

Kuo-Ping Chiu

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

11
papers

2,096
citations

8
h-index

11
g-index

11
ext. papers

2,278
ext. citations

6.7
avg, IF

3.51
L-index

#	Paper	IF	Citations
11	The Oct4 and Nanog transcription network regulates pluripotency in mouse embryonic stem cells. <i>Nature Genetics</i> , 2006 , 38, 431-40	36.3	1920
10	Long-read sequencing in deciphering human genetics to a greater depth. <i>Human Genetics</i> , 2019 , 138, 1201-1215	6.3	37
9	Concordant and discordant regulation of target genes by miR-31 and its isoforms. <i>PLoS ONE</i> , 2013 , 8, e58169	3.7	36
8	Palindromic sequence impedes sequencing-by-ligation mechanism. <i>BMC Systems Biology</i> , 2012 , 6 Suppl 2, S10	3.5	31
7	Loop-sequence features and stability determinants in antibody variable domains by high-throughput experiments. <i>Structure</i> , 2014 , 22, 9-21	5.2	22
6	Analysis of microbial sequences in plasma cell-free DNA for early-onset breast cancer patients and healthy females. <i>BMC Medical Genomics</i> , 2018 , 11, 16	3.7	17
5	Global assessment of <i>Antrodia cinnamomea</i> -induced microRNA alterations in hepatocarcinoma cells. <i>PLoS ONE</i> , 2013 , 8, e82751	3.7	10
4	Application of cell-free DNA sequencing in characterization of bloodborne microbes and the study of microbe-disease interactions. <i>PeerJ</i> , 2019 , 7, e7426	3.1	8
3	Improved in situ hybridization: color intensity enhancement procedure for the alkaline phosphatase/Fast Red system. <i>BioTechniques</i> , 1996 , 20, 964-6, 968	2.5	6
2	Single cell transcriptome analysis of MCF-7 reveals consistently and inconsistently expressed gene groups each associated with distinct cellular localization and functions. <i>PLoS ONE</i> , 2018 , 13, e0199471	3.7	6
1	Genomic sequencing of <i>Troides aeacus</i> nucleopolyhedrovirus (TraeNPV) from golden birdwing larvae (<i>Troides aeacus formosanus</i>) to reveal defective <i>Autographa californica</i> NPV genomic features. <i>BMC Genomics</i> , 2019 , 20, 419	4.5	3