Ali Navid

List of Publications by Year in descending order

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	840776		940533	
17	374	11	16	
papers	citations	h-index	g-index	
19	19	19	715	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	CITATIONS
1	A Beginner's Guide to the COBRA Toolbox. Methods in Molecular Biology, 2022, 2349, 339-365.	0.9	2
2	Curating COBRA Models of Microbial Metabolism. Methods in Molecular Biology, 2022, 2349, 321-338.	0.9	1
3	Collagen Turnover in Relation to Risk Factors and Hemodynamics in Human Intracranial Aneurysms. Stroke, 2020, 51, 1624-1628.	2.0	18
4	System-level analysis of metabolic trade-offs during anaerobic photoheterotrophic growth in Rhodopseudomonas palustris. BMC Bioinformatics, 2019, 20, 233.	2.6	18
5	Combining multiple functional annotation tools increases coverage of metabolic annotation. BMC Genomics, 2018, 19, 948.	2.8	35
6	Application of a Physiologically Based Pharmacokinetic Model to Study Theophylline Metabolism and Its Interactions With Ciprofloxacin and Caffeine. CPT: Pharmacometrics and Systems Pharmacology, 2016, 5, 74-81.	2.5	7
7	Do genomeâ€scale models need exact solvers or clearer standards?. Molecular Systems Biology, 2015, 11, 831.	7.2	68
8	D‣actate production as a function of glucose metabolism in <i>Saccharomyces cerevisiae</i> . Yeast, 2013, 30, 81-91.	1.7	21
9	Rapid Countermeasure Discovery against Francisella tularensis Based on a Metabolic Network Reconstruction. PLoS ONE, 2013, 8, e63369.	2.5	19
10	Development of Constraint-Based System-Level Models of Microbial Metabolism. Methods in Molecular Biology, 2012, 881, 531-549.	0.9	0
11	Syntrophic metabolism of a co-culture containing Clostridium cellulolyticum and Rhodopseudomonas palustris for hydrogen production. International Journal of Hydrogen Energy, 2012, 37, 11719-11726.	7.1	30
12	Genome-level transcription data of Yersinia pestis analyzed with a New metabolic constraint-based approach. BMC Systems Biology, 2012, 6, 150.	3.0	68
13	Applications of system-level models of metabolism for analysis of bacterial physiology and identification of new drug targets. Briefings in Functional Genomics, 2011, 10, 354-364.	2.7	13
14	Yeast Dynamic Metabolic Flux Measurement in Nutrient-Rich Media by Hplc and Accelerator Mass Spectrometry. Analytical Chemistry, 2010, 82, 9812-9817.	6.5	15
15	Genome-scale reconstruction of the metabolic network in Yersinia pestis, strain 91001. Molecular BioSystems, 2009, 5, 368.	2.9	42
16	Systems Biology of Microbial Communities. Methods in Molecular Biology, 2009, 500, 469-494.	0.9	4
17	Simulated complex dynamics of glycolysis in the protozoan parasite Trypanosoma brucei. Journal of Theoretical Biology, 2004, 228, 449-458.	1.7	10