

Rahul Shaw

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4863145/publications.pdf>

Version: 2024-02-01

12
papers

418
citations

1163117

8
h-index

1199594

12
g-index

12
all docs

12
docs citations

12
times ranked

530
citing authors

#	ARTICLE	IF	CITATIONS
1	Single-Cell Transcriptome Analysis in Plants: Advances and Challenges. <i>Molecular Plant</i> , 2021, 14, 115-126.	8.3	127
2	Responses to Light Intensity in a Genome-Scale Model of Rice Metabolism. <i>Plant Physiology</i> , 2013, 162, 1060-1072.	4.8	117
3	A Dynamic Multi-Tissue Flux Balance Model Captures Carbon and Nitrogen Metabolism and Optimal Resource Partitioning During Arabidopsis Growth. <i>Frontiers in Plant Science</i> , 2018, 9, 884.	3.6	43
4	A Genome-Scale Metabolic Model of Soybean (<i>Glycine max</i>) Highlights Metabolic Fluxes in Seedlings. <i>Plant Physiology</i> , 2019, 180, 1912-1929.	4.8	43
5	A mass and charge balanced metabolic model of <i>Setaria viridis</i> revealed mechanisms of proton balancing in C4 plants. <i>BMC Bioinformatics</i> , 2019, 20, 357.	2.6	25
6	Reconstruction of <i>Oryza sativa indica</i> Genome Scale Metabolic Model and Its Responses to Varying RuBisCO Activity, Light Intensity, and Enzymatic Cost Conditions. <i>Frontiers in Plant Science</i> , 2017, 8, 2060.	3.6	21
7	Multi-tissue to whole plant metabolic modelling. <i>Cellular and Molecular Life Sciences</i> , 2020, 77, 489-495.	5.4	18
8	Metabolic trade-offs between biomass synthesis and photosynthate export at different light intensities in a genome-scale metabolic model of rice. <i>Frontiers in Plant Science</i> , 2014, 5, 656.	3.6	10
9	Flux balance analysis of genome-scale metabolic model of rice (<i>Oryza sativa</i>): Aiming to increase biomass. <i>Journal of Biosciences</i> , 2015, 40, 819-828.	1.1	6
10	Metabolic Plasticity and Inter-Compartmental Interactions in Rice Metabolism: An Analysis from Reaction Deletion Study. <i>PLoS ONE</i> , 2015, 10, e0133899.	2.5	3
11	Integration of crop growth model and constraint-based metabolic model predicts metabolic changes over rice plant development under water-limited stress. <i>In Silico Plants</i> , 2021, 3, .	1.9	3
12	Random Weighting through Linear Programming into Intracellular Transporters of Rice Metabolic Network. <i>Lecture Notes in Computer Science</i> , 2013, , 662-667.	1.3	2