

Borjana Arsova

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2665693/publications.pdf>

Version: 2024-02-01

20
papers

1,762
citations

623734

14
h-index

713466

21
g-index

28
all docs

28
docs citations

28
times ranked

3013
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Modulators or facilitators? Roles of lipids in plant root-microbe interactions. Trends in Plant Science, 2022, 27, 180-190. | 8.8 | 45 |
| 2 | The molecular basis of zinc homeostasis in cereals. Plant, Cell and Environment, 2022, 45, 1339-1361. | 5.7 | 14 |
| 3 | Root Growth and Architecture of Wheat and Brachypodium Vary in Response to Algal Fertilizer in Soil and Solution. Agronomy, 2022, 12, 285. | 3.0 | 4 |
| 4 | N-dependent dynamics of root growth and nitrate and ammonium uptake are altered by the bacterium <i>Herbaspirillum seropedicae</i> in the cereal model <i>Brachypodium distachyon</i> . Journal of Experimental Botany, 2022, 73, 5306-5321. | 4.8 | 11 |
| 5 | Isolation of Novel Xanthomonas Phages Infecting the Plant Pathogens <i>X. translucens</i> and <i>X. campestris</i> . Viruses, 2022, 14, 1449. | 3.3 | 6 |
| 6 | Time-resolution of the shoot and root growth of the model cereal Brachypodium in response to inoculation with Azospirillum bacteria at low phosphorus and temperature. Plant Growth Regulation, 2021, 93, 149-162. | 3.4 | 10 |
| 7 | The Metabolic Response of Brachypodium Roots to the Interaction with Beneficial Bacteria Is Affected by the Plant Nutritional Status. Metabolites, 2021, 11, 358. | 2.9 | 8 |
| 8 | Transcriptional regulation of <i>ZIP</i> genes is independent of local zinc status in Brachypodium shoots upon zinc deficiency and resupply. Plant, Cell and Environment, 2021, 44, 3376-3397. | 5.7 | 9 |
| 9 | Dynamics in plant roots and shoots minimize stress, save energy and maintain water and nutrient uptake. New Phytologist, 2020, 225, 1111-1119. | 7.3 | 37 |
| 10 | Energy costs of salt tolerance in crop plants. New Phytologist, 2020, 225, 1072-1090. | 7.3 | 284 |
| 11 | MapMan4: A Refined Protein Classification and Annotation Framework Applicable to Multi-Omics Data Analysis. Molecular Plant, 2019, 12, 879-892. | 8.3 | 353 |
| 12 | Multilab EcoFAB study shows highly reproducible physiology and depletion of soil metabolites by a model grass. New Phytologist, 2019, 222, 1149-1160. | 7.3 | 55 |
| 13 | Plant genome and transcriptome annotations: from misconceptions to simple solutions. Briefings in Bioinformatics, 2018, 19, bbw135. | 6.5 | 62 |
| 14 | Monitoring of Plant Protein Post-translational Modifications Using Targeted Proteomics. Frontiers in Plant Science, 2018, 9, 1168. | 3.6 | 41 |
| 15 | Evolution of the Phosphoenolpyruvate Carboxylase Protein Kinase Family in C3 and C4 <i>Flaveria</i> spp. Plant Physiology, 2014, 165, 1076-1091. | 4.8 | 23 |
| 16 | The genome of the stress-tolerant wild tomato species <i>Solanum pennellii</i> . Nature Genetics, 2014, 46, 1034-1038. | 21.4 | 391 |
| 17 | Precision, Proteome Coverage, and Dynamic Range of Arabidopsis Proteome Profiling Using ¹⁵ N Metabolic Labeling and Label-free Approaches. Molecular and Cellular Proteomics, 2012, 11, 619-628. | 3.8 | 16 |
| 18 | The use of heavy nitrogen in quantitative proteomics experiments in plants. Trends in Plant Science, 2012, 17, 102-112. | 8.8 | 32 |

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|----|--|-----|-----------|
| 19 | Current status of the plant phosphorylation site database PhosPhAt and its use as a resource for molecular plant physiology. <i>Frontiers in Plant Science</i> , 2012, 3, 132. | 3.6 | 16 |
| 20 | Plastidial Thioredoxin <i>z</i> Interacts with Two Fructokinase-Like Proteins in a Thiol-Dependent Manner: Evidence for an Essential Role in Chloroplast Development in <i>Arabidopsis</i> and <i>Nicotiana benthamiana</i> . <i>Plant Cell</i> , 2010, 22, 1498-1515. | 6.6 | 281 |