

Yariv Brotman

List of Publications by Year in descending order

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Version: 2024-02-01

51
papers

2,355
citations

236925

25
h-index

223800

46
g-index

52
all docs

52
docs citations

52
times ranked

3082
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-wide association of the metabolic shifts underpinning dark-induced senescence in Arabidopsis. <i>Plant Cell</i> , 2022, 34, 557-578.	6.6	29
2	Bringing more players into play: Leveraging stress in genome wide association studies. <i>Journal of Plant Physiology</i> , 2022, 271, 153657.	3.5	11
3	Autophagy is required for lipid homeostasis during dark-induced senescence. <i>Plant Physiology</i> , 2021, 185, 1542-1558.	4.8	22
4	Multi-omics analysis of early leaf development in Arabidopsis thaliana. <i>Patterns</i> , 2021, 2, 100235.	5.9	24
5	When vegetation indicates reproduction: The affinity between leaf morphology and flowering commitment in the lily meristem. <i>Physiologia Plantarum</i> , 2021, 172, 2022-2033.	5.2	0
6	Mass spectrometry-based metabolomics: a guide for annotation, quantification and best reporting practices. <i>Nature Methods</i> , 2021, 18, 747-756.	19.0	403
7	The utility of metabolomics as a tool to inform maize biology. <i>Plant Communications</i> , 2021, 2, 100187.	7.7	17
8	Genomic basis underlying the metabolome-mediated drought adaptation of maize. <i>Genome Biology</i> , 2021, 22, 260.	8.8	44
9	It takes two: Reciprocal scion-rootstock relationships enable salt tolerance in 'Hass' avocado. <i>Plant Science</i> , 2021, 312, 111048.	3.6	3
10	Tomato Yellow Leaf Curl Virus (TYLCV) Promotes Plant Tolerance to Drought. <i>Cells</i> , 2021, 10, 2875.	4.1	19
11	Metabolomic Analysis of Natural Variation in Arabidopsis. <i>Methods in Molecular Biology</i> , 2021, 2200, 393-411.	0.9	0
12	Cytochrome respiration pathway and sulphur metabolism sustain stress tolerance to low temperature in the Antarctic species <i>Colobanthus quitensis</i> . <i>New Phytologist</i> , 2020, 225, 754-768.	7.3	32
13	The Acetate Pathway Supports Flavonoid and Lipid Biosynthesis in Arabidopsis. <i>Plant Physiology</i> , 2020, 182, 857-869.	4.8	35
14	Model-assisted identification of metabolic engineering strategies for <i>Jatropha curcas</i> lipid pathways. <i>Plant Journal</i> , 2020, 104, 76-95.	5.7	11
15	Multi-omics reveals mechanisms of total resistance to extreme illumination of a desert alga. <i>Nature Plants</i> , 2020, 6, 1031-1043.	9.3	33
16	Liquid Chromatography–Mass Spectrometry (LC–MS)–Based Analysis for Lipophilic Compound Profiling in Plants. <i>Current Protocols in Plant Biology</i> , 2020, 5, e20109.	2.8	16
17	Network-based strategies in metabolomics data analysis and interpretation: from molecular networking to biological interpretation. <i>Expert Review of Proteomics</i> , 2020, 17, 243-255.	3.0	70
18	Correlation-based network analysis combined with machine learning techniques highlight the role of the GABA shunt in <i>Brachypodium sylvaticum</i> freezing tolerance. <i>Scientific Reports</i> , 2020, 10, 4489.	3.3	13

