

Sahra Uygun

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7594319/publications.pdf>

Version: 2024-02-01

12
papers

415
citations

933447

10
h-index

1281871

11
g-index

14
all docs

14
docs citations

14
times ranked

874
citing authors

#	ARTICLE	IF	CITATIONS
1	Phase II Trial of Neoadjuvant Carboplatin and Nab-Paclitaxel in Patients with Triple-Negative Breast Cancer. <i>Oncologist</i> , 2021, 26, e382-e393.	3.7	27
2	Optimising the use of gene expression data to predict plant metabolic pathway memberships. <i>New Phytologist</i> , 2021, 231, 475-489.	7.3	3
3	Cis-Regulatory Code for Predicting Plant Cell-Type Transcriptional Response to High Salinity. <i>Plant Physiology</i> , 2019, 181, 1739-1751.	4.8	29
4	Recovery from N Deprivation Is a Transcriptionally and Functionally Distinct State in <i>Chlamydomonas</i> . <i>Plant Physiology</i> , 2018, 176, 2007-2023.	4.8	30
5	Regulatory Divergence in Wound-Responsive Gene Expression between Domesticated and Wild Tomato. <i>Plant Cell</i> , 2018, 30, 1445-1460.	6.6	23
6	Predictive Models of Spatial Transcriptional Response to High Salinity. <i>Plant Physiology</i> , 2017, 174, 450-464.	4.8	24
7	Diversity, expansion, and evolutionary novelty of plant DNA-binding transcription factor families. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2017, 1860, 3-20.	1.9	75
8	Utility and Limitations of Using Gene Expression Data to Identify Functional Associations. <i>PLoS Computational Biology</i> , 2016, 12, e1005244.	3.2	63
9	The Impact of the Branched-Chain Ketoacid Dehydrogenase Complex on Amino Acid Homeostasis in <i>Arabidopsis</i> . <i>Plant Physiology</i> , 2015, 169, pp.00461.2015.	4.8	74
10	Measuring semantic similarities by combining gene ontology annotations and gene co-function networks. <i>BMC Bioinformatics</i> , 2015, 16, 44.	2.6	39
11	Molecular Evidence for Functional Divergence and Decay of a Transcription Factor Derived from Whole-Genome Duplication in <i>Arabidopsis thaliana</i> . <i>Plant Physiology</i> , 2015, 168, 1717-1734.	4.8	28
12	Evaluating the effects of clustering methods in co-expression-based functional inference in <i>Arabidopsis thaliana</i> . <i>New Biotechnology</i> , 2012, 29, S149.	4.4	0