

Research Report

Open Access

The Relationship between Epigenetic Changes and Seasonal Changes in Rabbits

Mengshi Jiang ✉

Shangyu Anghua Biotechnology Co., Ltd., Shangyu, 312300, China

✉ Corresponding author email: 1624454379@qq.com

International Journal of Molecular Zoology, 2024, Vol.14, No.1 doi: [10.5376/ijmz.2024.14.0007](https://doi.org/10.5376/ijmz.2024.14.0007)

Received: 12 Jan., 2024

Accepted: 17 Feb., 2024

Published: 29 Feb., 2024

Copyright © 2024 Jiang, This is an open access article published under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

Preferred citation for this article:

Jiang M.S., 2024, The relationship between epigenetic changes and seasonal changes in rabbits, International Journal of Molecular Zoology, 14(1): 54-61 (doi: [10.5376/ijmz.2024.14.0007](https://doi.org/10.5376/ijmz.2024.14.0007))

Abstract This study explores the close relationship between epigenetic changes and seasonal changes in rabbits. Through in-depth analysis of the epigenome of rabbits in different seasons, the researchers found that there are significant differences in the epigenome of rabbits in winter and summer, indicating that seasonal changes may be a key factor driving epigenetic changes in rabbits. One. Further research revealed the impact of seasonal changes on rabbit gene expression, showing that the expression levels of genes related to adaptation to low-temperature environments increased in the cold season, while other genes showed different expression patterns in the hot summer. Research limitations are mainly reflected in the limited geographical scope and sample size. Future research can understand the epigenetic changes of rabbits in different seasons from a more comprehensive perspective by expanding the research scope and introducing more regional data. In addition, future research can also be expanded to spring and autumn, as well as under different climate conditions, to gain a deeper understanding of the impact of seasonal changes on rabbit epigenetics. Finally, this research is not only of great significance to understanding rabbit biology, but also provides some useful inspirations for ecology and agriculture, and provides theoretical support for animal protection and animal husbandry.

Keywords Rabbits; Epigenetic changes; Seasonal variations; Gene expression; Ecology

Epigenetics is a cutting-edge scientific field that studies gene expression and inheritance. It goes beyond traditional genomics and focuses on the exploration of non-coding genetic changes. Unlike changes in the DNA sequence itself, epigenetics focuses on non-coding changes that are closely related to DNA, including mechanisms such as DNA methylation and histone modifications. These epigenetic changes play a key role in regulating gene expression, thereby affecting an organism's traits and adaptability.

As a ubiquitous natural phenomenon on the earth, seasonal changes have a profound impact on the physiological processes of living organisms. Seasonal changes in climate, light, food supply, and ecosystems drive organisms to adapt to different seasonal environments. In this context, it becomes particularly important to study the relationship between seasonal changes and epigenetic changes. Many organisms show adaptive physiological and behavioral changes in the face of seasonal changes, and these changes may be realized through epigenetic regulation (An et al., 2021).

The main purpose of this study is to deeply explore the impact of seasonal changes on epigenetic regulation of organisms, using rabbits as the research object. As a mammal, rabbits have excellent reproductive capabilities and survival adaptability. They show obvious behavioral and physiological changes in different seasons, such as changes in reproductive behavior, food intake, hair color, etc. The relationship between these seasonal adaptations and epigenetic changes is a focus of researchers. By studying the epigenetic characteristics of rabbits in different seasons, this study hopes to reveal how seasonal changes regulate the expression of rabbit genes, thereby affecting their physiology, behavior and ecological adaptability. It is expected that this research will contribute to a more comprehensive understanding of the adaptation mechanism of organisms in seasonal environments and provide new theoretical foundations and application prospects for ecology, agriculture and wildlife conservation. By in-depth studying the epigenetic regulatory mechanism of rabbits as a model organism under seasonal changes, this study is expected to provide new insights into unlocking the mysteries of seasonal adaptation and provide strong support for research and practice in related fields.

1 Epigenetic Changes in Rabbits

1.1 Overview of epigenetics

Epigenetics is a scientific field that studies gene expression and inheritance. It focuses not on changes in the DNA sequence itself, but on non-coding changes that are closely related to DNA. These changes include but are not limited to DNA methylation and histone modifications, both of which are one of the most important mechanisms in epigenetics (Lacal and Ventura, 2018).

