

Joslynn S Lee

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2090194/publications.pdf>

Version: 2024-02-01

10
papers

12,266
citations

1305906

8
h-index

1526636

10
g-index

14
all docs

14
docs citations

14
times ranked

16788
citing authors

#	ARTICLE	IF	CITATIONS
1	Creb<scp>A</scp> increases secretory capacity through direct transcriptional regulation of the secretory machinery, a subset of secretory cargo, and other key regulators. <i>Traffic</i> , 2020, 21, 560-577.	1.3	20
2	Reproducible, interactive, scalable and extensible microbiome data science using QIIME 2. <i>Nature Biotechnology</i> , 2019, 37, 852-857.	9.4	11,167
3	Functional classification of protein structures by local structure matching in graph representation. <i>Protein Science</i> , 2018, 27, 1125-1135.	3.1	8
4	Ribbon regulates morphogenesis of the <i>Drosophila</i> embryonic salivary gland through transcriptional activation and repression. <i>Developmental Biology</i> , 2016, 409, 234-250.	0.9	7
5	DNA-binding sequence specificity of DUX4. <i>Skeletal Muscle</i> , 2015, 6, 8.	1.9	30
6	Beyond antioxidant genes in the ancient Nrf2 regulatory network. <i>Free Radical Biology and Medicine</i> , 2015, 88, 452-465.	1.3	74
7	Protein function annotation with Structurally Aligned Local Sites of Activity (SALSAs). <i>BMC Bioinformatics</i> , 2013, 14, S13.	1.2	15
8	Design and evaluation of xanthine based adenosine receptor antagonists: Potential hypoxia targeted immunotherapies. <i>Bioorganic and Medicinal Chemistry</i> , 2013, 21, 7453-7464.	1.4	10
9	Electrostatic Properties for Protein Functional Site Prediction. , 2011, , 183-196.		0
10	FUNCTIONAL CLASSIFICATION OF PROTEIN 3D STRUCTURES FROM PREDICTED LOCAL INTERACTION SITES. <i>Journal of Bioinformatics and Computational Biology</i> , 2010, 08, 1-15.	0.3	18