

Jun Lin

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5254028/publications.pdf>

Version: 2024-02-01

11
papers

253
citations

1478505

6
h-index

1125743

13
g-index

15
all docs

15
docs citations

15
times ranked

372
citing authors

#	ARTICLE	IF	CITATIONS
1	Methods for genetic transformation of filamentous fungi. <i>Microbial Cell Factories</i> , 2017, 16, 168.	4.0	118
2	Applications of sequencing technology in clinical microbial infection. <i>Journal of Cellular and Molecular Medicine</i> , 2019, 23, 7143-7150.	3.6	35
3	Study on the Effect of Capsaicin on the Intestinal Flora through High-Throughput Sequencing. <i>ACS Omega</i> , 2020, 5, 1246-1253.	3.5	29
4	CircRNA CircRIMS Acts as a MicroRNA Sponge to Promote Gastric Cancer Metastasis. <i>ACS Omega</i> , 2020, 5, 23237-23246.	3.5	18
5	Rapid detection of low-level HeLa cell contamination in cell culture using nested PCR. <i>Journal of Cellular and Molecular Medicine</i> , 2019, 23, 227-236.	3.6	14
6	Study of oral microorganisms contributing to non-carious cervical lesions via bacterial interaction and pH regulation. <i>Journal of Cellular and Molecular Medicine</i> , 2021, 25, 3103-3112.	3.6	5
7	Metagenomic Sequencing Revealed the Potential Pathogenic Threats of Banknotes. <i>ACS Omega</i> , 2021, 6, 3499-3507.	3.5	4
8	In vivo and in vitro study of co-expression of LMP1 and Cripto-1 in nasopharyngeal carcinoma. <i>Brazilian Journal of Otorhinolaryngology</i> , 2020, 86, 617-625.	1.0	3
9	Cas9 Protein Triggers Differential Expression of Inherent Genes Especially NGFR Expression in 293T Cells. <i>Cellular and Molecular Bioengineering</i> , 2020, 13, 61-72.	2.1	3
10	Metagenomic sequencing revealed the potential of banknotes as a repository of microbial genes. <i>BMC Genomics</i> , 2021, 22, 173.	2.8	1
11	rRNA Analysis Based on Long-Read High-Throughput Sequencing Reveals a More Accurate Diagnostic for the Bacterial Infection of Ascites. <i>BioMed Research International</i> , 2021, 2021, 1-8.	1.9	0