

Zhe Wang

List of Publications by Year in descending order

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Version: 2024-02-01

22
papers

734
citations

840776

11
h-index

713466

21
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22
all docs

22
docs citations

22
times ranked

843
citing authors

#	ARTICLE	IF	CITATIONS
1	Cabotegravir for HIV Prevention in Cisgender Men and Transgender Women. <i>New England Journal of Medicine</i> , 2021, 385, 595-608.	27.0	359
2	An Oncogenic Role for Alternative NF- κ B Signaling in DLBCL Revealed upon Deregulated BCL6 Expression. <i>Cell Reports</i> , 2015, 11, 715-726.	6.4	66
3	Adaptive Evolution of 5'HoxD Genes in the Origin and Diversification of the Cetacean Flipper. <i>Molecular Biology and Evolution</i> , 2008, 26, 613-622.	8.9	60
4	Prenatal development supports a single origin of laryngeal echolocation in bats. <i>Nature Ecology and Evolution</i> , 2017, 1, 21.	7.8	43
5	Digital gene expression tag profiling of bat digits provides robust candidates contributing to wing formation. <i>BMC Genomics</i> , 2010, 11, 619.	2.8	26
6	Unique expression patterns of multiple key genes associated with the evolution of mammalian flight. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2014, 281, 20133133.	2.6	25
7	Comparison of whole embryonic development in the duck (<i>Anas platyrhynchos</i>) and goose (<i>Anser</i>) Tj ETQq1 1 0.784314 rgBT /Overlook	3.4	25
8	Circular RNA Profiling Identifies Novel circPPARA that Promotes Intramuscular Fat Deposition in Pigs. <i>Journal of Agricultural and Food Chemistry</i> , 2022, 70, 4123-4137.	5.2	22
9	Molecular signatures and functional analysis of beige adipocytes induced from in vivo intra-abdominal adipocytes. <i>Science Advances</i> , 2018, 4, eaar5319.	10.3	18
10	Differential Expression of Meis2, Mab21l2 and Tbx3 during Limb Development Associated with Diversification of Limb Morphology in Mammals. <i>PLoS ONE</i> , 2014, 9, e106100.	2.5	17
11	Identification of Candidate Circular RNAs Underlying Intramuscular Fat Content in the Donkey. <i>Frontiers in Genetics</i> , 2020, 11, 587559.	2.3	16
12	Unveiling the Biogeography and Potential Functions of the Intestinal Digesta- and Mucosa-Associated Microbiome of Donkeys. <i>Frontiers in Microbiology</i> , 2020, 11, 596882.	3.5	11
13	Synergy between <i>MC1R</i> and <i>ASIP</i> for coat color in horses (<i>Equus caballus</i>). <i>Journal of Animal Science</i> , 2019, 97, 1578-1585.	0.5	9
14	Prolonged treatment with 3-isobutyl-1-methylxanthine improves the efficiency of differentiating 3T3-L1 cells into adipocytes. <i>Analytical Biochemistry</i> , 2016, 507, 18-20.	2.4	8
15	The Great Roundleaf Bat (<i>Hipposideros armiger</i>) as a Good Model for Cold-Induced Browning of Intra-Abdominal White Adipose Tissue. <i>PLoS ONE</i> , 2014, 9, e112495.	2.5	7
16	Pooled Sequencing Analysis of Geese (<i>Anser cygnoides</i>) Reveals Genomic Variations Associated With Feather Color. <i>Frontiers in Genetics</i> , 2021, 12, 650013.	2.3	6
17	Accelerated Evolution of Limb-Related Gene <i>Hoxd11</i> in the Common Ancestor of Cetaceans and Ruminants (Cetruminantia). <i>G3: Genes, Genomes, Genetics</i> , 2020, 10, 515-524.	1.8	4
18	Mucosal Microbiota and Metabolome in the Ileum of Hu Sheep Offered a Low-Grain, Pelleted or Non-pelleted High-Grain Diet. <i>Frontiers in Microbiology</i> , 2021, 12, 718884.	3.5	4

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19	A Rapid Embryonic Stem Cell-Based Mouse Model for B-cell Lymphomas Driven by Epstein-Barr Virus Protein LMP1. <i>Cancer Immunology Research</i> , 2015, 3, 641-649.	3.4	3
20	Whole Genome Sequencing Reveals Signatures for Artificial Selection for Different Sizes in Japanese Primitive Dog Breeds. <i>Frontiers in Genetics</i> , 2021, 12, 671686.	2.3	3
21	A missense mutation in <i>ASIP</i> is associated with light point variation in donkeys. <i>Animal Genetics</i> , 2020, 51, 629-629.	1.7	2
22	Adaptive Evolution of the Fox Coronavirus Based on Genome-Wide Sequence Analysis. <i>BioMed Research International</i> , 2022, 2022, 1-8.	1.9	0