## Maochun Wang

List of Publications by Year in descending order

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1478505 1199594 12 290 12 6 citations h-index g-index papers 15 15 15 508 docs citations times ranked citing authors all docs

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