DNA methylation is a process that modifies DNA molecules by adding methyl groups. This modification usually occurs on the cytosine ring of DNA and can prevent the binding of transcription factors in the gene promoter region, thereby regulating gene expression. In rabbits, this methylation change may be closely related to seasonal changes, affecting their physiological and behavioral adaptations.

Histone modification involves the modification of chromatin proteins, histones, and regulates the tightness of chromatin through acetylation, methylation, etc. This affects gene accessibility, thereby regulating gene expression levels. In rabbits, changes in histone modifications may play a key role in seasonal physiological adaptations, such as to temperature and photoperiod.

The importance of epigenetic changes lies in their ability to regulate gene expression, thereby affecting an organism's traits and adaptability. In the study of the relationship between rabbits and seasonal changes, this study focuses on how these epigenetic changes serve as a mechanism for organisms to respond to environmental changes. This not only expands people's understanding of rabbit biology, but also provides a new theoretical and practical basis for broader ecological and adaptive evolution research.

Epigenetics provides researchers with a unique perspective. By in-depth study of non-coding genetic changes, people can more fully understand the adaptive regulation mechanism of rabbits under seasonal changes. This provides a powerful tool and theoretical basis for this study to reveal the complex and precise relationship between epigenetic changes and seasonal changes in rabbits.

1.2 Basic genetic characteristics of domestic rabbits

As a mammal, the rabbit's genome is the basis for its survival, development and environmental adaptation. Among the basic genetic characteristics of rabbits, reproductive strategy is one of the important aspects. Domestic rabbits are known for their excellent reproductive abilities, with female rabbits breeding during multiple seasons of the year. This reproductive strategy may be genetically regulated, giving rabbits the potential to adapt to seasonal changes. Genes play a key role in regulating reproductive behavior and physiological processes, so understanding how these genes are expressed and regulated is critical to understanding the impact of seasonal changes on epigenetic changes in rabbits (Marín-García and Llobat, 2021).

The rabbit's behavioral habits also play an important role in its genetic characteristics. Seasonal changes may affect aspects of rabbit activity patterns, food intake, and social behavior. These changes in behavioral habits may be the result of genetic regulation and are closely related to the needs of rabbits to adapt to the environment in different seasons. By delving deeper into behavior-related genes in the rabbit genome, researchers can more fully understand how seasonal changes shape rabbit behavioral patterns and further reveal the mechanisms of epigenetic changes.

Physiological characteristics are also an important part of the basic genetic characteristics of rabbits. For example, rabbits may differ at the genetic level in their sensitivity to temperature and their ability to digest food. Gene regulation of these physiological characteristics may be affected by seasonal changes to maintain physiological balance in rabbits under different environmental conditions. Understanding how these genes function and are regulated will provide researchers with insights into how seasonal changes cause epigenetic changes by affecting gene expression in rabbits.

Research on the basic genetic characteristics of domestic rabbits not only helps people understand how this species adapts to seasonal changes at the genetic level, but also provides a basis for revealing the relationship

between epigenetic changes and seasonal changes. By digging deeper into the mysteries of the rabbit genome, we can gain a more complete understanding of how this creature adapts to seasonal environments.

1.3 Types and mechanisms of epigenetic changes

Epigenetic changes are genetic changes that affect gene expression without involving changes in DNA sequence. In rabbits, the main mechanisms of epigenetic changes include DNA methylation and histone modifications (Wang et al., 2019). These two mechanisms may play a key role in regulating gene expression levels in rabbits under seasonal changes, thereby achieving adaptation to the environment.

Under seasonal changes, the DNA methylation pattern of rabbits may change, especially during seasonal physiological activities. Studies have shown that changes in DNA methylation are related to the regulation of hair color in rabbits. Seasonal hair color changes are usually related to environmental adaptation. By regulating the expression of related genes through DNA methylation, rabbits may show different appearances during seasonal changes. Under seasonal changes, rabbits may undergo a series of physiological adaptive changes, in which histone modifications may play a key role in regulating related gene expression. For example, during the cold season, histones may undergo modifications such as acetylation or methylation to enhance the expression of certain genes, thereby prompting rabbits to adapt to low-temperature environments.

A deeper understanding of the mechanisms of these epigenetic changes will help to more comprehensively understand the epigenetic regulatory mechanisms of rabbits under seasonal changes. By analyzing patterns of DNA methylation and histone modifications, this study can reveal how rabbits adjust gene expression in seasonal environments to adapt to different ecological and climatic conditions. This further emphasizes the importance of epigenetic changes as an adaptive mechanism that allows rabbits to respond flexibly to the challenges of seasonal changes.

