

Joaquã-n Giner-Lamia

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7924939/publications.pdf>

Version: 2024-02-01

19
papers

713
citations

687363

13
h-index

839539

18
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21
all docs

21
docs citations

21
times ranked

1047
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Towards the biogeography of prokaryotic genes. <i>Nature</i> , 2022, 601, 252-256. | 27.8 | 85 |
| 2 | Extracellular vesicles as an alternative copper-secretion mechanism in bacteria. <i>Journal of Hazardous Materials</i> , 2022, 431, 128594. | 12.4 | 14 |
| 3 | GeCoViz: genomic context visualisation of prokaryotic genes from a functional and evolutionary perspective. <i>Nucleic Acids Research</i> , 2022, 50, W352-W357. | 14.5 | 9 |
| 4 | Structure-based analyses of <i>Salmonella</i> RcsB variants unravel new features of the Rcs regulon. <i>Nucleic Acids Research</i> , 2021, 49, 2357-2374. | 14.5 | 10 |
| 5 | A protease-mediated mechanism regulates the cytochrome <i>c</i> ₆ /plastocyanin switch in <i>Synechocystis</i> sp. PCC 6803. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, . | 7.1 | 18 |
| 6 | Genome analysis of <i>Salmonella enterica</i> subsp. <i>diarizonae</i> isolates from invasive human infections reveals enrichment of virulence-related functions in lineage ST1256. <i>BMC Genomics</i> , 2019, 20, 99. | 2.8 | 24 |
| 7 | ChIP-seq Experiment and Data Analysis in the Cyanobacterium <i>Synechocystis</i> sp. PCC 6803. <i>Bio-protocol</i> , 2018, 8, e2895. | 0.4 | 0 |
| 8 | Identification of the direct regulon of NtcA during early acclimation to nitrogen starvation in the cyanobacterium <i>Synechocystis</i> sp. PCC 6803. <i>Nucleic Acids Research</i> , 2017, 45, 11800-11820. | 14.5 | 82 |
| 9 | Draft Genome Sequence of <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Infantis Strain SPE101, Isolated from a Chronic Human Infection. <i>Genome Announcements</i> , 2017, 5, . | 0.8 | 10 |
| 10 | Extracellular Proteins: Novel Key Components of Metal Resistance in Cyanobacteria?. <i>Frontiers in Microbiology</i> , 2016, 7, 878. | 3.5 | 37 |
| 11 | The Transcriptional Landscape of the Photosynthetic Model Cyanobacterium <i>Synechocystis</i> sp. PCC6803. <i>Scientific Reports</i> , 2016, 6, 22168. | 3.3 | 47 |
| 12 | Ni interferes in the Cu-regulated transcriptional switch <i>petJ/petE</i> in <i>Synechocystis</i> sp. PCC 6803. <i>FEBS Letters</i> , 2016, 590, 3639-3648. | 2.8 | 5 |
| 13 | CopM is a novel copper-binding protein involved in copper resistance in <i>Synechocystis</i> sp. PCC 6803. <i>MicrobiologyOpen</i> , 2015, 4, 167-185. | 3.0 | 30 |
| 14 | The unfolded protein response and its potential role in Huntington's disease elucidated by a systems biology approach. <i>Frontiers in Molecular and Cellular Biosciences</i> , 2015, 4, 103. | 1.6 | 32 |
| 15 | The unfolded protein response and its potential role in Huntington's disease elucidated by a systems biology approach. <i>Frontiers in Molecular and Cellular Biosciences</i> , 2015, 4, 103. | 1.6 | 29 |
| 16 | Metals in Cyanobacteria: Analysis of the Copper, Nickel, Cobalt and Arsenic Homeostasis Mechanisms. <i>Life</i> , 2014, 4, 865-886. | 2.4 | 124 |
| 17 | Global Transcriptional Profiles of the Copper Responses in the Cyanobacterium <i>Synechocystis</i> sp. PCC 6803. <i>PLoS ONE</i> , 2014, 9, e108912. | 2.5 | 46 |
| 18 | The CopRS Two-Component System Is Responsible for Resistance to Copper in the Cyanobacterium <i>Synechocystis</i> sp. PCC 6803. <i>Plant Physiology</i> , 2012, 159, 1806-1818. | 4.8 | 88 |

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|----|---|-----|-----------|
| 19 | Redox control of copper homeostasis in cyanobacteria. <i>Plant Signaling and Behavior</i> , 2012, 7, 1712-1714. | 2.4 | 15 |