven females (33.33%) were carried to full term. More investigations are needed to confirm the fertility rate using frozen thawed semen of eventually large scale application of AI as important step of assisted reproductive techniques in camelids.

The monitoring of the ovarian follicular wave patterns of the female camel by transrectal ultrasound examinations is mandatory for AI success. Ovulation is induced by a single intravenous injection of GnRH (ex: 20 mg buserelin) accompanied by timed artificial insemination once the dominant follicle reaches 1.3–1.8 cm (Skidmore and Billah 2006b). Female camels inseminated 24 h after ovulation induction resulted in higher pregnancy rate (53.7–70.0%) (Skidmore and Billah 2006b; Al-Bulushi et al. 2019a, b) compared to insemination performed at the time of ovulation induction or 30 h avert induction (68.4 and 23.5%, respectively) (Al-Bulushi et al. 2019a, b).

The results of the insemination into the uterine body or the tip of uterine horn ipsilateral to the ovary containing the dominant follicle showed respective increasing conception rate of 0, 7, 53% and 7, 40, 43% according to increased sperm concentration per milliliter of 40, 80 and 150×10^6 (Skidmore and Billah 2006a). Deposition of semen in the uterine body resulted in lower pregnancy rates compared to deposition in the tip of the horn (35.3% versus 72.2%) (Al-Bulushi et al. 2019a, b). AI with 75×10^6 motile spermatozoa resulted in lower pregnancy rates compared to 150 and 300×10^6 motile spermatozoa doses (40.9% versus 65.2 and 70.0%, respectively) and pregnancy rate without significant correlation with semen extenders (Green buffer, Optixcell or Triladyl) (Al-Bulushi et al. 2019a, b). Indeed, the use of insemination with small sperm concentration requires a deep uterine insemination technique. If semen is deposited only in the uterine body rather than at the tip of the horn a substantial loss of spermatozoa occurs due to the semen's backflow through the short and open cervix during the female camel's oestrus.

The short, open cervix that occurs during oestrus of the female camel would lead to considerably loss of spermatozoa, due to backflow of semen through the cervix, when the semen is deposited just into the body of the uterus rather than at the tip of the horn (Skidmore and Billah 2006a; Al-Bulushi et al. 2019a, b). Uterine manipulation during the intrauterine insemination may simulate penile stimulation by male camel or aide sperm transport by a greater release of uterine hormones involved in uterine contractility (Martinez et al. 2002).

Form the given studies; it appears that large scale utilization of AI in camels faces many technical difficulties particularly those relative to oestrus synchronization, preservation of camel sperm and low fertility of females inseminated with frozen semen. In addition, many basic concepts relevant to AI application are still not available, such as infectious disease screening, biosecurity controls, male management, training procedures and hygienic collection and processing of semen (Monaco and Lacalandra 2019).

2.7 Conclusion

Good reproductive performances allow obtaining a high numerical productivity of camel herds and consequently increasing the information available on the siblings and shorten the average generation interval. However, in traditional reproductive management, increasing reproductive efficiency of camel herds faces certain constraints such as long gestation period, short breeding season and traditional reproductive and herd management methods leading to widespread venereal infections and reducing fertility. Oestrus synchronization and AI are simplest and cheapest reproductive technologies to overcome some of these problems. They have a significant influence on the genetic improvement rate. A higher reproduction rate means a lower number of breeding animals for a given population size and, thus, a greater selection strength. A greater number of offspring per breeder also promotes a more accurate estimate of genetic values. Another advantage is the faster diffusion of higher genetic material and international movement of genetic material. To date, it is difficult to accept large scale practice of artificial insemination in DC because of the technical and zootechnical difficulties associated with this biotechnology. Investigations on the technical aspects are still in progress; such as semen collection methods, semen liquefaction and cooling, deep freezing and thawing of camel spermatozoa to optimize AI protocols and standardize procedures. The main zootechnical constraint is the nature of the breeding systems of camel herds which are in continuous movement throughout the year in pastoral areas. This makes difficult the ultrasound monitoring of ovarian activity, ovarian activity synchronization followed by timed ovulation induction and AI or blind mating. In addition, there is the risk of inbreeding and the reduction in reproductive capacities by using a reduced number of breeding males over a large number of females during short breeding season.

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Chapter 3

Biotechnological Advancements in Livestock Production



Bhaskar Sharma, Dixita Chettri, and Anil Kumar Verma 

Abstract Animal-based agriculture is being used to surmount the food demand for the ever-increasing human population under limited land and water resources. The meat and milk consumption has severely grown over the past few decades with increasing population and the demand is expected to grow by double in the next three decades. The fulfillment of milk, meat, and related products demand has turned into a greater challenge for poor and developing countries which is leading to malnutrition. The traditional methods for livestock production are facing challenges due to limited offspring production, unstable genetic improvement, tedious methods, high cost, and disease susceptibility. Here we review the modern technological developments offering interesting solutions to the issues associated with conventional methods for animal production. The advanced biotechnological interventions such as *in vitro* fertilization, somatic nuclear transfer, embryonic stem cell technology, multiple ovulations, and embryo transfer could improve the nutritional composition and animal health. The capabilities of precise and desired interventions at the cellular and molecular levels renders unprecedented developments in animal reproduction. Molecular marker-assisted selection and transfer of useful and superior traits to the recipient line ensures improvements such as better growth, quality, and disease-resistance for several generations. The chapter covers biotechnological methodologies adopted in modern animal breeding, including marker-assisted selection methods for stable and improved livestock production.

Keywords Marker assisted selection · Somatic cell nuclear transfer · Artificial insemination · Embryo transfer · Stem cell technology

B. Sharma

School of Life and Environmental Sciences, Faculty of Science, Engineering, and Built Environment, Deakin University, Geelong, VIC, Australia

Department of Biotechnology, TERI School of Advanced Studies, New Delhi, India
e-mail: sharmabha@deakin.edu.au

D. Chettri · A. K. Verma (✉)

Department of Microbiology, Sikkim University, Gangtok, Sikkim, India
e-mail: dchettri02@sikkimuniversity.ac.in; akverma@cus.ac.in

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Abbreviations

AFLPs	Amplified fragment length polymorphisms
AI	Artificial insemination (AI)
BCWD	Bacterial cold-water disease
bFGF	Basic fibroblast growth factor
BLUP	Best linear unbiased prediction
BOEC	Bovine oviduct epithelial cells
CRISPR Cas9	Cluster regularly interspaced short palindromic repeats
DNA	De-oxy ribonucleic acid
eCG	Equine chorionic gonadotropins
ESC	Embryonic stem cell technology (ESC)
ET	Embryo transfer (ET)
FAO	Food and Agriculture Organization
FSH	Follicle-stimulating hormones
GAS	Gene-assisted selection
IETS	International Embryo Technology Society
IVF	In vitro fertilization (IVF)
JAK/STAT	Janus kinase and signal transducer and activator of transcription proteins
LD	Linkage disequilibrium
LE	Linkage equilibrium
LH	Luteinizing hormone
LIF	Leukemia inhibitory factor
MAI	Molecular marker-assisted introgression
MAS	Marker Assisted Selection
MOET	Multiple ovulation and embryo transfer (MOET)
NGS	Next-generation sequencing
PCC	Premature chromosome condensation
QTL	Quantitative Trait Locus
RAPD	Random amplified polymorphic DNA
RFLPs	Restriction fragment length polymorphisms
SCNT	Somatic cell nuclear transfer (SCNT)
SCR	Sire conception rate
SNP	Single nucleotide polymorphism
SSR	Simple sequence repeats
TALEN	Transcription activator-like effector nuclease
TGF-beta	Transforming growth factor beta
TGS	Third-generation sequencing

3.1 Introduction

Livestock production began with the domestication of sheep, chicken, pigs, goats, and other cattle about 8000–10,000 years ago (Porter 2020). Eventually, the need for the genetic improvement of livestock turned up to produce better characteristics through breeding. Later, with the rediscovery of Mendel's law (Mendel 1866) and the establishment of modern genetics, the selection and crossbreeding of livestock revolutionized genetic improvement and production. The growth rate of the commercial production of the pig, chicken, and other poultry was substantially enhanced in unprecedentedly lesser time (Muir and Aggrey 2003). Therefore, the development of genetic improvement tools led the increased livestock production in developed and developing countries. The global livestock production attributes to around 40% of total agricultural production. Livestock production contributes more than half in developed countries and approximates one-third in developing countries. Livestock demand is increasing in developing countries due to the larger population size and dietary habits. On the other side, the agricultural food consumption was decreased to 1.1% from 2.2% in the last 40 years. The annual per capita meat consumption was increased by 100% in developing countries between 1980 and 2002. The FAO (Food and Agriculture Organization) projected an increase of 60% in the global animal food production demand by 2050, considering the huge population growth (Alexandratos and Bruinsma 2012). There is an urgent requirement for the Livestock revolution to meet the population food demand while sustaining limited natural resources. The emerging trends in the livestock sector are influenced by increasing competition for grazing and water resources, enhanced usage of pigs and poultry, livestock production in hot, moist, and disease-prone habitat, and health risks due to industrialization and urbanization (Harrison et al. 2002). There is a need to introduce effective public policies for natural resource management, food safety, environmental pollution, and public health considering large transformation in the livestock sector (Harrison et al. 2002).

Recently, the demand for animal-based foods such as eggs, milk, and meat has increased with the growing population in developing countries. The share of total calories from animal-based food has risen in the past few decades, while the total calories from traditional staples and root-based products such as cassava, potatoes, and plantains have decreased by approximately 4%, making animal-based products the second major source for the calories after cereals in developing countries. The consumption of animal-based products is still low in developing countries compared to developed countries but continuously rising over time. The most organized structural organizations for livestock production and maintenance have been intensive establishments in urban areas for poultry and meat production in East Asia and Latin America (Seré C 1996). The production systems for the livestock have dramatically evolved because of a surge in the demand and increasing pressure on natural resources. The use of the grazing system contributes to a large proportion of total livestock production. However, the grazing systems are severely affected by climate change, land degradation, land enforcement, and increasing population. Similarly,

using limited land with crop-livestock production, where ruminant livestock is grown with crops in a mixed system, is a popular choice in developing countries (Seré C 1996). Crops are used to feed the animals and livestock that produce manure and food that generate diversified income for rural farmers. Such livestock production systems are critical for household food security and immediate income generation with limited resources. Another type of production system is an intensive industrial livestock production system that involves heavyweight machinery, limited land area, improved production, and low-cost production. It is particularly beneficial for meeting the demand of the growing population but could reduce the value of the livestock (Seré C 1996).

Biotechnology can be defined as a technological application using a biological system to produce or modify any method or product for human benefits. The animal, plants, and microorganisms are modified using a set of techniques to produce desired products for human use. The application of the modern knowledge of the techniques and methods results in the higher qualitative and quantitative improvement in the desired traits of the selected organisms at a lower cost. The discovery of the genetic code led to the origin of the new modern method known as 'biotechnology'. The first-ever product developed using biotechnology in the field of animal sciences was bovine somatotropin metabolite, commercially produced at a large scale and administered to more than half cattle in the United States of America. Such technologies are necessary for ensuring nutritive food for the growing population. The major concern with the use of biotechnology has been observed with regulatory policies where potential risks of releasing genetically modified materials in the environment have attracted the attention of the wider scientific community. Biotechnology largely deals with the modification of the genetic material that poses a potential threat to human health and the environment. Sometimes, the quality of the product is compromised when producing in large quantities. Given the safety concerns, industrial manufacturing must be regulated in a robust and organized manner to establish a solid livestock production system. Overall, biotechnological interventions have revolutionized animal-based production, making it a stable industry for feeding future generations.

3.2 Biological Methods in Animal Breeding

3.2.1 Somatic Cell Nuclear Transfer (SCNT)

The cellular ability to generate all cell types is termed as totipotency (Lu and Zhang 2015). In mammalian cells, the totipotency is observed in the blastomere of preimplantation embryos and eventually lost during embryo development to produce inner cell mass. The totipotency is a vital characteristic of somatic cell nuclear transfer (SCNT) that can be used to clone animals such as sheep (Wilmot et al. 1997a), cows (Cibelli et al. 1998), pigs (Onishi et al. 2000), and rabbits (Chesné

et al. 2002) from differentiated somatic cells (Gurdon 1962). The somatic cell nuclear transfer is a method where DNA from the somatic cell of the donor is transferred to the enucleated metaphase II oocyte of the acceptor for the production of genetically identical cells to the donor. The discovery of Dolly through SCNT led the researchers to think about cellular differentiation reprogramming. The differentiated nucleus are reversed to attain totipotency using nuclear reprogramming. The reprogramming of the nucleus through SCNT involves enucleation from recipient cells followed by injecting the nucleus from donor cells and finally activation of the reconstructed embryos. The nuclear membrane breaks down during an enucleation process to make a premature chromosome condensation (PCC) promoted by M-phase promoting factors (Campbell et al. 1996). PCC is necessary for initiating the reprogramming and in which the chromatin-bound proteins dissociate from the genome. After fertilization, the phospholipase from sperm triggers the oocyte activation through calcium signaling (Saunders et al. 2002). In the case of the absence of phospholipases, the activation is induced by either treatment of the strontium chloride or electro pulse treatment (Liu et al. 2014). Later, nuclear expansion occurs in the fertilized zygote to form a paternal and maternal pronucleus followed by DNA replication. The zygote further activates the genome by restoring transcription in the new genome that is known as a zygotic genome activation. Further, the SCNT process extends to chromatin, transcriptional, and epigenetic reprogramming that leads to the development of the clone. SCNT is an excellent tool for the study of genetic disease, growth regulation, gene therapy (Capecchi 2000), and genetic imprinting (Solter 2004). SCNT can be used to produce pharmaceutically important organisms, therapeutic proteins, and the preservative of endangered species. Despite the advancements, developmental abnormalities and low transformation efficiencies were observed that limit the widespread usage of the SCNT (Hill et al. 2000b; Kubota et al. 2000). The low transformation efficiencies varied from 0–10% and were predicted due to errors in the epigenetic signal reprogramming. The abnormalities could result in immunodeficiency, early death, respiratory defects, and obesity (Loi et al. 2016). Several factors can affect nuclear reprogramming, such as the origin and age of the donor and recipient cells (Wells et al. 2003; Hill et al. 2000b), epigenetic signals (Enright et al. 2003), tissue, and cell culture (Zakhartchenko et al. 1999), and drugs (Kubota et al. 2000). There were instances when the SCNT efficiency was enhanced through optimized experimental handling of the SCNT (Wilmut et al. 1997b). Overall, the SCNT method is highly suitable for cattle and other animals for genetic improvements. The method can undergo a wide variety of genetic improvements and produce quality transgenic lines without affecting animal biological properties. The somatic cells are very easy to collect, store, and transport that increases the success rate and stability of the method. SCNT is very useful in the biomedical applications and the therapeutic treatments of the animals. However, high cost and tedious process is limiting its extensive application. Moreover, poor oocyte recovery has been observed in few cases that need to address before successful implementation of the method (Ogura et al. 2002). There are several legal and ethical restriction imposed on the SCNT products in many countries (Kishigami

et al. 2006). The SCNT needs significant improvement to be easily accessible to most of the part of the world.

3.2.2 *Artificial Insemination (AI)*

Artificial insemination is a method in which the semen from a male donor is placed into the reproductive tract of the female recipient. The method does not use any natural method for mating or reproduction (Salisbury et al. 1978). Artificial insemination is a part of a group of technologies known as assisted reproduction technologies. The methodology allows the transfer of the conception products into females and the collection of semen, the development of reliable methods for inserting semen into the female reproductive organ is a prerequisite. Usually, the semen is collected into a sterilized insulated vessel using an artificial vagina consisting of suitable temperature and pressure for stimulation and ejaculation (Salisbury et al. 1978). Few other methods may include electroejaculation apparatus for collecting semen from animals with abnormalities or as an alternative for extreme conditions. Semen is sensitive to enzymatic and chemical exposure (Wani et al. 2008). The semen exists in liquid form and consists of various energy molecules such as protein, magnesium, calcium, zinc, and fructose. The semen plasma helps spermatozoa reach the fertilization site and survival for successful fertilization. The collected semen is used fresh or can be stored at 4 °C. The spermatozoa are sensitive to a higher temperature whereas the lower temperature minimizes metabolic activities and bacterial infection that increase the lifespan of spermatozoa. Cryopreservation can be opted for long-term preservation at a much lower temperature (Brinsko and Varner 1992). The semen sample is mixed with the sugar, lipoprotein, and cryopreserving glycerol for longer time preservation. Natural detection of estrus and ovulation is a critical task for timely insemination (Bó and Baruselli 2014). The ovulation can be induced using vasectomized boys or the administration of the using hormones. Artificial insemination has been improvised and upgraded continuously with the latest technological developments.

Artificial insemination is the most common method for improving domestic livestock such as dairy cattle, pigs, horses, and sheep (Table 3.1). The usage of artificial insemination has been extended to human fertility treatments. The physical contact between livestock was the major reason for the various disease transmission, which resulted in a shorter life span and loss in productivity. Artificial insemination avoids contact between livestock and retains the quality of the livestock for a longer time. The strength of artificial insemination is the genetic improvement and higher production yield using the semen of superior quality male for best female (Morrell 2011). The methodology is highly useful for the conservation of genetic material of the endangered species and the revival of the genetic merits of the animals. The breeding became possible for animals under physical challenges or abnormalities, and different relocation. Despite the advantages of technology, few limitations were observed for the artificial reproduction of the animals. The viral and bacterial

Table 3.1 Major advantages and limitations of biological methods in animal breeding

Sr. No.	Technology/ method	Advantages	Constraints/limitations	References
1	Somatic cell nuclear transfer (SCNT)	<ol style="list-style-type: none"> 1. Introduction of selective genetic enhancements 2. Capable of producing efficient transgenic offspring and transgenic animal lines 3. Wide range of genetic alteration to <i>in vitro</i> cultured cells 4. No major biological properties and nutritional quality alteration in cloned animals compared to non-cloned animals 5. Easy to obtain, culture, and cryopreserve somatic cells 6. Therapeutic and biomedical applications 	<ol style="list-style-type: none"> 1. Poor quality oocyte recovery 2. Abnormalities in generated clones such as defected phenotypes 3. Resource intensive and costly procedure 4. Several legal and ethical issues related to the use of SCNT products 	Pickering et al. (2005), Hill et al. (2000a), Ogura et al. (2002), and Kishigami et al. (2006)
2	Artificial insemination (AI)	<ol style="list-style-type: none"> 1. Limits the spread of disease and sterility 2. Early detection of semen defects ensures better breeding efficiency 3. Semen transportation, storage, and extensive uses are possible 4. Helpful to inseminate physically different sized animals and increased rate of conception 5. Reduce the number of sires maintained for insemination 	<ol style="list-style-type: none"> 1. Time-consuming, and tedious operations 2. Well trained staff needed, and special instruments required 3. Advanced hygiene and maintenance of the instruments 4. The spread of genetic defects is possible 5. Reduction of the effective population of breed and inbreeding can spread 	Noakes et al. (2018)
3	Embryo transfer (ET)	<ol style="list-style-type: none"> 1. Reduced risk of disease transmission due to limited transportation of live animals 2. Easy to exchange genetic material and promote international trading 3. Increased number of offspring per female 4. Storage and expansion of rare and endangered species genetic stock 5. Desired mating and faster genetic process 	<ol style="list-style-type: none"> 1. High cost involved in the method 2. Advanced expertise needed to conduct the embryo transfer 3. The success rate can be low sometimes 4. Spread of the disease through recipient is possible 	Bó and Mapletoft (2018)

(continued)

Table 3.1 (continued)

Sr. No.	Technology/method	Advantages	Constraints/limitations	References
4	Embryonic stem cell technology (ESC)	<ol style="list-style-type: none"> 1. Lesser animal testing, diagnosis, and research needed 2. Efficient preservation and revival of rare and endangered species 3. Targeted treatment and care for animals 4. Advanced genetic improvement is possible 	<ol style="list-style-type: none"> 1. Require skilled experts for the successful implementation of the technology 2. The costly method requires <i>in vitro</i> maintenance 3. Several legal and ethical issues associated with the use of ESCs 	Deckha and Xie (2008)
5	Multiple ovulation and embryo transfer (MOET)	<ol style="list-style-type: none"> 1. Improve the reproductive potential of the animals 2. Production of numerous offspring is possible 3. Easy collection, storage, and transportation of the embryos 4. No surgical methods are mandatory for the successful MOET 5. High yield and growth in a short period 6. MOET can be undertaken at farm area which reduces handling damages 	<ol style="list-style-type: none"> 1. Costly method and require skilled experts 2. Sometimes a longer waiting period for animals to become productive 3. Hormone dosage optimization to reduce the risk of abnormalities 	Bergstein-Galan et al. (2019)
6	In vitro fertilisation (IVF)	<ol style="list-style-type: none"> 1. Preservation of valuable genetic stocks for a longer period 2. Lesser hormonal treatments to animals 3. A shorter generation interval is needed 4. No impact of physiological abnormality on offspring 5. Semen cost is highly reduced 6. A range of donors can be used 	<ol style="list-style-type: none"> 1. IVF is a costly method and requires specialized instruments 2. Skilled and experienced staff needed 3. Special hygiene and maintenance required in every step 4. The success rate can decline if animals are facing oocyte inactivation 	del Collado et al. (2017) and Abdalla et al. (2009)

infection in the semen is a major threat to the successful implementations of artificial reproduction (Morrell 2011). The widespread use of artificial reproduction methods compromised the fertility of cattle and animals and eventual loss of genetic variation (Morrell 2011). Artificial insemination can adopt the latest biotechnological advancement and integrate new tools to improve the process of large-scale livestock production. Artificial insemination can be beneficial for insemination of different size animals, semen transportation, early detection of the disease, and limiting the disease transmission through animals (Noakes et al. 2018). However, the AI method cannot overcome with any possible genetic defect in donors or recipients. The AI has been used traditionally to breed animals at a larger scale and can be readily improved with new technological advancements.

3.2.3 Embryo Transfer (ET)

The embryo transfer has emerged as a complementary method of artificial insemination. The embryo transfer is an artificial breeding method where the embryos are retrieved from an elite donor female before implantation and transferred to a recipient female that serves as surrogate mother (Wilmut et al. 1985; Betteridge 1981). The embryos are recovered through ovulation by hormonal treatment. The embryo transfer methods spread the desired genetic traits from superior female animals (Table 3.1). Embryo transfer can be achieved surgically or non-surgically. In the surgical method, the uterus of the donor animal is accessed through an incision on the midline or the flank (Tan et al. 1990). The uterine horns are flushed by penetrating the uterine wall using a tube from one end and with a hypodermic needle from another end. The medium is collected in a dish usually forced by a syringe. The embryos are roughly found to be 0.2 mm in size in the 32-cell stage of development. The embryos are examined under a microscope and picked up with a small glass pipette attached to a micro-syringe. These embryos are directly transferred to the recipient female uterus by puncturing the uterine wall with a needle. Further, the glass pipette with an embryo is inserted into the reproductive tract of the animal and embryo is released into the uterine lumen with slight pressure. The embryos are transferred within 4–5 h of recovery to minimize the risk associated with temperature and pH. Additionally, it is necessary to note that the estrous cycle of the donor and recipient animals should be in synchronization. Moreover, many chemical agents can be used to induce estrous-synchronization or prostaglandins.

The non-surgical method is based on the use of the foley catheter for collecting the embryos (Wright 1981). The genital area of the animal is washed properly, and local anesthetic injection is given. Later, the catheter is inserted into the vagina and goes through the cervix followed by the pressure release of the catheter. The flushed fluid is collected, and embryos are filtered from the collection of the medium. The procedure of embryo transfer involves the selection and management of the donor and recipient female. The estrus cycle detection should be normal for the superovulation to happen on the time. The superovulation and embryo production occur

eventually followed by the embryo transfer event. The embryo transfer reduces disease transmission, circumvent infertility, induce genetic testing, rapid exchange of genetic material, and increase the offspring per female (Smith 1988; Betteridge 1981). The embryo transfer technique has been applied to most domestic animals. Overall, embryo transfer is another major technique developed to enhance the impact of superior genotype in the population. The latest developments are introduced to embryo transfer to establish an easy and less destructive method. The ET ensure faster genetic improvements in major animal species and comparatively higher offspring generation (Bó and Mapletoft 2018). The ET reduces the risk of disease transmission and allows better storage and transportation of the embryo which makes international trade easy.

3.2.4 Embryonic Stem Cell Technology (ESC)

Embryonic stem cells (ESCs) are self-renewable and pluripotent cells that can differentiate into any cell type of primary germ layer. They originate from the inner cell mass of the pre-implanted blastocyst. These characteristics make them ideal candidates for genetic modification, gene therapy, and cell regeneration. ESCs are categorized into naïve and primed (Nichols and Smith 2009). The mouse ESCs are naïve pluripotency while human ESCs are developed into an advanced primed pluripotent state (Hanna et al. 2010). The naïve pluripotent stage in the mouse is characterized as domed colonies, depends on JAK/STAT signaling, and contains reactivated X-chromosome. These cells can contribute to the chimera formation into the blastocyst and exhibit higher pluripotent homogeneity. The primed ESCs are however, the naïve ESCs have failed to further differentiate into functional cell types in the various cell culture medium, and protocol optimization and establishment of the stable embryonic stem cell line are necessary before attempting the differentiation (Talbot et al. 1995; Wang et al. 2005). The derivation of embryonic stem cells in domestic cattle has not yet been established. Several studies including different feeder layers (Cong et al. 2014), micro-drop culture (Kim et al. 2012), and 2i conditions (Furusawa et al. 2013; Maruotti et al. 2012), combinations of different culture medium LIF and bFGF (Gong et al. 2010; Maruotti et al. 2012) showed poor pluripotency and limited cell proliferation. The derivation of the stable bovine pluripotent has been challenging and does not qualify the in vivo teratoma assay, in vitro embryoid body formation, and chimera formation (Blomberg and Telugu 2012; Ezashi et al. 2016). The bovine ESCs often show poor proliferation and derivation capabilities, and the disappearance of pluripotency markers (Soto and Ross 2016; Wang et al. 2005; Talbot et al. 1995). Recently, a study demonstrated pluripotency marker gene expression, stable morphology, similar karyotype, and transcriptome profiles, and population doubling time. The cell-culturing conditions used were custom base medium lacking TGF-beta supplemented with bFGF and Antagonist I (Wu et al. 2015). Antagonist I is an inhibitor of the WNK- β catenin signaling pathway that decides the cell fate and stem cell pluripotency. The

conditions were the same as ‘region-selective’ primed type pluripotent stem cells sharing molecular functions with gastrula-stage-epiblast in human ESCs. Apart from the latest developments in cattle, similar trials are being attempted to establish stable ESCs in sheep and pigs. The establishment of the bovine embryonic stem cells will facilitate early development and biotechnological applications such as genome editing for the genetic improvement of the livestock. The full potential of ESC technology has not been experienced yet but this can improve stable genetic integration and treatments in the animals. ESCs can differentiate into any cell of the animal body which will help increase integration of desired genetic changes (Deckha and Xie 2008). However, legal, and ethical restrictions have made the technology inaccessible to large part of the world and will take longer time to be adoptable. Genome editing and livestock genetic improvement using embryonic stem cell technology have greater potential to revolutionize the livestock breeding.

3.2.5 Multiple Ovulation and Embryo Transfer (MOET)

Artificial insemination and embryo transfer methods for breeding have been popularly used for the improvement of animal characteristics. These methods are not well established in cattle and are usually expensive. The Multiple ovulation and embryo transfer (MOET) method is an advancement of the embryo transfer method where the fertilized eggs are removed from the super-ovulated female donors and transferred in the genetically different multiple surrogate mothers (Bergstein-Galan et al. 2019). The MOET can be used in the sheep, camel, goat, buffalo, and other cattle except for horses which are difficult to super-ovulate. This method produces significantly higher offspring in cattle with improved characteristics with lower cost compared to traditional methods. Thus, genetically superior donors can generate numerous progeny than it could earlier with other traditional methods such as artificial insemination and embryo transfer (Vettical et al. 2019). The screening of the health and breeding capabilities of the suitable donors and recipients is a crucial part of the process of embryo transfer (Seidel Jr. and Seidel 1991). Animals without any disease or infection and reproductive abnormalities such as ovarian-bursal adhesion should be selected for a successful embryo transfer and super-ovulation (Tibary and Anouassi 1997). The super-ovulation is achieved through various doses of follicle-stimulating hormones (FSH), progesterone, equine chorionic gonadotropins (eCG) for a few days (Anouassi and Tibary 2013). The optimization of hormone dosage and administration time is necessary for better stimulation (McKinnon et al. 1994). The synchronization of the embryo stages in the recipient is achieved through stimulating hormones based on their follicular size. Even a few days difference in synchronization often lead to decreased pregnancy (Skidmore et al. 2002). Finally, the embryo transfer is achieved using either surgical or non-surgical methods. The non-surgical methods commonly include insertion of catheter and embryo flushing media for the careful recovery of the embryos (Anouassi and Tibary 2013). The morphological and biochemical investigations of the recovered embryos are

performed to categorize the transferable and non-transferable embryos. Healthy and suitable embryos can be cryopreserved for future use. The method is versatile and frequently used for a better reproductive rate. MOET can be used to identify or screen the genetic defects by cross mating the parental and super-ovulated progenies and the transmission of the disease is largely eliminated from parental generation.

3.2.6 *In Vitro Fertilisation (IVF)*

In vitro fertilization is a method for embryo culture and manipulation without in-vivo maternal signaling. The embryo propagation is vulnerable to environmental factors such as temperature, oxygen availability, light, nutrition, and embryo-maternal signaling. In-vitro fertilization is a technique to generate an embryo in a suitable synthetic environment or external environment through the fertilization of oocytes and sperm cells. The process of in vitro fertilization involves the recovery of the oocytes, maturation, fertilization, and propagation of the embryo. The oocytes are removed using the castration method from ovaries of the domestic animals followed by in vivo maturation (de Collado et al. 2017). The immature oocytes are usually aspirated in vivo from unstimulated or mild stimulated ovaries and cultured in the standard maturation medium (Rose and Bavister 1992). During aspiration of the follicles from the ovaries, a granular, multilayered, tight, and rough cumulus is identified as cumulus-oocyte complexes (Yang and Lu 1990). The propagation media allow resuming of oocyte first meiotic division. The luteinizing hormone (LH) and follicle-stimulating hormones (FSH) are commonly added with the maturation medium. The growth of the oocyte can be affected by any alteration in the standard maturation conditions. The qualitative losses can be assessed during maturation through blastula and morula yield estimation (Khosla et al. 2001; Luciano and Sirard 2017). During maturation, the culture conditions affect developmental competence and epigenetic growth of the embryo. Several cytoplasmic changes undergo before nuclear maturation during the maturation process (Hyttel et al. 1997). In vitro maturation process involves the messenger RNA and protein synthesis, mobilization, and storage which is vital for the early development of the embryo (Motlik and Fulka 1986; Luciano and Sirard 2017). After maturation of oocytes, the medium is changed to induce sperm capacitation and fertilization which can last up to one day. The ejaculated semen is preferably used for fertilization that gives rise to more eggs than epididymal spermatozoa. The fresh semen is incubated with heparin or any high ionic strength medium for maximum oocyte penetration (Pavlok et al. 1988). A variable concentration of heparin, depending on the species, was effective for better fertility (Leibfried-Rutledge et al. 1989). The serum albumin, epinephrine, heparin, caffeine, calcium, and hypo-taurinechemicals can be used for sperm capacitation (Parrish 2014; Polisca et al. 2013). The co-culture of the oocyte and sperm cells and the fertilization time strictly depend on the type of animal species and conditions used for the early embryo development and maturation. The fertilized embryo is propagated in the enriched culture medium such as bovine oviduct epithelial cells (BOEC) or supplemented with serum, growth regulators, proteins, and hormones (Fukui and Ono 1989; Li et al. 2018). The chemical composition

and fluid properties are adjusted to mimic the *in vivo* environment of embryo development. The methods, protocols, and guidelines for embryo evaluation, classification, transfer, and maintenance are adapted according to the International Embryo Technology Society (IETS) (Bo and Mapletoft 2018; Stringfellow and Seidel 1998). The morula or early blastocyst stages of the embryo are suitable for transferring into the recipient and later blastocyst stages can be considered with special hygiene and following guidelines of IETS for the safe and disease-free transfer into the recipient (Stringfellow and Seidel 1998). The IVF methods in cattle face difficulties due to oocyte inactivation after injection of the spermatozoon (Abdalla et al. 2009). The inactivation could be a result of the difference in morphological characteristics of the oocytes or physiological deficiencies. IVF has great potential in animal reproduction and recently the integration of IVF with advanced genomic methods can provide precise and comprehensive information of the desired characteristics in animals. IVF can be used in the modern genome editing methods such as CRISPR Cas9 and TALEN for efficient genome editing to achieve desired genetic selection or modification in the animals.

3.3 Marker Assisted Selection (MAS) for Livestock Production

With technological progress in the identification of different genetic markers, research is being done for the development of breeding techniques using genetic markers. This technology is used to identify the genomic segments that control the variation seen in the quantitative trait of a species and their application in breed selection (Dekkers and Van der Werf 2007). Information about the identified Quantitative Trait Locus (QTL) is used for screening traits at the DNA level using an associated marker in the newly emerged technique of Marker Assisted Breeding (MAB) or Marker Assisted Selection (MAS). It is an indirect method of trait selection based on genotype composition, where the presence of a marker linked to a desired trait in the polymorphic loci is used as the criteria for selection and not the trait itself. The main objective of MAS is to collect the marker and QTL-related information and to complement it together with the phenotypic information to improve the selection process (Wakchaure et al. 2015). Since quantitative traits are targeted for various breeding programs and classical selection methods depend on phenotypic characteristics, the development of genetic markers has helped make a step forward in the development of strain improvement programs (Moniruzzaman et al. 2014). Integrated use of the classical method with this new genetically based approach will facilitate the identification and selection process in the early developmental phase.

The MAS has its utility for those traits which are expressed, for example, in the later stages of life, depends on the sex, have a low level of inheritance, or are regulated by a few genes. The traits with expensive measurement procedures can also be selected using this technique. An increase in the sensitivity of the selection response from -0.7% to 64% for a single trait is estimated when MAS is used along with

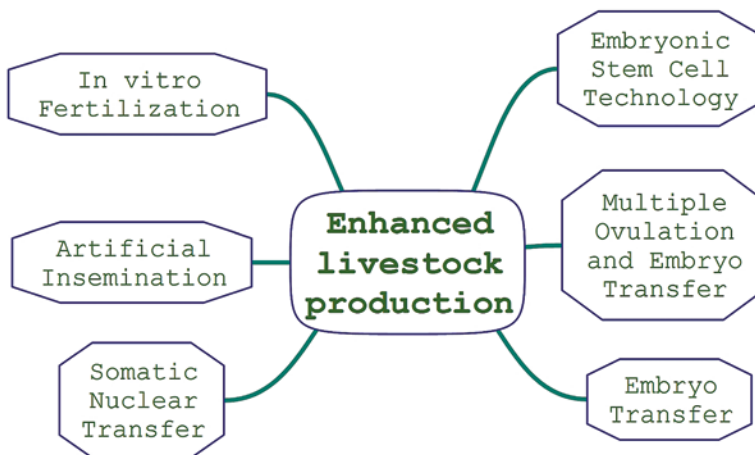


Fig. 3.1 Biotechnological strategies for enhanced livestock production

BLUP (a linear mixed model for predicting the random effect) than the BLUP alone. The trait studied using MAS is the amount and quality of carcass, longevity, milk production, and inheritance-related diseases (Ashraf et al. 2019) (Fig. 3.1).

3.3.1 Molecular Markers for MAS

Molecular markers are the point of reference in the genome. They do not necessarily play an important role but help identify specific DNA segments in the genome. Several genetic markers have been used to study the trait for their application in MAS, such as single nucleotide polymorphism (SNP), restriction fragment length polymorphisms (RFLPs), random amplified polymorphic DNA (RAPD), amplified fragment length polymorphisms (AFLPs), and simple sequence repeats (SSR) or microsatellites. Each of these markers has its limitations and disadvantages, however, the relationship between the marker and the target gene influences the success rate of MAS (Dekkers and Van der Werf 2007).

3.3.2 Properties of an Ideal Marker

The molecular markers are required to exhibit some unique characteristics such as codominance for easy identification of allelic nature of a diploid organism i.e. whether heterozygous or homozygous. Moreover, the molecular marker should be highly polymorphic, frequently appearing in the genomic compositions of the organism and no external factor should influence the selective nature of the marker. Other important features include high availability and duplicability, simple and rapid assays with ease of exchange of the data (Yadav et al. 2017).

