

Tsunglin Liu

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2886417/publications.pdf>

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#	ARTICLE	IF	CITATIONS
1	Novel potential functions of amoeboid cells in thraustochytrids revealed by <i>Aurantiochytrium limacinum</i> BL10. <i>European Journal of Protistology</i> , 2022, 82, 125860.	1.5	1
2	Nasal Microbiome Change During and After Exacerbation in Asthmatic Children. <i>Frontiers in Microbiology</i> , 2021, 12, 833726.	3.5	8
3	Dissecting efficiency of a 5â€™ TM rapid amplification of cDNA ends (5â€™ TM -RACE) approach for profiling T-cell receptor beta repertoire. <i>PLoS ONE</i> , 2020, 15, e0236366.	2.5	9
4	Joining Illumina paired-end reads for classifying phylogenetic marker sequences. <i>BMC Bioinformatics</i> , 2020, 21, 105.	2.6	18
5	A novel zebrafish model to emulate lung injury by folate deficiency-induced swim bladder defectiveness and protease/antiprotease expression imbalance. <i>Scientific Reports</i> , 2019, 9, 12633.	3.3	12
6	Expanding the mesodermal transcriptional network by genome-wide identification of Zinc finger homeodomain 1 (Zfh1) targets. <i>FEBS Letters</i> , 2019, 593, 1698-1710.	2.8	2
7	Expressional Profiling of Carpet Glia in the Developing <i>Drosophila</i> Eye Reveals Its Molecular Signature of Morphology Regulators. <i>Frontiers in Neuroscience</i> , 2019, 13, 244.	2.8	8
8	Hepatitis B virus surface gene preâ€‘S2 mutant as a highâ€‘risk serum marker for hepatoma recurrence after curative hepatic resection. <i>Hepatology</i> , 2018, 68, 815-826.	7.3	33
9	Rare Compound Heterozygous Frameshift Mutations in <i>ALMS1</i> Gene Identified Through Exome Sequencing in a Taiwanese Patient With Alstr�m Syndrome. <i>Frontiers in Genetics</i> , 2018, 9, 110.	2.3	8
10	TRlg: a robust alignment pipeline for non-regular T-cell receptor and immunoglobulin sequences. <i>BMC Bioinformatics</i> , 2016, 17, 433.	2.6	10
11	Complete Genome Sequence of Community-Acquired <i>Klebsiella pneumoniae</i> KP36, a Strain Isolated from a Patient with an Upper Urinary Tract Infection. <i>Genome Announcements</i> , 2016, 4, .	0.8	2
12	Highly diversified fungi are associated with the achlorophyllous orchid <i>Gastrodia flavilabella</i> . <i>BMC Genomics</i> , 2015, 16, 185.	2.8	19
13	Prediction of personalized microRNA activity. <i>Gene</i> , 2013, 518, 101-106.	2.2	2