

Maochun Wang

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5294243/publications.pdf>

Version: 2024-02-01

12
papers

290
citations

1477746

6
h-index

1199166

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.	5.7	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.	1.1	1
3	Genetic Dissection of Hypertrophic Cardiomyopathy with Myocardial RNA-Seq. <i>International Journal of Molecular Sciences</i> , 2020, 21, 3040.	1.8	26
4	Genome-Wide Analysis of MicroRNA-related Single Nucleotide Polymorphisms (SNPs) in Mouse Genome. <i>Scientific Reports</i> , 2020, 10, 5789.	1.6	11
5	A multiplex SNP genotyping by allele-specific PCR based on stem-loop and universal fluorescent primers of Chr1 dixin mice. <i>Electrophoresis</i> , 2019, 40, 1600-1605.	1.3	2
6	miR-505-3p is a repressor of puberty onset in female mice. <i>Journal of Endocrinology</i> , 2019, 240, 379-392.	1.2	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.	0.5	3
8	A multiplex sensitive quantification of microRNAs based on competitive PCR. <i>Biotechnology and Bioprocess Engineering</i> , 2017, 22, 95-99.	1.4	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.	1.0	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.	0.8	8
11	Electrospun gelatin nanofibers loaded with vitamins A and E as antibacterial wound dressing materials. <i>RSC Advances</i> , 2016, 6, 50267-50277.	1.7	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.	1.9	80