3.3.3 *Marker Types*

Three types of markers are used based on the linkage between the marker and genes for the trait of economic value:

- (i) Direct marker: The marker can be positioned within the target gene sequence. This relationship is highly favorable, and the MAS technique for such genes is termed gene-assisted selection (GAS). However, the difficulty in the identification of such markers makes their use less frequent.
- (ii) LD- marker: The marker may be in close association with the gene of interest so that they are always inherited together showing a high value of linkage disequilibrium (LD).
- (iii) LE markers: The marker may be in linkage equilibrium (LE) with the gene of interest. The random combination of these genes will have the same value as that of their independent expression showing zero linkage disequilibrium values.

Based on the techniques utilized for their identification, the DNA molecular markers can further be classified into three types (Yadav et al. 2017):

- (i) Hybridization-based DNA Markers: Radio-labelled or enzyme-conjugated DNA segments or oligonucleotides are used in this method for DNA hybridization that helps easy detection of the marker molecules. Restriction Fragment Length Polymorphisms (RFLPs) and Oligonucleotide fingerprinting are the two common approaches that use this technology.
- (ii) PCR-based DNA Markers: In this technique, the presence of the DNA molecule is determined by its PCR amplification using specific primers. The Random Amplified Length Polymorphic DNAs (RAPDs), Simple Sequence Repeats, or microsatellites (SSRs) and Amplified Fragment Length Polymorphisms (AFLPs) are the examples of this technique.
- (iii) DNA Chip and Sequencing-based DNA Markers: Development of rapid genome sequencing methods like Next-generation sequencing (NGS) and third-generation sequencing (TGS) has led to the discovery of single nucleotide polymorphism (SNP) based markers. These markers have a high frequency of occurrence with ease of access and other desirable features for their use in MAS (Singh and Singh 2015).

3.3.4 *Designing the MAS Program*

The MAS has been applied in various livestock breeding programs, from improvement of the traits to the traceability of products. The prerequisite for design of a successful MAS breeding program is a strong and close linkage between the targeted gene and a marker gene. The success of the MAS program also depends on the progress made in other techniques of genetics, namely (i) gene mapping: it is

Table 3.2 The technologies adopted for genetic improvement in a few livestock

Sr. No.	Animal/ livestock	Methodologies adopted for genetic improvement	References
1.	Cow/ buffalo	Somatic cell nuclear transfer, Artificial insemination, embryo transfer, embryonic stem cell technology, IVF, MOET	Cibelli et al. (1998), Chebel et al. (2004), Drost et al. (1986), Gugjoo et al. (2019), Thakkar, and Wei et al. (2010)
2.	Sheep/ goat	Somatic cell nuclear transfer, Artificial insemination, embryo transfer, embryonic stem cell technology, IVF, MOET	Wilmot et al. (1985), Evans and Maxwell (1987), Cognie (1999), Notarianni et al. (1991), Bari et al. (2003), and Catalá et al. (2012)
3.	Pig	Somatic cell nuclear transfer, Artificial insemination, embryo transfer, embryonic stem cell technology, IVF, MOET	Gupta et al. (2013), Knox (2016), Brüßow et al. (2000), Romagnuolo et al. (2019), and Hicks et al. (2020)
4.	Chicken	Somatic cell nuclear transfer, Artificial insemination, embryo transfer, embryonic stem cell technology, IVF	Gupta et al. (2013), Burrows and Quinn (1939), Jones et al. (2006), Laval and Pain (2010), and Davidson (2004)
5.	Rabbit	Somatic cell nuclear transfer, Artificial insemination, embryo transfer, embryonic stem cell technology, IVF, MOET	Chesné et al. (2002), Morrell (1995), Song et al. (2017), Wang et al. (2007), Arias-Álvarez et al. (2017), and Garcia-Dominguez et al. (2020)

necessary for the identification and mapping of the target gene and the associated marker, (ii) marker genotyping: it is used to study the genetic differences in large numbers of individuals based on a large number of markers available. It must be made cost-effective for processing many samples for detecting QTL for the application of MAS. (iii) QTL detection: for the identification of the gene and the associated gene marker of the economically valuable traits. The degree of association must also be estimated. (iv) Genetic evaluation: for the statistical analysis of genotypic and phenotypic data to estimate the breeding value of individuals and (v) MAS for the efficient use of information at the DNA level to develop strategies for successful breeding programmes (Moniruzzaman et al. 2014).

The design of MAS includes the selection of the parent as a critical initial step. The selected parent breeds should have contrasting characteristics so that the genome of the individual parent and the heritability of its segment in the different recombination formed in the next generation can be easily identified. To achieve this, the germplasm of the parent breed is screened. Once the parents have been selected, a breeding population is developed by crossbreeding between the parents. Finally, the DNA from successive generation is used to select traits based on MAS (Table 3.2).

3.3.5 Application of MAS

The MAS finds its application in various breeding programs for selecting different traits.

- (i) Molecular marker-assisted introgression (MAI): Molecular markers can also be used as a guide to identify the introgressive genes expressing individuals. While traditional breeding techniques require a series of backcrosses to identify individuals with introgressive genes, the use of MAI reduces the workload, time, and cycles of backcrossing. Growth, meat or milk quality, etc. are the traits targeted for breeding programmes (Fertilization). The most common application of MAI is its use in the introgression of disease resistance genes (Prajapati et al. 2017). The MAS has been studied for its accuracy in the identification of the Bacterial cold-water disease (BCWD) resistance-related genes in rainbow trout (*Oncorhynchus mykiss*). SNP-based marker was applied for the study and it was observed that MAS provided accuracy of about 0.5 which is higher than the classical approach (Liu et al. 2018).
- (ii) Progeny testing and multiple ovulation and embryo transfer (MOET) programs: The different molecular markers are linked to the gene of interest for QTL in the livestock industry and used as GAS or LD-MAS for trait pre-selection and breeding. For the accuracy of selection, progeny testing of the traits before widespread use is required for assuring accuracy of selection. However, the high interval between the generations reduces the genetic gain. The integration of MAS reduces the interval but maintains selection accuracy to increase the genetic gain. A significant gain of 65% in the traits of the carcass has been reported. Similarly, the response of milk production increased to 15% compared to the previous report of 6% when the MAS and MOET were used together (Wani et al. 2008).
- (iii) Product traceability: To ensure the quality of the various animal-based products and their food safety, molecular markers have been used as a trustworthy tool. The molecular markers have been used to identify the genotype and parentage of the various animal products other than their use in the selection process (Moniruzzaman et al. 2014).
- (iv) Assessment of undesirable genes: Like the selection of desirable traits, the MAS can be used for screening of the presence of undesirable genes, physical or genetic defects in breeding livestock. The MAS help identify the genes responsible for the various defects in the individual and their removal from breeding programmes (Fertilization).
- (v) Fertility study: The MAS can also be used for fertility studies in different breeding programs whereby the breeding soundness of the cattle involved can be evaluated. Since a large number of insemination dosages i.e. approximately three lakh doses are formulated from a single bull, it is essential to assure its fertility for successful conception and economic realization of artificial insemination (Raina et al. 2020). Conventionally the fertility of the bull is determined through phenotypic analysis which includes the study of sperm cell concentration, morphology, motility along with the absence of any physical or genital disease in the bull and Sire Conception Rate (SCR) (Morrell et al. 2017). However, all these parameters could not give a clear representation with reports indicating a continuous decline in the fertility rate of the bulls. Since the fertility trait of the bull has high variability and low heritability, the pheno-

typic selection is not competent enough for the identification of bulls with high fertility. Identification of molecular markers linked with the fertility traits like the SCR and quality of semen have been carried out for their application in MAS for differentiation and selection of bulls with high fertility for the process of artificial insemination (Raina et al. 2020).

3.3.6 Advantages and Limitation of MAS

The MAS has a major advantage of faster selection in the early stages of life cycle, as the phenotype can be calculated based on the genotype information. This helps reduce the generation gap for selection and expression of the desired trait. Since the genotype is considered, the traits that need to be transferred to the successive generation can be identified without taking environmental influences into account. Moreover, when the phenotypic selection is difficult or unfeasible to examine for the different features such as disease resistance, protein content of the product to be derived, the MAS has the lead. The traits, with low heritability, are sex-linked (milk or egg production) or can only be estimated after the slaughter of the animals (meat quality), can also be selected using MAS giving an advantage over the conventional methods (Wakchaure et al. 2015).

The major setback of MAS is the high cost of genotyping. Sample collection and gathering of genotypic information from the entire population is unfeasible for developing cattle breeding program. Missing markers or incomplete information on the genotype, lack of consistency of QTL data over some time, or varied population does not make the QTL a reliable source of information. Furthermore, the validity of the markers used must be continuously confirmed for each breeding population. Although the molecular marker may not be affected by environmental factors, its associated trait may get influenced and show different phenotypes. These factors must also be considered, and the marker analysis needs to be done in a varied environment making the process cumbersome. Finally, the availability of experts with the necessary technical knowledge is another obstacle to the application of MAS in developing countries (Moniruzzaman et al. 2014) (Table 3.3).

3.4 Conclusion

The current scenario of the world population and climate change indicate a higher food demand in the next two decades. Livestock production and consumption in the world has increased over time. Livestock will be one of the major sources of food for developing countries. The genetic improvements of this livestock through the latest biotechnological interventions and molecular marker-assisted selection of the beneficial traits and transfer into the recipient line can enhance the food quantity while maintaining the quality in a relatively shorter time. The latest technologies can provide increased accuracy and intensity of the improved trait selection for the

Table 3.3 Industrial application and advantage of MAS

Sr. No.	MAS applications	Advantages	References
1	Dairy cattle breeding schemes	Increased milk production	Meuwissen and Van Arendonk (1992)
2	Closed nucleus breeding schemes	Gain in carcasses	Goddard (1996)
3	Assessment of genetic defect	Removal of carriers of genes related to bovine leukocyte adhesion deficiency	Shuster et al. (1992) and Dekkers (2004)
4	Introgression of genes under NARI-Suwarna scheme	Introgression of the <i>FecB</i> mutation in sheeps for increased ovulation rate	Nimbkar
5	QTL mapping and introgression of genes	Increased meat quality of Qinchuan cattle breed	Abd El-Hack et al. (2018)
6	SNP marker assisted BLUP	Increase breeding value of Turbot	Lyu et al. (2019)
7	Marker linked sex identification	Selective production of faster growing (XX) females for mass production and (YY) super males for basic research in the Southern catfish (<i>Silurus meridionalis</i>)	Zheng et al. (2020)

livestock. The inception of the embryonic stem cell technology for livestock breeding has opened new avenues of the reprogramming genetic material for the in-vitro selection of the desired characteristics. The livestock sector requires aggressive research for the successful implementation of the existing technologies.

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Chapter 4

Applications of Stem cells Technology in Livestock Production



Vinay Bhaskar, Satish Kumar, and Dhruva Malakar

Abstract Urbanized population and a gigantic demand for livestock products are expected to rise dramatically in the coming decades. Initiatives have been geared up to intensify the livestock production. Decipher of large animal genomic sequence boost the genetic enhancement. Embryonic stem cells are efficient platforms for genome editing to produce transgenic animals. Moreover, stem cells opens a new avenue to insight into developmental processes. Stem cells increase the production efficiency by maintaining the health and welfare of the livestock. Conservation of elite species also adds advantage to use reprogrammed induced pluripotent stem cells in farm animals. However, uses of embryonic stem cells are bound with ethical concerns adult stem cells promises alternative approach. Based on the ease of availability and isolation, mesenchymal stem cells are extensively studied and widely used stem cells for therapeutic purposes. Multipotency, immunomodulation, homing properties of mesenchymal stem cells plays a vital role to alleviate diseases and increase the production. Stem cell technology used in large animals can also serve as a model for various human diseases. Hence conventional approaches to enhance the productivity and welfare of animals were reformed by the advent of stem cell technology.

Keywords Stem cells · Embryonic stem cells · Mesenchymal stem cells · Livestock production · Therapeutic applications

Abbreviations

ART	Assisted reproductive technology
BMP	Bone morphogenic protein
CD	Cluster of differentiation
CRISPR	Cluster of regularly interspaced short palindromic repeats
CXCR	Chemokine receptor type

V. Bhaskar · S. Kumar · D. Malakar (✉)
Animal Biotechnology Centre, National Dairy Research Institute, Karnal, Haryana, India

EC	Embryonic carcinoma cells
ES	Embryonic stem cells
FAO	Food and agriculture organization
GCTM	Germ cell tumor marker
HGF	Hepatocyte growth factor
HSC	Hematopoietic stem cells
IBD	Inflammatory bowel disease
IL	Interleukin
INF- γ	Interferon gamma
iPSC	Induced pluripotent stem cells
ISCT	International society for cellular therapy
LIF	Leukemia inhibitory factor
MHC	Major histo compatibility complex
MSC	Mesenchymal stem cells
NO	Nitric oxide
OCT 4	Octamer binding transcription factor 4
PGE2	Prostaglandin E2
QTL	Quantitative trait locus
SDF	Stromal derived factor
SSEA	Stage specific embryonic antigen
TALEN	Transcription activator like effector nuclease
TGDF	Teratocarcinoma derived growth factor
TGF	Transforming growth factor
TNF	Tumor necrosis factor
TRA-1	T cell receptor alpha locus
VCAM	Vascular cell adhesion molecule
ZFN	Zinc finger nuclease

4.1 Introduction

In this new millennium where we are exploring biological solutions for biological problems, We are on the brink of a paradigm change, emerging from basic mechanical care services to considering biological approaches for health promotion, evaluation of risk assessment, treatment, and even prognosis. Slow but steady attempts have made to understand regenerative capacities and underlying mechanisms. The stem cells set to transform the entire health care delivery. Past decade witnessed a significant flux of stem cells technology in human and veterinary research. The properties of stem cells fascinates the researchers to explore its wide range of applications. Stem cells are unspecialized cells and was first reported in bone marrow as self renewing cells in the year 1963. Stem cells have clonogenic capacity which can

give rise to same kind of genetically identical cells. Self renewing capacity (Asal and Güven 2020) and the ability to differentiate into wide variety of cell types even to germ cell type (Malik et al. 2020; Ota 2008) were some of the remarkable properties of stem cells. Embryonic stem cells (ESC), Adult stem cells, Induced pluripotent stem cells (iPSC) are some of the extensively explored stem cell types. These cells are primarily undifferentiated cells found in the body throughout the lifespan starting from early embryonic development. Embryonic stem cells first isolated from mouse blastocyst in 1981 then recently in 2006 induced pluripotent stem cells were produced for the first time by reprogramming skin cells to early pluripotent state. After their first discovery they are reported in many livestock species (Table 4.1). Based on their differentiation ability stem cells are categorized into totipotent- Stem Cells from early Embryo (1–3 days) are totipotent as each cell have the ability to develop into whole individual. Pluripotent- Some cells of blastocyst (5–9 days) are pluripotent they can give rise to all adult type of cells into body not placenta. Multipotent-Adult stem cells are multipotent which have capacity to differentiate into many tissues like hematopoietic cells. Oligopotent- Stem cells which have capacity to differentiate only few tissues like lymphoid or myeloid stem cells. Unipotent-Stem cells are unipotent which can give rise to only same kind of cells like skin cells are a good example of unipotent stem cells.

Table 4.1 Stem cells source, types and markers in livestock

Species	Source	Type	Marker	References
Cattle	Adipose, endometrium, bone marrow, mammary tissue, Wharton's jelly	MSC, ES, iPSC	Oct-4, Sox-2, Nanog, TRA-1-81, TRA-1-60, CD73, CD 90, CD 105	Hill et al. (2019)
Buffalo	Mammary tissue, adipose, placenta, endometrium, Wharton's jelly	MSC, ES, iPSC	CD73, CD 90, CD 105, Oct-4, Sox-2, Nanog	Gugjoo et al. (2019a, b)
Sheep	Adipose, umbilical cord Wharton's jelly	ES, iPSC	Oct-4, Sox-2, Nanog	Gugjoo and Amarपाल (2018)
Goat	Adipose, umbilical cord	ES, iPSC	Oct-4, Sox-2, Nanog	Gugjoo et al. (2020)
Pig	Bone marrow, adipose, umbilical, endometrium, Wharton's jelly	MSC, iPSC	Oct-4, Sox-2, Nanog, CD73, CD 90, CD 105	Bodek et al. (2015), Xu et al. (2019), and Vassiliev et al. (2010)
Horse	Bone marrow, adipose, umbilical cord, endometrium, Wharton's jelly	MSC, ES	Oct-4, Sox-2, Nanog, TRA-1-81, TRA-1-60, CD73, CD 90, CD 105	Burk et al. (2013) and MacDonald and Barrett (2020)
Dog	Adipose tissue, umbilical cord.	MSC, iPSC	Oct-4, Sox-2, Nanog, CD 73, CD 90, CD105	Kriston-Pál et al. (2017) and Baird et al. (2015)

4.2 Stem cell Types and Their Identification

Embryonic stem cells are pluripotent stem cells derived from ICM of blastocyst. They have ability to differentiate into all three germ layers. ESC derived from Humans, murines and primates shows peculiar cell cycle pattern they spends most of the time in S phase as it lacks G1 check point (White and Dalton 2005; Bárta et al. 2010). Different methods were adopted to isolate the embryonic stem cells like immunosurgery, mechanical dissection, Laser dissection, enzymatic method and mostly cultured on mitotically inactivated fibroblast feeder layers (Yu and Thomson 2014). Intrinsic and extrinsic pathways influence the self-renewing and pluripotency of stem cells. Both self-renewing and pluripotency processes are coordinated processes altered by different signaling pathways like LIF/stat3, wnt/b-catenin, FGF/ERK, TGF/SMAD and PKC signaling (Huang et al. 2015). Induced pluripotent stem cells are another type stem cells which have pluripotency properties. Due to ethical concerns of embryonic stem cells iPSC gain its popularity because of its technical simplicity (Liang and Zhang 2013). iPSCs are transcription factors mediated reprogrammed cells. Yamanaka described Oct4, Sox2, Nanog and c-myc are key transcription factors which helps in reprogramming of cells to pluripotent state (Takahashi and Yamanaka 2006). Oct4, Sox2, Nanog are important factors that maintain pluripotency in both types. Oct-4 is the master regulator of maintaining pluripotency Embryo restricts the development of ICM in the absence of Oct 4. Most of the cases pluripotent identity is maintained by Oct4 in associated with Sox2 form heterodimer and regulate Nanog expression (Wu and Schöler 2014).

Adult stem cells were found in bone marrow, dental pulp, retina, pancreas, spinal cord, cornea, heart, liver, blood etc. almost in all the body parts. They are clonogenic and maintain homeostasis (Nakada et al. 2011). They promote healing by replacing the deceased cells (Kanji and Das 2017). Adult stem cell differentiate into specialized cell type suitable to microenvironment (Tang et al. 2006). As per the standards set by ISCT cells positive for CD90, CD73, CD105 and negative for CD45, CD 34, CD79 are considered as mesenchymal stem cells (Dominici et al. 2006). Hematopoietic stem cells are positive for CD34 and lack in marker in mesenchymal stem cells. Expression of markers varies with species for example CD73 cell surface marker in dog showed negative while it is an established marker in humans. Stem cell surface markers in canines shows positive for CD105, CD90, CD44, STRO-1 (Amanda B.T. Hill et al. 2018). In equine CD105, CD90, CD44 were shown positive but umbilical derived mesenchymal stem cells expresses low levels of CD-90 (Barberini et al. 2014). Identified markers of stem cells so far were detailed in Table 4.2. Bovine mesenchymal stem cells shows positive for CD29, CD166, CD106, CD73, CD44, CD90 (Sampaio et al. 2015; Gao et al. 2014) (Fig. 4.1). Stem cells have great potential in terms of cells for therapeutic applications under different conditions, including metabolic, degenerative and inflammatory diseases, restoring and regenerating damaged or destroyed tissues, as well as a cancer treatment. Immunomodulatory, antimicrobial activity and migration of adult stem cells are discussed further in this chapter.

Table 4.2 Reported markers for stem cells

Markers	Type of stem cells	References
TRA-1-81	EC, ES	Andrews et al. (2005)
TRA-1-60	EC, ES	Zhao et al. (2012)
Telomerase	ES, EC	Hiyama and Hiyama (2007)
c-Kit	MSC,HSC,ES,EC	Lennartsson and Rönstrand (2012)
SSEA-4	EC, ES	Zhao et al. (2012)
SSEA-3	EC, ES	Andrews et al. (2005)
OCT-4	EC, ES	Pan et al. (2002)
Germ cell nuclear factor	EC, ES	H. Wang et al. (2016)
GCTM-2	EC, ES	Andrews et al. (2005)
TDGF	ES	Baldassarre et al. (1997)
CD133	HSC, neural stem cells	Zhao et al. (2012)
CD30	EC, ES	Zhao et al. (2012)
Alkaline phosphatase	EC, ES	Štefková et al. (2015)
Neurosphere	ES	Campos (2004)
Stro-1	MSC, HSC	Lin et al. (2011)
Lin	MSC, HSC	Seita and Weissman (2010)
Colony forming unit	MSC, HSC	Kaufman et al. (2001)
CD 44	MSC	Jiang et al. (2019)
BMPR	MSC	Mira et al. (2010)
CD 73	MSC	Maleki et al. (2014)
CD 34	MSC, HSC	AbuSamra et al. (2017))
Thy-1	HSC, MSC	Saalbach and Anderegg (2019)

4.3 Application of Stem cells in Livestock

Conventional livestock production should be optimized to reach the demand of daily needs of fast growing population. According to FAO, 60% of animal and crop output has to be increased to meet the requirement. The requirement for meat and milk will relatively increased by 70% by 2050 (Herrero et al. 2015). Advancement in genomic technologies during early twenty-first century led to the identification of hundreds of QTLs. This discovery opens a wide opportunity to improve the genomic selection in livestock across the globe. Advancements in Stem cell technology integrated with genomic selection offers a wide range of applications in livestock production.

Stem cells especially embryonic stem cell type increases the efficiency of cloning somatic cell nuclear transfer. It was reported that embryonic stem cells has 10–20 times higher cloning efficiency in comparison with cumulus cells (Singh 2019; van Thuan et al. 2010). Chimeric and transgenic animals can be produced, Desired trait carrying transgenic animal production is ramp by using embryonic stem cells as it was reported that higher genetic modifications is possible in ES than somatic cells. Homologous recombinations were higher in case of ES in comparison with somatic cells (Choi et al. 2020). Manipulating the live stock species

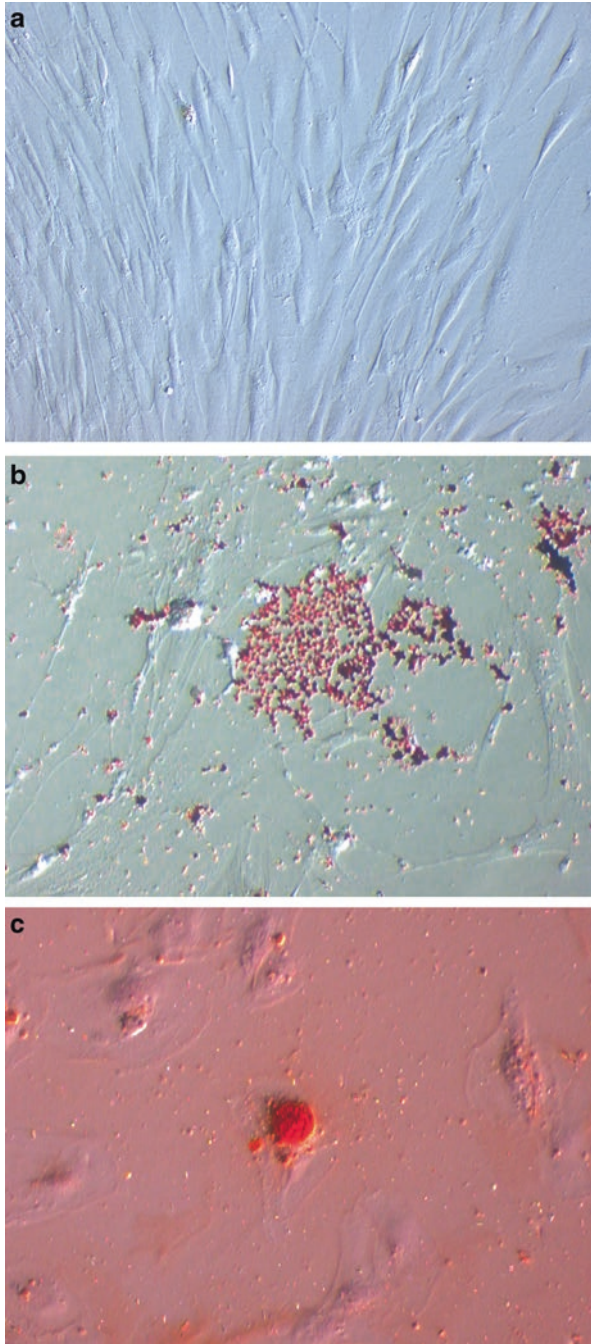


Fig. 4.1 Buffalo stem cells and their differentiation into bilineage (a) Fully confluent Buffalo adipose derived mesenchymal stem cells (b) Adipogenic differentiated stained with Oil red O staining. (c) Osteogenic differentiated stained with alizarin red S

genome could efficiently increase the animal production with desirable traits. Enhancing welfare of animal, increasing food production are the main objectives of gene editing. They also used in resolving the environmental issues. Embryonic stem cells were used as donor cells for gene editing further used in cloning. ZFN, TALEN, CRISPR Genome editing technologies enable production of disease resistance and high yielding milk livestock (Menchaca et al. 2020). Economically developed countries dairy industry facing huge economic loss due to mastitis. These cutting edge technologies can produce the mastitis resilient trait. Introgression of lysostaphin gene in cow (Wall et al. 2005) and lysozyme in goat (Maga et al. 2006) are reported to have resilient to mastitis. Likewise influenza resistant pig, PRRSV (porcine reproductive and respiratory syndrome virus) resilient porcine trait (Lu et al. 2017) and environmental friendly porcine trait which digest the phytate (Golovan et al. 2001) were produced using these technologies. Over the past few years around 300 edited cattle, sheep, pigs goats were produced (Tan et al. 2016).

ES also able to generate germ cells which has its application in alleviation of infertility in livestock. Although ARTs reduced the effectiveness of this problem, ethical issues, invasive, expertise handling and costly treatment hinder the progress of solving the issue. Stem cells offer a new approach of treatment. Oocyte generation, ovarian regeneration and dysfunctional sperm were possible and treated using stem cells. so that elite gene pool can be conserved (Wang et al. 2019). Moreover ES opens a new way for drug development. A vibrant framework ideal for defining innovative molecular targets and improving specific drugs is defined by stem cells. Pathological context of disease, testing of drugs and assess the effectiveness of drugs and its safety can be evaluated using stem cells. Desirable cells were obtained by iPSC then effect of drugs and chemicals were evaluated (Rubin and Haston 2011).

Due to its complexity developmental biology is less understood. Embryonic stem cells and induced pluripotent stem cells are versatile biological cells that open insight into developmental process. Embryonic stem cells have the capacity to differentiate into almost any kind of cells. The pathways involved in embryonic development, fate of cells and regulation of transcription factors can be revealed by exploring the stem cells (Zhu and Huangfu 2013). The list of endangered species increasing alarmingly. Conservation strategies are not sufficient to protect them from becoming extinct. A recent potential technology like iPSC attracts scientists to revive the lost ecosystem. Extinct animals can also be resurrected by using this technology. Still lots to explore its application in conserving the endangered species recent advances are encouraging (Stanton et al. 2019).

Ethical limitations of embryonic stem cells are overcome by adult stem cells. Mesenchymal stem cells are extensively studied stem cells. As they are easy to isolate and expand the popularity gained after the recognition of immune modulation, anti proliferative, migration abilities. Clinical use of stem cells in veterinary medicine was well reviewed by Voga et al. (2020).

4.4 Mesenchymal Stem Cells

MSCs are adult stem cells which have the ability to differentiate into multiple lineages. MSCs considered immunomodulatory since they lack type II MHC antigen and thus do not cause immune reactions. MSCs have recently proven to regulate endogenous tissue and immune cells, initially open the way as a regenerative therapy to reconstruct skeletal tissue. Preclinical action mechanism studies indicate that the potential effects of MSC transplantation are short-lived, and linked to active, paracrine interactions between MSCs and host cells. The shreds of evidence indicate that MSC's primary mechanism of action based on paracrine signalling that results in functional changes in the immune cells, such as natural killer cells, B-cells, T-cells, dendritic cells, and monocytes/macrophages. There have been studies of many factors contributing to MSC's immunomodulatory effects such as transforming growth factor β , prostaglandin E2 (PGE2), interleukin 10 (IL 10) indolamine 2,3 dioxygenase (IDO) and stimulated gene-6 (TSG-6) tumour necrosis factor- (TNF) (Voga et al. 2020). Moreover, MSCs secretes a variety of soluble factors and Antimicrobial peptides like cathelicidin, lipocalin, β -Defensins,hepcidin etc. (Marrazzo et al. 2019).

4.4.1 MSC Immune Modulation and Antimicrobial Properties

MSCs have anti-inflammatory properties and reduce scar formation. MSCs strongly inhibit T lymphocytes and b lymphocytes, by activating programmed death pathway and proliferation and subsequent cytokine signaling (Augello et al. 2005). MSCs suppress excess production of autoantibodies. However, B cell-MSc interactions are not well known. MSCs interact with other immune cells and inhibit B-cells, and at the same time, it induces bregs (Fan et al. 2016). MSCs interacts with B-cells and inhibit the proliferation of B-cells and other co-stimulatory factors (Rosado et al. 2014). MSCs do not produce any immunological reaction as it lacks MHC 1 (Krampera et al. 2003), MHC 2 (Nauta and Fibbe 2007), CD80 and CD86 molecules on its surface (Le Blanc et al. 2003). The proliferation of NK cells is associated with IL-2. MSCs inhibit IL-2 and NK cells. The activated NK receptors NKp30, NKG2D is responsible for the lysis of MSCs (Spaggiari et al. 2006). VCAM1 and galectin-1 (Ren et al. 2010; Gieseke et al. 2010) are also involved in MSC-cell mediated immunoregulation. Eventually, MSCs strongly upregulates the expression of galectin-9 upon activation INF- γ and tend to be an important proliferative mediator, thus act as an immunoregulatory marker (Ungerer et al. 2014). CD8+ induce apoptosis in MSCs, which offers clinical benefits as they completely lack the immune rejection. This shows the clinical efficiency of MSCs. Apoptosis is important for MSC's anti-inflammatory and regenerative events. In agreement, apoptosis stimulated before treatment of MSC increased its effectiveness (Galleu et al. 2017).

MSCs also secrete some soluble factors which are necessary for immune modulation. The key mediators of MSC's immunosuppressive activity in the presence of pro-inflammatory cytokines are, PGE2, cyclooxygenase 2 (COX-2) and indoleamine-pyrrole 2,3-dioxygenase (IDO) (Krampera et al. 2003). PGE2 has been associated directly in macrophage production of IL-10 (Tam et al. 2009) and in preventing the monocytes differentiation into DCs. (Spaggiari et al. 2009). IFN- γ regulates the immunomodulation of MSCs and TNF- α and produces PGE2 (Ren et al. 2008), IL-6 (Ghannam et al. 2010), TGF- β 1 (Nemeth et al. 2010), NO, LIF and HGF (Najar et al. 2010) are other crucial MSC immune regulatory molecules found till date. These factors interact with both innate and adaptive immunity and initiate the immune suppression. T- cells and NK cells suppression is mediated by IDO. Allograft kidney acceptance is mediated by the MSC secreted IDO via treg generation (Ge et al. 2010). IL-10 is secreted by interaction of MSCs with the Antigen presenting cells thus regulate the immune suppression irrespective of allogenic or naïve or pre activated T- cells (Yang et al. 2009). Critical role of nitric oxide secreted by mesenchymal stem cells is reported by Sato et al. 2007. Proliferation of T- cells is repressed by the phosphorylation of stat 5 in the presence of NO and MSC. Inhibition of Nitric oxide synthase of MSC tends to the proliferation of T-cells (Fig. 4.2).

The use of mesenchymal stem cells (MSC) in the treatment of bacterial infections has gained growing attention in recent years. MSCs have direct bactericidal activity. Thus, MSC administration could replace traditional antibiotic therapy (Kadam et al. 2019; Marrazzo et al. 2019) MSC also produces antimicrobial peptides (AMPs), usually found in neutrophils or epithelial cells. AMPs directly destroy bacteria by destroying the integrity of the microbial membrane (Krasnodembskaya et al. 2010) or by triggering the production of pro-inflammatory cytokines. Human MSC produces multiple AMPs, including cathelicidin peptide LL-37 (Krasnodembskaya et al. 2010), hepcidin (Alcayaga-Miranda et al. 2017), β -defensin 2, and lipocalin 2 (Gupta et al. 2012). MSC-produced AMPs which considered to be one of the critical components in the removal of bacterial infections, as investigated in multiple animal models (Johnson et al. 2017).

4.4.2 MSC Migration and Homing

MSC Homing is of two types based on the administration route (1) Systemic homing (2) Non-systemic homing. Systemic homing happened when MSCs administered intravenously or intraarterial. MSC enters the bloodstream and undergoes a multistep process to exit and blood stream and migrates to the targeted region. Non-systemic homing MSCs injected at target tissue MSCs travels to the damaged tissue. Some of the crucial stages of homing are the migration of MSCs in the bloodstream and trans endothelial immigration. MSCs thought to use the same mechanism as the movement of leukocytes to pass into the bloodstream. Adhesion molecules are essential for the trafficking of mesenchymal stem cells (Butcher and Picker 1996). These adhesion molecules include integrins, chemokine receptors, and selectins.

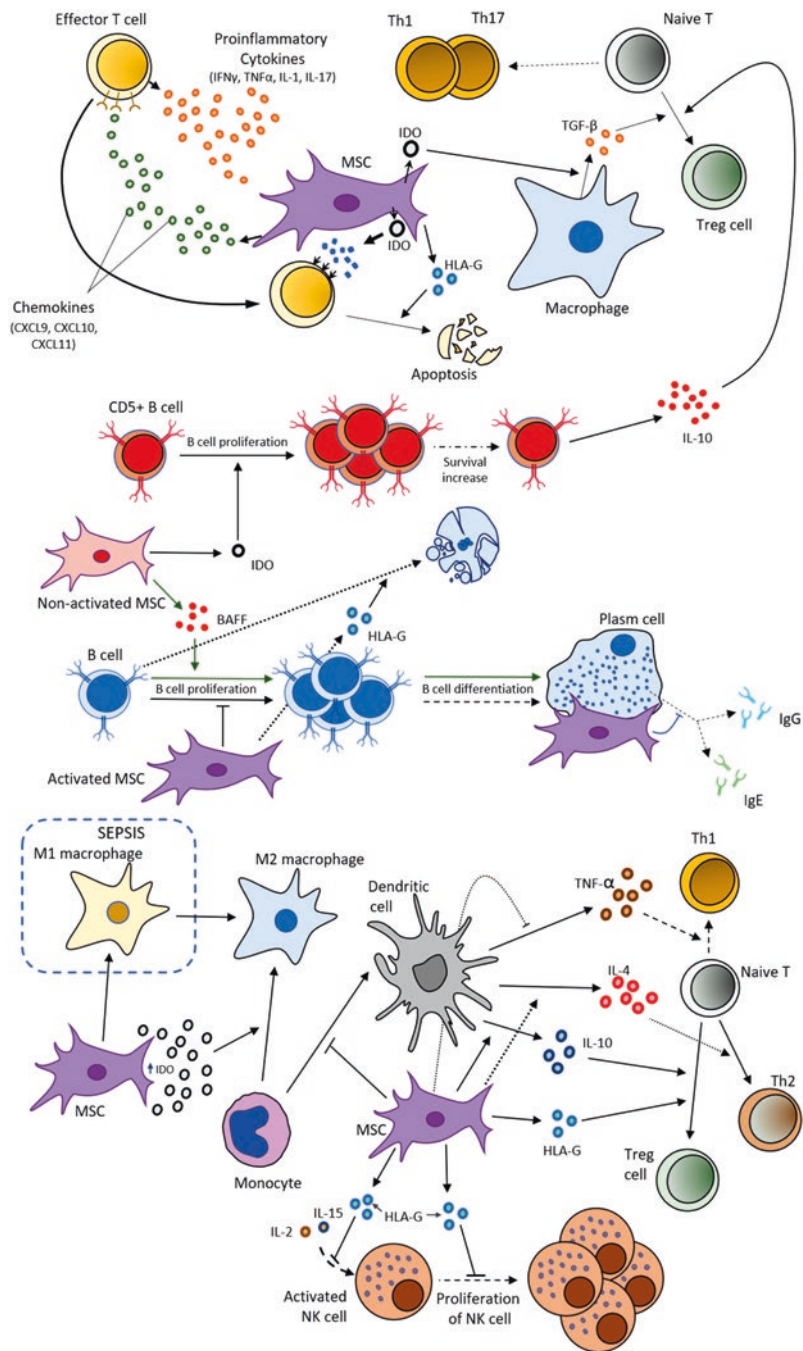


Fig. 4.2 Overview of mesenchymal stem cells immunomodulation. MSC interaction with adaptive and innate immune cells by releasing various soluble factors. (This figure is adapted from Jimenez-Puerta et al. 2020)

Primarily MSCs associate with endothelium by gathering and rolling. In the second, G-protein - coupled receptors activate cells, accompanied in the third step by integrin-mediated, activation-dependent arrest. Eventually, the cells transmigrate in the 4th phase via the endothelium (Ullah et al. 2019). MSCs will move to inflamed tissues reacting to chemokine and chemokine receptor signals induced according to inflammatory conditions (Kia et al. 2011). MSCs express a wide variety of receptors of chemokine and chemokine which helps in MSC homing. Some studies have shown chemokine receptor expression; namely, CCR1, CCR2, CCR4, CCR6, CCR7, CCR8, CCR9, CCR10, CXCR1, CXCR2, CXCR3, CXCR4, CXCR5, CXCR6 are present on MSC (Ghaffari-Nazari 2018). Reports have shown that the SDF-1/CXCR4 fulfil the essential function of MSC migration to the bone marrow. CXCR4 takes dose-dependent responses to SDF-1 (Kitaori et al. 2009) (Fig. 4.3).