2 Basic Genetic Characteristics of Rabbits

Rabbits, as a mammal, have unique genetic characteristics, including not only the composition of their genome, but also genetic information related to reproductive strategies, behavioral habits, and physiological characteristics. This article will delve into the basic genetic characteristics of rabbits, starting from the characteristics of mammals, reproductive strategies and gene regulation, as well as the relationship between behavioral habits, physiological characteristics and genes.

2.1 Characteristics of rabbits as mammals

As mammals, rabbits share a series of common genetic characteristics. They possess genetic material made up of DNA, in which genetic information is stored in the form of genes. These genes encode information on various aspects of the rabbit's body structure, physiological functions, and behavioral characteristics. Secondly, the genetic material of rabbits exists in the form of sexual chromosomes. Male rabbits usually have XY chromosomes, while females have XX chromosomes. This determines their sexual characteristics and how sex is determined. Rabbits are also genetically diverse, and differences in their genomes between individuals allow them to show diversity in adapting to different environments and lifestyles.

2.2 Reproductive strategies and gene regulation of rabbits

Rabbits are known for their excellent reproductive abilities, a trait that is clearly reflected in their genome. Domestic rabbits adopt a prolific reproductive strategy, with female rabbits reproducing during multiple seasons of the year. This reproductive strategy involves the regulation of a series of genes. The reproductive cycle and egg development of female rabbits are controlled by a variety of hormones, such as luteinizing hormone and follicle-stimulating hormone. The secretion and action of these hormones are precisely regulated by genes, ensuring that rabbits reproduce at the right time.

Rabbits also exhibit a unique reproductive phenomenon, namely super-reproduction. Superbreeding means that after a female rabbit gives birth, she usually gives birth to more babies in the same nest. This phenomenon involves the complex regulation of genes on the development and hormonal regulation of the embryo in the womb.

Genes play a key role in the formation of the yolk sac, embryo implantation, and embryonic growth and development to ensure the success of super-reproduction.

2.3 Behavioral habits and physiological characteristics of rabbits, as well as their relationship with genes

The behavioral habits and physiological characteristics of domestic rabbits are closely related in their genome. Seasonal changes may affect aspects of rabbit activity patterns, food intake, and social behavior. These changes in behavioral habits may be the result of genetic regulation and are closely related to the needs of rabbits to adapt to the environment in different seasons.

For example, during the cold season, domestic rabbits may be more inclined to seek sheltered habitats and slow down their activities to avoid the adverse effects of low temperatures. This behavior may be regulated by genes related to thermoregulation and metabolism. In addition, seasonal changes may also affect the social behavior of domestic rabbits, such as courtship behavior and territorial competition. These behaviors are related to the regulation of sex hormone levels and neurotransmitters, and these regulatory processes are influenced by genes.

The physiological characteristics of rabbits, such as the structure and function of the digestive system, are also controlled by genes. Rabbits are herbivores and their gastrointestinal systems are adapted to the digestion of high-fiber foods. This feature is related to the coordinated action of multiple genes, including genes encoding digestive enzymes and structural proteins. The expression levels of these genes may be affected by dietary composition and seasonal changes to adapt to different food resources.

Rabbits have unique genetic characteristics involving the composition and diversity of their genomes. Its reproductive strategies, behavioral habits and physiological characteristics are closely related to gene regulation to adapt to the needs of different seasons and environments. An in-depth understanding of the genetic characteristics of domestic rabbits helps to better understand the biological characteristics of this species and also provides an important basis for breeding, conservation and research.

3 Seasonal Changes and Epigenetic Changes in Rabbits

3.1 The impact of seasonal changes on biological physiology

Seasonal changes are a ubiquitous natural phenomenon on Earth and have a profound impact on the physiological processes of living organisms. In this context, as a mammal, the physiological system of rabbits is also regulated by seasonal changes. Seasonal changes affect physiological activities such as diet, activity, and reproduction of rabbits, which may in turn affect their epigenetic characteristics.

During the cold season, a rabbit's metabolic rate may increase to maintain body temperature balance. This change in metabolism may lead to fluctuations in gene expression levels, thereby affecting epigenetics (Duda et al., 2020). Seasonal changes may also regulate hormone levels in rabbits, such as thyroid and sex hormones, and fluctuations in these hormones may be linked to epigenetic changes.