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19	Towards model-driven characterization and manipulation of plant lipid metabolism. <i>Progress in Lipid Research</i> , 2020, 80, 101051.	11.6	28
20	Low-temperature tolerance of the Antarctic species <i>Deschampsia antarctica</i> : A complex metabolic response associated with nutrient remobilization. <i>Plant, Cell and Environment</i> , 2020, 43, 1376-1393.	5.7	21
21	Tomato yellow leaf curl virus (TYLCV)-resistant tomatoes share molecular mechanisms sustaining resistance with their wild progenitor <i>Solanum habrochaites</i> but not with TYLCV-susceptible tomatoes. <i>Plant Science</i> , 2020, 295, 110439.	3.6	13
22	Balancing the double-edged sword effect of increased resistant starch content and its impact on rice texture: its genetics and molecular physiological mechanisms. <i>Plant Biotechnology Journal</i> , 2020, 18, 1763-1777.	8.3	36
23	A Biostimulant Obtained from the Seaweed <i>Ascophyllum nodosum</i> Protects <i>Arabidopsis thaliana</i> from Severe Oxidative Stress. <i>International Journal of Molecular Sciences</i> , 2020, 21, 474.	4.1	62
24	Lipidomic and transcriptomic analysis reveals reallocation of carbon flux from cuticular wax into plastid membrane lipids in a glossy <i>Newhall</i> navel orange mutant. <i>Horticulture Research</i> , 2020, 7, 41.	6.3	23
25	Network Analysis Provides Insight into Tomato Lipid Metabolism. <i>Metabolites</i> , 2020, 10, 152.	2.9	10
26	Nano and Micro Unmanned Aerial Vehicles (UAVs): A New Grand Challenge for Precision Agriculture?. <i>Current Protocols in Plant Biology</i> , 2020, 5, e20103.	2.8	13
27	Large-scale metabolite quantitative trait locus analysis provides new insights for high-quality maize improvement. <i>Plant Journal</i> , 2019, 99, 216-230.	5.7	37
28	Branched-Chain Amino Acid Catabolism Impacts Triacylglycerol Homeostasis in <i>Chlamydomonas reinhardtii</i> . <i>Plant Physiology</i> , 2019, 179, 1502-1514.	4.8	26
29	The proof is in the bulb: glycerol influences key stages of lily development. <i>Plant Journal</i> , 2019, 97, 321-340.	5.7	21
30	An integrated multi-layered analysis of the metabolic networks of different tissues uncovers key genetic components of primary metabolism in maize. <i>Plant Journal</i> , 2018, 93, 1116-1128.	5.7	38
31	Unraveling lipid metabolism in maize with time-resolved multi-omics data. <i>Plant Journal</i> , 2018, 93, 1102-1115.	5.7	38
32	The metabolic (under)groundwork of the lily bulb toward sprouting. <i>Physiologia Plantarum</i> , 2018, 163, 436-449.	5.2	6
33	Salt tolerance of two perennial grass <i>Brachypodium sylvaticum</i> accessions. <i>Plant Molecular Biology</i> , 2018, 96, 305-314.	3.9	4
34	Mapping the Arabidopsis Metabolic Landscape by Untargeted Metabolomics at Different Environmental Conditions. <i>Molecular Plant</i> , 2018, 11, 118-134.	8.3	116
35	Differential lipidome remodeling during postharvest of peach varieties with different susceptibility to chilling injury. <i>Physiologia Plantarum</i> , 2018, 163, 2-17.	5.2	27
36	Quantitative Trait Loci Analysis Identifies a Prominent Gene Involved in the Production of Fatty Acid-Derived Flavor Volatiles in Tomato. <i>Molecular Plant</i> , 2018, 11, 1147-1165.	8.3	63

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37	Metabolome and Lipidome Profiles of <i>Populus</i> $\tilde{\text{A}}$ — <i>canescens</i> Twig Tissues During Annual Growth Show Phospholipid-Linked Storage and Mobilization of C, N, and S. <i>Frontiers in Plant Science</i> , 2018, 9, 1292.	3.6	18
38	Guidelines for Sample Normalization to Minimize Batch Variation for Large-Scale Metabolic Profiling of Plant Natural Genetic Variance. <i>Methods in Molecular Biology</i> , 2018, 1778, 33-46.	0.9	13
39	Molecular Mechanisms Preventing Senescence in Response to Prolonged Darkness in a Desiccation-Tolerant Plant. <i>Plant Physiology</i> , 2018, 177, 1319-1338.	4.8	26
40	Cucumber ovaries inhibited by dominant fruit express a dynamic developmental program, distinct from either senescence- or fruit-setting ovaries. <i>Plant Journal</i> , 2018, 96, 651-669.	5.7	8
41	Interorganelle Communication: Peroxisomal MALATE DEHYDROGENASE2 Connects Lipid Catabolism to Photosynthesis through Redox Coupling in <i>Chlamydomonas</i> . <i>Plant Cell</i> , 2018, 30, 1824-1847.	6.6	51
42	Integrated genomics-based mapping reveals the genetics underlying maize flavonoid biosynthesis. <i>BMC Plant Biology</i> , 2017, 17, 17.	3.6	34
43	Canalization of Tomato Fruit Metabolism. <i>Plant Cell</i> , 2017, 29, 2753-2765.	6.6	47
44	Combined Use of Genome-Wide Association Data and Correlation Networks Unravels Key Regulators of Primary Metabolism in <i>Arabidopsis thaliana</i> . <i>PLoS Genetics</i> , 2016, 12, e1006363.	3.5	67
45	Broadening Our Portfolio in the Genetic Improvement of Maize Chemical Composition. <i>Trends in Genetics</i> , 2016, 32, 459-469.	6.7	25
46	Omic Relief for the Biotically Stressed: Metabolomics of Plant Biotic Interactions. <i>Trends in Plant Science</i> , 2016, 21, 781-791.	8.8	76
47	Using lipidomics for expanding the knowledge on lipid metabolism in plants. <i>Biochimie</i> , 2016, 130, 91-96.	2.6	39
48	The maize leaf lipidome shows multilevel genetic control and high predictive value for agronomic traits. <i>Scientific Reports</i> , 2013, 3, 2479.	3.3	29
49	Trichoderma-Plant Root Colonization: Escaping Early Plant Defense Responses and Activation of the Antioxidant Machinery for Saline Stress Tolerance. <i>PLoS Pathogens</i> , 2013, 9, e1003221.	4.7	299
50	Transcript and metabolite analysis of the Trichoderma-induced systemic resistance response to <i>Pseudomonas syringae</i> in <i>Arabidopsis thaliana</i> . <i>Microbiology (United Kingdom)</i> , 2012, 158, 139-146.	1.8	172
51	Trichoderma. <i>Current Biology</i> , 2010, 20, R390-R391.	3.9	85