4.5 Transplantation Modes and Therapeutic Applications of Stem Cells

Based on source of stem cells for transplantation possible treatment methods are (1) Autogenic transplantation, (2) Allogenic transplantation and (3) xenogenic transplantation. Autogenic transplant refers to the stem cells that are isolated from the animal's own tissues like adipose tissue, cord blood etc. In allogenic transplantation stem cells were isolated from another donor of same species. Stem cells were isolated from different species for transplantation in xenogenic transplantation which involves the risk of rejection. Stem cells are introduced into animal's body intravenously or locally injected based on the condition and severity of disease. Stem cells are reported to be more tolerant to oxidative stress as animals undergo stress in order to increase the productivity.

4.5.1 Bone, Cartilage and Ligament Injuries

Equines and canines are more susceptible to different injuries. Stem cells have the ability to differentiate into osteocytes and myocytes. Reports increasing on the reparative effect of mesenchymal stem cells in different species and recent studies quoted the effectiveness of MSC treatment in tendon, cartilage and bone injuries (Renzi et al. 2013; Canapp et al. 2016; Gibson et al. 2017). Adequate supply of blood and absence of nerve and lymph to cartilage bounded with reparative limitation favorable results were reported for cartilage repair by stem cells in rabbit and equine (Agung et al. 2006). Although bones have regeneration capacity, deformity caused by the devastating events can't be compensated in some critical cases. Reproductive inefficiency, loss of meat and loss in production of milk were some results of bone fracture in livestock which increases the proportion to lose the elite

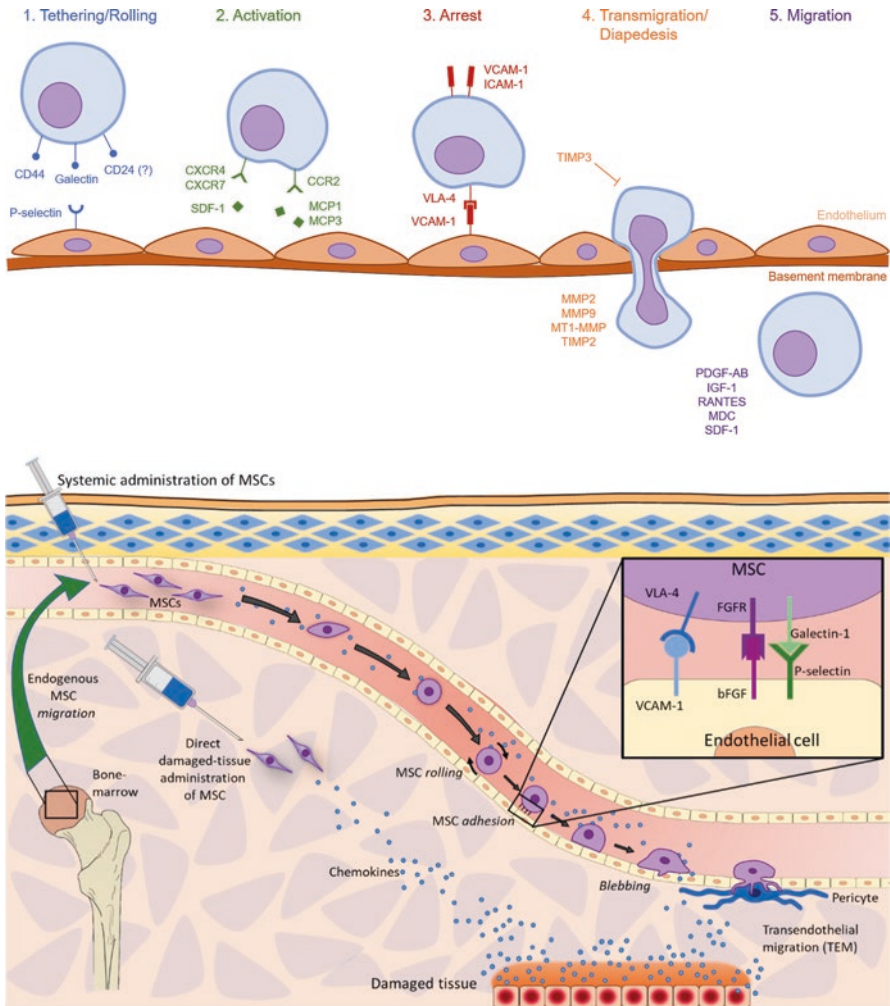


Fig. 4.3 (a) Step wise mechanism of homing of Mesenchymal stem cells. (This figure is reproduced with the permission of Ullah et al. 2019) (b) On administration migration of MSC to site of infection. (This figure is adapted from Jimenez-Puerta et al. 2020)

species (St. Jean and Anderson 2014). Changed trends in the society to conserve the economically productive animals reformed the treatment methods. Stem cells enhance the regeneration of bone (Iaquinta et al. 2019). Stem cells not only recuperate the animal’s structure but also stabilize its functionality (Moritz and Ambrosio 2017).

4.5.2 Degenerative Diseases

Stem cells can differentiate into osteocytes and chondrocytes and possess immune modulatory effect to treat certain degenerative diseases. Degenerative diseases like Osteoarthritis effects the quality of semen (D. F. Wolfe 2018). Promising results of stem cell injection to osteoarthritis condition significantly improves the local cartilage and synovial fluid viscosity in equines (Broeckx et al. 2019; Brehm et al. 2014 & Frisbie et al. 2009). Reports in canines showed infusion of stem cells in cartilage regeneration also infer a reliable approach (Shah et al. 2018; Sasaki et al. 2019). Most common progressive neurodegenerative disease in dogs is degenerative myelopathy (DM). Recent Success in mice model promises to use stem cells in canine suffering with DM (Hoffman and Dow 2016).

4.5.3 Inflammatory Diseases

Inflammatory diseases like Mastitis, laminitis, IBD, metritis, dermatitis in bovine, ovine, canine and equine species cause huge economical loss and has detrimental effect on animals health. Mastitis cause huge loss to the dairy industry. Resistance of bacteria to antibiotics makes its treatment difficult. Stem cells offers an alternative therapy for mastitis becoming popular now a days. Recent reports quote the safety of mesenchymal stem cells for the treatment of mastitis. Intra mammary infusion of mesenchymal stem cells decreases the bacterial population in the milk (Peralta et al. 2020). Rise of inflammatory bowel disorder cases over past decade and failure of present therapies to treat this condition attracts stem cell in order to maintain homeostasis in the gut. IBD is a gastrointestinal inflammatory disease and its etiology is unknown till date. Sub-mucosal and intravenous infusion of stem cells showed the significant improvement in reconstruction of disruptive mucosal tissue after the infusion of stem cells (Shi et al. 2019). Lamella tissue inflammation cause laminitis frequently found in cattle and horse. Use of MSC shown promisable treatment for laminitis in equine and cattle (Gugjoo et al. 2019a, b) Metritis is a reproductive disorder characterized by inflammation of endometrial tissue followed by bacterial infection. Antibiotics are the most common method to treat these diseases. However positive selection of bacteria with antibiotic treatment making microbes drug resistance and yielding huge loss to bovine and equine industry (Lara et al. 2018). Endometrial stem cells holds promise able treatment for this condition. Mambelli et al. (2013) observed the homing of stem cells to injured endometrial tissue. Falomo et al. (2015) observed the increase in the expression of IL10 and decrease in the expression of TNFA indicating the down regulation of chronic inflammation of disease in equine. Canine atopic dermatitis is the most common multifunctional disorder occurs in dogs in which less antimicrobial peptides were produced exposing animal to bacteria. Immunomodulatory and antimicrobial activities of MSCs holds promising treatment for this disease (Dias et al. 2019).

4.5.4 *Animal Models for Human Diseases*

To reduce the gap between clinical practice and translational research animal models are required to alleviate the human diseases. As small animal models failed to reproduce the results for human use large animal models can be alternative and more appropriate models to explore the use of stem cell technology for human welfare. Pigs, Dogs, cattle, goat, rabbit, and buffalo can be alternative sources to explore the stem cells full spectrum of action. Less data availability on safety and efficacy of the stem cells has to be addressed by proving the stem cells ability in large animal models (Harding et al. 2013). The mystery riddled around development process in human can also be untangled with the anatomical similar large animal models. Xenotransplantation of organs from swine to human, transgenic pigs to understand the pathophysiology of cystic fibrosis are some of examples of the stem cell large animal models for the welfare of mankind (Wolfe 2009; Klymiuk et al. 2010). Human Xenogenic implantation of MSC showed decrease in pulmonary edema and enhance oxygenation in acute respiratory distress syndrome model of ovine (Asmussen et al. 2014). Stem cell therapeutic animal model for osteoarthritis, inflammatory bowel disease, crohns disease, ulcerative colitis, diabetes to understand the effect and safety of infusion. Regime of biologics, administration route, and therapy results can be quickly transferred to human use.

4.6 Conclusion

Tremendous properties of stem cells hold a reliable source to solve extensive problems now our community is facing. The impact of stem cells on production and welfare of the livestock is revolutionary. On administration migrating to the site of infection, restoring the damaged tissue and boosting its immunity made stem cells a robust technology to use in farm animals. No optimized protocol to isolate the stem cells from different tissues across different species, effect of age on stemness, contradictory results on therapeutic efficacy and its safety needs extensive research.

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Chapter 5

Metabolomics and Proteomics Signatures in Feed-Efficient Beef and Dairy Cattle



Ahmed A. Elolimy, Mohamed Zeineldin, Mohamed Abdelmegeid, Alzahraa M. Abdelatty, Abdulrahman S. Alharthi, Mohammed H. Bakr, Mona M. M. Y. Elghandour, Abdelfattah Z. M. Salem, and Juan J. Loor

Abstract Feed accounts for 40–60% of total expenses of beef and dairy cattle production costs. Therefore, feed-efficient cattle have a great potential to reduce production costs without compromising meat or milk production levels, resulting in a greater profit margin for producers. Many approaches for measuring feed efficiency are available with residual feed intake being one of the most common. The residual feed intake is defined as the difference between actual dry matter intake and expected dry matter intake based on animal size and production level. Therefore, compared with a least-efficient animal, the most-efficient animal would have a negative resid-

A. A. Elolimy (✉)

Department of Animal Production, National Research Centre, Giza, Egypt

M. Zeineldin

Department of Animal Medicine, College of Veterinary Medicine, Benha University, Banha, Egypt

M. Abdelmegeid

Department of Animal Medicine, College of Veterinary Medicine Kafr-Elsheikh University, Kafrelsheikh, Egypt

A. M. Abdelatty

Department of Nutrition and Clinical Nutrition, Faculty of Veterinary Medicine, Cairo University, Giza, Egypt

A. S. Alharthi

Department of Animal Production, College of Food and Agriculture Sciences, King Saud University, Riyadh, Saudi Arabia

M. H. Bakr

Animal Production Department, Faculty of Agriculture, Cairo University, Giza, Egypt

Mona M. M. Y. Elghandour · A. Z. M. Salem (✉)

Facultad de Medicina Veterinaria y Zootecnia, Universidad Autonoma del Estado de Mexico, Toluca, Mexico

J. J. Loor

Department of Animal Sciences, University of Illinois, Urbana, IL, USA

ual feed intake coefficient value, indicating that it consumed less dry matter intake while maintaining the same level of production. Recent studies have focused on investigating changes in key metabolites and proteins that would shift metabolic pathways to support better feed efficiency. Recent reports highlighted that in most-efficient cattle metabolic pathways associated with energy, vitamins, and amino acid metabolism in rumen and skeletal muscle are upregulated to provide extra energy, thus, allowing for a similar level of production despite lower dry matter intake. Other studies demonstrated that most-efficient cattle reduce protein turnover in skeletal muscle including upregulation of key protein synthesis pathways, such as mechanistic target of rapamycin signaling, and the downregulation of key proteins in protein degradation such as ubiquitin-proteasome pathway, resulting in greater protein deposition in muscle. In this chapter, we discuss applications of novel comprehensive techniques for protein and metabolite profiling in rumen, intestine, blood, liver, and skeletal muscle to elucidate adaptive biological functions that support better feed efficiency in beef and dairy cattle.

Keywords RFI · Cow · Calves · Metabolomics · Proteomics · Rumen · Blood · Liver · Muscles · Hindgut

5.1 Introduction

Feed costs in the beef and dairy cattle industry are the most expensive inputs, and represent on average 40–60% of total expenses (Montaño-Bermudez et al. 1990). Therefore, enhancing feed efficiency would dramatically decrease overall costs, raise producer profitability, and increase animal protein availability for consumers (Clemmons et al. 2020). Hence, discovering robust biomarkers for selecting the most-efficient beef and dairy cattle is crucial.

Residual feed intake (RFI) is a commonly-used measurement of feed efficiency in beef and dairy cattle (Li et al. 2020; Zhang et al. 2020). The RFI is defined as the divergence of predicted dry matter intake (DMI) for maintenance and meat or milk production from the actual DMI after adjusting DMI for the level of production through a linear regression model (Xi et al. 2016). The predicted DMI is calculated as a function of changes in body weight (BW) and production level (Potts et al. 2017). The regression model defines which animal is below (negative) or above (positive) the predicted DMI (Durunna et al. 2011). The fact that RFI is a trait independent of body size and production level renders it a reliable measurement of feed efficiency (Gomes et al. 2012). Most-efficient (M-eff) cattle, i.e. with the favorable negative RFI coefficient, utilize less DMI than predicted to cover both maintenance and production requirements (Potts et al. 2017). Therefore, M-eff cattle are biologically- and economically-efficient compared with their least-efficient (L-eff)

counterparts having an undesirable positive RFI coefficient (Gomes et al. 2012; Lawrence et al. 2013). Despite the proven value of the RFI, the underlying biology associated with this trait is still not well known, hence, supporting the use of modern technologies in an effort to uncover putative biomarkers.

Metabolomics profiling provides a novel approach for rapidly-identifying M-eff animals. These techniques focus on detecting, identifying, and quantifying available free metabolites in a given biological sample (Fontanesi 2016). The origin of these metabolites could be either endogenous (i.e. derived from the animal) or xenobiotic (i.e. metabolites from plants or microbes) (Fontanesi 2016). Metabolites are intermediates or products of metabolic pathways (Clemmons et al. 2020) involved in energy, protein, and vitamin metabolism, all of which are of particular interest for feed efficiency divergence in cattle (Nafikov and Beitz 2007; Ferrell and Jenkins 1984). Liquid chromatography-mass spectrometry (LC-MS) techniques have been used to detect differences in ruminal, intestinal and circulating metabolite abundance between beef and dairy cattle divergent in RFI (Clemmons et al. 2017; Elolimy et al. 2020), with several metabolites identified as having a key contribution to better feed efficiency (summarized in Table 5.1). These alterations in metabolome profiles between M-eff and L-eff cattle could be used as predictive biomarkers for feed efficiency (Clemmons et al. 2019b; Artegoitia et al. 2017; Novais et al. 2019).

The study of protein profiles in a biosample collected at a certain physiological state is called “proteome” (Reinhardt et al. 2012). Proteomics data provide a wealth of information that gene expression analysis could not. Therefore, proteomics analyses provide a better understanding of the biological functions in cattle. Few studies in recent years have applied proteomic approaches to unmask alterations in protein profiles in blood and skeletal muscle associated with RFI divergence that would help our understanding of the biology behind this trait (summarized in Table 5.2).

5.2 Metabolomic Signature in Feed-Efficient Cattle

In a recent study, Clemmons et al. (2020) used DionexUltiMate 3000 ultra-high performance liquid chromatography (UHPLC) system with an Exactive Plus Orbitrap MS to identify the metabolic signature in ruminal fluid between M-eff and L-eff beef steers. Authors used 50 purebred Angus steers at the age of 7 months and 264 ± 2.7 kg of BW fed individually using the GrowSafe System for 70 days to evaluate RFI divergent groups. At day 70 of the trial, the authors selected the extreme M-eff ($n = 14$) and the top L-eff ($n = 15$) steers. Ruminal fluid samples were collected on day 70 of the study using stomach tubing. Results revealed 8 DEM between M-eff and L-eff steers, with M-eff steers having greater succinate. In the rumen, bacteria such as *Selenomasruminantium* metabolize succinate in the presence of several bacterial enzymes and coenzyme A (CoA) to propionate (Wirth et al. 2018), a vital volatile fatty acid (VFA) used in the liver for gluconeogenesis (Wirth et al. 2018). Therefore, the greater ruminal succinate in M-eff group indicates a better capacity for hepatic gluconeogenesis to maintain similar meat

Table 5.1 Metabolomics studies in residual feed intake (RFI) divergent beef and dairy cattle

Breed type	Breed name	Sex	Age (months)	Average body weight (kg)	Sample type	Approach and platform	Changes in M-eff	References
Beef	Angus	Steers	7	264	Rumen fluid	Untargeted metabolomics (UHPLC-MS)	↑ succinate ↑ Uracil ↑ Thymine ↑ Hypoxanthine ↑ Pyridoxic acid ↑ Citraconate ↑ Xylose	Clemmons et al. (2020)
Beef	Nellore	Bulls	16-20	376	Blood (serum)	Untargeted metabolomics (LC-MS)	↓ Retinal ↓ Progesterone ↓ Stearic acid ↓ Vomifolol, 2,3 ↓ Dihydroflavone ↓ Limonoate ↓ Phytanic acid	Novais et al. (2019)
Beef	Angus	Steers	7	264	Blood (serum)	Untargeted metabolomics (LC-MS)	↑ Pantothenate	Clemmons et al. (2019a, b)
Beef	Nellore	Cows	36	484	Blood	Targeted metabolites (enzymatic kits)	↑ Cholesterol	Broleze et al. (2020)
Beef	Nellore	Bulls	7	207	Blood (serum)	Targeted metabolites (enzymatic kits)	↓ Cortisol	Bonilha et al. (2017)
Dairy	Holstein	Heifers	At birth	43	Feces	Untargeted metabolomics (LC-MS)	↑ Cholesterol ester ↑ Biotin ↑ L-Tryptophan	Eloimy et al. (2020)
Dairy	Holstein	Cows	584	Lactating	Blood (serum)	Targeted metabolites (enzymatic kits)	↑ NPY ↓ Leptin ↓ NEFA	Xi et al. (2016)
Dairy	Murrah buffalo	Calves	4-6	70	Blood (plasma)	Targeted metabolites (enzymatic kits)	↑ IGF-1 ↑ Triiodothyronine T3 ↓ Thyroxin	Sharma et al. (2016)

Table 5.2 Proteomics studies in residual feed intake (RFI) divergent beef and dairy cattle

Breed type	Breed name	Sex	Age (months)	Average body weight (kg)	Sample type	Approach and platform	Changes in M-eff	References
Beef	Red Angus	Steers and heifers	Finishing stage	836	Ruminal epithelium	Targeted proteins (western blot)	↑ p-EEF2K ↑ p-EEF2K:EEF2K ↑ p-EIF2A:EIF2A ↓ UBA1 ↓ NEDD4 ↓ STUB1 ↓ MDM2	Elolimy et al. (2019)
Beef	Nellore	Bulls	7	239	Skeletal muscles	2D-PAGE	↑ HSPB1 ↓ 14-3-3 epsilon	Carvalho et al. (2019)
Beef	Nellore	Bulls	24-26	557	Liver	2D-PAGE	↓ HBB ↓ Aldehyde dehydrogenase ↓ Aspartate aminotransferase ↓ Glycine amidinotransferase	Baldassini et al. (2018)

production levels relative to L-eff animals. In support of this notion, Myer et al. (2015) reported that M-eff cattle had more succinate- and propionate-producing bacteria in the rumen such as *Succiniclasticum* spp. Similarly, M-eff steers had a greater concentrations of pantothenate in the serum, a precursor of CoA (Clemmons et al. 2017). These results indicate greater hepatic energy production in M-eff steers, which likely contribute to maintain growth in M-eff steers despite lower DMI (Fan et al. 2015).

Clemmons et al. (2020) also reported that M-eff steers had more abundant nucleic acids and nucleic acid derivatives in the rumen including uracil, thymine, and hypoxanthine, indicating an increased production of microbial protein in M-eff steers (Leng and Nolan 1984). Additionally, Clemmons et al. (2020) highlighted that M-eff steers had greater pyridoxic acid, a byproduct of vitamin B6 catabolism (Linkswiler and Reynolds 1950), suggesting better protein and muscle accretion in the M-eff group (Clemmons et al. 2020). Other key metabolites associated with carbohydrate metabolism such as citraconate was more abundant in ruminal fluid in M-eff steers (Clemmons et al. 2020). Citraconate is a metabolite generated through TCA cycle activity suggesting higher energy production took place in the rumen of M-eff steers (Clemmons et al. 2020). Collectively, data indicate that M-eff cattle have greater capacity for energy production in the rumen and liver to maintain similar growth performance despite lower DMI.

A recent study used untargeted metabolomics via liquid chromatography-mass spectrometry (LC-MS) to uncover differences in serum metabolomic profiles between young M-eff and L-eff Nellore bulls (Novais et al. 2019). In this study, serum samples from 98 Nellore bulls at 16–20 months of age and 376 ± 29 kg BW were collected 21 days before the start of a 70 day RFI evaluation period (Novais et al. 2019). Authors detected 7808 DEM between M-eff and L-eff groups (Novais et al. 2019). Seven metabolites had lower concentrations in the M-eff group including retinal, progesterone, stearic acid, vomifoliol, 2,3 dihydroflavone, limonoate and phytanic acid (Novais et al. 2019). Retinal is involved in the retinol pathway, previously reported to be downregulated in feed-efficient beef cattle (de Almeida Santana et al. 2016). Similarly, progesterone (a key metabolite in steroid hormone biosynthesis) was suppressed in the liver of M-eff Jersey cows (Salleh et al. 2017). Interestingly, this study revealed metabolites exclusively produced by bacteria or plants including vomifoliol, 2,3 dihydroflavone, limonoate and phytanic acid that were lower in the M-eff group (Novais et al. 2019), likely due to lower DMI.

Clemmons et al. (2019a) conducted a study to discover differences in serum metabolome between M-eff and L-eff beef cattle. In this study, they used LC-MS analysis for untargeted metabolomics of serum samples collected from M-eff ($n = 14$) and L-eff ($n = 15$) weaned Angus steers at 7 months old and 264 ± 2.7 kg of BW. The GrowSafe system was used to monitor individual DMI for each steer during the 70-day RFI trial (Clemmons et al. 2019a). Weekly, 9 mL of blood samples were collected via venipuncture from the coccygeal vein to separate serum. Results indicated that pantothenate was greater in the M-eff group (Clemmons et al. 2019a). Pantothenate, a substrate for CoA synthesis, is produced by ruminal bacteria such as *Flavobacteriathen* absorbed via ruminal epithelium to reach the

circulation (Clemmons et al. 2019a). Interestingly, Clemmons et al. (2019a) reported that Flavobacteriia were more abundant in M-eff steers, a result that is in line with previous studies revealing a better capacity for energy production from lower DMI in M-eff cattle.

Using commercial enzymatic kits, Broeze et al. (2020) evaluated differences in targeted metabolites in blood between M-Eff and L-eff beef cows. In this study, DMI of 53 primiparous Nellore beef cows at 36.8 ± 1.23 months of age and 484 ± 40.9 kg of BW was monitored individually using the GrowSafe System for 168 days between 22 and 190 days in milk (DIM; early-to mid-lactation stage) to calculate RFI coefficients for each cow (Broeze et al. 2020). Blood samples were collected from all animals for analysis of glucose, cholesterol, triglycerides, and β -hydroxybutyrate (Broeze et al. 2020). Cholesterol was the only metabolite that differed in concentration, being greater in M-eff beef cows (204 mg/dL vs. 192 mg/dL) (Broeze et al. 2020). Because another study reported increased plasma cholesterol in feed-restricted dairy cows (Gross et al. 2015), authors suggested that lower DMI in M-eff cows (consumed 11.5% DMI) partly explained the response observed (Broeze et al. 2020). In another study, Bonilha et al. (2017) used commercial enzymatic kits to evaluate differences in specific serum metabolites including insulin, non-esterified fatty acids (NEFA), insulin-like growth factor I (IGF-I), and cortisol between M-eff ($n = 13$) and L-eff ($n = 12$) Nellore bulls at 210 days of age and 207 kg BW monitored for individual DMI for 70 days. Cortisol, a biomarker of stress, was lower in M-eff bulls indicating a lower degree of systemic stress (Bonilha et al. 2017) and providing support to previous studies reporting lower circulating cortisol in M-eff cattle (Gomes et al. 2013; Richardson et al. 2004).

In a recent study, Elolimy et al. (2020) detected shifts in hindgut metabolomics profiles between M-eff and L-eff dairy preweaned calves. In this study, DMI in 26 neonatal Holstein heifer calves was individually monitored from birth to weaning at 42 days of age. Calves were retrospectively classified into two groups: M-eff ($n = 13$) and L-eff ($n = 13$) heifers based on individual RFI coefficient (Elolimy et al. 2020). Fecal samples were collected every two weeks throughout the study to perform untargeted metabolomics using an LC-MS approach (Elolimy et al. 2020). At birth, M-eff calves had an enrichment of metabolites belonging to energy-producing pathways including pyruvate metabolism, gluconeogenesis, TCA cycle, and biotin suggesting a greater availability of energy for growth and development during the preweaning period (Akram 2014; Vailati-Riboni et al. 2016; Elolimy et al. 2020). Furthermore, the M-eff group upregulated vitamin (biotin metabolism), fatty acid (arachidonic acid metabolism), and amino acid (alanine metabolism) related pathways that would likely enhance gut function and development (Elolimy et al. 2020; León-Del-Río 2019). During the preweaning period, Elolimy et al. (2020) demonstrated that M-eff calves had greater supply of B vitamins in the hindgut including vitamins B6, B7 (biotin) and B9 (folate). Vitamin B6 is essential for metabolism of fatty acids, amino acids, and glucose (Rodríguez-Melendez and Zemleni 2003). Vitamin B7 is important for mucosal immune responses (Jenkins et al. 2017). M-eff calves also had greater capacity for metabolism of amino acids

such as tyrosine, tryptophan, and phenylalanine (Elolimy et al. 2020) likely contributing to the similar growth achieved (Elolimy et al. 2020).

Another study measured the concentrations of leptin, prolactin, neuropeptide Y (NPY), insulin-like growth factor 1 (IGF-1), ghrelin, insulin, β -hydroxybutyrate, glucose, NEFA and growth hormone (GH) in serum between M-eff and L-eff dairy cows (Xi et al. 2016). The authors selected 29 lactating Holstein cows from a total of 84 based on their RFI coefficients to end up with two groups including M-eff (n = 15) and L-eff (n = 14) dairy cows (Xi et al. 2016). Blood samples were collected from all 29 cows through jugular venipuncture on day 1, 25 and 50 during the feeding period (Xi et al. 2016). The authors reported no differences in serum prolactin, IGF-1, ghrelin, insulin, β -hydroxybutyrate, glucose and GH (Xi et al. 2016). However, M-eff cows had greater concentration of NPY and lower leptin and NEFA (Xi et al. 2016) indicating lower propensity for fat mobilization, suggesting that M-eff cows likely had sufficient energy supply and experienced a lesser degree of negative energy balance.

Sharma et al. (2016) calculated RFI coefficients for 18 growing male Murrah buffalo calves at 4–6 months old and 70 ± 1.0 kg of BW after 99 days of a feeding trail, resulting in M-eff (n = 7) and L-eff calves (n = 11). Blood samples were collected at the start and end of the feeding trial by venipuncture of the anterior *vena cava* to evaluate plasma content of IGF-1, GH, creatinine, insulin, albumin, hydroxyproline, triiodo-thyronine (T3), thyroxin (T4), and total protein using commercial kits (Sharma et al. 2016). No differences in plasma concentrations of creatinine, insulin, albumin, hydroxyproline and total protein were detected (Sharma et al. 2016). However, M-eff calves had greater plasma IGF-1, and T3, but lower T4 (Sharma et al. 2016).

Overall, the above findings of alterations in several metabolites associated with better feed efficiency in beef and dairy cattle provide a list of robust biomarkers that could be studied in the future for their potential as physiological indicators predictive of feed-efficient cattle.

5.3 Proteomics Signature in Feed-Efficient Cattle

Elolimy et al. (2019) investigated changes in ruminal epithelium protein abundance between M-eff (n = 6) and L-eff (n = 6) Red Angus heifers and steers using the western blot approach. In this study, Elolimy et al. (2019) evaluated 29 proteins involved in protein synthesis (MTOR signaling) and degradation (ubiquitin-proteasome pathways) in ruminal epithelium collected after slaughter at the end of 70 days of a feeding trail. The M-eff group had greater abundance of proteins crucial for cellular protein synthesis such as phosphorylated eukaryotic elongation factor 2 kinase (p-EEF2K), phosphorylated eukaryotic elongation factor 2 kinase:total eukaryotic elongation factor 2 kinase (p-EEF2K:EEF2K, and phosphorylated eukaryotic translation initiation factor 2A:total eukaryotic translation initiation factor 2A (p-EIF2A:EIF2A)(Elolimy et al. 2019). On the other hand, M-eff cattle had

lower abundance of proteins involved in protein degradation pathways such as total ubiquitin like modifier activating enzyme 1 (UBA1), total neural precursor cell expressed, developmentally downregulated 4, E3 ubiquitin protein ligase (NEDD4), total STIP1 homology and U-box containing protein 1 (STUB1), and total MDM2 proto-oncogene (MDM2) (Elolimy et al. 2019). No differences were detected in plasma insulin and ruminal epithelium insulin signaling proteins (Elolimy et al. 2019). These data indicated that M-eff beef cattle have a greater rate of protein synthesis relative to protein degradation in ruminal epithelium. These changes likely result in better growth of ruminal epithelium to absorb more VFA produced from the anaerobic microbial fermentation of plant fiber in M-eff beef cattle.

Carvalho et al. (2019) employed a proteomics approach to unmask differences in key proteins associated with energy metabolism in skeletal muscle of RFI divergent beef cattle. Daily DMI was recorded for 129 young Nellore bulls at 7 months old and 239 ± 30.1 kg of BW during 98 days of RFI evaluation period (Carvalho et al. 2019). At the end of the study, Carvalho et al. (2019) selected 9 bulls for M-eff group and another 9 bulls for L-eff group. After slaughter, longissimus muscle was sampled for protein profiling using a two-dimensional electrophoresis (2D-PAGE) with mass spectrometry (ESI-MS) (Carvalho et al. 2019). Heat shock protein beta 1 (HSPB1), a key protein for cellular development and differentiation (Zhang et al. 2014; Carvalho et al. 2014) and inhibitor of protein degradation in muscle fibers, was greater in the M-eff group (Carvalho et al. 2019) suggesting greater protein synthesis. Therefore, this adaptation likely contributed to better feed efficiency in M-eff cattle through decrease protein turnover in skeletal muscle.

Another proteomics study conducted by Baldassini et al. (2018) used 2D-PAGE and ESI-MS techniques to profile hepatic proteins in RIF divergent Nellore bulls. In this study, Baldassini et al. (2018) used 18 Nellore bulls at 24–26 months of age during the finishing period (M-eff = 9 and L-eff = 9). After slaughter, liver samples were collected from the 18 animals for protein extraction and proteomic profiling (Baldassini et al. 2018). Results indicated that hemoglobin subunit beta protein (HBB) was downregulated in the M-eff group (Baldassini et al. 2018) likely from lower numbers of red blood cells since hemoglobin binds to oxygen to form oxyhemoglobin inside the red blood cells (Hsia 1998). The data indicated a need to determine blood hemoglobin concentrations between M-eff and L-eff groups in future studies. Baldassini et al. (2018) reported that the M-eff group had lower abundance of oxidative stress-associated proteins such as aldehyde dehydrogenase, aspartate aminotransferase, and glycine amidino transferase proteins, highlighting a lower degree of hepatic oxidative stress and less reactive oxygen species (ROS) content in liver of feed-efficient cattle.

Davis et al. (2016) investigated differences in oxygen uptake by mitochondria in muscle and respiratory chain proteins between M-eff and L-eff beef cattle. They calculated RFI coefficients in 92 Hereford-crossbreed steers in 63-day feeding period using the individual feed intake system. The top 10 M-eff and top 8 L-eff steers based on RFI coefficient ranking were used for subsequent analysis. Mitochondrial complex I (CI), II (CII), and III (CIII) protein concentration in M-eff and L-eff groups was assessed using bicinchoninic acid colorimetric procedures.

Lymphocytes were isolated from blood samples collected via jugular venipuncture from both groups (Davis et al. 2016). Results indicated that M-eff steers did not differ in CI, CII, and CIII protein concentration. Therefore, mitochondrial proteins do not seem to play a key role in the RFI divergence between M-eff and L-eff steers.

5.4 Conclusions

Comprehensive metabolome and proteome studies revealed associations between RFI divergent cattle and shifts in metabolome/proteome profiles that might explain the biology behind superior feed efficiency in cattle. Overall, M-eff cattle are characterized by metabolite and protein profiles that would provide extra energy and nutrients to help maintain similar levels of production despite a lower DMI. Further studies are warranted to expand our understanding of the biological contribution of key metabolites and proteins to RFI divergence in cattle at different production stages. Additionally, the relationship between metabolome/proteome profiling among different tissue in the same individual animals such as rumen, blood, skeletal muscles and milk should be examined in order to provide a more holistic overview into feed-efficient cattle.

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Chapter 6

Biotechnological Applications in Dairy Products and Safety



Ayushi Kapoor, Monica Yadav, Aparna Verma, and Kiran Ambatipudi

Abstract From harvest to distribution, the food industry's application is as old as the industry itself. Conventional technologies such as pasteurization, homogenization, and direct microscopic count are well established and practiced mainly in the dairy industries. However, there is a need to evolve these technologies due to food security, changing consumer needs, environmental concerns, economic viability, and policy reforms. Apart from this, today's food industry is also facing several challenges related to food safety, preservation, nutrition, and allergies. Therefore, the presently used technologies in the production, processing, packaging, and safety of various dairy products need to evolve further to overcome these challenges. Here we review the potential applications of food safety tools like hazard analysis and critical control point, sustainable packaging materials like polylactic acid, and non-conventional technologies such as ultrasound to solve current problems. This chapter discusses the role of various biotechnologies in food packaging, processing, and safety, focusing on different dairy products, from indigenous to by-products. The food industry also needs to identify and overcome socioeconomic challenges to proceed with these non-conventional, relatively new technologies from laboratory to industry.