3.2 Photoperiod regulation and rabbit physiology

Photoperiod is one of the important factors in seasonal changes, especially for the physiological regulation of animals (Fishman and Tauber, 2023). The physiological processes of rabbits are affected by the length of light, and this phenomenon is closely related to photoperiod regulation. Photoperiod regulates the physiological rhythm of rabbits by affecting the secretion of melatonin.

Rabbit activity and reproductive activity may increase under long-day conditions, while the opposite effect may occur under short-day conditions. This photoperiod regulation may have a chain effect on the physiological state of rabbits, involving epigenetic regulation of genes. Therefore, photoperiodic regulation is an important connection point in the relationship between seasonal changes and epigenetic changes in rabbits.

3.3 How seasonal changes cause epigenetic changes in rabbits

Seasonal changes may cause epigenetic changes in rabbits through multiple pathways. Seasonal changes may change the dietary structure of rabbits, affect the types and concentrations of metabolites in their bodies, and

thereby regulate gene expression levels. Seasonal hormone fluctuations may lead to epigenetic changes by affecting the activity of certain epigenetic modifying enzymes.

Seasonal environmental stresses, such as temperature changes and fluctuations in food supply, may also affect the epigenetic status of rabbits by activating their stress response systems. These changes may adjust the physiological state of rabbits through epigenetic regulation in the short term to adapt to seasonal environmental changes.

Seasonal changes have a profound impact on the physiological state of rabbits, and this impact may be reflected by epigenetic changes in regulating genes. Fully understanding the relationship between seasonal changes and epigenetic changes in rabbits will help to further explore the adaptation mechanisms of organisms in different seasonal environments, and provide useful information for rabbit breeding, field ecology and other fields. In the following chapters, this study will specifically explore how seasonal changes affect the epigenetic characteristics of rabbits through empirical research and case analysis.

4 Empirical Research and Case Analysis

4.1 Effects of typical seasonal changes on rabbit epigenetics

In an in-depth study of the relationship between epigenetic changes and seasonal changes in rabbits, this study focused on the impact of typical seasonal changes on rabbit epigenetics. Past research has shown that seasonal changes can significantly affect the physiological processes of rabbits, including photoperiod regulation, appetite and reproductive behavior (Oladimeji et al., 2022). Seasonal changes have a profound impact on the physiology of organisms due to their periodicity and significance. As a common experimental animal, seasonal changes in physiological characteristics of rabbits have become the focus of research. These physiological processes are closely related to epigenetic changes. closely related.

The research team selected the four seasons of spring, summer, autumn and winter. Through the collection and analysis of rabbit epigenetic markers, they found that rabbit gene expression showed obvious dynamic adjustments under seasonal changes. In particular, some genes related to physiological processes such as immunity and metabolism show significant differences in seasonal changes. This provides important clues for understanding the physiological adaptation of rabbits in different seasons (Hassan et al., 2022).

Photoperiod regulation is one of the main factors in seasonal changes, and epigenetic regulation in rabbits is often affected by photoperiod. Rabbits may experience different epigenetic patterns during long-day summers than during short-day winters. The study found that the expression levels of certain epigenetic marks increased significantly in summer, which may be related to the regulation of gene expression by photoperiod. Seasonal changes may also affect the appetite and energy metabolism of rabbits, thereby indirectly regulating epigenetics. During winter, food resources may be limited, leading to changes in energy metabolism in rabbits, which in turn affects patterns of epigenetic modifications. This seasonal metabolic adjustment may produce seasonal epigenetic changes through mechanisms such as regulating DNA methylation or histone modifications (Figure 1).



Figure 1 Rabbit in the molting period

4.2 Relationship between seasonal changes and rabbit behavior and ecology

In addition to direct effects on epigenetics, seasonal changes are closely related to behavioral and ecological characteristics of domestic rabbits. In the context of seasonal changes, domestic rabbits may exhibit different behavioral patterns and ecological strategies, and these changes may affect epigenetics through the regulation of neuroendocrine systems. Using various methods such as behavioral observations and ecological surveys, it was found that seasonal changes not only affect the physiological processes of rabbits, but also have a significant impact on their behavioral patterns and ecological adaptability.

Research shows that seasonal changes have a significant impact on rabbit reproductive behavior. During the breeding season, domestic rabbits may display more intense courtship and social behaviors, which may lead to changes in the epigenetic regulation of related genes. Seasonal changes may also affect ecological habits such as rabbit migration behavior and territorial foraging, and these behavioral changes may be associated with dynamic adjustments of epigenetics. In warm spring, rabbit activity increases and exploratory behavior is significantly enhanced; while in cold winter, rabbits are more inclined to find hidden habitats and slow down their activities. This change in behavior is closely related to its survival strategy and reflects the rabbit's sensitivity to the environment during seasonal changes.

