

Bohao Zhao

List of Publications by Year in descending order

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| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Exosomal miRNA-181a-5p from the cells of the hair follicle dermal papilla promotes the hair follicle growth and development via the Wnt/ β 2-catenin signaling pathway. <i>International Journal of Biological Macromolecules</i> , 2022, 207, 110-120. | 7.5 | 24 |
| 2 | miR-129-5p Participates in Hair Follicle Growth by Targeting HOXC13 in Rabbit. <i>Genes</i> , 2022, 13, 679. | 2.4 | 1 |
| 3 | Promoter Methylation Changes in KRT17: A Novel Epigenetic Marker for Wool Production in Angora Rabbit. <i>International Journal of Molecular Sciences</i> , 2022, 23, 6077. | 4.1 | 2 |
| 4 | Characterization and functional analysis of SMAD2 regulation in hair follicle cycle in Angora rabbits. <i>Gene</i> , 2021, 770, 145339. | 2.2 | 1 |
| 5 | Bacitracin Methylene Disalicylate Improves Intestinal Health by Modulating Its Development and Microbiota in Weaned Rabbits. <i>Frontiers in Microbiology</i> , 2021, 12, 579006. | 3.5 | 2 |
| 6 | Deubiquitination of MITF-M Regulates Melanocytes Proliferation and Apoptosis. <i>Frontiers in Molecular Biosciences</i> , 2021, 8, 692724. | 3.5 | 8 |
| 7 | GNAI2 Promotes Proliferation and Decreases Apoptosis in Rabbit Melanocytes. <i>Genes</i> , 2021, 12, 1130. | 2.4 | 5 |
| 8 | A Treatment Combination of IGF and EGF Promotes Hair Growth in the Angora Rabbit. <i>Genes</i> , 2021, 12, 24. | 2.4 | 17 |
| 9 | A Genetic Evaluation System for New Zealand White Rabbit Germplasm Resources Based on SSR Markers. <i>Animals</i> , 2020, 10, 1258. | 2.3 | 3 |
| 10 | Characterization and functional analysis of Krtap11-1 during hair follicle development in Angora rabbits (<i>Oryctolagus cuniculus</i>). <i>Genes and Genomics</i> , 2020, 42, 1281-1290. | 1.4 | 5 |
| 11 | Characterization of HTATIP2 and its role during hair follicle cycles in Angora rabbit. <i>Genome</i> , 2020, 63, 179-187. | 2.0 | 9 |
| 12 | Characterization and functional analysis of SIAH1 during skin and hair follicle development in the angora rabbit (<i>Oryctolagus cuniculus</i>). <i>Hereditas</i> , 2020, 157, 10. | 1.4 | 4 |
| 13 | Analysis of Genome DNA Methylation at Inherited Coat Color Dilutions of Rex Rabbits. <i>Frontiers in Genetics</i> , 2020, 11, 603528. | 2.3 | 6 |
| 14 | Characterization of POU2F1 Gene and Its Potential Impact on the Expression of Genes Involved in Fur Color Formation in Rex Rabbit. <i>Genes</i> , 2020, 11, 575. | 2.4 | 7 |
| 15 | KIT is involved in melanocyte proliferation, apoptosis and melanogenesis in the Rex Rabbit. <i>PeerJ</i> , 2020, 8, e9402. | 2.0 | 10 |
| 16 | Morphological Characterization and Gene Expression Patterns for Melanin Pigmentation in Rex Rabbit. <i>Biochemical Genetics</i> , 2019, 57, 734-744. | 1.7 | 11 |
| 17 | Characterization and Establishment of an Immortalized Rabbit Melanocyte Cell Line Using the SV40 Large T Antigen. <i>International Journal of Molecular Sciences</i> , 2019, 20, 4874. | 4.1 | 18 |
| 18 | Systematic Analysis of Non-coding RNAs Involved in the Angora Rabbit (<i>Oryctolagus cuniculus</i>) Hair Follicle Cycle by RNA Sequencing. <i>Frontiers in Genetics</i> , 2019, 10, 407. | 2.3 | 57 |

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|----|---|-----|-----------|
| 19 | Slc7a11 Modulated by POU2F1 is Involved in Pigmentation in Rabbit. International Journal of Molecular Sciences, 2019, 20, 2493. | 4.1 | 15 |
| 20 | miR-218-5p regulates skin and hair follicle development through Wnt/ β -catenin signaling pathway by targeting SFRP2. Journal of Cellular Physiology, 2019, 234, 20329-20341. | 4.1 | 57 |
| 21 | MicroRNAs Profiling Identifies miR-125a and Its Target Gene Wnt2 in Skins of Different Haired Rabbits. Frontiers in Genetics, 2018, 9, 628. | 2.3 | 10 |
| 22 | RNAi-mediated SLC7A11 knockdown inhibits melanogenesis-related genes expression in rabbit skin fibroblasts. Journal of Genetics, 2018, 97, 463-468. | 0.7 | 6 |
| 23 | Identification and profiling of microRNA between back and belly Skin in Rex rabbits (<i>Oryctolagus</i>) Tj ETQq1 1 0.784314 rgBT /Overloc | 0.6 | 1 |
| 24 | Gene expression profiling analysis reveals fur development in rex rabbits (<i>Oryctolagus cuniculus</i>). Genome, 2017, 60, 1060-1067. | 2.0 | 11 |
| 25 | Impacts of diarrhea on the immune system, intestinal environment, and expression of PGRPs in New Zealand rabbits. PeerJ, 2017, 5, e4100. | 2.0 | 12 |