Keywords Food industry · Pasteurization · Dairy products · Food safety · Food Packaging · Nutrition · Non-conventional technologies · Non-thermal processing · Policy reform · Biotechnological applications

A. Kapoor · M. Yadav · A. Verma · K. Ambatipudi (✉)
Department of Biosciences and Bioengineering, Indian Institute of Technology Roorkee,
Roorkee, Uttarakhand, India
e-mail: akapoor@bt.iitr.ac.in; myadav@bt.iitr.ac.in; averma@bt.iitr.ac.in;
kiran.ambatipudi@bt.iitr.ac.in

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6.1 Introduction

Food technology deals with various applications used in the production, packaging, safety, storage, and distribution of food products. Such technologies enhance the shelf life of perishable food, convert the comparatively bulky food to a convenient form, offer more food choices and make food available during off-seasons, thus fulfilling consumer demand. Once harvested from agricultural fields, post-harvest technologies such as threshing, drying, cleaning, and milling are applied to the agricultural produce to protect it until further processing begins. From the age-old techniques such as smoking, salting, drying, and pickling to recent developments like irradiation, canning, freeze-drying, today's food industry uses all sorts of technologies for processing. With technological advancements, it's not only the main product processed, but several by-products are also manufactured with further processing. The large-scale production and availability of food are also advantageous in ensuring food security, especially for developing nations. Biotechnological applications like enzyme and fermentation technology are responsible for increasing the shelf life, enhancing the flavor, texture, and nutritive value of food products (FAO International Technical Conference, 2010). Fermentation technology is highly employed in the dairy industry, especially in yogurt, cheese, cultured buttermilk, kefir, and sour cream. More recently, the kefir grains have been used to ferment and develop soybean beverages supplemented with inulin (dos Santos et al. 2019). Similarly, enzyme technology is well established in fruit juice for liquefaction and clarification. For example, the addition of protease in cherry juice and immobilizing polygalacturonase in calcium alginate microspheres in apple juice reduces juice's turbidity levels (Pinelo et al. 2010; Deng et al. 2019). The advances taking place in these technologies from time to time further improve the product, thus meeting evolving market demand requirements.

Food packaging is one of the most crucial manufacturing steps since it is responsible for protecting its content, conveying information through labels, appealing to consumers, and maintaining product quality. The food packaging industry has come a long way from coating cheese in grease and lard to biobased packaging material and edible films. For instance, active food packaging with polymer material is functionalized with synthetic peptides to form antimicrobial packaging (Agrillo et al. 2019). The packaging is closely associated with the safety aspect of food. The food safety tools are implemented to control physical, chemical, and biological hazards and assure that food upon consumption shall not cause harm. Therefore, it becomes equally essential to ensure that processing methods must comply with food safety norms. Not just nutritious, food must be safe and fit for consumption to prevent foodborne illness and outbreak. Food scares are the repercussions of pandemics and public health emergencies as they cause safety concerns among consumers, and the COVID-19 pandemic is a recent example. Even though the researchers reported no relation between the spread of the virus through food, the pandemic has created awareness and raised significant concern for safe food consumption.

The dairy industry is an integral part of the whole food system, with annual global milk output of 843 million tons in 2018, and India as the largest producer and consumer. In recent times, various dairy by-products have started being explored for

their functional properties, for example, phospholipids isolated from buttermilk, butter serum, and cheese whey confer health benefits in humans (Verardo et al. 2017). Utilizing dairy industry by-products is economical and adds to the existing products, and reduces the environmental load. The dairy industry, in particular, uses most of the applications, which are used in food industries. This includes, but is not limited to, fermentation technology, membrane filtration, microbial testing, etc. The chapter consists of various dairy industry biotechnological applications, especially in dairy products (e.g., cheese, yogurt, paneer, and infant formula). The chapter starts with the dairy industry description and further explains various production, processing, packaging, and safety technologies applied in the dairy industry. The current limitations of food industries and their potential solutions are also discussed, including socioeconomic constraints and possible solutions.

6.2 Historical Perspective

Due to limited knowledge, technological advancements, and demand, most dairy products prepared until the mid-nineteenth century were farm-level with the challenges such as non-uniformity in product quality, low quality of raw material, lack of proper market, improper packaging. But with increasing population and subsequently increasing demand, rapid industrialization, improving transportation facilities, and development of technology, dairy production transited from farm to factory level production (De 1980). Pasteurization was invented in 1862 for processing wine, but in the early 1900s, it found application in the dairy industry for pasteurizing milk (Smith-Howard 2017). From vat pasteurization, high-temperature short-time pasteurization around the 1930s to ultra-high-temperature pasteurization in 1948, the technology has evolved because of the advancements of the core concept of killing unwanted microorganisms. Cooling milk is a crucial step in milk processing since it is highly responsible for keeping milk quality. The cooling cans used earlier were replaced due to the advent of mechanical refrigeration in the mid-twentieth century. Homogenization, the pressure-driven process of reducing the size of milk fat globule, was first used in commercialized milk in 1919 but gained consumer acceptance around 1940–1950 (Cano-Ruiz and Richter 1997; Weimar 1994).

It was not just the technological development in terms of processing, but the food safety norms also progressed with time. Processing methods are highly interlinked and have a significant impact on product quality and microbial safety. This relation could be better understood, for example, pasteurization kills pathogenic microorganisms, and refrigeration of milk helps in keeping the microbial load low, therefore, processing methods like these directly contribute to ensuring food safety. Similarly, penicillin, whose discovery was unrelated to the dairy industry, impacted milk processing (Gaynes 2017). Even the packaging of dairy products has come a long way. For example, earlier cheese was packaged in lard, cheese grease, or paraffin, which then shifted to plastic materials like polyethylene terephthalate and now progressing towards edible films and coatings. The commercialized milk was earlier available in returnable quart bottles, then in plastic-coated milk cartons, and later in

plastic containers. With rising environmental concerns, plastic is being replaced with environment-friendly and sustainable solutions such as biobased packaging material. The industrial concern is transitioning from mere food preservation to safety, food security, and sustainability. This is apparent with stricter regulations where nutrition labeling was introduced and emphasized in the 1970s, acts such as the food safety modernization act are the need of the hour. Prevention of Food Adulteration Act, 1954 was a prominent act in the Indian food system until 2011, when food safety and standards rule came into force. Prevention of Food Adulteration Act and various such acts were repealed. This itself exemplifies the shift from food adulteration to food safety. The rising concern of the environmental impact of food processing, waste treatment, and the quest for greener and cleaner solutions is shifting scientists' focus on long-term food processing methods. Novel food processing methods such as high hydrostatic pressure and pulsed electric fields offer great potential in this regard (Fig. 6.1).

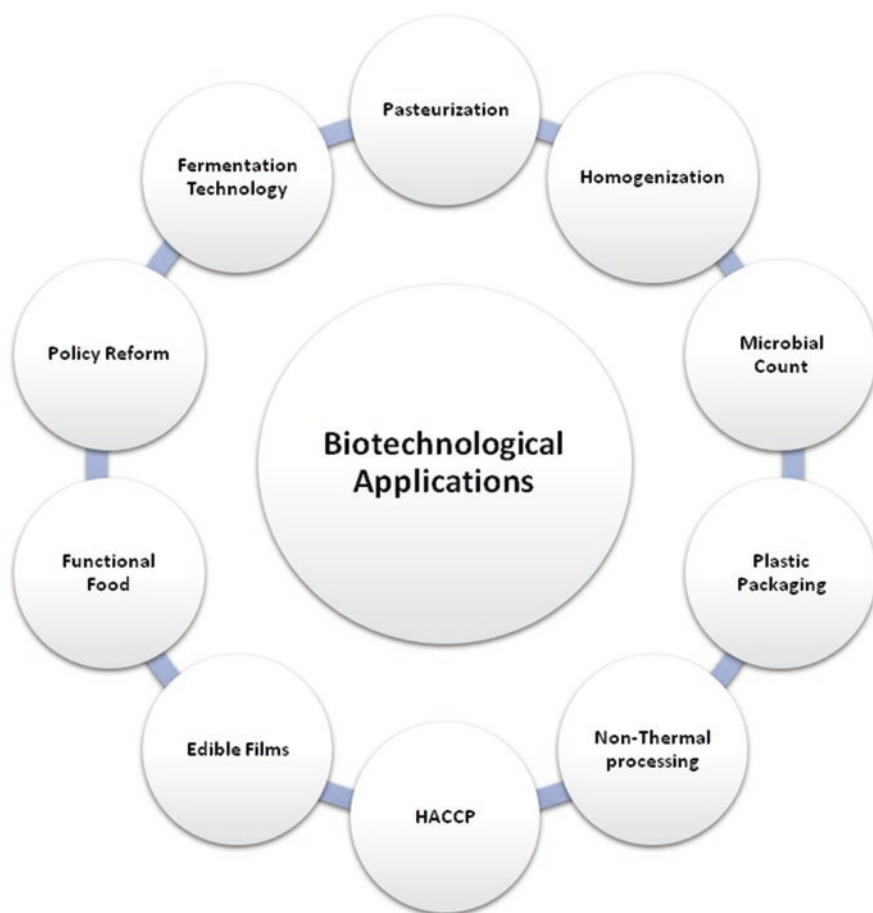


Fig. 6.1 Schematic overview of biotechnological applications in the food industry

6.3 Dairy Products

6.3.1 Cheese

Cheese is made from milk using rennet enzymes and/or lactic acid bacteria for casein coagulation followed by separation of developed curd. Initially, it was made at the farmhouse level in small quantities with low milk quality as the major challenge in large-scale production (Johnson 2017).

6.3.1.1 Production and Processing

The advancement of microbiology and technology in refrigeration, commercial starters, and pasteurized milk has improved cheese quality and initiated factory level production in the latter half of the nineteenth century (Johnson 2017; De 1980). The early challenges in cheese production were overcome gradually by applying various processing methods such as milk in cans, methods to cool milk, direct microscopic counting of bacteria, and, most importantly, by pasteurization. These processing steps contributed to bringing uniformity to the end product, promoting commercial production. Milk naturally contains many bacteria, such as coliforms hindering the fermentation process, resulting in undesirable acidification and gassy cheese. In contrast, pasteurization kills most bacteria that ferment lactose and/or cause uncontrolled fermentation thus giving more control over the cheese production. Even though pasteurization interferes with the flavor development of cheese and a costly technique, it had an advantage in quality improvement with higher cheese yield (Price 1927; Wilson et al. 1945). Presently, a high-temperature short-time pasteurization method is used where milk is heated using a heat exchanger to 72 ° C for 15 s, followed by cooling (Singh and Heldman 2001). This heating-holding-cooling method enables the production of cheese with uniform quality.

Milk composition and cheese yield are the key factors that determine profit, milk price, and cheese making efficiency, with yield primarily depending on moisture, fat and casein content, and retention of each component during the processing (Wedholm et al. 2006; Lopez-Fandino et al. 1996). Variation in milk composition is a relatively common phenomenon, attributed to alterations across seasons, breed, feed, health conditions of the animal, etc. Therefore, to maintain uniformity in commercial cheese production, it is imperative to ensure consistency in the raw milk used to produce cheese. Membrane technology (e.g., microfiltration, nanofiltration, and osmosis) is a separation process where pressure is applied to facilitate the separation of individual components in the form of retentate while allowing the permeate to flow through. This pressure-driven process is used in cheese manufacturing because of its role in improving the nutritive value, cheese yield and consistency, whey utilization, and relative reduction in the use of rennet and starter culture (Kumar et al. 2013). Membrane filtration has been strongly advocated to remove lactose and water for concentrating the milk fat and casein. With the membrane pore

size ranging from 1 to 100 nm, the widely used ultrafiltration technology is primarily used to eliminate lactose and increase fat and casein content to improve cheese yield. The ultrafiltration technology is further categorized based on the concentration factor: low, medium, and high-concentration ultrafiltration. The low-concentration ultrafiltration is mainly used to standardize milk protein, while medium-concentration factor ultrafiltration is primarily used to concentrate milk (Soodam and Guinee 2018). The microfiltration process efficiently removes bacteria and spores from milk due to its membrane pore size of 0.2 to 2 μm . Thus, the microfiltration of pre-treated milk improves curd firmness and reduces the number of additives such as calcium dichloride to promote ripening (Caron et al. 1997).

Membrane technology is also applied in the processing of whey, the nutritionally rich by-product obtained during cheese production. The large-scale manufacturers retrieve the different nutrients such as lactose, protein, and minerals from whey, while small-scale processors consider it as a waste as further processing for valorization is regarded as uneconomical, especially in developing nations (Chen et al. 2019). Membrane filtration of whey leads to the production of high-quality products such as whey protein isolates (>90% protein) and concentrates (30–80% protein) for foaming, gelling and emulsifying agents (Kumar et al. 2013). Recently, Samtlebe et al. (2017) designed a tubular membrane filtration system for controlling the phage before processing cheese whey to develop different whey products. A 0.1 μm membrane filtration model was found better suited for phage reduction as well as protein transmission. The application of membrane technology has become a well-established technology in the cheese industry, including the processing of its by-product whey with fouling as one of the significant limitations of the technology. Fouling occurs mainly due to pore blockage, particle adsorption on pores, the formation of cake and protein, and mineral deposition (James et al. 2003), resulting in a significant drop in membrane efficiency and economic loss. Therefore, fouling is reduced by regularly cleaning membranes, using low-fouling membranes, and applying high pressure.

Fermentation is an integral part of cheese making, and commercial starters proved to be a milestone in factory level cheese production. Pasteurization of raw milk kills most of the bacteria, resulting in unwanted fermentation, enabling to better control cheese's fermentation characteristics, and developed acidity is due to added starter culture to maintain cheese quality. The discovery of plasmid DNA encoding proteolytic activity and lactose fermentation was a significant advancement in starter cultures (McKay et al. 1976; Romero and Klaenhammer 1993). For example, commonly used starters in cheese consist of mesophilic lactic acid bacteria, predominantly *Lactococcus lactis* species which is responsible for acidification, proteolytic activity, flavor production, and phage insensitivity. Starter cultures are also associated with flavor development in cheese. Ayad et al. (2000) reported wild strains of *Lactococci* spp. in developing new flavors in cheese. Additionally, permeabilization, autolysis, and intracellular enzyme release of starter culture are also impacted by the alterations in cheese components such as fat and salt content (Yanachkina et al. 2016). Rennet is a set of enzymes used to coagulate milk during cheese making, and chymosin is the active enzyme in rennet. Before 1980, most of

the rennet was extracted from calf stomach, but the advent of calf genes' insertion led to chymosin production from molds. And today, the majority of the rennet used is fermentation-produced chymosin (Johnson 2017). Various advances have been made in recent studies researching the parameters associated with the rennet in cheese making. For instance, the milk-clotting enzyme BY-1 developed from *Bacillus subtilis* PNG27 exhibited better flavor and acceptability in cheese, indicating its potential use in cheese production (Meng et al. 2018). More recently, Dussault-Chouinard et al. (2019) reported the enhanced cheese-making characteristics due to lowered renneting pH. Additionally, processing of the milk is performed using ultrasound, different inoculation methods, starter cultures, and ripened molds to influence and enhance the rennet coagulation properties and sensory qualities during cheese making (Ragab et al. 2020; Galli et al. 2016). Cheese ripening is the development of flavor, texture, and other characteristics, such as aroma and specific taste. It is the enzymes and microbiota in specific which contributes to the flavor development of cheese. Flavor in cheese plays a crucial role and often distinguish one variety of cheese from the other, for example, without the addition of lipase enzyme, the flavor of Provolone and Romano would be similar to mozzarella (Johnson 2017).

6.3.1.2 Packaging

Initially, cheese was coated with cheese grease, paraffin, or lard to prevent drying, but with the large-scale production and increasing demand, newer ways were developed to protect the product and prevent contamination. Significant challenges in cheese commercialization are the losses due to contamination by yeast, moulds, and bacteria during storage, often leading to off-flavor and reducing the quality (Costa et al. 2018). The improper packaging causes moisture loss in a few cheese varieties, resulting in increased hardness and low organoleptic properties. The majority of the currently used food packaging materials are composed of plastic materials, including polyethylene terephthalate, polystyrene, polypropylene, polyvinylchloride, and polyethylene, which is primarily associated with environmental concerns because they cannot be recycled entirely or biodegraded (Piscopo et al. 2019). Thus, researchers focus on developing newer technologies such as edible films and coatings, nanoparticle embedded packaging materials, engineered nanofibers for packaging, and treating food with pulsed light to prevent post-process contamination. Biobased packaging materials, such as polylactic acid, are primarily derived from renewable resources produced through chemical synthesis of biomass monomers, have relatively low water vapor barrier properties limiting their use in cheese packaging. For instance, polylactic acid in packing semi-hard cheese resulted in moisture loss and surface drying (Holm et al. 2006). The edible films and coatings comprising polysaccharides, proteins, and lipids are another recent advancement in food packaging where films can be consumed with the food, thereby avoiding waste generation. To curb the bacterial contamination in cheese, researchers have recently developed nanofibers containing nisin and loaded with poly-g-glutamic acid/

chitosan and moringa-oil chitosan nanoparticles, which has shown promise as a potential active packaging material in cheese preservation (Cui et al. 2017; Lin et al. 2019). Pulsed light treatment, a non-thermal processing technology, also finds a potential application in reducing cheese contamination. More recently, it has been reported that pulsed light's post-packaging application could be used successfully to minimize post-processing contamination in cheese.

6.3.1.3 Food Safety

The quality of the milk used for cheese production has drastically improved over time, mainly due to the application of technologies such as pasteurization. Today, the bacterial count in pasteurized milk is significantly fewer (e.g., 100 bacteria/mL) compared to the previous cheesing making procedure (5 billion CFU/mL) (Kelly 1939). As cheese is highly susceptible to contamination at various processing and post-processing steps, it demands strict cleaning and sanitation facilities. For instance, water activity, pH of cheese, and competitiveness with other bacteria determine the growth and survival of pathogens present due to contamination during manufacturing, storage, and cheese ripening. Similarly, few surface-ripened kinds of cheese like Limburger and mold-ripened cheese such as Camembert and Brie are more prone to pathogenic growth because of their high-water activity and tendency to lose acidity during ripening (Johnson 2017). Although pasteurization has proved to produce better quality cheese, negligence in proper cleaning and sanitization and prolonged use of the equipment has often led to developing a complex microbial consortium called biofilms. Recently, Lee et al. (2017) reported the biofilm-producing ability of *Listeria monocytogenes* on stainless steel and polystyrene microplates in cheese plants. Antibiotics residue is another primary safety concern in cheese manufacturing as it could inhibit acid production by starter cultures that leads to technological and economic loss (Chiesa et al. 2020). In contrast, Silva et al. (2020) reported no effect of fermentation and physicochemical characteristics of Minas Frescal cheese by an antibiotic (e.g. monensin) residue in milk. Nonetheless, antibiotic residues remain a matter of concern due to their effect on product and consumer health.

6.3.2 Paneer

Paneer is an indigenous dairy food product rich in animal protein and possesses a high biological value of 80–86. It is obtained by heat treatment of milk, followed by coagulation, where the end product has a mildly acidic flavor, spongy texture, and whitish color. Both cow and buffalo milk are suitable to prepare paneer; however, the latter is preferred because of its high fat, minerals, and calcium content and whitish appearance, as desirable. In contrast, paneer made from cow milk has a soft texture with a yellowish appearance and does not appeal to consumers. Paneer has

been classified as one of the soft cheese varieties highly used in culinary dishes (Kumar et al. 2014).

6.3.2.1 Production and Processing

To maintain uniformity in the final product, it is necessary to ensure consistency in the raw materials obtained, processing conditions, quality analysis of the product, proper storage, and optimum conditions throughout the supply chain. For instance, membrane technology, the widely used filtration technology, has been extensively used to concentrate the standardized milk for paneer preparation. The ultrafiltration concentrates standardized milk to 27% total solids, resulting in more whey proteins for paneer manufacturing, leading to the total solids recovery up to 95% and higher paneer yield. Kanawjia and Rizvi (2000) reported that adding 0.15% calcium chloride to the membrane filtration retentate improves the paneer's textural and organoleptic properties. Reverse osmosis has also been used for concentration, but its main advantage lies in low energy consumption. The recent advancement of utilizing reverse osmosis to obtain milk solids from dairy rinse water is an efficient environmental strategy for the dairy industry (Brião et al. 2019). In paneer processing, milk is coagulated after filtration, heating, and cooling. The most commonly used coagulants in paneer preparation are lactic acid, citric acid, alum, tartaric acid, or sour whey, including several coagulants that impart different properties to the quality of the final product. The use of citric and tartaric acid at a 2% concentration has been reported to make paneer with the most suitable properties from reconstituted milk (Khan et al. 2014).

In contrast, malic acid was found as the most suitable coagulant for better flavor and soft body, while calcium lactate for higher yield and whiter color, and citric acid for overall acceptability of paneer prepared from cow milk (Kumar et al. 2019; Shanaziya et al. 2018). Recently, Amini et al. (2019) reported fermented rice milk as a novel coagulant for paneer preparation. Sharma et al. (2002) reported that solids-not-fat, total solids, and paneer yield were highest in winter (8.983%, 13.639%, 15.5.1% respectively), summer, and rainy season, emphasizing not only coagulants but seasons also influencing paneer composition and yield. Interestingly, sour fruit juices such as lemon and amla (Indian gooseberry) juice have also been used as paneer coagulants. The sensory evaluation to analyze the appearance, flavor, body, and texture found sour juice coagulated paneer as a suitable product (Ahmed and Bajwa 2019). Like further processing, paneer whey, similar to cheese whey, is utilized as a potential product for human consumption. Lactose, also called milk sugar, accounts for approximately 70% of total solids present in paneer whey. Recovering lactose post further processing is economical and environment friendly since lactose reduces the biochemical oxygen demand of whey by 80%. Bund and Pandit (2007) studied sonocrystallization to fasten lactose recovery from paneer whey and optimized the recovery rate at 90.3% in 20 min. Paneer whey has also been explored to prepare functional pineapple beverages and has been reported to exhibit higher mineral content and better shelf life than cheese whey (Baba et al.

2016). Additionally, paneer whey has also been reported for inulinase production as a cost-effective substrate (Singh et al. 2019).

6.3.2.2 Packaging

Food packaging is mainly done to enhance the product's shelf life, maintain quality, provide mechanical, chemical, and microbial protection, appeal customer, and provide details on labels. Paneer is a perishable product, highly susceptible to physico-chemical and microbial contamination, and therefore, packaging correctly is critical for consumption. The most commonly used packaging material for paneer includes coextruded films, polythene sachets, laminates, parchment paper, etc. (Kumar et al. 2014). Similarly, modified atmospheric packaging is a food packaging technique with a modified atmosphere inside the package to enhance the shelf life and maintain food quality. This is achieved either by flushing desirable gases inside the package, or it occurs due to food's respiration process. The application of modified atmospheric packaging in paneer packaging is relatively new, but the technology has gained significance since the primary concern associated with paneer packaging is its shelf life. The modified atmospheric packaging has been studied in paneer packaging to enhance the product shelf life (along with hurdle technology for paneer processing), to evaluate paneer quality based on its color, and to study its influence on chemical properties of paneer such as moisture, pH, titratable acidity and free fatty acid (Thippeswamy et al. 2011; Shrivastava et al. 2013; Rai et al. 2008). The application of sequential treatment such as supercritical-carbon dioxide in association with food-grade acetic acid works as a sustainable, non-thermal method to enhance paneer shelf life up to 30 days (Kapoor et al. 2013). Edible films and coatings are also a potential packaging material for paneer as it not only increases paneer's shelf life but also fortifies it, thereby enhancing its nutritional value. This includes the usage of whey protein concentrate-based edible coatings consisting of ferrous sulfate heptahydrate, starch, and carboxymethyl cellulose-based edible film treated with clove oregano essential oil (Jotarkar et al. 2018; Karunamay et al. 2020). Vacuum packaging is another advancement in food packaging where the air is removed from the package and sealed hermetically. The primary advantages include shelf life extension, flexible film packages like shrink film, and oxygen reduction to limit aerobic bacteria's growth. Paneer is usually vacuum packed in co-extruded and laminated films. The linear low-density polyethylene and the binding agent have been proved as an effective packaging combination for vacuum packing and paneer tikka storage for 40 days (Ahuja and Goyal 2013).

6.3.2.3 Food Safety

Sensory evaluation is a discipline that evaluates consumer food products by using human senses such as smell, sight, touch, taste, and hear. The food should taste not only acceptable but also look appealing. For example, no matter how nutritious

orange juice is, consumers will not accept the product unless it appears to be 'orange.' Therefore, the sensory evaluation of products is conducted to analyze the overall acceptability. Since sensory evaluation is all about perceiving human senses, mostly human panels are involved as trained and consumer panels. Schlossareck and Ross (2020) evaluated the intensity of aftertaste and liking of spicy paneer by employing a consumer panel and observed that the aftertaste stayed was directly proportional to the spice level of paneer. Nonetheless, error and inaccuracy are still high with human panelists in sensory evaluation because of each panelist's varying experience. Therefore, more recently, the electronic tongue, an analytical tool, has drawn attention for better accuracy as a potential tool in qualitative analysis of food products like paneer.

Paneer is a highly perishable food product with short shelf life, and the high moisture content of approximately 70% increases the chance of microbial contamination. Although refrigeration is suggested to store it, there are microorganisms like *Escherichia coli* that jeopardize the safety of food. Sharma et al. (2019) demonstrated the use of phytochemicals, including gallic acid, piperine, menthol, quercetin, and eugenol, with antimicrobial properties against *Escherichia coli* in paneer. Another critical parameter in paneer preservation is maintaining a low temperature. The cold-chain facilities in India are still in the developing phase where most areas lack proper electricity supply, thus hampering the product quality and limiting the shelf life of various dairy products. The paneer's shelf life is a significant restriction in its proper utilization, and therefore, efforts are made to enhance the paneer's shelf life. One such attempt was made by Pandey et al. (2019), where paneer whey was used for the production of paneer making powder with bio-preservative potential. The paneer obtained exhibited a lower count of the total plate, *Staphylococcus*, lactic acid bacteria, and coliforms, thereby increasing the shelf life to 75% compared to control.

6.3.3 Yogurt

Milk is the most desired food consumed to fulfill the essential nutritional requirement of humans. The dairy products such as yogurt and other fermented dairy products (e.g., cheese, kefir) impart high nutritional value, sensory properties, and overall health (Cuevas-Gonzalez et al. 2020). The incorporation of lactic acid bacteria in fermented products due to their beneficial health benefits has led to probiotic's evolution, with the ingestion of live microorganisms to restore or improve the host's gut microflora. Yogurt (popular fermented dairy product) is consumed worldwide because of its crucial role in improving intestinal health, immune system (Weerathilake et al. 2014; Pei et al. 2017), maternal health, and pregnancy outcomes (Kok C. R and Hutkins R 2018; He A et al., 2020). It acts as a suitable carrier for delivering functional ingredients like probiotics and prebiotics (Castro et al. 2015; Hill et al. 2017). Yogurt is relatively more nutritious than milk as an excellent source of calcium, magnesium, potassium, milk fat, and vitamins (e.g., B₂, B₆, B₁₂) (Ayar

et al. 2014), including milk proteins and amino acid with a higher biological value for sustaining good health (Lourens-Hattingh 2001). Ndife et al (2014) reported a crucial role of yogurt in improving lactose tolerance, immune enhancement, metabolic, and gastrointestinal disorders (e.g., inflammatory bowel disease). Furthermore, several non-nutrient components present in yogurts, such as sphingolipids, conjugated linoleic acid, and butyric acid, have been reported to have anti-cancerous properties (Sanders et al. 2007; Gahruie et al. 2015). Interestingly, fortified yogurt consumption with iron, fibers, and seed oil has been reported to effectively ameliorates nutritional diseases such as osteoporosis, rickets, malnutrition, and atherosclerosis (Gahruie et al. 2015). Yogurt has also found an important place in the older population's diet due to its accessibility and beneficial role in improving their nutrition and health profiles (El-Abbadi et al. 2014). Thus, due to increased awareness of the numerous beneficial effects of yogurt, there has been a steady rise in the demand and popularity of yogurt globally, and in turn, has driven scientists to develop innovative strategies to improve further the quality and quantity of yogurt production (Chandan et al. 2017). In the market, multiple yogurt varieties such as plain, stirred/blended, vanilla flavored, smoothies/whips/mousses, frozen, organic, and natural are available for consumption. These yogurts are incorporated with different additives such as stabilizers, colors, exopolysaccharides, flavors, fruits, preservative, and sweetening agents to improve their sensory and texture characteristics (Fazilah et al. 2018).

6.3.3.1 Production and Processing

Bacterial exopolysaccharides are natural stabilizers and have gained momentum as safe during yogurt manufacturing, thereby eliminating the commercial stabilizers (Gawai et al. 2017). To generate novel functional food, yogurt-based products have been enriched with plant-derived sterols/olive or carrot juice, including many other products (e.g., cereal yogurt-like beverages, synbiotic yogurt) to impact human health positively (Buyuktuncer et al. 2017; Zoidou et al. 2017). To this end, emerging fields of biotechnology, microbiology, and food bioengineering have contributed significantly to improve the overall quality and safety of different dairy products. For example, biotechnological techniques have been used to incorporate microorganisms in food to enhance the aroma, taste, flavor, shelf life, and nutritional value of foods. Additionally, biotechnological tools have been used to produce a range of products (e.g., enzymes, vitamins, and food ingredients), including selection and manipulation of microorganisms to upgrade the product quality, safety, consistency, and yield (Ghoshal et al. 2018). Yogurt has been defined as a mixture of milk (whole, reduced-fat, low fat) produced by culturing lactic-acid producing bacteria such as *Lactobacillus delbrueckii* spp. *Bulgaricus* and *Streptococcus thermophiles*. Additionally, *Acidophilus* spp. culture has also been reported to produce yogurt and considered as "Generally recognized as safe" (Weerathilake et al. 2014). The microbial cultures decrease the pH and coagulate the milk proteins via lactic acid obtained through lactose fermentation, providing a viscous gel-like

structure. The milk components converted to volatile and non-volatile acids (e.g., acetone, acetaldehyde) within the yogurt give a characteristic flavor (Beshkova et al. 1998; Dan T et al. 2019).

Starter culture strains such as *Streptococcus thermophiles* zlw TM11 secrete a high amount of exopolysaccharide to improve the texture and viscosity (Folkenberg et al. 2006). Of note, flavor, texture, and taste vary depending on the lactic acid bacteria starters and metabolites secreted by the bacteria. Subsequently, the different manufacturing processes (e.g., homogenization, heat treatment, pasteurization) and types of ingredients (e.g., fruit pieces, stabilizers, bacterial cultures, flavors, and colors) have been reported to alter yogurt's quality (Nagaoka et al. 2019). The most commonly available yogurts are the set and strained, but frozen and drinking yogurt has gained significant popularity more recently. To date, yogurt manufacturing has been mainly from bovine milk (e.g., sheep, goat, water buffalo), but other mammals' (e.g., camels) milk has recently aroused interest. These dairy products are carriers of probiotic bacteria, which enter the human intestine by maintaining their survivability through their stomach and modulating intestinal microflora. It requires a thorough understanding to increase their usage and application in dairy science and medicine (Elizaquivel et al. 2011). The manufacturing of yogurt begins with the standardization based on varying fat amount in milk, and proteins (5–15%), followed by physical treatment such as heat (90–95°C, 3–7 min), homogenization (20–25 Mpa at 70°C), deaeration (70°C), cooling (42°C), and inoculation of microbial culture (10^6 – 10^7 CFU/ml). Subsequently, after fermentation, yogurt production proceeds with harvesting, post-treatment, and packaging. The formation of a few yogurts such as set-type, stirred, drinking, and concentrated depends on the steps (e.g. number) during the manufacturing influences the yogurt's final quality (Lopes R P et al. 2019). After the final packaging, yogurt is stored at low temperature (4°C or 5°C) to limit the post-acidification and improve safety. The fermentation of yogurt is a batch process, but few operations (e.g., treatment, cooling, packaging) are designed and carried out in a continuous and semi-continuous process. Acceptable and good manufacturing practices are followed to maintain control of industrial manufacturers' microbial risk (Corrieu and Beal 2015).

6.3.3.2 Genetic improvement of starter cultures

The Fermentation bioprocess makes use of microbial inoculants to produces a wide range of microbial products that functions as preservatives, stabilizers, flavoring, and coloring agents. Several traditional and molecular-based approaches have been applied to improve microorganisms' metabolic properties and quality found in fermented food.

Traditional approaches

Traditional methods such as mutagenesis and natural gene transfer methods have formed the basis for developing bacterial starter cultures. For instance, mutants are produced by exposure of the microbial strain to UV rays or mutagenic chemicals (e.g., alkylating and deaminating agents) by inducing changes in their genomic sequence in classical mutagenesis. Subsequently, the strains with improved quality are selected based on flavor or ability to resist bacterial viruses. In contrast, gene transfer methods such as transduction, conjugation, and transformation have led to a better understanding of microbial starter cultures' genetics and used for strain improvement (Harlander et al. 1992). Additionally, the introduction of DNA into bacteria through electroporation has revolutionized bacteria's transformation (Chassy et al. 1988; Hisatsune et al. 2016). Recently, Welker et al. (2019) reported that multi-pulse electroporation increases the transformation efficiency in bacteria (e.g., *Lactococcus lactis*) compared to that obtained with single-pulse electroporation (Welker et al. 2019).

Molecular Approaches

The different molecular approaches used to improve starter cultures are genetic modification, characterization, and genomics-based platforms. For example, recombinant DNA technology has been extensively used to modify bacteria, yeast, and mould to either promote, alter, or inactivate the expression of a specific gene (Shinde et al. 2018). To make this approach more successful, specifically in the application of food processing, requires the construction of food-grade cloning vectors with plasmids devoid of antibiotic-resistant marker carrying DNA sequence from generally recognized as a safe organism (Landete et al. 2017). Additionally, the research work on the genetic characterization of microbial organisms has accelerated due to the progress made in information collection of the genome sequence, high-throughput RNA and protein analysis, and high-throughput comparative genome bioinformatics tools. For instance, molecular diagnostic techniques such as polymerase chain reaction (PCR) and fluorescent in situ hybridization (FISH) tools have been extensively used to identify, detect and characterize different microbial strains to improve the fermentation process. Similarly, the application of high-throughput approaches such as the Next-Generation Sequencing (NSG) tool, Loop-Mediated Isothermal Amplification (LAMP), and microbial marker development could be applied to understand food biotechnological processes better. For instance, the techniques above have been used to study the ecological communication between bacterial strains thoroughly, appearance, competition, prevalence, food-microbe, microbe-microbe interaction in the gut, and development of the final product (De Vos et al. 2001). The era of microbial genomics also holds an important place in food biotechnology for the conversion of raw materials to edible form. As microorganisms are an integral part of food-grade ingredients and fermented food due to their interaction within the intestine, functional and comparative genomics of

food-grade organisms are crucial to developing functional and safe food. More recently, functional genomics has been used to regulate the gene expression pattern and interactions based on the availability of the complete genome sequence of a particular organism, such as understanding the response of microorganisms to environmental influences (Parkhill et al. 2000). As we move to proteins from a gene level, proteomics' application has led to the identification and characterization of protein expression and protein-protein interaction in a microorganism to improve the fermentation technology (Parkhill et al. 2000). Similarly, genetic and protein engineering techniques have also been used to increase enzyme production (e.g., chymosin) in the dairy industry to manufacture fermented products such as cheese (Teuber et al. 1993).

6.3.4 Buttermilk

Buttermilk is the liquid phase by-product that is obtained during the processing of butter. It has high nutritional value and is explored as a functional food. When the cream is churned in butter making, it yields 2 streams – butter and buttermilk. Upon churning, the milk fat globule membrane breaks down, and while most of the milk fat goes to butter, phospholipids present on the milk fat globule membrane (MFGM) fragment migrates to the buttermilk. This liquid phase primarily consists of water-soluble components, including lactose, milk proteins, and minerals, making it a nutritionally rich by-product (Morin et al. 2007).

6.3.4.1 Production and Processing

When the cream is churned to produce butter, a stream of buttermilk is obtained. Although ideally, the cream should yield 50% butter and 50% buttermilk, various conditions such as fat content of the cream, churning temperature, churning speed, aging of cream, etc., are responsible for the yield of both products. Therefore, just like any other dairy product, processing parameters are crucial in buttermilk yield as well. Not just the churning but the initial steps in buttermilk processing also affects the composition of the product. Morin et al. (2007) studied the effects of processing such as cream pasteurization and spray drying on the buttermilk composition and microstructure. Cream pasteurization was a pivotal step in determining higher recovery and composition of milk fat globule membranes, whereas spray drying had a pronounced effect on phospholipids content. Buttermilk has applications in various other dairy products, such as yogurt, cheese, fermented beverage, etc. The development of fermented drink based on buttermilk, where barley and fructooligosaccharide were used as a functional ingredient, is an example of buttermilk utilization in manufacturing other products (Sheth and Hirdayani 2016). Incorporating buttermilk in the preparation of fermented milk beverages reduced excessive post-acidification during storage, thus highlighting the acceptability of buttermilk in a

fermented beverage (De Bassi et al. 2012). Using buttermilk in the making of a variety of cheese has been reported by several studies. Poduval and Mistry (1999) prepared Mozzarella cheese using 0.5% fat milk with various combinations of cream and ultrafiltered sweet buttermilk and found alterations in functional properties such as spongy and open protein matrix due to the addition of sweet buttermilk. A similar pizza cheese was also studied for its compositional changes due to the inclusion of unconcentrated sweet cream buttermilk and was found responsible for improved cheese yield (Govindasamy-Lucey et al. 2006). On the other hand, utilizing buttermilk powder in the processing of cheese similar to that of Cheddar was a successful attempt since the process resulted in softer cheese with better quality and potential health benefits (Hickey et al. 2018).