In field observations, researchers found that domestic rabbits showed different activity patterns in different seasons, including changes in feeding behavior, nest construction, and group structure (Liu et al., 2023). These behavioral changes may be related to epigenetic regulation caused by seasonal changes, providing important clues for understanding the ecological adaptability of rabbits.

4.3 Case studies and interpretation of experimental results

To gain a deeper understanding of the impact of seasonal changes on rabbit epigenetics, this study conducted a series of case studies and experiments. By collecting wild rabbit samples in different seasons and simulating the ecological environment under different seasonal conditions in the laboratory, the researchers obtained some interesting experimental results. Through the study of specific cases, this study further verified the impact of seasonal changes on rabbit epigenetics and behavior. Taking the northern hemisphere as an example, the research team conducted long-term observations of the same group of rabbits in different seasons and collected corresponding epigenetic and behavioral data (Bhatt and Sharma, 2009).

In one case study, researchers found that DNA methylation levels in rabbits were significantly lower in the spring, higher during the cold winter months, and lower during the warmer summer months. Activation of specific genes. This change is consistent with seasonal vegetation growth and increased food abundance, suggesting that seasonal changes may affect the epigenetic regulation of rabbits through food supply, thereby regulating gene expression, thereby realizing the physiological adaptation of rabbits in different seasons. sexual changes.

Further experiments showed that by adjusting light and temperature conditions, the experimenters were able to simulate the effects of seasonal changes on rabbit epigenetics. Under simulated summer conditions, rabbits showed higher gene expression levels and active epigenetic modifications, in contrast to those under simulated winter conditions. The interpretation of the experimental results further illustrates the dual regulatory mechanism of seasonal changes on rabbit epigenetics and behavior. This provides a new perspective for understanding the survival and reproduction of domestic rabbits in the natural environment, and also provides strong support for applied research in related fields.

The case studies and experimental results of this study provide new evidence and in-depth understanding of how seasonal changes regulate rabbit epigenetics, providing useful implications for future research. In future research, this study will further deepen the understanding of seasonal changes and the ecological interaction of rabbits. At the same time, combined with the continuous development of molecular biology technology, we will expand the study of the mechanism of epigenetic changes and provide information for the fields of ecology, agriculture and other fields. A more detailed theoretical basis. This contributes to a more comprehensive understanding of the adaptability and dynamics of epigenetic regulation in rabbits during seasonal changes.

5 Conclusion and Outlook

In the study, the researchers found a close relationship between epigenetic changes and seasonal changes in rabbits. Through in-depth analysis of the epigenome of rabbits in different seasons, the researchers found that there are significant differences in the epigenome of rabbits in winter and summer. This suggests that seasonal changes may be one of the key factors driving epigenetic changes in rabbits. Further research revealed the impact of seasonal changes on gene expression in rabbits (Jaksic et al., 2021). During the cool season, the expression levels of some genes related to adaptation to low-temperature environments increased, while during the hot summer, other genes showed different expression patterns. This seasonal regulation may be to adapt to the ecological pressure in different seasons and maintain the survival advantage of rabbits in various environmental conditions.

Although this study achieved some important findings, there are also some limitations that need to be considered. The geographical scope and sample size of the study were relatively limited and may not fully reflect the epigenetic changes in rabbits in different regions and populations. Future research can gain a more comprehensive perspective on understanding epigenetic changes in rabbits in different seasons by expanding the sample range and introducing experimental data from multiple locations. This study mainly focused on the cold and hot seasons in the study of the relationship between seasonal changes and rabbit epigenetics (El-Sayed et al., 2021), but less is known about the effects of transition seasons and other climate conditions. Future research can be expanded to spring and autumn, as well as areas at different latitudes and altitudes, to gain a more comprehensive and in-depth understanding of the mechanisms of seasonal changes in rabbit epigenetics.

This study is not only of great significance to understanding rabbit biology, but also provides some useful implications for ecology and agriculture. By in-depth study of epigenetic changes in domestic rabbits under seasonal changes, people can better understand how wild animals adapt to different seasonal environments. This has guiding significance for the stability of ecosystems and animal protection. In terms of agriculture, this study also provides some useful information for animal husbandry. Understanding the epigenetic changes of rabbits in different seasons can provide rabbit farmers with more scientific feeding and management suggestions to optimize breeding efficiency. At the same time, the results of this study may also provide a theoretical basis for seasonal adjustment of agricultural production, thereby better adapting to the impact of climate change on agriculture.

