

Creation and analysis of biochemical constraint-based models v.3.0

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Citation Report

#	ARTICLE	IF	CITATIONS
1	A systematic assessment of current genome-scale metabolic reconstruction tools. <i>Genome Biology</i> , 2019, 20, 158.	3.8	150
2	Predicting gastrointestinal drug effects using contextualized metabolic models. <i>PLoS Computational Biology</i> , 2019, 15, e1007100.	1.5	16
3	Genome-scale metabolic model of the rat liver predicts effects of diet restriction. <i>Scientific Reports</i> , 2019, 9, 9807.	1.6	10
4	Metabolic Model of the <i>Phytophthora infestans</i> -Tomato Interaction Reveals Metabolic Switches during Host Colonization. <i>MBio</i> , 2019, 10, .	1.8	23
5	Machine and deep learning meet genome-scale metabolic modeling. <i>PLoS Computational Biology</i> , 2019, 15, e1007084.	1.5	194
6	Integration of probabilistic regulatory networks into constraint-based models of metabolism with applications to Alzheimer's disease. <i>BMC Bioinformatics</i> , 2019, 20, 386.	1.2	13
7	Lysine harvesting is an antioxidant strategy and triggers underground polyamine metabolism. <i>Nature</i> , 2019, 572, 249-253.	13.7	99
10	Integrated Analyses of Microbiome and Longitudinal Metabolome Data Reveal Microbial-Host Interactions on Sulfur Metabolism in Parkinson's Disease. <i>Cell Reports</i> , 2019, 29, 1767-1777.e8.	2.9	102
11	Rewiring carbon metabolism in yeast for high level production of aromatic chemicals. <i>Nature Communications</i> , 2019, 10, 4976.	5.8	177
12	Genome-Scale Identification of Essential Metabolic Processes for Targeting the Plasmodium Liver Stage. <i>Cell</i> , 2019, 179, 1112-1128.e26.	13.5	92
13	High-resolution ¹³ C metabolic flux analysis. <i>Nature Protocols</i> , 2019, 14, 2856-2877.	5.5	132
14	Synthetic methylotrophy: Strategies to assimilate methanol for growth and chemicals production. <i>Current Opinion in Biotechnology</i> , 2019, 59, 165-174.	3.3	51
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17	Arterio-venous metabolomics exploration reveals major changes across liver and intestine in the obese Yucatan minipig. <i>Scientific Reports</i> , 2019, 9, 12527.	1.6	14
18	MOOMIN – Mathematical exploration of Omics data on a Metabolic Network. <i>Bioinformatics</i> , 2020, 36, 514-523.	1.8	15
19	Metabolic Modeling of Human Gut Microbiota on a Genome Scale: An Overview. <i>Metabolites</i> , 2019, 9, 22.	1.3	66
20	Integration of Metabolomic and Other Omics Data in Population-Based Study Designs: An Epidemiological Perspective. <i>Metabolites</i> , 2019, 9, 117.	1.3	47

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22	Human Systems Biology and Metabolic Modelling: A Review-From Disease Metabolism to Precision Medicine. <i>BioMed Research International</i> , 2019, 2019, 1-16.	0.9	56
23	Local convergence of the Levenberg-Marquardt method under Hölder metric subregularity. <i>Advances in Computational Mathematics</i> , 2019, 45, 2771-2806.	0.8	20
24	Systematic assessment of secondary bile acid metabolism in gut microbes reveals distinct metabolic capabilities in inflammatory bowel disease. <i>Microbiome</i> , 2019, 7, 75.	4.9	215
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38	<sc>Gapsplit</sc>: efficient random sampling for non-convex constraint-based models. <i>Bioinformatics</i> , 2020, 36, 2623-2625.	1.8	11

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