As far as yogurt is concerned, adding buttermilk is responsible for enhancing reduction in undesirable sourness and an increase in odor and flavor, thereby improving the product's overall acceptability (Zhao et al. 2018). Similarly, employing buttermilk powder for the milk fat globule membrane's isolation and then making yogurt indicated functional product development (Le et al. 2011). Apart from being nutritionally rich, buttermilk is also known to have good emulsifying properties. Wong and Kitts (2003) explained how buttermilk solids exhibit antioxidant properties. Therefore, it is a potential ingredient for value addition in stabilizing food matrix against lipid peroxidation reactions (Fig. 6.2).

6.3.4.2 Food Safety

Being an integral part of the food industry, food safety plays a crucial role, especially in dairy products like buttermilk, which are highly susceptible to spoilage. Buttermilk is a perishable product, and therefore, its proper packaging is of utmost importance whether it is meant to be on the shelves of supermarkets or transported to another industry to be used in processing. The food product itself, but the manufacturing facility is also responsible for ensuring food safety. Manufacturing units where food safety tools such as hazard analysis and critical point are not implemented were found to produce cultured buttermilk contaminated with microorganisms such as *Escherichia coli* and coliforms (Paraffin et al. 2019).

6.3.5 Milk

Milk is a significant source of proteins, carbohydrates, vitamins, and minerals (potassium, magnesium, phosphorous, calcium, iodine, sodium). Milk comprises total solids (12–13%) and water (87%). Total solids consist of fat (4%) and solids-not-fat (9%), e.g., proteins, lactose, and minerals and vitamins. The fat amount is the primary factor that categorizes milk drinks into different types. On average, fresh milk has a milk fat of 3.8%, while low-fat milk has a fat of less than 1.5%. Milk is further categorized based on processing such as pasteurization and

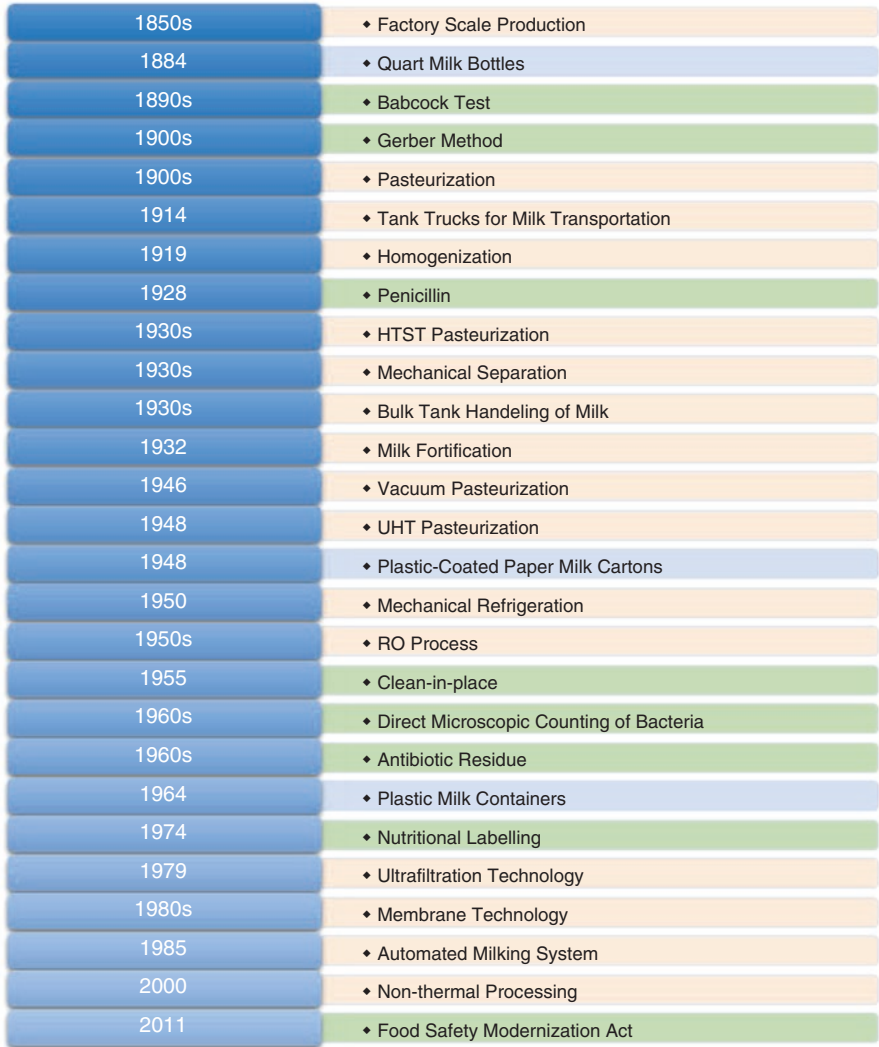


Fig. 6.2 Timeline diagram of major technological advancements of the dairy industry. The pink color indicates production and processing; blue color indicates food packaging; and green color indicates food safety

Foot note:

Production and Processing	Food Packaging	Food Safety
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homogenization. Pasteurized kills any potentially harmful bacteria, while homogenized milk reduces the size of MFGM and uniformly blends to prevent them from rising with cream. Similarly, fortified milk with extra nutrients such as calcium, vitamin D, or omega-3 fatty acids and modified milk like lactose-free milk are also popular products. Furthermore, according to usage, milk has been categorized into Grades A and B. For instance, grade A milk is produced under appropriate sanitary conditions for usage as fluid milk, while grade B milk is acceptable for manufactured products such as certain cheeses and requiring further processing. In contrast, certified milk is produced under exceedingly high sanitary standards and sold at a higher price than Grade A milk.

Besides, specialty milk includes flavored milk, such as chocolate milk with a flavoring syrup. Furthermore, concentrated milk products have varying degrees of water removed from evaporated milk, condensed milk, and dry milk. Biotechnological intervention can add more value to milk by extending the shelf life, applying better packaging, storage facilities, sterilization, and processing procedures. Milk plants process raw milk collected from farms into marketable milk, requiring standard and clean processing procedures. Maintaining aseptic and clean processes directly affect the quality of milk. Furthermore, the application of rapid detection methods would prevent spoilage of milk, thus making sure the safety and quality of milk.

6.3.5.1 Extending Shelf Life (ESL) Milk

Processing technology and packaging material can extend milk's shelf life. Limited shelf life has been a matter of concern for dairy retailers and producers, and investing in technology that can extend the shelf life can enhance the longevity of dairy products allowing milk to be transported longer distances. The three steps to extend the shelf life of milk include:

Reducing bacterial count

Microbial contamination in raw milk occurs from milk-secreting alveoli cells, the udder's exterior surface, and the surface of milking equipment (Guerra et al. 2013), causing undesirable changes affecting the physicochemical properties of milk (Pereda et al. 2007). In contrast, healthy udder, hygienic procedures for milking, aseptic storage, significant, and transportation practices significantly affect milk's microbial quality. Thus, bacterial count in milk should be reduced to minimal to make heat treatment methods more efficient. There are two ways to reduce bacteria: bactofugation and microfiltration.

Bactofugation

This technique separates the bacteria from the milk by centrifugation. The technique is done in an airtight mode, thereby eliminating any bacteria entering into milk throughout the bactofugation process and preventing oxidation to preserve milk quality. Bactofugation reduces the microbial milk load to 89.66% by eliminating pasteurization-resistant microorganisms and psychrotrophic bacteria like *Lysinibacillus fusiformis*, *Bacillus invictae*, *Enterococcus faecalis*, and *Kurthia gibsonii*. Milk psychrotrophics, e.g., Gram-negative (e.g., *Pseudomonas*, *Aeromonas*) and Gram-positive (e.g., *Bacillus*, *Clostridium*), are majorly responsible for milk deterioration (Sørhaug and Stepaniak 1997) and significantly reduce the quality and shelf life of fluid such as milk by producing lipases and proteases. (Dogan and Boor 2003). Bactofugation has proven to be an efficient way to remove spores and vegetative forms of bacteria from milk, thereby prolonging the shelf life by 3–5 days.

Microfiltration

This technique is very efficient in removing bacteria (99.999 %) and preserving the shelf life of milk. In this technique, pressure is used to push the milk through a membrane for the particles to go through it, while majority of bacteria are left behind. More recently, ceramic membranes have been used for microfiltration to reduce the mesophilic microflora to 4 logs and 2 logs, thereby improving the quality of ovine and bovine milk (Panopoulos et al. 2020).

Preventing Re-introduction of Bacteria

After removing bacteria, additional steps should be taken as precautionary measures to avoid recontamination of milk, which can happen during handling and pre-processing activities. Residual milk deposited on the surfaces of milking tools and equipment supports the growth of various microorganisms such as coliforms, pseudomonas, and streptococci (Poghossian et al. 2019, Naing et al. 2019). Additionally, improper refrigeration conditions allow psychrotrophs to multiply rapidly before milk is processed (Paludetti et al. 2018). Thus, to ensure recontamination, a well-designed milk plant with no exposure to air and proper cleaning methods will prevent bacterial growth making it more suitable for preservation.

Preservation

Milk components make it a suitable medium for microorganisms to grow. Pathogenic and spoilage bacteria (e.g., *Escherichia coli*, staphylococcus, streptococci) cause the spoilage of milk by releasing the extracellular and intracellular enzymes, which makes it unfit for consumption. Enzymes (e.g., oxidases, proteases, lipases) cause degradation of milk components, facilitating bacteria's growth (Poghossian et al.

2019). Thus, preserving milk and screening it to be microbe-free in all becomes an essential step. Preserving the milk can be done either by heat treatment (Pasteurization) or chemical treatment.

Pasteurization

Pasteurization's main objective is to eliminate pathogenic bacteria and enzymes that cause spoilage and extend the milk's shelf life. Pasteurization has been classified in four ways:

High-Temperature Short Time (HTST) Method

This is the most popular way of preserving milk. In this process, milk is heated to 72° C (or 161.5°F) for 15 s to kill any pathogenic bacteria. It is also known as Flash pasteurization or continuous flow pasteurization. Milk pasteurized by this method is labeled as HTST milk and has a refrigerated shelf life of two or three weeks.

Ultra-High Temperature (UHT)

In this method, milk is heated at 138 °C (280 °F) for at least 2 s. The UHT pasteurized milk has a refrigerated shelf life of a month and is the fastest and most economical way of preserving milk.

Vat or Batch Pasteurization

It is the longest and expensive method of pasteurization. Temperature controlled closed vats are used in batch pasteurization, with the milk heated to 71° C (160°F) for 30 min, followed by cooling to 4°C (40°F).

Solar Milk Pasteurization

This method is based on the utility of solar intensity as a source of energy for pasteurization. The solar pasteurizing unit consists of a parabolic solar dish, energy conversion system, heat exchanger, and cooling unit. The system can pasteurize milk at 73°C for 30 min and subsequently transferred it to the cooling unit. However, this method is directly dependent on the availability of solar intensity. Development of novel solar pasteurization systems with advancements in higher temperature using photovoltaic integrated concentrator (Meraj et al. 2021) and portable pasteurization plants (Sur et al. 2020) for remote areas can be an efficient alternative for pasteurization.

Chemical Treatment

Chemical preservatives have been added to milk to improve shelf-life; however, they are a challenge to milk safety. Chemicals like hydrogen peroxide, salicylic acid, benzoic acid, mercuric chloride, and formalin. These preservatives are bactericidal and have a health/environmental hazard, including unsuitable for storage. Studies conducted to preserve raw milk using hydrogen peroxide (H₂O₂) showed an absence of coliform (e.g., *Escherichia coli*) growth in the treated milk and was found safe for human consumption. Furthermore, the application of transglutaminase enhanced the micellar stability in skim milk on the addition of ethylene diamine tetraacetic acid (EDTA) and did not affect the dynamics between salts and serum phase. Additionally, the use of thyme-free nano emulsified essential oil is also known for inhibiting microbes' growth in milk (Jemma et al. 2017).

Proper Packaging

The longer shelf life of dairy products is achieved through packaging in a sealed and sterilized environment. Milk can be packed into different types of packages such as cartons, glass, pouches, Polyethylene terephthalate (PET) bottles, C-enamelled tinfoil cans, etc. The introduction of different plastic materials, combined with paper, was developed for packaging dairy products. Petrus et al. 2010 studied storage temperature evaluation on pasteurized milk with different packaging materials like polyethylene (PE) monolayers specked with titanium dioxide (TiO₂). The study showed better storage of pasteurized milk when high-density polyethylene (PE-HD) bottles were used compared to low-density polyethylene (PE-LD) pouches. Similarly, Polyethylene terephthalate (PET), another packaging material, has shown better results by protecting milk from light-induced lipid oxidation (Kontominas 2010; Van Aardt et al. 2001).

The growing demand for safe and fresh products has led to the development of technologies that assure food quality and improve shelf life through cold chain distribution (Rysstad and Kolstad 2006). The enhanced shelf life of dairy products depends on the biochemical interactions between dairy products, packaging, and the existing environment (Haghighi-Manesh, 2017). Therefore, modern food preservation methods must ensure packaging to enhance the shelf life, fresh characteristics, and quality of food. Newer technologies used in packaging include nanotechnology, modified atmosphere, active and intelligent/smart packaging. The active form of packaging has been introduced to the market to enhance products' safety and quality. It is an innovative food packaging system that includes additives or agents that absorb oxygen, moisture, carbon dioxide, ethylene, flavors, and odours or release carbon dioxide, antimicrobial agents, and antioxidants flavours (Wong and Goddard 2014). Some popular active packaging systems include moisture-scavenging (desiccation), oxygen-scavenging, carbon dioxide-absorbing, and antimicrobial systems (Jalilzadeh et al. 2015). Furthermore, non-thermal food preservation technologies

such as pulsed electric fields (PEF) and high-pressure technology (HPP) are also useful in extending the shelf life of the pasteurized product (Abida et al. 2014).

Novel Preservation Technologies

The use of increasing temperatures in pasteurization impacts the taste of milk and is not sufficient enough as an extended shelf life product. Different technologies have been adopted by milk plants to extend the shelf life without impacting the taste of pasteurized milk. New technologies include non-thermal technologies like ultrasound, irradiation, and ultraviolet, while thermal technologies include microwave, ohmic heating, and radiofrequency pasteurization methods. Similarly, microwave and radiation processing have also evolved as novel pasteurization techniques. These new technologies exhibit great potential in maintaining the pasteurized milk product's shelf life and nutritional properties.

Ultrasound Technology

In this technique, low-frequency ultrasound (power ultrasound) waves are applied to milk, causing vibration of molecules and the generation of bubbles, which disrupts the cells, leading to inactivation of microorganisms. Application of low-frequency ultrasound waves (20–100 kHz) generates micro-currents of implosion and explosion, causing physical damage to cell membrane and breakage of cell walls (Carcel et al. 1998). These waves have a lethal effect on cells, thereby inactivating the pathogenic and spoilage bacteria (Bermudez-Aguirre et al. 2009). A study reported by Noci et al. (2009) thermo-sonication of milk and the pulsed electric field application enhanced the inactivation of *Listeria innocua* in milk significantly. This technology renders shorter processing time and imparts significant enhancements to final milk quality after processing. For instance, the milk color turns whiter due to fat globules' homogenization (Bermudez-Aguirre et al. 2009).

Irradiation Technology

Irradiation (transfer of energy) in milk is used to inactivate pathogenic microorganisms (Osaili et al. 2008). Microorganisms like Salmonella, Listeria, and *Escherichia coli* have been inactivated with irradiation technology (Crawford and Ruff 1996). The ionizing radiation includes gamma rays, X-rays, and electron beams (Satin 1996) and generates secondary electrons causing ionization, dissociation, and excitation. The transfer of energy kills the pathogenic bacteria; however, changes the physical and chemical properties of milk components. For example, in a study, lower vitamin A and carotene contents were observed on the application of gamma-irradiation with thermal treatment to pasteurized milk ((Naghmouh et al. 1983).

Ohmic Heating

Ohmic heating (electro-pasteurization) involves the application of electrical current to pasteurize milk. The technique of Ohmic heating is based on electroporation with pore-forming mechanisms leading to the inactivation of cells. The current generation generates heat of low frequency (50–60 Hz), mainly responsible for bacterial death. Nevertheless, some studies showed permanent membrane damage on the application of high voltage discharges, while the killing with lower voltages needs further investigation. As a novel technology, there are few studies reported on electro-pasteurization of milk. In a study reported by Sun et al. 2008, ohmic heating pasteurization inactivated *Streptococcus thermophilus* faster than thermal treatment processing.

6.3.6 Infant Formulas

Human milk is a complete food for the development, nourishment, and protection of infants. The mother's milk's unavailability has driven a new platform for manufacturing artificial formulas to nurture developing infants. Although inferior to natural milk, these formulas are designed to provide early nutrition to infants and long-lasting metabolic influence during their life (Thompson et al. 2012). The most commonly used breast milk substitute is cow or soya milk, which is marketed as ready-to-feed, liquid, and powder forms. The specialized formulas depending on the infant's health conditions (e.g., gastrointestinal disorders, allergy), are also designed and mentioned as "safe to use" (Green corkins et al. 2016). The government closely monitors infant formulas' composition, and the manufacturers should adhere to strict guidelines to maintain nutrients' self-life (Martin et al. 2016). The recent advances in biotechnology have opened up numerous unique food industry opportunities to produce formulas with bioactives, targeting the infant's gut, nutritional, and functional composition to simulate human milk. For example, the production of recombinant milk proteins (e.g., lactoferrin) through genetic engineering techniques to enhances the nutritional value of synthetic infant formulas and human milk fortifiers have improved the diet and feeding of very-low-birth-weight and premature infants (Gaul 2000; Hartmann et al. 2008; Aggarwal and Aggarwal 2020). Notably, a functional ingredient such as human milk oligosaccharides, long-chain polyunsaturated fatty acids (PUFA), nucleotides, probiotic bacteria, synbiotics, and fat globule membrane is being incorporated to humanize infant formula to provide nutrition, including enhancing immunity, microbiome, and cognitive development (Pande and Akoh 2016; Ahern et al. 2019;). Formulas with prebiotics and probiotics could generate epigenetic modification by modulating gut microbiota's composition and activity (Lemaire et al. 2018).

Human milk is dynamic in composition, especially during the initial six months of lactation compared to the static composition of infant formulas, making it noteworthy to prepare age-based formulas to meet the infant's desirable nutritional

requirement of the infant (Lönnerdal et al. 2016; Spalinger et al. 2017). Additionally, during formula preparation, it is essential to maintain a hygienic condition and acceptable manufacturing practices to avoid microbial contamination and adulteration due to the neonates' fragile gut and immature immune systems. In the future, efforts should be directed to develop fast and effective technologies to increase the prepared formulas' microbiological safety, such as bacteriophage therapy, as natural weapons to control pathogens in the food industry (Gutierrez et al. 2016). Similarly, natural antimicrobial (e.g., lactic acid) has effectively prevented microbial spoilage (Al-Holy et al. 2010, 2015; Yemiş G. P and Delaquis P 2020). Furthermore, strategies to increase milk's shelf life and reduce microbial spoilage without hampering the bioactive milk components have been employed through different techniques such as microfiltration, ultrafiltration, microfiltration, and reverse osmosis (Ahern et al. 2019). The use of partially fermented formulas has recently gained momentum in restoring infant health with components derived from a microorganism such as microbial cells, metabolites (postbiotics), and non-viable cells (parabiotics) (Wegh et al. 2019; Salminen et al. 2020). These postbiotics create an appropriate environment for the proper colonization and development of intestinal microbiota, vital for the infant's future well-being. Recent studies have also supported the application of postbiotics as suitable and safest alternative agents than other 'biotics' to improve infant's health with fewer challenges to product storage and shelf life (Deshpande et al. 2018; Rad et al. 2020). Most importantly, these unique features make postbiotics a promising tool for food, biotechnology, and the pharmaceutical industry to promote health benefits as a functional food (Cuevas-González et al. 2020).

The new era of 'biotic' research (e.g., postbiotics, parabiotics) has been increasingly used for the development of biotechnology-related products (e.g., Bactistatin®, Cytoflora®, Lacteol® Fort) with functional elements (e.g., metabolites, cell lysate, non-viable cells) for the nutraceutical industry. The evolving concept of postbiotics brings food, microbiology, and personalized therapies closer (Olle et al. 2013). A particular focus on the impact of host-gut microbe interaction on potential benefits and risk factors is: essential. Therefore, future studies based on bioinformatics, epigenetic and omics-based approaches (e.g., metatranscriptomics, metabolomics, lipidomics) will provide an insight to understand better the complex interplay between postbiotics, host biology, and the gut microbiome, including measurement of the composition and functions of gut microbiota (Wegh et al. 2019). Despite these biotics' tremendous positive impact, researchers are looking forward to developing novel bioengineered postbiotics with desirable features (Cuevas-González et al. 2020). Interestingly, the postbiotics could be tailored towards the prevention and treatment of pandemic disease such as SARS-CoV-2 as the changes in structure and metabolic activity of microbiota residing in the intestine from a healthy state may lead to the occurrence of a microbial biomarker for the prediction of coronavirus (Gou et al. 2020). Taken together, such progress may pave the route to reproduce future infant formula mimicking human milk. Of note, postbiotics' mechanism and signaling pathways need further studies in promoting health, including their usage in infant-based formulas.

6.4 Biotechnology in Food Diagnosis and Safety

Milk production has tremendously increased in India, with sufficient quantities available to prepare different dairy products with acceptable microbial load and quality. For human consumption, the safety of milk and processed products (e.g., yogurt) are ensured through the use of different molecular techniques such as gene probing and immunological assays (e.g., monoclonal and polyclonal antibodies) to detect the foodborne pathogens (Sundarraaj et al. 2018). The characterization of individual pathogens and their products is crucial to identify the pathogen source involved in diseases. The application of molecular techniques provides valuable information to understand how pathogens pass through the food chain and cause infections (Macori and Cotter 2018). For example, molecular techniques such as Polymerase Chain Reaction (PCR), an extremely versatile, sensitive, and reliable technique, have been extensively used to rapidly detect, identify, and determine pathogens. Additionally, they have been extensively used in DNA cloning, genetic fingerprinting, and detection and diagnosis of infectious diseases. Through this technique, the 16S rRNA gene of bacteria is amplified using universal or specific primers, followed by sequencing to identify not known or novel bacteria species (Magistrado et al. 2001). Furthermore, modified PCR technique such as Multiplex PCR uses multiple primers within the one PCR mixture in order to rapidly detect, identify and differentiate various bacterial species and strains (Shi et al. 2010), leading to amplification, more than one target sequence in a reaction and for different DNA sequences.

Similarly, modified PCR technique such as real-time PCR (qPCR) amplifies and quantifies the target DNA simultaneously within a reaction. It uses a specified primer set, one or two probes, or fluorescent dye for improving the detection signals (Dhanasekaran et al. 2010). As the reaction progresses, the amplified DNA is detected rather than towards the end and shortens detection time compared to standard PCR, including resolving the relative or an absolute number of bacteria across various samples (Shi et al. 2010; Agrimonti C et al. 2019). This technique has also been used to detect *Enterobacter* in the infant formula and *Clostridium* spore quantification in dairy products (Liu et al. 2006; Lopez-Brea 2018). It even encompasses the capability for quantifying target organisms in complex matrices and hence is regarded as a promising tool in improving the safety and quality of food (Martínez et al. 2011). Various studies have recently studied the utilization of droplet digital polymerase chain reaction (ddPCR) that involves eight individual TaqMan™ reactions for the simultaneous detection of, without selective enrichment of *Listeria* spp, *L. monocytogenes*, *Salmonella* spp., and *Campylobacter* spp. in cheese. The ddPCR is a third-generation PCR that quantifies target DNA without standard curve and higher sensitivity than qPCR (Cremonesi et al. 2016; Pan et al. 2020). Additionally, High Resolution Melting quantitative-PCR (HRM-qPCR), a development of PCR technique, has been implemented to identify foodborne pathogens (Forghani et al. 2015; Agrimonti C et al., 2019). In contrast, in the reverse-transcription PCR, the

target RNA is reverse transcribed to complementary DNA (cDNA) to detect only viable pathogenic cells (Choi S.H. and Lee S. B, 2011).

The molecular typing methodologies such as restriction fragment length polymorphism (RFLP), random amplified polymorphic DNA (RAPD), and pulsed-field gel electrophoresis (PFGE) has been used to monitor, characterize, detect, and diagnose spoilage flora (Tabit F T 2016; Maryam et al. 2017). The PFGE is classified as a third-generation technique and is considered the 'gold standard' typing method by many researchers to investigate foodborne pathogens and other epidemiological studies (Alonso R et al., 2005; Parizad EG et al., 2016; Neoh HM et al., 2019). It enables research on the entire food supply chain and also traces, including estimating the strain responsible for food contamination, from animal raising to foodborne illness in a human. The PFGF is applied to correlate the strain isolated from different sites to resolve the relationship between strains (Frasao, B. D. S et al. 2017). The RAPD-based method is used for rapid detection of target DNA using arbitrary primers (typically 10-mer primers) under low-stringency PCR conditions (Wassenaar T.M. and Newell D. G, 2000). Furthermore, RFLP uses variation in homologous DNA sequences for bacteria's characterization (Foley SL et al., 2009; Adzitey F et al., 2013). Recently, biosensor-based nanomaterial has been gaining interest to detect allergenic and toxicant compounds and food authentication (Gomez-Arribas et al. 2018; Nikoleli et al. 2018). The application of such approaches and methods has significantly improved human health and hygiene (Meshram et al. 2018; Patel et al. 2016). Newer techniques such as high-resolution melting HRM-qPCR, ddPCR, Loop-mediated isothermal amplification (LAMP), and Next-generation sequencing (NGS) have been used for food labeling and authentication due to their speed, specificity, sensitivity, and multiplexing (Bohme et al., 2019). Notably, the applications of culture-independent nucleic acid-based food microbiology techniques have solved newer emerging problems associated with culture-dependent methodologies. Furthermore, the application of foodomics (e.g., metagenomics, metatranscriptomics) has been extensively used to analyze genes linked with characteristics properties and functionalities such as flavor formation, taste, and probiotic activity, including understanding the behavior of microbes in the food ecosystem (Ceuppens et al. 2014). The membrane-based technologies that are non-thermal environment-friendly, greener, and energy-saving technologies have gained broader applications in the food and dairy sector (Dhinesh Kumar et al., 2017). Different types of membranes, such as microfiltration, ultrafiltration, nanofiltration, and reverse osmosis, have been used to enhance shelf life, yield, texture, compositional properties, and quality of milk products (Kumar et al., 2013). Therefore, it is essential to continuously develop superior membranes that could be extensively used to improve other dairy products for human consumption (Kumar et al., 2013). From the burgeoning data, it is evident to acknowledge biotechnology's immense contribution to reshaping the dairy industry. However, we envision and confident that newer technologies will be developed to improve the quality, nutrition, and safety of different dairy products to impact human health and introduce newer products in the market.

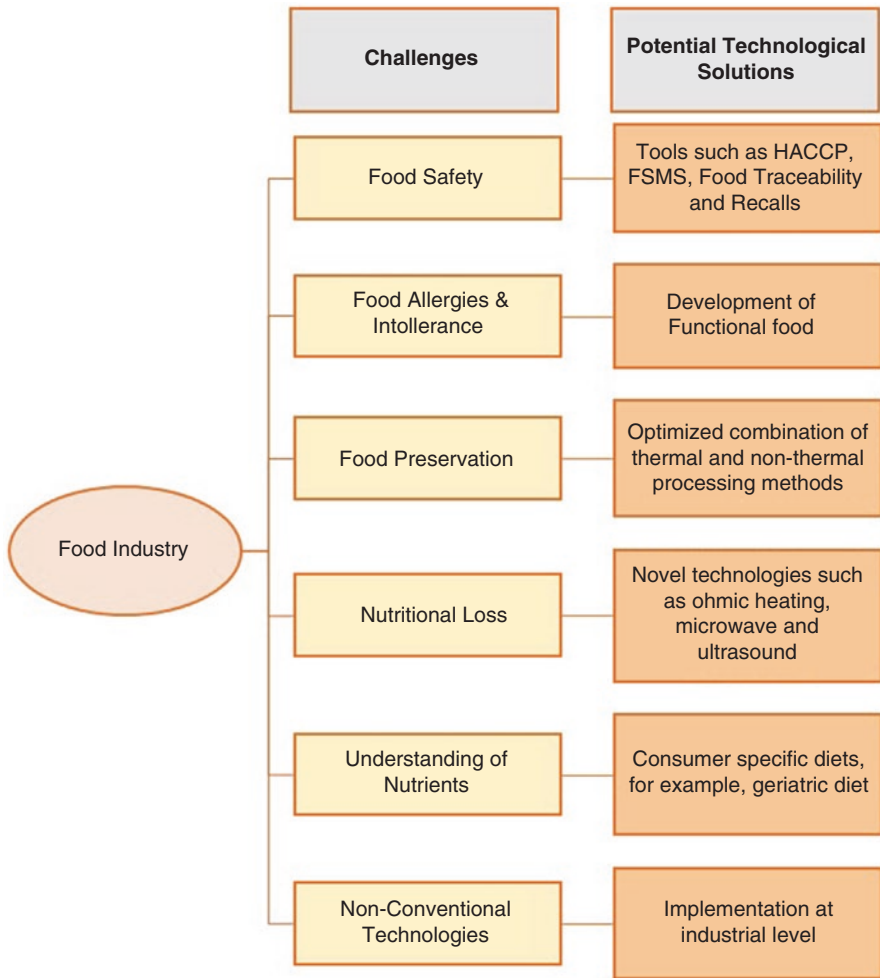


Fig. 6.3 Overview of challenges and potential technological solutions. The present challenges of the food industry could be resolved by respective technological solutions, as presented. HACCP: Hazard analysis, critical control point. FSMS: Food safety management system

6.5 Challenges in Application of Technologies in the Food Industry

Food is an indispensable requirement for life, and therefore, the food industry is always in demand. The food industry has evolved from using fire for cooking to thermal processing, sprinkling salt for preservation to brine, sun drying to solar drying, the focus has shifted from merely satiating the hunger to safe food consumption. Nevertheless, the food industry has been experiencing multiple challenges due

to the expanding market, growing competition, volatile demand, and changing trends. Food technologists and researchers worldwide are trying to solve these challenges and have come a long way. Some of the significant challenges are listed as follows and presented in Fig. 6.3.

6.5.1 Food safety

Food safety is an assurance that the food has been prepared, handled, and stored in a way to avoid any food-borne illness and suitable for consumption. The concerns about safe food consumption have been there since the advent of factory-level food production. Although processing technologies such as pasteurization were implemented at the commercial level in the late nineteenth and early twentieth century, food safety has remained a significant concern even today primarily because of the development of new products, and knowledge about microorganisms keeps on emerging. Various tools and techniques in assuring food safety have focused on detecting, monitoring, and preventing physical, chemical, and biological hazards. For instance, hazard analysis critical control point (HACCP) is a systematic approach to identify, evaluate, and control food safety hazards (FDA HACCP Guidelines, 2017). From heating, chilling, drying, and fermentation, and HACCP can be successfully implemented in all cases. Food traceability and recall, the capability to follow the development of food at any stage, and the action of removing food at any stage from the market, respectively, are other fundamental approaches in managing risk as a response to food safety events (FAO, Traceability and Recalls). Food scares are the repercussions of pandemics and public health emergencies as they cause safety concerns among consumers. Although there is no evidence of COVID-19 transmission by food, the pandemic has raised concern about safe food consumption among consumers and increased attention on zoonotic disease, antimicrobial resistance, and food fraud among scientists (FAO, Food Safety and Quality, 2020).

6.5.2 Food Allergies and Intolerances

Allergens and intolerances are a significant challenge for the dairy industry and the whole food industry. The main concern associated with food allergies is the industry's limited knowledge of dealing with them, mainly to avoid cross-contamination (Augusto 2020). Cow milk allergy arising from a particular immune response is undoubtedly the most common type of milk allergy that affects non-exclusively breastfed infants (Vandenplas 2017). Another dairy-associated allergy is casein sensitivity, where the body mistakenly produces antibodies to fight against the milk protein. The third most common dairy intolerance is lactose intolerance, which is often wrongly referred to as an allergen. Lactose intolerance is associated with the

inability to digest milk sugar and lactose. Avoiding milk intake due to allergies and intolerances is undoubtedly not the solution since it may lead to adverse health effects such as nutritional rickets, low bone mineral density, and increased fracture risk (Heine et al. 2017). Therefore, functional food development is a potential solution; for example, fermented dairy products consisting of probiotics reduce lactose intolerance symptoms (Hasan et al. 2014).

6.5.3 Food preservation

It is the technique of preserving food from physical, chemical, biological, and biochemical spoilage. The main objectives of food preservation include, but not limited to, are making food safe for consumption, enhancing its shelf life, improving texture, flavor, and consistency, and increasing its acceptability. The age-old practice of drying, curing, pickling, fermentation, and newer technologies of canning, irradiation, and freezing are all food preservation forms. Despite food preservation being practiced for long, various gaps still exist, such as a lack of thorough understanding of microbial and enzymatic inactivation. Thermal inactivation of microbial spores, which forms the basis of calculating thermal preservation processes' safety, rarely follows first-order kinetics (Peleg and Cole 1998). Another case is enzymatic inactivation in commercially available coconut water. The drink's thermal processing damages its flavor, while non-thermal treatment such as high-pressure processing cannot inactivate the peroxidase and polyphenol oxidase, resulting in pink coloration. Therefore, an optimized combination of thermal and non-thermal processing methods could be a potential solution (Augusto 2020).

6.5.4 Nutritional Loss

Over-processing food items lead to the nutritional loss. Conventional thermal processing such as pasteurization, sterilization, and canning are widely used techniques to control microbial spoilage that heavily contributes to the loss of nutrients, including loss of color, texture, and appearance of the food products. There are two approaches to avoid nutritional loss during processing and/or enhancing food's nutrient content. Firstly, through a process known as food enrichment, lost nutrients during processing are added to the food that comes with an added cost and is not feasible in every case. However, a more reliable long-term solution is in novel thermal techniques like ohmic heating, microwave, non-thermal technologies, for example, irradiation and high-pressure processing, and pre-treatments like an ultrasound for food drying. Novel thermal processing, for example, ohmic heating, microwave, radio-frequency heating, is efficient in minimizing nutrient and quality loss due to heating (Ling et al. 2015). Employing an electron beam for pasteurization is another advancement in non-thermal milk processing affecting raw milk's

nutrient and aroma profile. Rojas et al. (2019) studied ultrasound technology for incorporating microencapsulated nutrients such as iron and carotenoids during pre-treatments of food drying. Surprisingly, it has been reported that thermal treatments responsible for the loss of natural antioxidants in tomato and coffee derivatives also aids in the formation of antioxidants due to Maillard reaction products (Nicoli et al. 1997).

6.5.5 Better Understanding of Nutrients

Although food safety and quality are often the prime focus while developing a new product or improving existing ones, its nutritional requirement throughout the processing is still overlooked. Also, it is equally crucial to understand the presence of nutrients in food and bioavailability. Verkempinck et al. (2020) reported how various food processing methods could enhance nutrient accessibility. For example, non-thermal techniques like high-pressure homogenization and ultrasound can be used to improve *in vitro* carotenoids. Similarly, the physicochemical properties of food matrices such as hardness, compactness, elasticity responsible for nutrient absorption of dairy products like yogurt (semi-solid), milk (liquid), and cheese (solid) are significantly affected (Fardet et al. 2019). Another essential aspect to understand is a customer requirement, which has to evolve due to consumer awareness, purchasing power, health issues, personal choices, etc. For instance, the nutrient requirement of the elderly is entirely different from a young person as it affects their mental, physical and social health (Kaur et al. 2019). Whey protein consists of β -lactoglobulin, bovine serum albumin, lactoferrin, α -lactalbumin are nutritionally rich and confer benefit in geriatric nutrition (Sreeja et al. 2013). Therefore, food technologists must work closely with nutrition scientists for wholesome development.