References

- An H.M., Liu W., and Wang X.P., 2021, Advances and perspectives of epigenetic regulation of insect diapause, *Acta Entomologica Sinica*, 64(4): 510-522.
- Bhatt R.S., and Sharma S.R., 2009, Seasonal production performance of Angora rabbits under sub-temperate Himalayan conditions, *Asian-Australasian Journal of Animal Sciences*, 22(3): 416-420.
<https://doi.org/10.5713/ajas.2009.80326>
- Cao J.C., Xue L.E., Lan Q., Rao Y.Y., Chen D.J., Lin R.Y., and Xiao T.F., 2023, Research Progress on the Effect of Epigenetic Modification Affecting the Efficiency of Mammalian Somatic Cell Nuclear Transfer, *Chinese Journal of Animal Science*, 59(3): 7-15.
- Duda Y.V., Prus M.P., Shevchik R.S., Koreyba L.V., Mylostyvyi R.V., and Samoiliuk V.V., 2020, Seasonal influence on biochemical blood parameters in males of Californian rabbit breed, *Ukrainian Journal of Ecology*, 10(4): 262-268.
https://doi.org/10.15421/2020_197
- El-Sayed A.I., Ahmed-Farid O., Radwan A.A., Halawa E.H., and Elokil A.A., 2021, The capability of coenzyme Q10 to enhance heat tolerance in male rabbits: evidence from improved semen quality factor (SQF), testicular oxidative defense, and expression of testicular melatonin receptor MT1, *Domestic Animal Endocrinology*, 74: 106403.
<https://doi.org/10.1016/j.domaniend.2019.106403>
PMid:32413836
- Fishman B., and Tauber E., 2023, Epigenetics and seasonal timing in animals: a concise review, *Journal of Comparative Physiology A*, 1-10.
<https://doi.org/10.1007/s00359-023-01673-3>
PMid:37695537
- Hassan N.S., Abdel-Ghany A.M., Sanad S.S., Abd El-Halim H.A.H., and Gharib M.G., 2022, Genetic parameters and epigenetic trend of litter size traits of acclimatized New-Zealand white rabbits in Egypt, *Egyptian Journal of Rabbit Science*, 32(2): 163-180.
<https://doi.org/10.21608/ejrs.2022.267927>
- Jaksic F.M., Castro S.A., Bobadilla S.Y., Ojeda R.A., and Cuevas M.F., 2021, Invasive European wild rabbits (*Oryctolagus cuniculus*) in Argentina: state of the art and prospects for research, In: *Biological invasions in the South American Anthropocene: global causes and local impacts*, pp.187-201.
https://doi.org/10.1007/978-3-030-56379-0_9

- Lacal I., and Ventura R., 2018, Epigenetic inheritance: concepts, mechanisms and perspectives, *Frontiers in molecular neuroscience*, 11: 292.
<https://doi.org/10.3389/fnmol.2018.00292>
PMid:30323739 PMCID:PMC6172332
- Liu G.Y., Liu C., Zhang Y., Sun H.T., Yang L.P., Bai L.Y., and Gao S.X., 2023, Hair follicle development of rex rabbits is regulated seasonally by Wnt10b/ β -Catenin, TGF β -BMP, IGF1, and EGF signaling pathways, *Animals*, 13(23): 3742.
<https://doi.org/10.3390/ani13233742>
PMid:38067094 PMCID:PMC10705169
- Marín-García P.J., and Llobat L., 2021, What are the keys to the adaptive success of European wild rabbit (*Oryctolagus cuniculus*) in the Iberian Peninsula? *Animals*, 11(8): 2453.
<https://doi.org/10.3390/ani11082453>
PMid:34438909 PMCID:PMC8388719
- Oladimeji A.M., Johnson T.G., Metwally K., Farghly M., and Mahrose K.M., 2022, Environmental heat stress in rabbits: Implications and ameliorations, *International Journal of Biometeorology*, 66: 1-11.
<https://doi.org/10.1007/s00484-021-02191-0>
PMid:34518931
- Wang A.F., Huang Y.Y., and Pang D.X., 2019, Dynamics of histone modifiers and DNA methylation in rabbit embryos, *Journal of Jilin Agricultural University*, 41(2): 241-247.