

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

687335

13
h-index

888047

17
g-index

20
all docs

20
docs citations

20
times ranked

3265
citing authors

#	ARTICLE	IF	CITATIONS
1	Disentangling the impact of environmental and phylogenetic constraints on prokaryotic within-species diversity. ISME Journal, 2020, 14, 1247-1259.	9.8	74
2	Extensive transmission of microbes along the gastrointestinal tract. ELife, 2019, 8, .	6.0	313
3	proGenomes: a resource for consistent functional and taxonomic annotations of prokaryotic genomes. Nucleic Acids Research, 2017, 45, D529-D534.	14.5	131
4	MOCAT2: a metagenomic assembly, annotation and profiling framework. Bioinformatics, 2016, 32, 2520-2523.	4.1	172
5	Durable coexistence of donor and recipient strains after fecal microbiota transplantation. Science, 2016, 352, 586-589.	12.6	461
6	Temporal and technical variability of human gut metagenomes. Genome Biology, 2015, 16, 73.	8.8	143
7	Molecular interaction networks for the analysis of human disease: Utility, limitations, and considerations. Proteomics, 2013, 13, 3393-3405.	2.2	17
8	Disturbed proteinâ€“protein interaction networks in metastatic melanoma are associated with worse prognosis and increased functional mutation burden. Pigment Cell and Melanoma Research, 2013, 26, 708-722.	3.3	12
9	A Multidimensional Matrix for Systems Biology Research and Its Application to Interaction Networks. Journal of Proteome Research, 2012, 11, 5204-5220.	3.7	5
10	A Genome-Wide Screen in Yeast Identifies Specific Oxidative Stress Genes Required for the Maintenance of Sub-Cellular Redox Homeostasis. PLoS ONE, 2012, 7, e44278.	2.5	40
11	Visualization of the interactome: What are we looking at?. Proteomics, 2012, 12, 1669-1686.	2.2	10
12	Time-Course Proteome Analysis Reveals the Dynamic Response of <i>Cryptococcus gattii</i> Cells to Fluconazole. PLoS ONE, 2012, 7, e42835.	2.5	17
13	Visualization and Analysis of the Complexome Network of <i>Saccharomyces cerevisiae</i> . Journal of Proteome Research, 2011, 10, 4744-4756.	3.7	25
14	Fourâ€“dimensional visualisation and analysis of proteinâ€“protein interaction networks. Proteomics, 2011, 11, 2672-2682.	2.2	16
15	Sequencing and Validation of the Genome of a <i>Campylobacter concisus</i> Reveals Intra-Species Diversity. PLoS ONE, 2011, 6, e22170.	2.5	32
16	Proteins Deleterious on Overexpression Are Associated with High Intrinsic Disorder, Specific Interaction Domains, and Low Abundance. Journal of Proteome Research, 2010, 9, 1218-1225.	3.7	18
17	The Interactorium: Visualising proteins, complexes and interaction networks in a virtual 3â€“D cell. Proteomics, 2009, 9, 5309-5315.	2.2	8