6.5.6 Potential of Non-conventional Technologies

Non-conventional technologies such as ultrasound, ohmic heating, pulsed electric fields, etc. offer great potential, as they are relatively environmentally friendly, economical, and consume less energy for operations. Nonetheless, these food-processing methods are used in industrial production. Although the concrete reasons behind this are unclear, ohmic heating has been studied for its application in improving the fermentation process to develop the probiotic dairy products like yogurts, fermented milk, and dairy-based drinks (Pereira et al. 2018). Similarly, non-conventional cell disruption technologies (e.g., high-pressure homogenization, microwave-assisted extraction, pulsed electric fields) have been reported to offers sustainable chemistry solutions in the extraction of food additives (Barba et al. 2015).

6.6 Socioeconomic Constraints and Potential Ways to Overcome Them

The local government's role in developing core sectors like food and agriculture cannot be neglected. The policies and regulations must be drafted to promote biotechnological developments in the food and dairy industry. The government's support in the form of favorable, flexible policies plays a crucial role in encouraging technological advancements. At the same time, it is necessary to ensure that the developed policies follow the global system. The flexibility of trade flow plays a crucial role in determining economic progress. Fostering startups and micro, small and medium enterprises could prove beneficial in the long run, especially when these businesses could quickly be promoted by providing tax incentives, elasticity in loans, etc. Since the developing nations have limited resources, they need to identify the key areas and prioritize their technological development work plan. Food safety is one such domain that should be the prime concern for every country, however, the extent and approach to deal with this issue could vary.

Food security is still an obstacle yet to overcome for many low-income countries. In most of these cases, agricultural production is sufficient to meet the population's demand. However, due to the poor food supply chain, the food wastage while reaching from farm to fork is countless. Another major drawback in the food supply chain is the proper maintenance of the cold chain due to disrupted power supply. Therefore, a consistent power supply needs to be achieved. There is a strong need to boost intellectual property rights since most indigenous products' traditional knowledge is passed through cultural channels. Disseminating information regarding patents and other intellectual property rights would promote and inculcate a sense of motivation among researchers and scientists. Another crucial aspect that is often overlooked while drafting policies is the consumer's needs, which could be achieved through effective communication between various stakeholders. The information and opinions must be shared between consumers and food industry stakeholders. Public participation in decision-making is crucial for implementing new biotechnological applications and working towards safe food consumption.

6.7 Conclusion

Biotechnological applications offer the food industry potential solutions to food processing, production, packaging, and safety challenges. The age-old pasteurization process is still relevant with certain modifications and finds its application in almost all dairy products. Notably, technologies such as biobased packaging material are prospective solutions towards a greener and sustainable environment. The utilization of dairy industry by-products such as whey and buttermilk are economical and confer health benefits.

Presently, the food industry faces multiple challenges covering various aspects of the industry, including food safety, allergy and intolerances, preservation, nutritional loss, and unexplored potential of non-conventional. Technology advances with time, and it is worthwhile to envision existing technologies to evolve based on requirements and demand. For instance, recent technologies such as non-thermal food processing need to find their way from the laboratory scale to the industry. This will further help meet various challenges that the food industry is facing today. Technology alone cannot be implemented without the support of policies and regulations. With evolving technology, the regulatory structures have to evolve to support the widespread adoption and rollout of the technology in the food industry. Technology, policy, and regulatory infrastructure should sync and complement each other to develop the food market. The ability of biotechnological applications in various food industry domains has enormous potential and further needs proper implementation.

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Chapter 7

On-Farm Point-of-Care Diagnostic Technologies for Monitoring Health, Welfare, and Performance in Livestock Production Systems



Mohamed Zeineldin, Ahmed A. Elolimy, P. Ravi Kanth Reddy, Mohamed Abdelmegeid, Miguel Mellado, Mona M. M. Y. Elghandour, and Abdelfattah Z. M. Salem

Abstract The occurrence of infectious diseases has a significant adverse effect on livestock health and production efficiency. Current diagnostic approaches in veterinary practice are primarily focused on the observation of changes in physical, clinical, behavioral, or performance of individual or groups of animals. In recent years, these diagnostic approaches have markedly improved livestock profitability during their production cycle. This is mainly achieved by using reliable and conveniently available on-farm and point-of-care diagnostic technologies for the rapid and accurate management of animal health. The availability of on-farm and point-of-care technology is rapidly changing decisions for bovine practitioners on both individual sick cows and herd health level. Early detection of infectious diseases using quick,

M. Zeineldin

Department of Animal Medicine, College of Veterinary Medicine, Benha University, Benha, Egypt

A. A. Elolimy

Department of Animal Production, National Research Centre, Giza, Egypt

P. R. K. Reddy

Veterinary Dispensary, Taticherla, Andhra Pradesh, India

M. Abdelmegeid

Department of Animal Medicine, College of Veterinary Medicine, Kafr-Elsheikh University, Kafrelsheikh, Egypt

M. Mellado

Department of Animal Nutrition, Autonomous Agrarian University Antonio Narro, Saltillo, Mexico

M. M. M. Y. Elghandour · A. Z. M. Salem (✉)

Facultad de Medicina Veterinaria y Zootecnia, Universidad Autonoma del Estado de Mexico, Toluca, Mexico

e-mail: salem@uamex.mx

effective, low-cost, automated technologies will allow timely detection of infected animals, thus reducing the economic loss and associated abuse of antimicrobial therapy. Here we review the currently available on-farm and point-of-care diagnostic technologies for health surveillance and disease detection in livestock production systems. We additionally review the advantages and disadvantages of each methodology, concerning their possible effect on the improvement of animal welfare and productivity of farm animals.

Keywords Bovine · Disease · Diagnosis · On-farm · Point-of-care · Technology

7.1 Introduction

Infectious diseases are a major source of livestock production inefficiency and cause significant economic losses to the global livestock industry due to associated morbidity, mortality, and treatment costs (Craft 2015). The presence of infectious agents, and related therapeutic and antibiotic residues, has serious consequences for the public perception of livestock production, food safety, and human health (Daszak et al. 2000). Although disease control and prevention are critical elements for mitigating the effect of disease on animal health and production efficiency, disease-free production systems are unrealistic (Schwabe 1982). Since diseases are an unavoidable component of modern livestock production systems, on-farm and point-of-care diagnostics for the accurate detection and diagnosis of affected animals are vital for minimizing morbidity, optimizing recovery, and maintaining profitability. Early and accurate disease detection plays an important role in reducing mortality, preventing disease transmission, and avoiding the long-term effects of disease on production, welfare, and profitability (Bisson et al. 2015). Early detection and diagnosis of infectious diseases in livestock also help to avoid irreversible pathologies, prevent the production of antibiotic-resistant bacteria, reduce public health concerns and optimize the action of antimicrobial therapy (Hennessy and Wolf 2018).

Compared to human medicine, current diagnostic approaches in veterinary practice have primarily focused on observation of changes in the physical, clinical, behavioral, or performance status of individual or groups of animals. Although these findings have historically been made by animal caretakers and animal health professionals, there has been an increased interest in the use of cow-side diagnostic tests to improve the sensitivity, precision, and timeliness of disease detection (Berckmans 2014a). The investment in rapid, selective, accurate, and cost-effective tests and the recognition among clinicians and animal stakeholders of the value of quick and accurate diagnosis of disease have resulted in rapid progress in point-of-care and on-farm diagnostics for rapid on-site testing of animals (Helwatkar et al. 2014). Similarly, technology adaptation and implementation are being used to

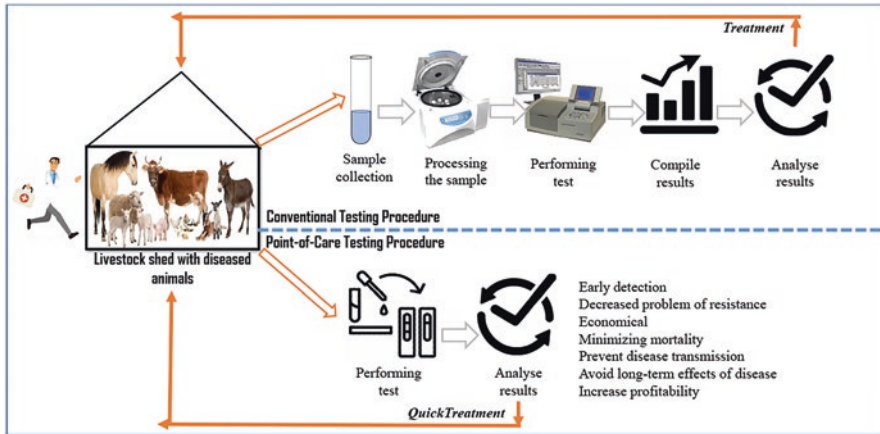


Fig. 7.1 Advantages of point-of-care diagnostic techniques compared to conventional diagnostic methods

create more robust systems for measuring animal growth, production, and the physical environment of animals, all to improve individual or group performance and production efficiency (Lokhorst and Ipema 2010). To date, most of the available diagnostic tests in humans and farm animals focus on the rapid identification of non-infectious conditions (e.g. blood glucose, β -hydroxybutyrate, and hemoglobin concentration, diagnosis of cardiac markers, detection of blood gas and electrolytes, etc.) (Tschandl et al. 2019; Sargeant and O'Connor 2020). On-farm and point-of-care diagnostics also provide an alternative for monitoring livestock health, which has a high potential for increasing the performance of livestock production systems, eliminating subjectivity the management decision-making, and maximizing the efficacy of unskilled or inexperienced labor in animal health management (Helwatkar et al. 2014). Advantages of point-of-care diagnostic techniques compared to conventional diagnostic methods are shown in Fig. 7.1.

In this chapter, we review the currently available on-farm and point-of-care diagnostic technologies for health and performance monitoring in livestock production systems. Additionally, we review the advantages and disadvantages of each approach, in particular regarding its possible effect for improving animal health and productivity.

7.2 Characteristics of an Ideal Point-of-Care and On-Farm Diagnostic Technologies

The list of available point-of-care and on-site diagnostic tests in both human and animal medicine has expanded steadily over the last few years. The global diagnostic market for veterinary diagnostic testing is projected to increase at an average

annual growth rate of 8.6% from 2016 to 2021, to hit US\$ 6.71 billion by 2021 (Commission 2011). This massive market potential is driven by the availability of quick, portable veterinary diagnostic devices to track animal health disorders (Helwatkar et al. 2014). Ideal technology should be developed as durable, low-cost, reliable, and effective explaining the intrinsic biological process (Frost et al. 2003). Moreover, it should be easily converted into a professionally tested and easy-to-maintain relevant action (Bolboacă 2019). The selection of the appropriate point-of-care and on-farm diagnostic technology is affected by many factors, including the frequency and type of measurement, the labor needed to conduct the test, the conditions of the livestock management system, and the costs of the technology (Berckmans 2014a). The idealized diagnostic technologies are challenged by the sample volumes of the complex structure of biological materials, the processing time of the sampling, and the diagnostic capacity and accuracy of the instrument (Gubala et al. 2012). The design of an ideal point-of-care and on-farm diagnostic should therefore depend not only on the development of technology but also on the design of the biological measurement goals and the surrounding livestock farming system.

7.3 Categories of Point-of-Care and On-Farm Diagnostic Technologies

Successful management of animal diseases depends on the timely identification of clinically infected animals through laboratory and clinical examination, along with an understanding of the factors influencing the epidemiology of the causative agent (Bisson et al. 2015). There is a wide variety of easier and more cost-effective diagnostic techniques, including point-of-care technologies for disease surveillance in both human medicine and veterinary practice (Helwatkar et al. 2014; Yanase and Triantaphyllou 2019; Mohr et al. 2020). Generally, each diagnostic tool measures a specific parameter related to the physiological state of the animal, performance or existence of infectious agents in an individual or group of animals, a change that entails an intervention to alleviate the disorder detected (Lokhorst and Ipema 2010). Besides, there may be relevant algorithms and computer programs necessary to understand an animal's health status, which may need to be coupled with other data (e.g. financial input) before an appropriate and effective decision can be taken (Das et al. 2015). This obstacle can be better tackled by the advancement of point-of-care and on-farm automated technology in a revolutionary way to provide fast and effective identification of animal health threats (Neethirajan et al. 2017). In this context, point-of-care diagnostics are valuable tools for detecting small sample volumes and low concentrations of biological components and infectious agents (Theurer et al. 2013). Practically, all the available technologies for the livestock industry fall into one of four broad categories related to their proximity (attached or non-attached) and association (invasive or non-invasive) with the animals themselves (Helwatkar

et al. 2014). Attached technologies include those that can be fixed to the outside the animal's body (non-invasive) (Krieger et al. 2019) or those that are fitted inside the body (invasive) (Rose-Dye et al. 2011). This category includes some of the most reliable technologies for continuously monitoring animal health throughout the day and comprises accelerometers, pedometers, vibration sensors, thermometers, and rumen temperature bolus. Non-attached technologies are those that animals pass by, over, or through for health monitoring (Stone et al. 2017). These devices are often set at fixed locations in the animal's environment (e.g. surveillance cameras (regular or thermographic), and video and audio recording systems). Similarly, point-of-care diagnostic technologies are also achieved with the use of unattached, transportable, and hand-held equipment and test kits to obtain blood samples and provide results in a very short time, so that decisions are taken very rapidly.

7.4 Molecular Diagnostic Technologies for Monitoring Animal Health and Disease

Traditionally, diagnostic approaches in veterinary practice have focused primarily on clinical evaluation of the behavior of individual animals or groups of individuals and the reaction to certain conditions of the disease (Van Veen 1997). These conventional approaches rely on animal history and visual clinical signs and can only be used to diagnose a small percentage of sick animals. Because of the multifactorial disease control and the wide variety and complexity of the livestock farming environment, these conventional methods are not sufficient to improve the sensitivity, precision, and timeliness of disease diagnosis (Saegerman et al. 2011). The usefulness of molecular and high-performance sequencing technologies for tracking the health and disease of livestock has been highlighted for accurate disease control (Reuter et al. 2015; Kumar et al. 2019). The recent application of molecular diagnostic approaches to livestock has shown the importance of these platforms for improving livestock management systems and limiting the spread of diseases (Shirley et al. 2010). Also, the existing use of molecular diagnostic techniques has demonstrated the complexity of disease-causing agents (Zeineldin et al. 2017a, b) and proven associations between early disease detection and livestock health and productivity (Naqvi 2007). Numerous molecular-based techniques are presently used to determine disease-causing agents, including immunohistochemistry, fluorescent in situ hybridization (FISH), marker-assisted selection, cloning, flow cytometry, RNA dot blotting hybridization, ELISA, and quantitative real-time PCR assays (Walker and Subasinghe 2000). Meanwhile, the field of molecular diagnostics has shown dramatic developments over the last few years in the use of low-cost high-density single-nucleotide polymorphism (SNP) technology in the genotyping of individuals at the SNPs level (van Arendonk 2011). Among these technologies, immunohistochemistry and fluorescent in situ hybridization have been used to show the presence and characteristics of species in complex biological samples (Cheon

and Chae 2000). Also, qPCR and FISH have been the most effective approaches used to amplify and quantify particular DNA or RNA sequences in complex biological samples from different hosts (Elelu et al. 2016). While these molecular techniques are appropriate for a plethora of different biological samples and provide reliable information on some agents, they do not provide the direct sequence of the infectious agent. Also, the use of these methods includes advanced awareness of the causative agents. These shortcomings eventually led to the development and widespread implementation of high-performance sequencing technologies in veterinary diagnostic reference laboratories, which can sequence all genomic material present in biological samples and produce thousands of sequences of previously unknown biological materials (Zeineldin et al. 2019b). In addition, next-generation technology has the ability to provide an unbiased sequencing platform for several microbial genomes and their antibiotic resistance genes in near-real time, which bring several possible advantages to create diagnostic reference laboratories for rapid diagnostic decision making (Zeineldin et al. 2019a). Recently, the current high-throughput sequencing technologies used in veterinary practice are primarily Illumina sequencing platforms (San Diego, CA, USA), Nanopore technology, and 454 pyrosequencing platforms (Ambardar et al. 2016; Singh et al. 2019). In the coming years, the cost of this sequencing technology is expected to be reduced and sequencing platforms will be available and used on a wide scale in livestock. The entire genome sequencing technology has recently been applied in a variety of studies to improve diagnostic methods accuracy in the veterinary field (Glaser et al. 2016). For example, the use of sequencing technologies for the genetic characterization of viral infectious agents has provided a much deeper and more detailed view of viral capsid and virus genetic material (Chappell et al. 2019). This knowledge is important for a deeper understanding of the genotypes of viruses and will help to identify viruses and the future production of vaccines. More recently, developments in mass spectrometry science, including metaproteomics, metabolomics, and metatranscriptomics, have improved diagnostic technologies to enhance animal health (Elolimy et al. 2020). Although valuable information is given by these techniques, the use of this technology in veterinary practice is minimal. Due to the associated high costs, access to these advanced technologies in developing countries is limited (Zumla et al. 2014). In addition, some enhancements are still required before the next generation and mass spectrometry technologies are used as a point-of-care and on-farm diagnostic testing.

7.5 Electrochemical Point-of-Care Biosensor Technology

The term point-of-care biosensors include instruments that can measure the physiological, immunological, and behavioral responses of animals, as well as the monitoring of the animal environment (Wang 2006). Electrochemical point-of-care is a flexible biosensor with a point-of-care functionality used to generate an electrochemical signal measured using a detector and a data analyzer (Dai and Liu 2019).

These instruments are not only extremely precise and sensitive to the parameters being analyzed, but are also accurate and easy to use and can improve the clinical assessment (Wang 2006). The emerging use of point-of-care biosensors in livestock management offers major benefits and applications in the monitoring of animal productivity, health monitoring, and disease detection, as well as monitoring of animal physiological conditions (Gattani et al. 2019; Robinson et al. 2020). The implementation of this technology is expected to enhance animal health and productivity of the livestock industry in the future, as well as to reduce the impact of the livestock industry on the environment (Gattani et al. 2019). In addition, the implementation of point-of-care biosensors in the livestock sector would contribute to social competitive advantage and organizational benefits for the global economy. The latest existing point-of-care biosensors in the veterinary field focus on the use of the available knowledge of animal physiology, nature, biology, nutrition, and the environment, and the incorporation of this knowledge into an effective and real-time diagnostic method (Neethirajan et al. 2017). Different established sensors and their locations in farm animals are presented in Fig. 7.2.

The key point-of-care biosensing devices rely mainly on identifying the desired biological biomarker (e.g. DNA, RNA, enzymes, hormones, metabolites, tissue, blood, cells, etc.) unique to a particular biological agent using a bioreceptor sensor (Wang 2006). The bioreceptor sensor is a key structure in the point-of-care biosensor device since it is crucial for the distinction between various biomarkers and molecules present in the same biological samples. The association between the bioreceptor sensor and the specific biomarker results in specific signals that allow qualitative and quantitative biomarker measurements (Vidic et al. 2017). A detailed overview of the theory and function of the electrochemical point-of-care biosensors has been reviewed elsewhere (Wang 2006; Dai and Liu 2019; Sun and Hall 2019).

The criteria for an ideal point-of-care biosensor in the veterinary sector is similar to that commonly used in human medicine but may vary concerning the particular needs of livestock farming (Kumar et al. 2020). Although designed for human application, respiratory rate sensors were evaluated successfully in cattle to measure thoracic and abdominal movements (Neethirajan 2020). Laser distance sensor while milking is another system for monitoring the respiratory rate of cows (Pastell and Kujala 2007). Sensor-equipped halters (belt-like device on an animal's chest) have been tested to measure the heart rate in dogs (Lahdenoja et al. 2019). Similarly, den Uijl et al. (2017) analyzed the results of behavioral traits such as headshake, walk, sleep, trot, canter, eat, and drink, using a neck collar-embedded with an accelerometer and validated through clinical setting. Several point-of-care diagnostic platforms are currently available in veterinary practice (Table 7.1), and these tools provide automatic readout results greatly reducing the time of diagnosis. However, the complete adoption and application of such innovations and their effective use in diagnostic procedures in livestock clinical practice imply further progress in the robustness of bioassay production and biomarker detection.

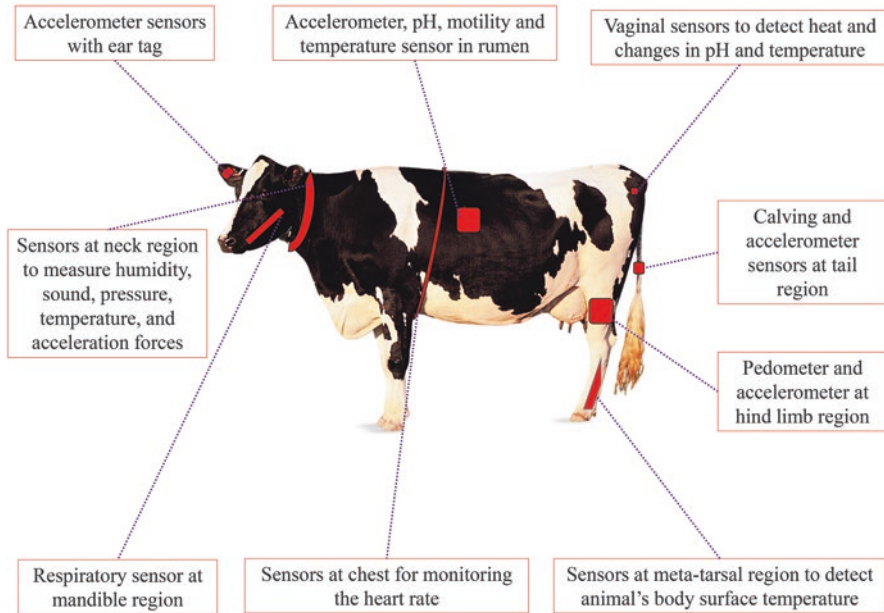


Fig. 7.2 Different established sensors being used in farm animals

7.6 On-Farm Diagnostic Technologies

7.6.1 On-Farm Video Surveillance and Digital Images Analysis Technologies

The frequency of animal behavior changes is often correlated with stress, anxiety, and pathological conditions (Von Holst 1998). These changes in behavior are cost-effective and easy to track via careful observation of individual animals or through the use of video surveillance devices on the farm (Zin et al. 2016). Simultaneous monitoring of these behavioral changes that occur on more than one animal for 24 h a day may be difficult to interpret for the farm observer (Ishiwata et al. 2007). The use of video monitoring systems to track these behavioral activities enables researchers to examine animal activity at their location in real-time (Zin et al. 2016). Further, the availability of digital imaging and the latest innovation of computer programs that can interpret and analyze these data has expanded the use of this technology in the livestock industry to track animal health without distracting the animals (Nilsson et al. 2015). For example, on-farm video monitoring and digital image processing tools have been commonly used for recording particular areas used by individual animals as they get up or lay down to make suggestions on stall size (Ceballos et al. 2004). Moreover, it was also used to track the thermoregulatory clumping of the animal to determine the efficacy of the pen temperature (Shao and Xin 2008). Another example of the use of on-farm video surveillance and digital image

Table 7.1 Some examples of point-of-care diagnostic technologies currently used in livestock husbandry system

Point-of-care test	Health status	Metabolites and agents	Animal species	Sample type	Reference
Protein G-based milk dipstick	Brucellosis	Brucella	Dairy cattle	Blood	Revathi Poonati et al. (2020)
Indirect enzyme-linked immunosorbent (iELISA) assays	Brucellosis	Brucella	Dairy cattle	Blood	Revathi Poonati et al. (2020)
Visual color-based serum phosphorous detection kit	Hypophosphatemia	Phosphorous	Cattle	Blood	D'souza (2020)
Handheld portable lactate analyzer	Bovine respiratory disease	L-lactate concentration	Cattle	Blood	Zeineldin et al. (2017a, b)
Handheld portable lactate analyzer	Healthy	L-lactate concentration	Cattle	Blood	Karapinar et al. (2013)
Portable blood lactate analyzer	Bovine respiratory disease	Plasma lactate	Cattle	Blood	Coghe et al. (2000)
β -hydroxybutyrate hand-held meter	Ketosis	β hydroxybutyrate	Dairy cattle	Blood	Iwersen et al. (2009)
Portable glucometer	Healthy	Blood glucose	Cattle and sheep	Blood	Katsoulos et al. (2011)
Lactate meters	Healthy	L-lactate concentration	Cattle	Blood	Karapinar et al. (2013)
Electrochemical glucose meter	Periparturient healthy cattle	Glucose	Dairy cattle	Blood and plasma	Megahed et al. (2015)
Portable ion-selective electrode meters	Periparturient healthy cattle	Potassium	Dairy cattle	Blood, plasma, milk, and abomasal fluid	Megahed et al. (2016)
Portable somatic cell count-based test	Healthy and sub-clinical mastitis	Somatic cell count	Dairy cows	Milk samples	Iraguha et al. (2017)
FreeStyle Precision Neo™	Healthy	β -hydroxybutyrate and glucose	Dairy cow	Blood samples	Macmillan et al. (2017)

(continued)

Table 7.1 (continued)

Point-of-care test	Health status	Metabolites and agents	Animal species	Sample type	Reference
β -hydroxybutyrate electrochemical meter	Periparturient healthy cattle	β hydroxybutyrate	Dairy cattle	Blood and plasma samples	Megahed et al. (2017)
Point-of-care glucometer	Healthy	Glucose	Goat	Blood samples	Quandt et al. (2018)
Epocal point-of-care analyzer	Healthy	Ionized calcium	Dairy cattle	Blood samples	Mahen et al. (2018)
Erythrocyte osmotic fragility assessment at field level	Stress-related disorders	Erythrocyte membrane	Sheep and goats	Blood samples	Reddy et al. (2019)
Cobalt chloride-impregnated device	Stress-related disorders	Moisture	Cattle	Sweat samples	Pereira et al. (2010)
Leukocyte esterase (LE) test strips	Subclinical endometritis	Leukocyte esterase	Dairy cow	Vaginal discharge	Van Schyndel et al. (2018)
Brix refractometry	Subclinical endometritis	Total solid percentage	Dairy cow	Vaginal discharge	Van Schyndel et al. (2018)
Colorimetric ammonium point-of-care test	Periparturient Holstein-Friesian cows	Ammonium	Dairy cow	Urine sample	Megahed and Constable (2020)

processing technology is the tracking of lameness problems in dairy cows by measuring the shift in animal gait characteristics as they pass to the milking parlor (Berckmans 2014b). The key drawbacks of on-farm video monitoring and digital image processing technologies are the difficulty in detecting the behavior of a large number of animals at the same point over a long period (Von Holst 1998). The challenge of recording in low ambient light and the labor needed to evaluate and display the frequency of such behaviors in individual animals over a long period impedes the application of this technology on a wide scale (Fleishman et al. 1998). Despite current limitations, on-farm video surveillance and digital image processing technologies may be used as a reference framework for other behavior tracking technologies in livestock management system.

7.6.2 On-Farm Audio Surveillance Systems for Sound Detection

Animal vocalization has been commonly used as a significant predictor of animal health status (Manteuffel et al. 2004). Sound monitoring technology has been implemented in the livestock industry to track and regulate animal health and welfare (Wathes et al. 2005; Handcock et al. 2009). The animal sound provides details not only about the abnormal state of the animal but also about the personality of the animal (Manteuffel et al. 2004). Automatic detection of irregular animal behaviors using bioacoustics, such as cough sounds, achieved maximum efficiency and minimal costs at a precision level of over 94% in the livestock industry (Anderson et al. 2011). For example, the use of sound recognition devices in the pig and dairy cattle industry provided an earlier diagnosis of respiratory distress (Exadaktylos et al. 2008; Ferrari et al. 2010). Respiratory diseases in the livestock sector have resulted in significant economic losses due to high mortality and morbidity rates (Zeineldin et al. 2016). Coughing is the primary clinical symptom of respiratory diseases in both cattle and pigs (Carpentier et al. 2018). Monitoring coughing as an indicator of respiratory disease can identify early clinical cases before the occurrence of serious complications and thereby reduce the risk of respiratory diseases in the livestock industry (Ferrari et al. 2010). It is well recognized that existing diagnostic methods for respiratory diseases in livestock are expensive and not very precise. Several attempts have therefore been made to classify the coughing characteristics in various animal species as a sign of respiratory diseases. Since coughing sound during respiratory distress has unique characteristics, the clinical use of on-farm audio surveillance systems to investigate cough sounds may be beneficial for real-time monitoring (Chung et al. 2013). Several algorithms have been validated for various cough sounds characterization in the livestock industry (Berckmans 2014b).

The advantage of using on-farm audio monitoring systems for sound detection is that the non-invasive nature of this device does not interfere with the daily activities of animals (Ferrari et al. 2008). In addition, this technology may be used to track multiple individuals at the same time. However, this may result in overlapping and less coherent sounds in a commercial setting (Carpentier et al. 2018). Despite the usefulness of this technology in early disease detection and tracking animal sounds, more consideration needs to be given to the realistic application and avoid drawbacks of these technologies in the veterinary field.

7.6.3 On-Farm Accelerometers and Pedometer for Walking and Standing Behavior Monitoring

Several automated technologies have been used in the livestock industry to track animal walking and standing activity as well as to record step frequency, including the use of accelerometers and pedometers placed on animals employing various

wearable devices such as collars, ear tags, leg or tail bands (Rushen et al. 2012). Over the last few years, accelerometers are probably the most promising technology for providing accurate data on activity tracking, body orientation, and complex posture behaviors (Sala et al. 2011). These instruments have been used to remotely assess animal walking distance and animal orientation (Rothwell et al. 2011; Ringgenberg et al. 2010; Enstipp et al. 2011). Besides, accelerometers have been used to track lying down activity, rumination, feeding, sleep pattern, approaching parturition, and lameness in dairy cows (Martiskainen et al. 2009). Similarly, pedometers are an inexpensive, simple, portable electronic system used to measure the number of distance traveled and the status of the animal's behavior using a mechanical sensor (Shepley et al. 2017). Recently, pedometer devices have also shown strong predictability when used to classify estrus behavior (Løvendahl and Chagunda 2010). Furthermore, a pedometer can assess several cow activities, including locomotion behavior (Alsaad and Büscher 2012) and approaching calving time (Felton et al. 2013). The application of accelerometers and pedometers in livestock involves using three main components: the main device, the receiving system, and the software program. The receiving system collects animal activities from the attached sensor and transmits these data in real-time through the receiving system to the software program (Brehme et al. 2008). The advantage of on-farm accelerometers and pedometer technologies is the continuous monitoring of animals in commercial systems, particularly when paired with other portable techniques (Moreau et al. 2009; O'Leary et al. 2020). The pitfalls of using these instruments for behavioral tracking include the expense of accessing sensor data and data processing. The risk of damage to sensor boxes due to continuous movements of the animal in the barn and growing labor demand for fixing and removing equipment from livestock are another disadvantage (Norling 1991). However, further modification including increasing battery life, memory capacity, and reducing the device size to be conveniently attached to the animal can efficiently transform on-farm accelerometers and pedometer technologies into functional behavioral measurements technologies.

7.6.4 On-Farm Global Positioning Systems (GPS) for Position Monitoring

On-farm GPS technology is designed to monitor the animal's location and remotely track the movement of the animal within a given area of the farm (Moen et al. 2001). The detailed methodology of GPS collars functioning is presented in Fig. 7.3. The GPS is designed to additionally inform the animal keeper of the distribution of animals on pastures (Griffin 2009). It can also track animal movement using a digital map for easy monitoring and submit information on animal status, such as animal behavior, delivery time, and position (Tang and Abplanalp 2014). Recently, low-cost GPS positioning sensors have been used in veterinary practice to assess the

position status of animals (Godsk and Kjærgaard 2011). GPS sensors are attached to the animal collars to detect possible changes in animal behavior, such as feeding, walking, lying down, and standing, which may be associated with stressful situations or changes in animal health (Tang and Abplanalp 2014). Advances in GPS technology have produced lighter and more precise receivers, but the identification of numerous animals in various geographical regions is still difficult (Davis et al. 2011; Foley and Sillero-Zubiri 2020). The drawbacks of GPS technology include costs, battery life, and the frequency of animal location updates. Current technology enables the animal’s location to be changed every second, but this rate exceeds the capacity resources available in most animal tracking barns (Tomkiewicz et al. 2010). These limitations restrict the ability to use GPS systems in behavioral tracking for longer periods and limit their use in small local zones. Furthermore, attaching the GPS to the animal collars may have an adverse effect, such as decreased power, tissue damage, and the death of the animal. Therefore, there is still much research needed to improve this technology concept and components used in these devices to be widely used on large scale.

7.6.5 On-Farm Automatic Milking Robot System for Monitoring Leg Health

The milking robot technology was used to track the leg health of the dairy animal in which the sensors were attached to the amplifier and the data was collected on a computer using advanced processing software (Pastell and Kujala 2007). This

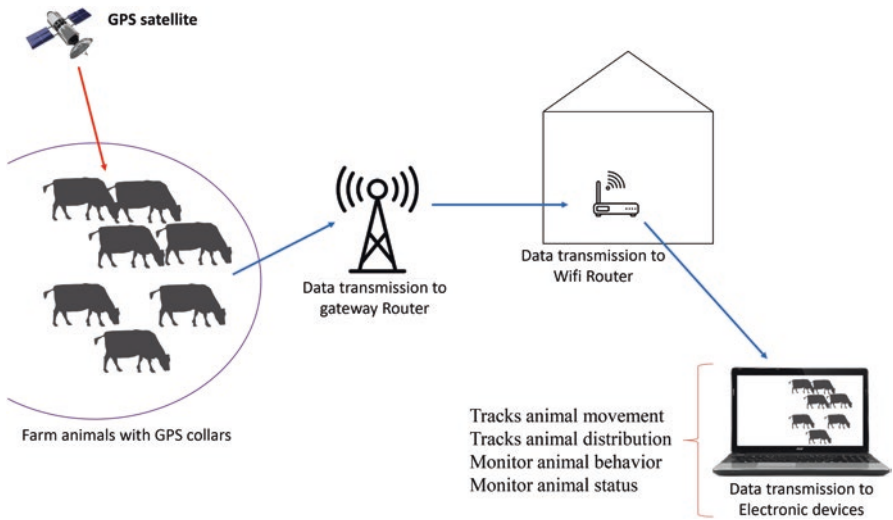


Fig. 7.3 Methodology of GPS collars functioning

technology can be used to automatically detect kick frequency during milking, which warns the operator of potential hoof diseases and other leg problems, but more work is required to improve accuracy (Borderas et al. 2008). Automated milking robots automatically collect data during the milking process and track poor health conditions in a regulated, confined barn up to three times a day. For example, lameness in dairy cows is evident in a reduced frequency of visits to robot system technology (Bach et al. 2007), but this appears to have a low specificity as many other cows are not lame (Borderas et al. 2008). Research focused on the weight distribution between the limbs and the probabilistic neural network showed that automated weight distribution measurements could classify lame cows markedly faster than those obtained by visual inspection applying gait score systems (Pastell et al. 2006). Moreover, these weight distribution steps are responsive to the degree of discomfort associated with lameness (Visen et al. 2002; Chapinal et al. 2011), but the relationship between input and output values is difficult to translate into mathematical representation (Visen et al. 2002). Therefore, the impact of different management factors and their interaction with the robot utilization should be carefully considered together to improve the system efficiency in livestock.

7.6.6 On-Farm Automated Feeder for Monitoring Feeding Behavior

Farm animals are increasingly housed in groups where disease is challenging to identify, leading to increased morbidity and care costs. Automatic detection of feed, water, and frequency of particular behaviors of animals, typically through radio frequency recognition, can be used for early identification of animals that are ill and can provide insight into potential improvements in the pain conditions of animals (Weary et al. 2009). Sick animals spend more time in the feed bunk than healthy ones, particularly the newly received animals (González et al. 2010). Data from such feeders can help recognize morbid animals from healthy ones based on differences in feeding behaviors (Sowell et al. 1999). Recording variations in feed behavior can also help track dairy cows suffering from periparturient diseases such as metritis, ketosis, or lameness (Proudfoot et al. 2010). Current limitation in monitoring the animal feeding behavior is the setup requirement, system maintenance, and the associated cost (Brown-Brandl and Eigenberg 2011). All of these possible drawbacks must be addressed when assessing automated feed intake and behavioral monitoring systems.

7.6.7 On-Farm Physical Inspection for Monitoring Health and Stress-Related Disorders

Any health and stress-related problems are reflected in the physiological parameters such as rectal temperature, pulse rate, and respiratory rate. Simple evaluation of these variables with physical or visual inspection for few minutes may explain the underlying health-related issues. However, quantification of these parameters involves animal contact, restraint, and human-interface (Jorquera-Chavez et al. 2019). Besides, monitoring these activities in large flocks is time-consuming, labor-intensive, and subjective, being inappropriate at the field level. Reddy et al. (2019) projected panting score as the facile measure for quantifying health problems, especially respiration-related issues. They attributed the technique's eminence to its non-invasive functioning of direct observation from two or three meters away from the animal (Nejad and Sung 2017). Therefore, digitalization of diagnostic technology to stress-related disorders through precision livestock farming technologies has the potential to address many aspects of animal health and related to animal welfare and public health.

