## Matthew DeJongh

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	High-throughput generation, optimization and analysis of genome-scale metabolic models. Nature Biotechnology, 2010, 28, 977-982.	17.5	974
2	KBase: The United States Department of Energy Systems Biology Knowledgebase. Nature Biotechnology, 2018, 36, 566-569.	17.5	955
3	Toward the automated generation of genome-scale metabolic networks in the SEED. BMC Bioinformatics, 2007, 8, 139.	2.6	120
4	The ModelSEED Biochemistry Database for the integration of metabolic annotations and the reconstruction, comparison and analysis of metabolic models for plants, fungi and microbes. Nucleic Acids Research, 2021, 49, D575-D588.	14.5	119
5	CytoSEED: a Cytoscape plugin for viewing, manipulating and analyzing metabolic models created by the Model SEED. Bioinformatics, 2012, 28, 891-892.	4.1	12
6	Computing and Applying Atomic Regulons to Understand Gene Expression and Regulation. Frontiers in Microbiology, 2016, 7, 1819.	3.5	7
7	A Bayesian Framework for the Classification of Microbial Gene Activity States. Frontiers in Microbiology, 2016, 7, 1191.	3.5	3
8	Implementing and evaluating a Gaussian mixture framework for identifying gene function from TnSeq data. , 2018, , .		1