

Maochun Wang

List of Publications by Year in descending order

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Version: 2024-02-01

12
papers

290
citations

1478505

6
h-index

1199594

12
g-index

15
all docs

15
docs citations

15
times ranked

508
citing authors

#	ARTICLE	IF	CITATIONS
1	Targeting macrophagic SHP2 for ameliorating osteoarthritis via TLR signaling. <i>Acta Pharmaceutica Sinica B</i> , 2022, 12, 3073-3084.	12.0	21
2	Candidate Regulators of Dyslipidemia in Chromosome 1 Substitution Lines Using Liver Co-Expression Profiling Analysis. <i>Frontiers in Genetics</i> , 2020, 10, 1258.	2.3	1
3	Genetic Dissection of Hypertrophic Cardiomyopathy with Myocardial RNA-Seq. <i>International Journal of Molecular Sciences</i> , 2020, 21, 3040.	4.1	26
4	Genome-Wide Analysis of MicroRNA-related Single Nucleotide Polymorphisms (SNPs) in Mouse Genome. <i>Scientific Reports</i> , 2020, 10, 5789.	3.3	11
5	A multiplex SNP genotyping by allele-specific PCR based on stem-loop and universal fluorescent primers of Chr1 ddx1 mice. <i>Electrophoresis</i> , 2019, 40, 1600-1605.	2.4	2
6	miR-505-3p is a repressor of puberty onset in female mice. <i>Journal of Endocrinology</i> , 2019, 240, 379-392.	2.6	4
7	Characterization of the dynamic change of microRNA expression in mice hypothalamus during the time of female puberty. <i>Genes and Genomics</i> , 2018, 40, 295-304.	1.4	3
8	A multiplex sensitive quantification of microRNAs based on competitive PCR. <i>Biotechnology and Bioprocess Engineering</i> , 2017, 22, 95-99.	2.6	4
9	Sequence analysis of chromosome 1 revealed different selection patterns between Chinese wild mice and laboratory strains. <i>Molecular Genetics and Genomics</i> , 2017, 292, 1111-1121.	2.1	2
10	Genome Sequencing of Chromosome 1 Substitution Lines Derived from Chinese Wild Mice Revealed a Unique Resource for Genetic Studies of Complex Traits. <i>G3: Genes, Genomes, Genetics</i> , 2016, 6, 3571-3580.	1.8	8
11	Electrospun gelatin nanofibers loaded with vitamins A and E as antibacterial wound dressing materials. <i>RSC Advances</i> , 2016, 6, 50267-50277.	3.6	127
12	A novel three-round multiplex PCR for SNP genotyping with next generation sequencing. <i>Analytical and Bioanalytical Chemistry</i> , 2016, 408, 4371-4377.	3.7	80