7.6.8 On-Farm Monitoring Change in Core Body Temperature

Body temperature is the most useful measure of animal reactions to physical-environmental stimuli, occurrence of diseases, and physiological functions such as diet, lactation, and reproduction (Galan et al. 2018). Remote temperature monitoring may help to rapidly detect diseases, minimizing associated treatment costs, and improving animal efficiency (Liang et al. 2013). Core body temperature is determined by many factors such as health status, physiological status, environmental temperature, humidity, water consumption, and feed intake (Davis et al. 2003). Core body temperature can be measured using a variety of methods, such as rectal probe, tympanic membrane, implanted transmitter, infrared thermography, and rumen temperature bolus (Firk et al. 2002). These approaches reflect mechanized technologies, but we must ensure that they are the most reliable, practical, and economical (Firk et al. 2002). Rectal probe is an alternative to conventional rectal thermometers since thermometers are labor-intensive and time-consuming (Godyń et al. 2019). Measurement of body temperature using a rectal probe is useful by continuous measurement of changing temperature over time; however, existing measurement methods are only suitable for single or short-term continuous measurements (Reuter et al. 2007). The rectal probe approach causes the loss of certain data while removing it out of the rectum. Hence, the probe must be connected to the animal in a manner that is not feasible for large-scale use. On the other hand, tympanic temperatures are a reliable indicator of heat stress in cattle, with scales somewhat close to rectal temperatures (Gasim et al. 2013). To measure the tympanic temperature, thermistor is connected to the data logger and then inserted several cm down the ear canal until

the tip is located near the tympanic membrane; nevertheless, the implementation of these techniques requires trained staff (Mader et al. 2002). The tympanic temperature procedure can only be used for limited numbers of animals and should not be used for long periods (Goodwin 1998). Animals with tympanic membrane recorders should also be closely monitored for infection at the insertion site (Davis et al. 2003).

Other methods for tracking core body temperature are infrared thermography (IRT) and rumen temperature bolus. Thermography imaging was used to detect nasal mucosal temperatures in a non-invasive manner (Willatt 1993) and corneal surface temperatures (Stewart et al. 2008). In bovine medicine, IRT is used mainly for diagnostic purposes, animal health evaluation, and feed quality monitoring (Montanholi et al. 2010). It has also been used for the prediction and early diagnosis of mastitis and lameness in dairy cattle (Hovinen et al. 2008; Alsaad and Büscher 2012; Stokes et al. 2012). Additionally, IRT has been used for bovine respiratory disease diagnosis at an earlier stage (Schaefer et al. 2007). The key drawbacks of these technologies are that images must be obtained in direct sunlight, and animal hair coats must be free of soil, moisture, and foreign material (Stewart et al. 2005). Other drawbacks of infrared thermography include the cost of infrared cameras, size, angle of measurement, and animal activity (Johnson et al. 2011). Therefore, IRT requires some adjustment before it becomes practical for the livestock industry.

Developing rumen temperature bolus for tracking core body temperature can be the most appropriate way for producers to monitor body temperature. These boluses could also periodically collect data at different time intervals to monitor sudden or subtle changes in the ruminal environment (Edwards 2010). The benefit of rumen temperature bolus over other approaches is the quick administration without side effects to the animal and the potential to track the large flock at once (Ipema et al. 2008). Rumen bolus is delivered by a baling gun and can be safely recovered after slaughter and does not irritate the epithelium of the reticulo-rumen wall (Ghirardi et al. 2006). This approach may have the longest duration in the body. Nonetheless, it may present some challenges for recovery in commercial harvesting facilities and the dynamics of rumen makes it susceptible to changes in the temperature from water intake and the type of diet due to fermentation (Timsit et al. 2011; Makinde 2020).

7.7 The Future Direction and Challenges of On-Farm and Point-of-Care Diagnostic Technologies

Advances in on-farm and point-of-care diagnostic technology have been linked to acceleration of mechanization, which could promote diagnostic validation, strengthen veterinary surveillance systems, increase livestock efficiency and improve animal welfare (Jones et al. 2019). Therefore, the potential effect of using this diagnostic technology depends on the number of livestock farming systems and the number of animals in each system to which these technologies will apply. These

new devices will encourage investment and will support our society (Abuelo and Alves-Nores 2016). The biggest challenge for global economies is to bring new products that meet customer needs in a cost-effective and productive manner. Other critical problems currently faced by on-farm diagnostic technologies are the sluggish usage of these technologies on a broad commercial scale and lack of harmonization between the use of these technologies and the large-scale implementation of data analysis (Busin et al. 2016). The key explanation for this is the abundant data produced and the available technology's inability to promptly turn the ample data into useful knowledge that could be used in the decision-making process. Additional work on the system using these technologies will also contribute to implementing the next generation of an economical and appropriate health monitoring system, which will capture additional behavioral changes related to the level of operation and connect the collected animal data to predict animal health events. The new technology has now reached a stage where its application to biological processes has become practical. Moreover, the livestock industry requires a large number of animals and procedures, making it possible to manufacture personalized, applied low-cost technology (Berckmans 2014a). Of course, no major advancement in precision livestock medicine come without disadvantages. Point-of-care and on-farm diagnostic technologies are still in the early stages of implementation particularly in veterinary medicine, and several major issues will need further research before these techniques become practical with a wide application in livestock management systems.

7.8 Summary and Conclusion

This chapter explains basic on-farm and point-of-care diagnostic technologies as a valuable way of monitoring animal health, welfare, and disease in the livestock industry. The promise of using this technology is valid; however, some drawbacks need to be tackled. Advances in on-farm diagnostic technology will increase the market's efficiency and make these technologies more competitive in the livestock sector. This chapter outlines the attached and unattached type of technologies for tracking physiological and behavioral responses that are important for evaluating animal welfare. The choice of the appropriate diagnostic method depends on the anticipated gain relative to the cost of the device used. Behavioral evaluation must also be selected based on their importance to animal welfare rather than their ability to be reported automatically. Further work on the system using these types of technologies will contribute to creating a next-generation, feasible, and cost-effective animal health monitoring system to detect other critical activities that will enable producers to introduce prevention and treatment protocols against diseases at earlier times.

Conflicts of Interest The authors declare no conflict of interest.

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Chapter 8

Biotechnological Applications in Poultry Farming



S. M. Lutful Kabir and S. K. Shaheenur Islam

Abstract Global meat consumption is growing exponentially because of rapid urbanization, population growth including lifestyle changes in the developing countries. Amongst the sources of meat consumption, maximum proportion comes from poultry birds as a white meat. Advancement of biotechnology revolution triggers a numerous opportunities in poultry industry to mitigate such emerging demand of meat and meat products. Biotechnology in poultry that never be regarded only for better farming, but its multifaceted impacts in the poultry production food systems in recent times. Here we review the activities related with the poultry sector, for example, breed development, health, nutrition, disease prevention and control, preservation of poultry origin meat and meat products along with the food safety are mostly achieved enormous improvements. The biotechnological tools based on modern techniques offer enormous prospective for the production of biologics and vaccines, medicinal products including disease diagnostic assays as a part of disease control and prevention. The advancement and application of these technologies are mostly used in the developed countries, however, these are being used moderately in developing countries in different segments of poultry science. Both conventional and modern biotechnologies in poultry have contributed enormously to enhance productivity and optimize income generation activities and reduce poverty, lessen the burdens of diseases and guarantee to environmental sustainable livestock production in developing countries.

Findings

In conformity with in-depth literature review on the above captioned title, we may conclude that biotechnology is attending to the upgradation of veterinary science especially in the light of poultry production and processing of poultry origin

S. M. L. Kabir (✉)

Department of Microbiology and Hygiene, Bangladesh Agricultural University,
Mymensingh, Bangladesh
e-mail: lkabir79@bau.edu.bd

S. K. S. Islam

Department of Microbiology and Hygiene, Bangladesh Agricultural University,
Mymensingh, Bangladesh

Department of Livestock Services, Krishi Khamar Sarak, Dhaka, Bangladesh

food that entwined with food safety issues. There is a need to publicize these issues since the approaches are almost new, and to provide handsome training for laboratorians from both private and government sectors targeted to getting best benefits on disease diagnostic areas. It is pertinent to instituting of inclusive and effective regulatory framework considering the country perspectives and the safe application of technologies regarding biosafety, biosecurity and to disperse the confusion among the mass people in this evolving discipline. The government with the partners of the food systems, and the biotechnology industry should completely utilize this science-based devise to corroborate food safety, and to stance at the back of public health decisions.

Keywords Poultry production and processing · Poultry origin food · Food safety · Food systems · Biotechnology · Disease diagnostic techniques · Regulatory framework · Biotechnology industry · Developing countries

Abbreviations

AI	Artificial insemination
AMR	Antimicrobial resistance
AMU	Antimicrobial uses
BAC	Bacterial artificial chromosome
BLAST	Basic local alignment sequence tool
BULP	Best linear unbiased prediction
CCAC	Canadian council of animal care
CCPS	Combination of crossbred and purebred selection
CLA	Conjugated linoleic acid
DNA	Deoxyribonucleic acid
EBU	Expected breeding value
EMO	Effective microorganism
EST	Expressed sequence tag
FASTA	Fast alignment
FCR	Feed conversion ratio
GAH	Good animal husbandry
GEBV	Genomic estimate breeding value
GPH	Good personal health
HA	Hemagglutinin
LBM	Live bird market
LD	Linked disequilibrium
MAS	Marker assisted selection
NA	Neuraminidase
NCBI	National Center for Biotechnology Information
NGS	Next generation sequencing

PCR	Polymerase chain reaction
PLS	Pure-line selection
QTL	Quantitate trait loci
QTL	Quantitate trait locus
SNP	Single nucleotide polymorphism
SRRC	Sheep red blood cell
TVC	Total viable count
US	United States
WGS	Whole genome studies

8.1 Introduction

As a diverse domain of scientific discipline, biotechnology has been associated with the human development when the life has advanced. Archaeologically, this was evolved when humans used microbes to prepare foods like bread and beverage, it may be dated before 4000 BC. The word “Biotechnology” is usually specified as the ability to use living microbes or their ingredients to promotion or to modernize a product, to develop fauna or flora or to change microorganisms for desired objectives (Raju et al. 2015; Vijayakumar and Sasikala 2012). Principally, this makes association among microorganism, human and animals, and with the effect of technology for the betterment of human life. The exploration in cell biology, agriculture, plant sciences, animal sciences, food science, medicine and environmental sciences are some paramount areas where biotechnology acts a key player (Sugumaran and Ponnusami 2017). A broad range of tools for the genetic upgradation of animal species, vaccines and biologics for disease control, and advancement of rapid test kits to diagnose the disease of poultry and livestock, including other animal species. Therefore, this discipline has already had a substantial influence on poultry production, control and prevention of diseases, and processing of poultry origin food (Pal et al. 2017) and also be linked with the safety standards of food and food products (Fig. 8.1).

As a part of global poultry meat production, 18% comes from Unites States (US) as a top producer, followed by China, Brazil and the Russian Federation. Nevertheless, considering global egg production, 42% is supplied by China as a top producer, followed by US and India. More than 60% of global egg production comes from Asia as considered to be the highest egg producing region in the World. From 1961 to 2017, the global poultry production increased about 14 times more (122 million tonnes) to mitigate incessant demand, however, egg production increased approximately 6 times (87 million tonnes) with 37% global contribution at present time. Interestingly, world egg production has been increased by more than 150% during past three decades. Enormous growth has been witnessed in Asia due to supply of cheapest animal source protein requirement for huge population in this region, where production increased nearly fourfolds. Approximately, 80% rural

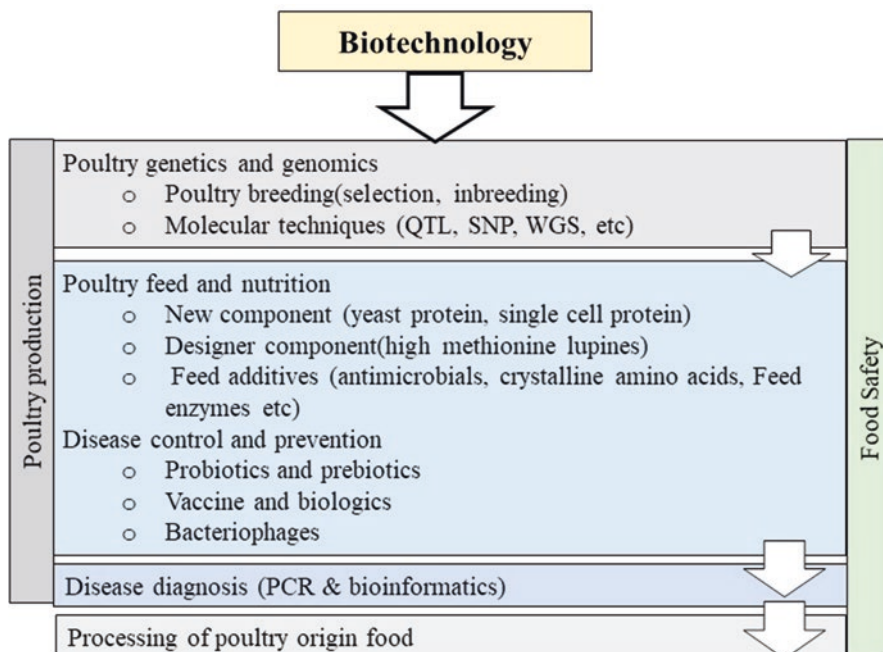


Fig. 8.1 Biotechnological application at poultry production, processing of poultry origin food and interlinking with food safety. (Adapted from Pal et al. 2017)

households rear poultry as their nutrition demand and livelihoods (FAO 2020) with biotechnologically developed high-yielding commercial layer or broiler poultry and low performance dual-purpose breeds in developing countries (Pym 2013). However, proportions in these two categories fluctuate widely as indigenous poultry shares around 90% of the poultry population in these countries (Pym et al. 2006). Moreover, this figure may vary due to rapid growth of the commercial poultry industry in developing countries at present time.

Nowadays, biotechnology is extensively applied in animal production along with a plentiful other potential uses in many countries in the world. This could have applied for enhance animal performance via better nutrition, upgraded production traits or better health condition through integrating antibodies or vaccines into feeds. However, this can efficiently defend the animals against infectious diseases and to enhance animal productivity economically (Daniel 2016). Over the last 75 years, a remarkable influence on poultry production through selection of production traits has been noticed. Moreover, genetic improvement of production parameters in poultry to be optimized in the next 20 years due to the emerging challenges of global food security (Mohammed et al. 2016). Nowadays, more emphasis on the demand of the producers and consumer like productivity, quality, food safety and animal welfare in addition to the necessities of the breeders on economically feasible options like low cost of feed and other production inputs (vaccine, biologics and

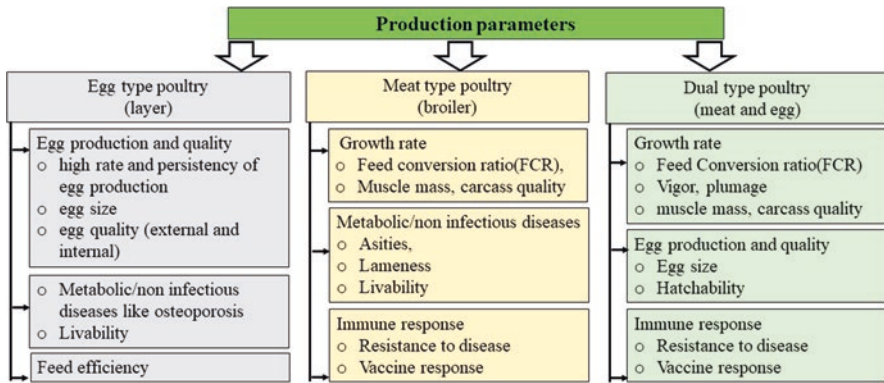


Fig. 8.2 Characteristics associated with the production parameters for selection of targeted poultry birds. (Adapted from Burt 2002, Engormix 2008, and Leeson and Summers 2010)

medicine) and higher fertility are the pivotal areas as prerequisite of poultry industry (Fig. 8.2). To obtain desire level of production, conventional genetic selection is unable to fulfill the targeted objectives unless the application of genomics to be functional. The poultry genomics has been influenced a lot as the progress of genetics technology of human and model organisms. At present, a number of techniques are well documented viz., quantitative trait locus (QTL), genetic markers and physical mapping, comparative mapping, bacterial artificial chromosome (BAC) resources, expressed sequence tag (EST) are tailoring to benefit to poultry industry (Burt 2002). The scope of this paper will glimpse on biotechnological approaches are being used for production and processing and to enable safety standard of poultry origin food. The paper is also focus on the potential limitation and challenges to be addressed to operationalize the biotechnological approaches in low resource settings. Moreover, this paper is also outlined the way forwards to mitigate the challenges as a long term goals.

8.2 Historical Perspectives

Biotechnology is a technology based biology through which the scientific and engineering principles applied in living animals for the production of biological materials targeted for the benefit of human and animal health. The animal biotechnology has been commenced approximately 8000 years before by means of animal domestication and artificial selection process. Nearly 48,000 mammalian species, less than 20 were successfully tamed in human captivity (Diamond 1999). Those animals were successfully tamed, and subsequently reared in captive condition, they shared and displayed a distinctive mixture of traits (Diamond 1999; Hale 1969). They were found to be comparatively quiet, and cooperative in their feed requirements, and could be raised and go maturity rapidly on natural available fodders, and could be

easily reproduced in the human confinement. These animals had hierarchical societal norms and permitted humans to restrain and they were adapted to living within big group.

The paradigm shifting in animal production had been taken place in the earlier decades via the techniques, namely artificial selection process to accumulation production traits, immunization to prevent and control of animal diseases, and artificial insemination technique to increase production of food animals. Nevertheless, in early 1960s, a contemporary biotechnology was emerged followed the genetic code confirmation (NRC 2002). Actually, the updated breeds of livestock differ very distinctly from their ancestors is mirroring of how rapidly the breeding took place. The dairy industry of contemporary United States where modern Holstein Frisian dominates, however, a little similarity to its descendants of about a 50 years ago. From 1945 to 1995, milk production increased each cow nearly three times (Majeskie 1996) due to breeding program of selected proven cattle. They had undertaken significant adjustment in traits over a short duration of time from an intense selection process and to bring the diversity observed in present breeds as revealed a huge disparity in pool genetics in species (Wayne and Ostrander 1999). The selection method formed the farm animal species (cattle) that facilitated to the development of the modern farmed poultry breeds of chickens, ducks, turkeys and geese raised for meat, eggs, and other uses like feathers. Thus the broiler poultry was established that gained at the market weight by around 40 days (Pisenti et al. 1999) through feeding of less quantity feed (Lacy 2000). The use of probiotics supplementation has a significant impact on live weight gain, increase carcass weight including high output of cut up meat and immune response including heat tolerance capacity of poultry, and to ensure the safety of poultry origin food (Roy et al. 2015; Kabir 2009a, b; Kabir et al. 2004). This has been established as probiotics inhibit harmful microorganisms through living in the inner side of the intestine (Kabir et al. 2005, 2014) and the prebiotics used in feed eliminate the harmful organism from gut and found to be better growth (FAO 2013).

Different of types of genetically modified animals have been developed principally for food, and others mainly for nonfood materials like pharmaceuticals, fibers, vaccines, and other high value products. Introduction of biotechnology have an important influence on livestock sector as vitamins, nutraceuticals for nutrition, vaccines and antibiotics for animal health, therapeutic proteins for transgenic animals and breeding for genomics (Mishra and Singh 2013). A rapid progresses in genomics with the important impetus of human genome project observed during last 15 years (Lander and Weinberg 2000). Research on livestock genetics has gained from this advancement, along with the formation of comprehensive genetic marker trait-genes through genetic mapping (Georges and Andersson 1996). Notwithstanding, quality food production, fuel synthesis from various raw products, and selective breeding program are the paramount areas of biotechnology.

The key biotechnological applications namely tissue culture, DNA finger printing, fermentation, selective breeding, and recombinant DNA technology are being used nowadays (Ledoux and Antunes 2018; Krasznai et al. 2018; Nitschke and Silva 2018, Kamle et al. 2017; Lucarini et al. 2016). However, biotechnology can

diagnose of various hereditary disorders including infectious diseases through complete genetic analysis and threat them accordingly (Šuster et al. 2017; Hamad et al. 2017; Lao et al. 2017; Kavousipour et al. 2017; Calvo-González 2016).

8.3 Biotechnology in Poultry Industry Through Genetics and Genomics

The great expansion of poultry industry was observed since last several decades due to scientific and technological progressions. The genetically improved superior stocks adept with higher productivity, even under extreme climate has shifted from rural farming to a complete form of poultry industry (Saxena and Kolluri 2018). The parameters were evaluated in layers birds with better feed conversion rate including other traits as high rate and persistent egg production, egg size and quality, and livability. In meat type poultry birds, growth rate, carcass quality, muscle mass, feed conversion rate and livability are essential parameters for consideration (Fig. 8.2). Since these traits are up gradated genetically in birds and achieved via selection process like crossbred and purebred selection (CCPS) that enhanced productivity per bird and production volume (Saxena and Kolluri 2018; Engormix 2008). In poultry, growth characters are connected with polymorphisms in various hormones and associated receptors like lambr1 ghrelin, growth hormone, growth hormone receptor, MC3R, MC4R, TGF- β and IGF-II, (Fang et al. 2007; Huang et al. 2007; Jiang et al. 2002; De Vries et al. 1998; Grobet et al. 1997; Feng et al. 1997). In animal production, genomic estimate of breeding value (GEBV) of an individual animal is anticipated through the whole genome as “genomic selection” was developed and considered to be as an innovative revolutionary technology using SNPs. (Saxena and Kolluri 2018). The impact of genetics and genomics at meat and egg type poultry production are as follows:

8.3.1 Impact on Broiler (Meat Type Poultry) Production

The past decade witnessed strong growth in the global per capita poultry consumption increased by 16% whereas the per capita beef and veal consumption decreased by nearly 5% between 2008 and 2017 (OECD 2018). At present nearly 100 million tonnes of poultry meat and 73 million tonnes of egg produced annually throughout the World (FAO 2016). However, backyard poultry contributes 2% of poultry meat demand, and 8% egg demand globally. The proportions of global poultry meat supply from broiler and layer poultry as 92% and 6% respectively (Mottet and Tempio 2017). The poultry especially broiler is playing a key role as the source of improved value protein and fats for the world people (Tavárez and Solis de los Santos 2016). Both in developing and developed countries the consumption of poultry meat is

gradually increasing since the past decade due to its cheaper price as the broiler industry is capable to attain remarkable production and efficiency (Scanes 2007). A broiler bird weighed approximately 600 g at 42 day old where a feed conversion ratio (FCR) was recorded as 2.8 in 1957, however, today the FCR is deduced to 1.7 and weighed 2900 g at the same age (Zuidhof et al. 2014) or more favorite FCR is observed in recent time. Due to development of production techniques that included enhanced genetics and breeding, good knowledge of nutrition and feeding along with total better management practices are responsible for such advancement (Tavárez and Solis de los Santos 2016). Nowadays, maximum production enhances (80–90%) observed due to impact of improved genetics and breeding (Havenstein et al. 2003a, b).

8.3.2 Impact on Egg (Layer Type Poultry) Production

At present, global demand of eggs is 75 million tonnes where yearly growth of 1 million tonnes observed. To minimize the escalating demands, approximately 50 million layer hens to be included per year. A good management environments to support the genetic traits for 20 kg egg mass/each hen with an age of 20–76 weeks (Preisinger 2018). Constant upgrading of egg production quality in hen is the best selection parameter in layers. Annual increment of around two to three eggs/year in a production cycle with a duration of 13 months could be predictable and confirm through field results on genetic trend (Fig. 8.3). Simultaneously, feed efficiency had been enhanced relatively with the egg production mass in brown layers, moreover, daily feed requirement were significantly lessened over the years (Fig. 8.4).

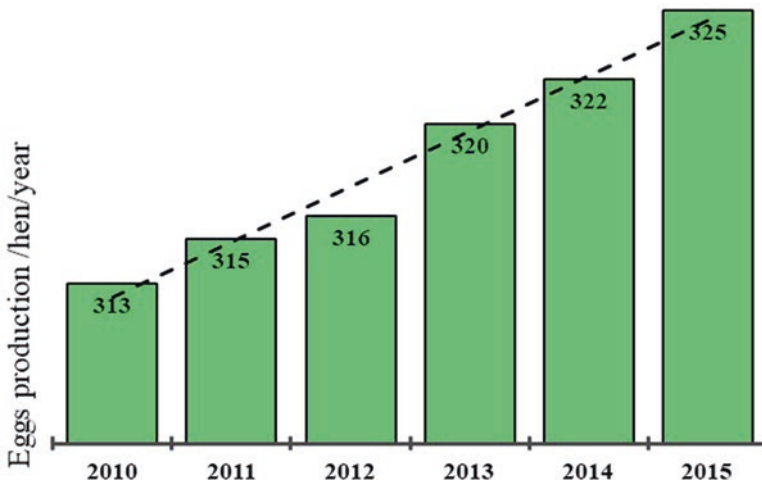


Fig. 8.3 Recent genetic trend with egg production output/year. (Adapted from Preisinger 2018)

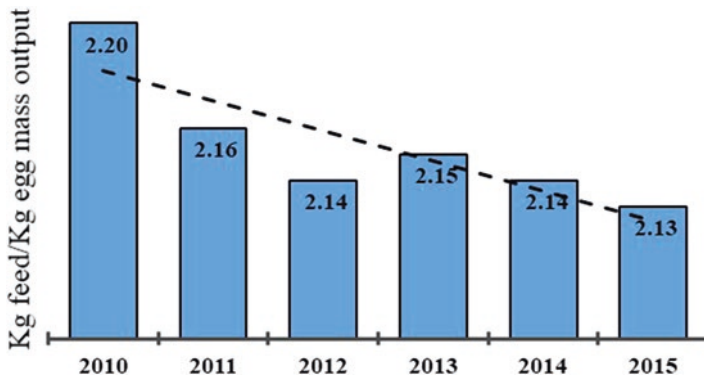


Fig. 8.4 Recent genetic trend with feed efficiency (kg feed/kg egg mass output). (Adapted from Preisinger 2018)

8.3.3 Biotechnology in Poultry Breeding

The genetic improvement in poultry is achieved through implementation of selection program from phenotypes of expected breeding Value (EBV) assessment. However, a lots of drawback are also challenged as capability to precisely and judiciously documenting the phenotypes on candidates and their near relatives; the expenses of documenting the data; and at beginning of major of the production traits in late life impedes genetic improvement each unit time. The heritability estimates of production parameters are important in determining the technique of selection. The additive genetic difference retains decreasing over the generations effecting in lowered evaluation of genetic upgradations and heritability for each generation (Saxena and Kolluri 2018).

8.3.3.1 Selection Process

The selection process in poultry breeding program has been adjusted as per the updated knowledge and the requirements. The birds those fulfilled desired traits were retained and the other birds eliminated from the flocks as a part selection process in early 1940s. After taken maize development program during 1980s, the knowledge of 2, 3, or 4 way breeding process was tailored in poultry birds. Interestingly, the purebred birds were also switched by hybrid commercial birds by terminal cross and the particular egg and or meat type poultry birds transformed to both type poultry birds. The adverse relationship among production and reproduction traits requires the improvement of particular male and female lines both in broiler and layer poultry birds. Above improvements enabled to established specialized lines in meat type (Chambers 1990) and egg type (Fairfull 1990) commercial poultry and also specific male and female lines characteristically with distinct basis of genetic source (O'Sullivan et al. 2010). The poultry breeding at present time

includes combination of crossbred and purebred selection (CCPS) and pure-line selection (PLS) (Saxena and Kolluri 2018).

Development of Specialized Lines Through Pure-Line Breeding

An exceptional selection method centered on distinct set of characteristics of father and mother lines by which desired father and mother lines are developed. The mother lines are tactfully selected for reproductive parameters, viz., egg production, egg size and weight, egg-shell quality, age of sexual maturity, hatchability along with growth at young age. On the contrary, the growth rate, configuration of body, FCR, carcass texture, and fertility are considered predominately for selection of father lines. Thus developed commercial broiler poultry birds associated with particular lines had been achieved to lessen the production cost. The consequences of the gene recombination in crossing among the genetically diverse lines resulting a heterotic outcome in progeny with economic characters. A control stock having the same improvements in inbreeding in the targeted poultry population could be conserved to evaluate of impacts.

Egg Type Poultry (Layer)

The primary objective of layer selection is to gain highest number of marketable eggs/hen per raised by reduced feed cost/egg or kg egg mass. Further, the eggs should be with better inner and outer conditions. The layer stocks should be with minimum mortality and maximum adaptableness at diverse conditions. However, layer parent stock are selected to enhance more than 30 characteristics are vital for profitable egg production. Breeders (grandparent stock) are selected for the sexual maturity age, laying rate, viability, body weight, weight of egg, feed conversion ratio (FCR), color of shell, shell strength, albumen height, presence of inclusion body in egg (meat and or blood spots), and adaptability. The selection approaches to enhance egg yields comprises the consistency of lay, clutch duration, FCR, skeletal deformities like osteomalacia and osteoporosis (Thiruvankadan et al. 2010).

Meat Type Poultry (Broiler)

The selection strategies in broiler poultry are focused on fast growing and carcass features. The best experienced strategy for broiler pure-line selection is related to be considered the commercial weight selection that gains the market weight and the age of selection complies relatively earlier as growth capacity improves. A salable age and period selection are also practiced for the selection broilers. Different biotechnological methods through breeding and selection approaches like trap nesting, mass selection, hybridization, artificial insemination, family feed conversion testing, selection index, individual feed conversion testing, Best Linear Unbiased Prediction (BLUP) breeding value estimation including DNA markers were adopted

since early 1900 for the genetic development of poultry (Thiruvankadan and Prabakaran 2017).

Evaluation of Breeding Program Under Particular Climatic Conditions

Since the phenotypic traits depends on genotype (G) and environmental factors (E), the environmental impacts also are necessary to emphasize during choosing the poultry stocks, and to confirm breeding policy. The vital objective of a breeding scheme is to establish a profitable crossbred animals are able in production at specific climatic conditions. For this reason, the output of breeding program requires to be focused under unlike climatic situation before its profitable use. The interaction between microenvironment and macro-environment and the genotypes is very crucial among inter-population and intrapopulation and. However, feeds, temperature, and climate changes between locations and seasons are the components of macro-environments. Moreover, unusual environmental variations are considered as a part of micro-environments within a population (Thiruvankadan and Prabakaran 2017).

Artificial Insemination and Cryopreservation

Application of conventional biotechnology method like artificial insemination (AI) in poultry production has supported to a quick transfer of hereditary traits from a lesser quantity of superior males to a greater quantity of females where AI could ensure excellent fertility in poultry relating to natural service. Effective uses of this method requirements better quality semen to be inseminated to the exact location of sperm preservation tubules in the female reproductive organ to ensure chances of best fertility in chicken (Mohan et al. 2018). However, liquid nitrogen can be used as cryopreservation in domestic birds for future using in AI. Since the last decade of the twentieth century, appropriate methods of cryo-storage of chicken semen of different breeds have developed. The techniques using the cryoprotectants glycerol, sulfoxide, dimethyl acetamide and dimethyl, slow or fast freezing-thawing procedures, pellets/vials/straws packaging have been established (Blesbois 2007).

8.3.3.2 Molecular Techniques Used in Poultry Production

The introduction of the era of molecular genetics in 1970s providing novel prospects to improve breeding programs via the application of DNA markers related to the traits of interest. As advance of whole genome studies (WGS) in genomics science is used in many biological systems like poultry breeding (Lander and Weinberg 2000). Various tools namely quantitative trait loci (QTL), candidate gene technique and high-density SNP genotyping for whole-genome selection, identification and genome wide scans, are widely applied nowadays in poultry genomics (Saxena and Kolluri 2018; Burt 2002).

QTL Classification and Genome Wide Scans

A QTL is defined as the genetic control of quantitative characters is foreseeable to be scattered all over the genome especially various locations of the genome that regulate the quantitative characters of interest. The specified experimental crosses use to confirm the QTLs particularly designed for the scope. The documentation of QTL and formulation of DNA tests were the key phases at applied uses of QTL through marker-assisted selection (MAS) as the selection based on an arrangement of information resultant from genetic markers linked with QTL and the indication of phenotype. Above these QTL examinations had accomplished by 200–350 MS markers and mating between diverse breeds, such as heavy meat type poultry birds with specific inbred lines or lighter egg producing birds (Fulton 2014).

Candidate Gene Technique

The candidate genes are those genes with direct and great impact on the character of importance. Formation of past information of certain candidate gene could be expected to be accountable for an identified main hereditary impact. The dissimilarity in sequences in target gene are documented and finally different alleles are attached with deviation of trait(s). The genes are straightly associated with production characters namely growth hormone (cGH) gene, growth hormone receptor (cGHR) gene, insulin-like growth factor-1 (IGF-1), IGF-1R, myostatin, etc. had assayed candidate genes and genetic markers like InDel/Dels and SNPs identified (Nie et al. 2005; Fritz et al. 2004; Lau and Leung 2002; Amills et al. 2003; Feng et al. 1998). Moreover, three physiological candidate genes viz. growth hormone genes for cGH, neuropeptide Y (NPY) and gonadotropin-releasing hormone receptor (GnRHR) were studied to detect their association with few production parameters like egg production/bird, dual yolk egg, and age at first laying egg (Ngu et al. 2015; Xu et al. 2007).

High-Density SNP Genotyping for Whole-Genome Confirmation

Advancement of NGS techniques and high performance of genotyping platforms has directed to the construction of high-density SNP array as an upgraded tool applicable for analyses of genetics and genomics in domestic animals. The important applications of SNP arrays in agriculture for upgrading of significant economic traits through genomic selection (Fan et al. 2010). An innovative type of marker-assisted selection (MAS) focuses on total markers in whole genome (Calus 2010; Hayes and Goddard 2001). It exactly forecasts the breeding values of animals using the data linked to the location of plentiful SNPs throughout the genome with a hypothesis that these SNPs are linked disequilibrium (LD) relations between QTL and SNPs. An abundance of SNPs basically needed for the outline and structure of arrays and could be retrieved by various techniques and platforms, for example,

genome sequencing and HapMap project, and reduced representation library (RRL) generate predicted SNPs sequencing, (Amaral et al. 2011; Matukumalli et al. 2009) via extracting SNP data from dbSNP of NCBI (<http://www.ncbi.nlm.nih.gov/projects/SNP/>).

8.4 Biotechnology in Poultry Feed and Nutrition

Innovation in biotechnology since the last decades has provided new prospects for improving the output and efficiency of animal production through ensuring improved feed and nutrition. At present, biotechnology have a comprehensive uses in animal feed and nutrition or formulation of feed-grade enzymes for availability of nutrients (Havenstein et al. 1992). Use of beneficial microorganisms (prebiotics and probiotics) made by recombinant DNA technology has an immense influence on poultry nutrition and production. However, food fermentation technologies ensure various nutrients, complete proteins/essential amino acids or to increase the digestibility of feeds that enhance animal production. As well as, recombinant bacteria produce improve nutrients through production of particular enzymes and hormones that can improve animal productivity including reduce environment pollution. Besides, uses of fiber degrading enzymes to increase animal production and decrease environmental impact (FAO 2010). A few of the aforementioned approaches are currently being used shown in Table 8.1. However, others are identified as likely to be potential, due to technical glitch, and food safety and public health issue are not applied at wider scale (Table 8.2).

