

Simone S Li

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2429471/publications.pdf>

Version: 2024-02-01

17
papers

1,497
citations

777949

13
h-index

993246

17
g-index

20
all docs

20
docs citations

20
times ranked

3604
citing authors

#	ARTICLE	IF	CITATIONS
1	Disentangling the impact of environmental and phylogenetic constraints on prokaryotic within-species diversity. <i>ISME Journal</i> , 2020, 14, 1247-1259.	4.4	74
2	Extensive transmission of microbes along the gastrointestinal tract. <i>ELife</i> , 2019, 8, .	2.8	313
3	proGenomes: a resource for consistent functional and taxonomic annotations of prokaryotic genomes. <i>Nucleic Acids Research</i> , 2017, 45, D529-D534.	6.5	131
4	MOCAT2: a metagenomic assembly, annotation and profiling framework. <i>Bioinformatics</i> , 2016, 32, 2520-2523.	1.8	172
5	Durable coexistence of donor and recipient strains after fecal microbiota transplantation. <i>Science</i> , 2016, 352, 586-589.	6.0	461
6	Temporal and technical variability of human gut metagenomes. <i>Genome Biology</i> , 2015, 16, 73.	3.8	143
7	Molecular interaction networks for the analysis of human disease: Utility, limitations, and considerations. <i>Proteomics</i> , 2013, 13, 3393-3405.	1.3	17
8	Disturbed proteinâ€“protein interaction networks in metastatic melanoma are associated with worse prognosis and increased functional mutation burden. <i>Pigment Cell and Melanoma Research</i> , 2013, 26, 708-722.	1.5	12
9	A Multidimensional Matrix for Systems Biology Research and Its Application to Interaction Networks. <i>Journal of Proteome Research</i> , 2012, 11, 5204-5220.	1.8	5
10	A Genome-Wide Screen in Yeast Identifies Specific Oxidative Stress Genes Required for the Maintenance of Sub-Cellular Redox Homeostasis. <i>PLoS ONE</i> , 2012, 7, e44278.	1.1	40
11	Visualization of the interactome: What are we looking at?. <i>Proteomics</i> , 2012, 12, 1669-1686.	1.3	10
12	Time-Course Proteome Analysis Reveals the Dynamic Response of <i>Cryptococcus gattii</i> Cells to Fluconazole. <i>PLoS ONE</i> , 2012, 7, e42835.	1.1	17
13	Visualization and Analysis of the Complexome Network of <i>Saccharomyces cerevisiae</i> . <i>Journal of Proteome Research</i> , 2011, 10, 4744-4756.	1.8	25
14	Fourâ€“dimensional visualisation and analysis of proteinâ€“protein interaction networks. <i>Proteomics</i> , 2011, 11, 2672-2682.	1.3	16
15	Sequencing and Validation of the Genome of a <i>Campylobacter concisus</i> Reveals Intra-Species Diversity. <i>PLoS ONE</i> , 2011, 6, e22170.	1.1	32
16	Proteins Deleterious on Overexpression Are Associated with High Intrinsic Disorder, Specific Interaction Domains, and Low Abundance. <i>Journal of Proteome Research</i> , 2010, 9, 1218-1225.	1.8	18
17	The Interactorium: Visualising proteins, complexes and interaction networks in a virtual 3â€“D cell. <i>Proteomics</i> , 2009, 9, 5309-5315.	1.3	8