

Fernando Garcia-Alcalde

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/12119043/publications.pdf>

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12
papers

4,066
citations

1040056

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1199594

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docs citations

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times ranked

9564
citing authors

#	ARTICLE	IF	CITATIONS
1	A previously uncharacterized gene, PA2146, contributes to biofilm formation and drug tolerance across the <i>E. coli</i> -Proteobacteria. <i>Npj Biofilms and Microbiomes</i> , 2022, 8, .	6.4	6
2	Comparative evaluation of rRNA depletion procedures for the improved analysis of bacterial biofilm and mixed pathogen culture transcriptomes. <i>Scientific Reports</i> , 2017, 7, 41114.	3.3	55
3	Heteroaryldihydropyrimidine (HAP) and Sulfamoylbenzamide (SBA) Inhibit Hepatitis B Virus Replication by Different Molecular Mechanisms. <i>Scientific Reports</i> , 2017, 7, 42374.	3.3	103
4	Qualimap 2: advanced multi-sample quality control for high-throughput sequencing data. <i>Bioinformatics</i> , 2016, 32, 292-294.	4.1	1,362
5	InFusion: Advancing Discovery of Fusion Genes and Chimeric Transcripts from Deep RNA-Sequencing Data. <i>PLoS ONE</i> , 2016, 11, e0167417.	2.5	62
6	Association study between mannose-binding lectin haplotypes and X gene mutation of hepatitis B virus from treatment naïve patients. <i>Aging</i> , 2016, 8, 2862-2870.	3.1	4
7	<i>Helicobacter pylori</i> Infection Causes Characteristic DNA Damage Patterns in Human Cells. <i>Cell Reports</i> , 2015, 11, 1703-1713.	6.4	114
8	Evidence for a crucial role of a host non-coding RNA in influenza A virus replication. <i>RNA Biology</i> , 2014, 11, 66-75.	3.1	90
9	CisMiner: Genome-Wide In-Silico Cis-Regulatory Module Prediction by Fuzzy Itemset Mining. <i>PLoS ONE</i> , 2014, 9, e108065.	2.5	10
10	Qualimap: evaluating next-generation sequencing alignment data. <i>Bioinformatics</i> , 2012, 28, 2678-2679.	4.1	799
11	Transdifferentiation of MALME-3M and MCF-7 Cells toward Adipocyte-like Cells is Dependent on Clathrin-mediated Endocytosis. <i>SpringerPlus</i> , 2012, 1, 44.	1.2	5
12	Differential expression in RNA-seq: A matter of depth. <i>Genome Research</i> , 2011, 21, 2213-2223.	5.5	1,456