Table 8.1 Biotechnological approaches are currently used in animal nutrition adapted from FAO (2013)

Approaches/ technologies	Objectives
(1) New ingredients	Produce microbial proteins as a novel feed source for livestock feed and nutrition, for example protein derived from single cell and yeast
(2) Designer ingredients	Improve nutrition status e.g. high-oil enriched maize, high-methionine enriched lupins Decrease the status of anti-nutritive ingredients in a common feed stuff, like low-phytate maize.
(3) Feed additives	
(a) Antimicrobials	Diminish the growth of pathogenic bacteria and to support to create a desired equilibrium of gastro intestinal flora like antimicrobials.
(b) Crystalline amino acids	Increase the feed supply of particular amino acids and to enhance the protein balance in formulations of animal ration.
(c) Feed enzymes	Improve the accessibility of nutrients like energy, phosphorus, amino acids, etc. in feed components by lessening the detrimental effects of anti-nutritive ingredients like xylanases actions on arabinoxylans in wheat and microbial phytases actions on phytate,

Table 8.2 Biotechnological approaches are future potential in animal nutrition adapted from FAO (2013) and Harlander (1990)

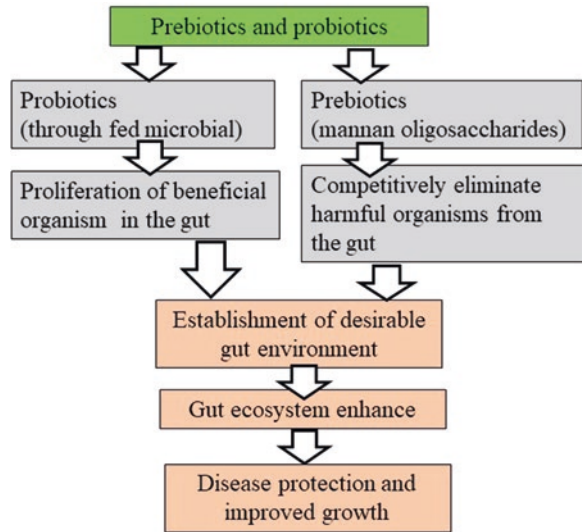
Approaches/ technologies	Objectives
(1) Adaptation of gut microbes	Modify microorganisms genetically the existing naturally into gastro intestinal tract, to improve their ability for desired functions or to include new functions
(2) Inclusion of novel gut microbes or bacteriocins	Inclusion novel genre of microbes into the gastro intestinal tract. Broad spectrum bacteriocins could be produced through engineering of nonpathogenic <i>Salmonella</i> spp. or new gut microbes. These strains could be included in feed and water for biological control of poultry pathogenic of <i>Salmonella</i> spp. or microbes in the gut.
(3) Bioactive peptides	Enhanced growth and efficiency like growth hormone-releasing peptides, better gut function, immunomodulation
(4) Antimicrobial replacers	Antimicrobial enzymes (like lysozyme), to deliver specific antibodies via spray-dried plasma and egg substances
(5) Transgenesis	Alter nutrient metabolism and enhance growth effectiveness by removal of genes.

8.4.1 Probiotics and Prebiotic

The role of biotechnology in poultry nutrition is enormous nowadays, as it shows the pivotal role in the poultry feed production. Since a wide reports on emergence of antimicrobial resistance (AMR) among the pathogens, the antimicrobial uses (AMU) as a protective measurement found to be flawed. In this situation, impact of antibiotics uses as growth promoters or therapeutic agents has created the alternative options where both consumer and producers will be gained. Therefore, probiotics was identified to cover this space. Nowadays, many farmers use them as antibiotics alternatives (Griggs and Jacob 2005; Nava et al. 2005; Trafalska and Grzybowska 2004). The probiotics organisms are used as gut ecosystem enhancers that improve to create desirable gut environment through proliferation of beneficial organism (fed-microbial). The prebiotics organisms direct eliminate harmful pathogens competitively (FAO 2013) and to improve FCR (Midilli et al. 2008) and improvement of host's health by particularly accelerating the growth and/or function of probiotic bacteria prevail in the intestinal tract (Borroto 2009) like Mannan oligosaccharides (Fig. 8.5).

Poultry nutritionists are constantly toiling their attempts into better feed formulation considering economical point of view. A “good feed” only not fulfill the target but also ensure substantial utilization. Nutritional modifications along with healthy feed can impact on equilibrium of the microflora in the intestine that prompting indigestion. Therefore, an optimum balanced ration with energy and nutrients components is needed to keeping a healthy gut. A lot of concerns have recently been highlighted from veterinary consultants and poultry nutritionists relating to proper utilization of nutrients along with uses of probiotics for growth promotion in poultry feed (Kabir 2009a, b).

Fig. 8.5 Prebiotics and probiotics act as a gut ecosystem enhancer adapted from FAO (2013)



Probiotic is a broad term, are the products may contain culture bacteria, yeast cells, or together that influence microorganisms to adjusting the gastrointestinal tract related to health condition and enhance feed efficiency (Dierick 1989). In many ways these can affect in poultry are comprised of (1) keeping standard intestinal microflora by antagonism and competitive exclusion process (Kizerwetter-Swida and Binek 2009; Kabir et al. 2005; Schneitz 2005; La Ragione and Woodward 2003; Stern et al. 2001; Fritts et al. 2000; Netherwood et al. 1999; Jin et al. 1998; Line et al. 1998; Nisbet et al. 1998; Owings et al. 1990; Fuller 1989; Nurmi and Rantala 1973; Rantala and Nurmi 1973), (2) modifying metabolism through intensifying digestive enzyme activity and lessening ammonia production and bacterial enzyme action (Yoon et al. 2004; Han et al. 1999; Chiang and Hsieh 1995; Jonvel 1993; Cole et al. 1987), and (3) enhancing feed intake and digestion capacity (Awad et al. 2006; Horniakova 2005; Kumprecht and Zobac 1998; Yeo and Kim 1997; Tortuero and Fernandez 1995; Nahashon et al. 1993; Nahashon 1992; Dierick 1989), and finally (4) boosting the immune system of the host (Apatá 2008; Brisbin et al. 2008; Nayebpor et al. 2007; Mathivanan and Kalaiarasi 2007; Haghighi et al. 2006; Haghighi et al. 2005; Dalloul et al. 2005; Kabir et al. 2004; Huang et al. 2004; Koenen et al. 2004). Therefore, the uses of probiotics in commercial broiler production even in low level of biosecurity in rural settings (Fig. 8.6).

8.4.1.1 Impact on Intestinal Microbiota and Intestinal Morphology

Probiotic supplementation in poultry feed have been observed as beneficial effect on growth performance at the different stages of age without relatedness of vaccination status (Awad et al. 2009; Sahin and Yardimci 2009; Roshanfekr and Mamooee



Fig. 8.6 (a and b) Uses of probiotics for safe broiler production that enables to no or minimum antibiotic uses under sector 3 (small scale commercial) poultry production system in rural Bangladesh with low/minimum level of biosecurity standard

2009; Willis and Reid 2008; Apata 2008; Willis et al. 2007; Nayebpor et al. 2007; Mountzouris et al. 2007; Khaksefidi and Ghoorchi 2006; Timmerman et al. 2006; Kamruzzaman et al. 2005; Gil De Los Santos et al. 2005; Hossain et al. 2005; Kabir et al. 2004; Islam et al. 2004; Kalavathy et al. 2003; Zulkifli et al. 2000; Jin et al. 1998). However, some changes in probiotics through inactivation by a high-pressure homogenizer that have a decisive impact on growth performance of broiler chicken as additives in concentrated feed (Huang et al. 2004). Use of selective probiotics like FM-B11 in turkeys was found to increase the usual daily weight gain and salable body weight to be considered as a financial benefit to better turkey production (Torres-Rodriguez et al. 2007).

8.4.1.2 Impact of Probiotics on Immune Status

The immune response was found to be significantly higher in probiotic administered birds than other birds. The variation of the weights of different organs like bursa and spleen in probiotic use and usual feed in broiler production could be confirmed at the different stage of antibody production in reaction to sheep erythrocytes (SRBC) (Kabir et al. 2004). In the same way, it was confirmed as antibody titer was suggestively higher after post-immunization (PI) at 5 and 10 days with supplementation of probiotic 50 mg/kg group, related to control (Khaksefidi and Ghoorchi 2006). The use of probiotics improves serum and intestinal usual antibodies against a few foreign antigens in poultry (Haghighi et al. 2005). A several studies confirmed the possible impact of probiotic on adjustment of immune response at a desired level in poultry (Apata 2008; Nayebpor et al. 2007; Mathivanan and Kalaiarasi 2007; Haghighi et al. 2005; Dalloul et al. 2005; Koenen et al. 2004; Zulkifli et al. 2000).

8.4.1.3 Impact on Improve Water Quality and Improve Microbial Balance

Probiotics organism directly play role to improve water quality and microbial balance as their presence in the poultry gut (Lalloo et al. 2007, Hill et al. 2009, Picchiatti et al. 2009) Supplementation of probiotics in broiler feed enhances quality of meat both at pre and post freezing storage time (Kabir 2009a, b; Kabir et al. 2005). Additionally, the physical appearance includes consistency, lusciousness and overall appropriateness of poultry meat substantially improves through probiotic (Lacto-Sacc) feeding as it declines total viable count (TVC) in poultry meat compared to untreated birds (Mahajan et al. 2000) or even meat tenderness could be enhanced by application *Saccharomyces cerevisiae* extract or whole yeast or (Zhang et al. 2005).

8.5 Biotechnology in Poultry Vaccine and Biologics

Immunization is an important method for maintaining animal health targeted for prevention and control of animal disease has been practiced for centuries and confirmed as a potential devise to lower animal suffering including the economic benefit for animal producers. A wide number of factors like capacity of national disease eradication programs and international trade policies along with cost-effectiveness of livestock and poultry production are responsible on availability of vaccines and biologics for disease control program. At present time, the techniques used in the development of vaccines have extended quickly due to improved understanding of the mechanisms at which protection is achieved, and the accessibility of genomic information on both hosts and pathogens (OIE 2010). Globally around 900 patent applications of poultry vaccine claims annually involving molecular techniques. The combined development of new tools in the area of molecular biology and immunology has a great influence on the formulation of novel vaccines including their quality. This is able to model of vaccines designed for the control and eradication of particular disease considering national, regional, and international aspects. As a cost-effective measure to control infectious diseases, vaccination has been proved widely for eradication of animal diseases. Nevertheless, a limited recombinant vaccines has been produce for commercial use, their utilization is insignificant in developing countries.

8.5.1 Reverse Genetics

As development of a reverse genetics system the area of virology through different techniques namely designed mutations, deletions and insertions into the viral genome of live viruses of different DNA and RNA viruses has been modernized. This is applied in a range of applications include attenuation of virus, the host

specificity alteration and the generation of replication-deficient viruses. These protocols have been used to the development of new vaccines and are extensively used in the characterization of the function and structure of distinct viral genes and their sequences. The production of a cloned copy of complementary DNA (cDNA) from RNA by reverse transcription *in vitro* is widely used in reverse genetics; subsequently transfection in the permissive cells with the cloned DNA(s). The tool was first recognized in a positive-strand RNA virus termed as bacteriophage Q-Beta (Taniguchi 1978) and also has been used for formation of RNA viruses with segmented/unsegmented negative sense genomes, e.g. vaccine development against avian influenza virus (AIV) included an engineered haemagglutinin (HA) gene from H5N1 virus and a neuraminidase (NA) gene from a H2N3 virus, with in a H1N1 skeleton (Meeusen et al. 2007). Thus inactivated H5N3 virus vaccine was produced that confer full protection in birds against H5N1 challenge.

8.5.2 Recombinant Vector Technologies

Progresses in genomics, reverse genetics, and proteomics have enabled to the documentation of process of pathogenicity, host-pathogen action along with protective antigens from many organisms, and the advance of appropriate vector for transfer of these antigens to the host animal. The availability of bacterial and viral genome sequences has improved the quick structure of distinct deletions in the genomes of a diverse range of pathogens is not only the effect of attenuation, but also it makes place for the insertion of imported antigens coding genes for of heterologous microorganisms (OIE 2010). Normally, for virulence, deletion of genes attenuate bacterial vectors is mandatory for the important metabolic processes of organisms. *Salmonella* utilized as a role of vector for vaccine antigens delivering and finally applied for the production of live vaccine strains to be used poultry (Babu et al. 2004). There are numerous bacterial vectors are developed using commensal microorganisms like *Lactococcus*, *Lactobacillus*, *Streptococcus*, and *Staphylococcus*, etc. However, pathogenic organisms are being used through attenuation namely *Shigella*, *Vibrio*, *Yersinia*, *Bacillus*, *Bordetella* and *Cornebacteria*, etc. as these are all evaluated for the capacity of production immunity at protection level.

The viral vectors were developed using those virus causing mild or no disease or even pathogenic viruses through deletion of virulence genes attenuation. Replication capacity of virus vectors can replicate progeny virus together with defective virus vectors replication and assessed as vaccine delivery vehicles. There are numerous commercial vaccines have been developed stand on DNA virus vectors technology and effectively approved for using in livestock and poultry practice like poxviruses and herpesviruses (Gerdts et al. 2006). However, a few viral vectors namely vaccinia virus, fowl pox virus, canary pox virus, and turkey herpes virus. Moreover, a number of viral vectors have been advanced or further evaluation is under way towards improvement. The information of certain virulence factor(s) of a pathogen and the accessibility of recombinant DNA technology has enabled the creation of

particular gene-deleted organism for the application as a live vaccine. The method is used for marking and assaying of defined gene deletions benefits in lowering the virulence without affecting their immunogenic potentiality. Such organisms could be used as vaccines as they have the immunogenic characteristics, however they could not perform as causative agents of disease (OIE 2010). A several live attenuated vaccine strains of bacterial disease have been developed through this approach and found to be genetically potent, safe for use and make better immunity compare to killed vaccines, for example gene-deleted *Salmonella enterica* serovar *enteritidis* and serovar *typhimurium* and vaccines in poultry (Meeusen et al. 2007; Babu et al. 2004).

Subunit vaccines Since 1980s, using recombinant DNA technology semi-pure or purified proteins subunit vaccines produced and available commercially as (Ulmer et al. 1995; Rhodes et al. 1994; Cohen 1993; Ulmer et al. 1993). However, this vaccine do not uses live recombinant vector techniques that will deliver the recombinant proteins in vivo. Using the conventional biochemical or recombinant DNA technologies the subunit antigens are produced (OIE 2010). A range of prokaryotic and eukaryotic expression systems of insect cell, yeast, and plants are used (Chichester and Yusibov 2007) via a variety of combined or transient expression system. Biochemical methods are still helpful in some cases however the recombinant expression is not suitable, such as antigens required intricate assembly (like fimbriae), or at the stage of post-translational modification.

New vaccination strategy for immunization with DNA is founded on a clear-cut idea. The vaccines can be described as antigen-encoded bacterial plasmids are able to make specific immune reactions after inoculation into an appropriate host. Immunization is completed through the receipt of purified plasmid in the host cells and it remains outside of the chromosome.. From a technical perspective, DNA vaccines can be easy to engineered, produced and purified, structured and evaluated in animal models within months be. DNA vaccines are considered to be very stable for a longer time or even carried without a cool chain to a long distance. The safety of this vaccine has been evaluated in various field trials in different animal models as well as in humans (Kim et al. 2001; Bagarazzi et al. 1998).

8.6 Biotechnology in Disease Diagnosis

Biotechnologies have applied in livestock health practices to increase the accuracy of diagnosis of disease and treatment. Immunology based diagnostic conventional method are being used monoclonal antibodies like enzyme-linked immunosorbent assays (ELISA) and radioimmunoassays (RIA) along with microscopic examinations are not sufficient to distinguish between species and subspecies of disease producing pathogens (Maharana et al. 2016). However, these techniques may not suitable for distinguishing of vaccinated from infected animals (DIVA). Molecular diagnostic assays including single DNA sequences give high levels of sensitivity,

specificity and even reliability in tests. Polymerase chain reaction (PCR) technique permits significantly higher level of specificity (Salih and Trifonov 2015).

The approaches in molecular biology discriminate targeted DNA sequences are now well familiar tools for detection of infectious agents (virus and bacteria) in poultry and other animals and widely used throughout the world even in developing countries. Mostly, PCR-based diagnostic assays are progressively used for timely diagnosis of causal agent of diseases, however, these are being used primarily in the labs of research organization and central national diagnostic laboratories due to incompatible facilities for diagnostic test (FAO 2010).

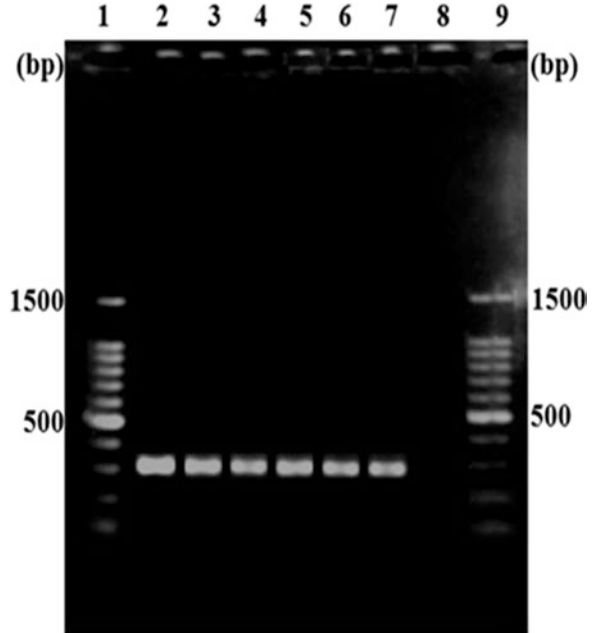
8.6.1 Polymerase Chain Reaction (PCR)

The PCR has an enormous impact on avian health. PCR can be used in diagnostic tools because of its higher specificity to detect microbial pathogens in clinical samples and epidemiological studies (Yamamoto 2002; Nagy 2001). This is an effective, simple and economical technique applied to amplify the preferred DNA sequence or sequences of into billions of identical copies. This assay is basically used for checking Short Tandem Repeats (STRs) and Single Nucleotide Polymorphisms (SNPs) in different livestock breeds, genetic mutation including parentage identification. The key component of the PCR are: (1) DNA template, (2) primers, (3) Taq polymerase enzyme, and (4) the PCR machine (Thermocycler) that devises the correct temperature for each step in every cycle. The PCR technique involves three major phases in each cycle that include; (1) denaturation, (2) annealing and (3) extension or elongation (Bartlett and Stirling 2003). To confirm as the PCR produced anticipated DNA fragment, agarose gel electrophoresis is often used for the confirmation of the sizes of PCR amplicons. The size of PCR amplicon is estimated by connection with a molecular weight marker (DNA ladder), that contains DNA fragments of known size, run on the gel quickly with the PCR amplicon (Jebakumar et al. 2012). In our most recent study, we had used *cdtC* gene-based multiplex PCR assay for detection of *Campylobacter coli* contamination in broiler meat samples (Neogi et al. 2020) (Fig. 8.7).

8.6.2 Bioinformatics to Diseases Diagnosis

In bioinformatics, biological data are organized and understand using upgraded computing techniques (Akunaedozi 2014). Bioinformatics techniques and relevant algorithms have now been developed with an aim to use the sequence of biological information. Through the bioinformatics tools viz. BLAST (Basic Local Alignment Sequence Tool), Fast Alignment (FASTA) and Clustal W solutions for sequence search and analysis, together with other techniques enable to a cost-effective and time reduction strategy to obtain vital data on gene and protein levels, however,

Fig. 8.7 *cdtC* gene-based multiplex PCR assay used for the detection of *Campylobacter coli* isolated from broiler meat samples of live bird market (LBM). Lanes: 1 and 9, 100 bp DNA ladder (Promega, USA); lane 2, positive control (*C. coli* ATCC33559); lanes 3–7, representative positive isolates; lane 8, negative control



these could not be achievable by other methods (Kaikabo and Kalshingi 2007). The phylogenetic analysis of the amplified nucleic acid sequences gives new information on the evolution of organisms and assists the studies of molecular epidemiology. Proof for interchange of DNA among different microorganisms with evolutionary ladder is reconsidering our knowledge on pathogens and reconfirming the classification. Nonetheless, the view of using this technology depends greatly on the accessibility of bioinformatics information. Present studies in Belgium, Western Europe confirmed the value of Next Generation Sequencing (NGS) technologies to veterinary science, with special emphasis for using in diagnosis of diseases and treatment capacities (Van Borm et al. 2015).

8.7 Biotechnology Application in Food Processing

Food processing sectors nowadays are using biotechnological tools for improvement of food products. The impacts of these techniques comprise increase food production, enrich the nutrition parameters, different food products produced through fermentation process, production of important enzymes, increase shelf life of the food, enhancing the organoleptic characteristics of food, and to improve safety compliances in food and food products (Lokko et al. 2018; Nguyen et al. 2017).

8.7.1 Increase Food Production

The use of transgenic technology is expanding poultry productions. The huge financial involvement of \$38 billion in the United States at the commercial sector importance of biotechnology is evident. The aim for transfer of gene in commercial poultry is mostly to improve hereditary superior poultry birds for production of meat and egg, which has been found to be the same target of conventional selection programs taken during last 50 years (Petitte and Mozdziaik 2014). The conventional selection program has been discussed elaborately in the subsection “selection process” under section biotechnological approaches in poultry breeding programs.

8.7.2 Enhance Nutritional Parameters

The idea of biotechnology to boost nutritional quality in food was first introduced in Japan during 1980s as targeted to certain health benefits spaced out from its own nutritional quality (Cruickshank 1934). There have been continuous interest in poultry biotechnology for changing the egg constitutes through manipulations of genetic and nutritional traits for human health advantages. This could be done by transforming cholesterol composition and its concentrations, fatty acids, lipids, amino acids, and minerals or by inclusion of therapeutic pharmaceutical elements (Alagawany et al. 2018). The designer egg approach was first introduced in 1934 by Cruickshank, who affirmed the adjustment of fatty acid contents in egg yolk by feed interventions (Arai 1996). Nowadays, designer eggs offer for vegetarian as immunity stimulant, safe or organic foodstuffs, better vitamins and minerals with adjusted ratio of fatty acids namely omega-6 and omega-3, lower level of total cholesterol, boost of antibodies including essential pigments like carotenoids (Alagawany et al. 2018). However, researchers had advanced to a designer egg with rich of antioxidants and omega-3 fatty acids (Sim and Sunwoo 2002). Further improvements in designer egg which were confirmed with enriched conjugated linoleic acid (CLA) (Raes et al. 2002), vitamin A and β -carotene content (Jiang et al. 1994). Chicken, beef and pork enriched with selenium (Se) can also be produced through dietary feed supplementation of organic Se in food animals and birds (Fisinin et al. 2009). Currently, the designer food or functional foods are receiving greater attention due to their role in health benefits and disease prevention (Fisinin et al. 2009).

8.7.2.1 Fermentation Process to Food Processing

Since hundred years ago, genetic upgradated microorganisms like bacteria, yeasts and molds have been used for the production of fermented foods. Conventional strain improvement techniques related to mutation and selection are erroneous and unmanageable. However, it is difficult to screening for all mutations that happen

and screening their process is challenging. Therefore, gene engineering keeps a tool to overcome several of these challenges as it allows for the selection to transfer of a single, particular defined traits from basically any living organism in an exact, manageable way. Suppression of pathogens and spoilage microorganism through 'Bacteriocin' production in poultry meat sausage is a unique example of genetic improvement of food-grade microorganisms in fermented food (Harlander 1990).

To yield different value added fermented food staffs through commercial fermentation processes, the starter cultures have been created for using either a single or mixed strains of organisms presented to be beneficial (Holzapfel 2002). Inhibitory action of these microorganisms was observed for the creation of one or several products like bacteriocins, hydrogen peroxide, organic acids and diacetyl (Hutkins 2006). Moreover, *Lactobacillus* strains can be applied as essential probiotics for the processing of meat and dairy products, and having abundance of health benefits (Pennacchia et al. 2006). Therefore, the biotechnological technologies can be fitted to obtain better-quality strains of microorganisms (bacteria, yeast, and moulds) and could be used for the processing of meat and meat products through fermentation.

8.7.3 Increase Shelf Life of Food

The shelf life of beverages and food can be prolonged through bacterial fermentation process (Brochu 2018; Paramithiotis 2017). Most of the food fermentations include lactic acid that is produced from sugars by the lactic acid producing bacteria under the genera of *Lactococcus*, *Streptococcus*, *Lactobacillus* and *Pediococcus*. However, nowadays, *Lactobacilli* have been considered due to the production of bacteriocins (Collins et al. 2017). These substances could be used food industry as natural preservatives to increase shelf life. The uses bacteria producing lactic acid and of their metabolites are recognized as safe (Patel and Prajapat 2013). Nisin is a bacteriocin used officially in the food industry and recognized globally (Kaškonienė et al. 2017).

Numerous processed food products in refrigerated vacuum-packaged of meat, fish, dairy, and vegetable category contain usually gram-positive bacterial strains under the genera *Brochothrix*, *Lactobacillus*, *Clostridium*, *Leuconostoc*, *Carnobacterium* (Marth 1998). They can able to spoilage of the product as they multiply at refrigerated temperature. By adding could be lessen. Through addition of *Lactobacillus sakei* culture in chilled raw ground meat and chilled cured pasteurized sliced vacuum-packed meats the *Listeria* count could be decreased, therefore, this facilitates to increase shelf life (Devi and Halami 2011; Hugas et al. 1998; Ennahar et al. 1998; Yang and Ray 1994; Kroeckel and Schmidt 1994).

8.7.4 Enhance Organoleptic Characteristics of Food

The food organoleptic properties can substantially have influence on acknowledgment of food and food products by the consumers. The genetics and genomics methods (discussed in Sect. 8.3.3) play a key role in the development of quality of product primarily for the nutritional and user friendly technological aspect and some for organoleptic properties (Smaldone et al. 2017). The cultured microbes used in food production are generally termed as starter cultures that also improves the organoleptic character of different foods stuffs. An earlier study has confirmed that above 100 commercial aroma chemicals are produced through biotechnological approach (Berger 2009). The foods produced through fermented process are found to be more value addition with higher nutrients with extend shelf life, and easy digestible as these are more appropriate for the digestive tract (Smaldone et al. 2017). The organoleptic properties of such meat are greater regarding taste, flavor, aroma and also color (Singh et al. 2012). Some aroma producing volatile compounds like acetic, acetaldehyde, diacetyl, ethanol, acetone, etc. were formed from carbohydrate during fermentation process of sausage (Awan et al. 2003).

8.8 Biotechnology and Food Safety

At present time biotechnology as a tool to surveillance on food safety, to avoid food-borne illnesses and to confirm traceability of the foods. In most present time a wide biotechnological developments have set common accessibility of tools of identification with rapid and economic than the conventional methods PCR and ELISA based methods are nowadays used for the detection of key food-borne microorganisms (Velusamy et al. 2010). Biotechnology has involved with the food safety in poultry production system in following areas:

8.8.1 Biotechnology Critics Get Priority to Food Safety

The outputs of biotechnology as new food animals, animal origin foods or processed food to fulfill certain desired goals. The animal biotechnology research emphases on producing transgenic poultry will use feed effectively, gain growth to slaughter weights at a minimal age, and can able to prevent a wide range of diseases. Therefore, farmers get more benefits from lower production costs, improved productivity with higher quality products. Consumers get benefit as farmers can provide leaner meat and poultry grown with a less dependency on vaccines, medicines. Additionally, consumers prefer to get the products at a lower price as the farmers paid less in a shorter period of time. The biotechnologically animal products and their associated events are exciting, however, likelihoods of public concerns and

worries on food safety issues are enormous. The concern for all foods or food products regarding to free of hazards like chemical or biologic that may affect safety of food for the human consumption. Consequently, food safety has got priority as an easy focus area for biotechnology critics. Food safety has been adopted in most recent time in developing countries, conversely, the consumers as well as mass people are not completely confident on poultry and poultry origin food as the production and processing along the value chain associated with biotechnical tools that creates public misperception. So, this is to be needed to take remedial actions properly through generating public awareness and motivation towards building confidence on biotechnologically poultry or poultry goods.

8.8.2 Food Safety Measures in Poultry Production

8.8.2.1 Protect Poultry Flock with Good Biosecurity

Implementation of good biosecurity measures to prevent the entry of pathogens into the farm and prevent the micro-organisms from transmission within the farm or from the farm if it enters and infects the poultry (Yap 2015). The biosecurity measures include the facility of fencing around the farm, netting in farm sheds, vehicles are cleaned and disinfected before entry of the farm, footwear to be clean before entry in the farm (Mridha et al. 2020; Akhter et al. 2018; Ratananakorn and Wilson 2011). Use safe production inputs like quality raw material, feed and day old chicks (DoC) and water that should be free of biological hazards (*Salmonella*) for production of safe poultry and poultry products.

8.8.2.2 Good Animal Husbandry (GAH) Practices

Good animal husbandry practices should be maintained at the farm level that include, vaccination against poultry diseases considering the country context, practice separation of age group e.g. should ensure buffer zones of >30 m, or with all-in-all-out practice in broiler rearing, suitable stocking density to minimize stress to poultry (1.6–2.0 sq.ft. per bird considering size of broiler). Use of probiotics for competitive exclusion to prevent the establishment of pathogens (discussed in Sect. 8.4.1) and to isolate or quarantine any sick poultry for treatment and observation, veterinary inspection of all poultry before marketing to certify on good health status. However, proper record keeping and documentation of farm activities are very much needed (Yap 2015).

8.8.2.3 Good Personal Hygiene (GPH) Practices

Practice good personal hygiene (GPH) to be maintained through ensuring that workers do not transmit any infectious disease e.g. *Salmonella* to poultry and vice versa, cut and wounds may be infected with *Staphylococcus aureus*, washing of hands after toilet use and handling live poultry or poultry waste, and before entering poultry sheds; use of personal protective equipment (PPE) like boots, gloves and mask during working in poultry sheds or cleaning poultry sheds; and poultry attendants who are sick, having cuts or wounds or immunological sick should not work in the farm.

8.8.2.4 Poultry Waste Management

Environmental friendly poultry waste management is very much needed to subsequent exposure disease producing pathogen from poultry to poultry or human. These practices involve good poultry waste & environment management practices e.g. bury poultry waste with lime, compost or incinerate and practice good pest control (Yap 2015). Poultry waste can be used for biogas production by large digesters can be an important source of energy to generate electricity or green energy for vehicle. Outside energy generation, the digester is a good source of biomaterials. A feather-degrading bacterium and the enzyme keratinase present in this biomaterials is beneficial to decompose feather meal and to enhance feed digestibility and facilitates to damage prion proteins (source of bovine spongiform encephalopathy) (Shih 2012). However, in low resource settings, poultry manure and including other organic waste materials can be transformed through anaerobic condition using effective microorganism (EMO) through Bokashi method to a more stable form within shorter period of time (2 weeks) to use as a soil improvement as organic source of fertilizer in vegetable gardening or crop cultivation (Fig. 8.8). The process



Fig. 8.8 (a) Demonstration of preparing compost of poultry manure using EMO in the course of an environment friendly procedure; (b) the compost is ready for use within shortest period of time (2 weeks) for vegetable gardening

is called ‘composting’ and the end product that is produced via composting is called ‘compost’ (Earth Care 2019). These technologies could be considered to utilize poultry waste through environment friendly waste management at low resource settings since the poultry drooping creates a huge nuisance nowadays among people as these are been dispose at open field without treatment.

8.8.2.5 Biotechnology Directs Food Safety

Unpredicted and unintentional composition variations occur with all forms of biotechnological foods. The bacteria utilized for feed production or other purpose as part of biotechnical process might be a hazard to human and animal health due to presence of resistance genes (EFSA 2007). To keep as satisfactory standard of food quality along with safety is completely obligatory for ensuring protection for consumers and to ease trade. Surveillance on microbial adulteration in the finished goods including the different segments of production value chains, cleaning and sanitation practices are the utmost priority factors of the manufacturing process in food technology and also in biotechnology (Van Belkum et al. 2017). More sensitive and specific tools to be applied for determination of microbial food impurities and their toxins by state-of-art technologies of proteomics and genomics. Several commanding techniques of biotechnology have now set massive advances like genetic engineering, PCR, random amplified polymorphic DNA, rDNA technology, amplified fragment length polymorphism, matrix associated laser desorption ionization-time of flight mass spectroscopy (Naveena et al. 2017). These techniques can also support to meat verification. However, development of new technique for rapid confirmation of high-risk food borne emerging pathogens in livestock food commodities is an appealing requirement in food safety connection.

8.9 Constraints and Limitations

Using new tools and techniques for disease prevention and control activities like novel and effective diagnostic methods, drugs including vaccine and biologics in low resource settings are not considered to be bottlenecks rather than to ensure their availability to the rural communities is important. However, there are several issues to be converged as limitations of biotechnology, and these are to be overcome to maximize optimum benefits from this discipline are as follows:

8.9.1 Animal Welfare and Ethical Issues

Biotechnology is considered to be indispensable part of the upcoming veterinary medicine, and husbandry practices. However, these novel technologies have involvement with animal species deserves an ethical obligation for an effective uses and the recognition. There are possible unexpected risks may result with the incredible benefits. Thus, ethical approval and animal welfare issues can arise at different phases of multiplication and longevity of a particular genetically transformed animal. A theory of “Three Rs” (reduce, refine, and replace alternatives in the use of animals in experiments) had been undertaken by the Canadian Council of Animal Care (CCAC) with an agreeable ethics of animal (Avey and Griffin 2016). Firstly, they do not consent for challenges to a scientist’s purpose in operating research with animals, even if that purpose is doubtful. A second point, gaps in the Animal Welfare Act have permitted researchers to keeping aside application of the Three R’s in action. Finally, the Three R’s have no application to novel and emerging issues of biomedical research that have potential to use of animals in experiments, comprised of research of stem cell, genetic modification, cloning, bioterrorism defense and xenotransplantation (Ibrahim 2006).

8.9.2 Lack of Available Data Like Traceability, Risk Assessment of Animal Biotechnological Foods

Available enriched foods to market necessitates the logical guarantee on safety concerns. Animal products derived through biotechnology are likely to be safe, however, that safety should be confirmed and documented as traceability to the safeguard of public health and support to public confidence (Crawford 1990). A system is to be developed concerning how biotechnology in poultry affect the meat and poultry industry will be scrutinized and assessed, and to share the findings/recommendations all scientific aspects have been taken into consideration. This is to be needed to building trust and confidence among mass people on biotechnologically developed poultry goods.

8.9.3 Lack of Consciousness Among the Mass People

The public concern for all foods or food products relating to free of agents like chemical or biological agents that may affect safety of the food for the human that got priority as an easy focus area for biotechnology critics. Food safety has been adopt in most the countries, conversely, the consumers as well as mass people are not completely confident and creates public misperception on poultry and poultry origin food as the production and processing along value chain has gone through the

biotechnical approaches (Borroto 2009). So, this is to be needed to address remedial approaches properly towards generating public awareness and building trust on biotechnologically poultry goods.

8.9.4 Legal Consideration and Government Inspections

Due to lack of inappropriate legal issues like relevant regulations along with government inspection to confirmation of safety standard in biotechnologically foods that has declined consumer confidence (Borroto 2009). The government inspection streamlines to intensify food product testing is the single best technique to improve food safety (Crawford 1990). To overcome food safety bottlenecks, government, consumers and industry should recognize their individual accountabilities to ensure safe food. The inspection facilities must be suitable and science based, and propose for their assessment of biotechnology products.

8.9.5 Lack of Resources

Due to lack of adequate resources in terms of skilled manpower, advance laboratory facilities in developing countries, biotechnological approach do not utilize its full potentials, for example, advance disease diagnosis tools (RT-PCR) are being utilized at the central level (Ruane and Zimmermann 2001). However, government facilities are not adequate to inspect the food safety parameters poultry origin biotechnologically improved food. Therefore, resource allocation to be needed to optimize the benefits of biotechnology as well to ensure food safety.

8.10 Conclusion

As a new technology has commenced at a time when consumers are worried on the application of technology in the food production system. General people desire natural products are devoid of synthetic chemical substances. They also like nutritious, appropriate, enhance quality, good-packaged foods that require technology. Biotechnology will help give consumers what they want. Therefore, food safety choices must be stand on the state of art science based knowledge. Psychological and socioeconomic parameters are crucial but could not confirm the safety in biotechnology products. There are several key issues to review the current insights in this paper on assessing the safety of poultry origin biotechnology poultry, poultry products and genetically modified poultry that bring a desired gene along with ingredients applied to poultry meat and meat products with a biotechnologically made enzyme/flavoring/other food additive, or use of prebiotics/probiotics in

production of poultry as an alternatives of antibiotics or growth promotors. This current review confirms that the substance used in production and processing of poultry meat and its related food items are found to be harmless and effective in its scientific and intended use.

In conclusion, we have delved from an in-depth review that biotechnology will serve as modernization in veterinary science especially poultry production and processing of poultry origin food that interlinked with food safety parameters where applicable. Biotechnology will benefit not only to fight against bacteria that contaminate food but also this will be helpful to the appropriateness of the meat products through inclusion of fat components and to add nutrients to other products. Now it is clear that biotechnology is encouraging the people as its involvement in food production systems and to ensure food safety and security. Further, backing to strengthen of national abilities in the development and suitable application of pro-poor animal biotechnologies through intervention of international and donor agencies is demanding. The government with other partners of the food system and the biotechnology industry should utilize completely the science based technology to corroborate food safety standards and to stance at the back of public health decisions.

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