

# Gothandapani Sellamuthu

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9741771/publications.pdf>

Version: 2024-02-01

18  
papers

592  
citations

1163117

8  
h-index

940533

16  
g-index

18  
all docs

18  
docs citations

18  
times ranked

875  
citing authors

| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 1  | Reduced apoplastic barriers in tissues of shoot-proximal rhizomes of <i>Oryza coarctata</i> are associated with Na <sup>+</sup> sequestration. <i>Journal of Experimental Botany</i> , 2022, 73, 998-1015.  | 4.8 | 2         |
| 2  | Diversity of Sodium Transporter HKT1;5 in Genus <i>Oryza</i> . <i>Rice Science</i> , 2022, 29, 31-46.   | 3.9 | 3         |
| 3  | Proto Kranz-like leaf traits and cellular ionic regulation are associated with salinity tolerance in a halophytic wild rice. <i>Stress Biology</i> , 2022, 2, 1.  | 3.1 | 4         |
| 4  | Rifampicin Increases Expression of Plant Codon-Optimized <i>Bacillus thuringiensis</i> $\delta$ -Endotoxin Genes in <i>Escherichia coli</i> . <i>Protein Journal</i> , 2022, , 1.   | 1.6 | 3         |
| 5  | Comparative Analysis of Root Na <sup>+</sup> Relation under Salinity between <i>Oryza sativa</i> and <i>Oryza coarctata</i> . <i>Plants</i> , 2022, 11, 656.  | 3.5 | 7         |
| 6  | Unravelling the physiological basis of salinity stress tolerance in cultivated and wild rice species. <i>Functional Plant Biology</i> , 2022, 49, 351-364.  | 2.1 | 12        |
| 7  | Identifying optimal reference genes for gene expression studies in Eurasian spruce bark beetle, <i>Ips typographus</i> (Coleoptera: Curculionidae: Scolytinae). <i>Scientific Reports</i> , 2022, 12, 4671.   | 3.3 | 6         |
| 8  | Targeting delta-endotoxin (Cry1Ac) of <i>Bacillus thuringiensis</i> to subcellular compartments increases the protein expression, stability, and biological activity. <i>International Journal of Biological Macromolecules</i> , 2022, 205, 185-192. | 7.5 | 0         |
| 9  | Reference Gene Selection for Normalizing Gene Expression in <i>Ips Sexdentatus</i> (Coleoptera:) Tj ETQq1 1 0.784314 rgBT /Overlock 10 T 752768.  | 2.8 | 11        |
| 10 | To exclude or to accumulate? Revealing the role of the sodium HKT1;5 transporter in plant adaptive responses to varying soil salinity. <i>Plant Physiology and Biochemistry</i> , 2021, 169, 333-342.   | 5.8 | 20        |
| 11 | Distinct Evolutionary Origins of Intron Retention Splicing Events in NHX1 Antiporter Transcripts Relate to Sequence Specific Distinctions in <i>Oryza</i> Species. <i>Frontiers in Plant Science</i> , 2020, 11, 267.                                 | 3.6 | 16        |
| 12 | Microhair on the adaxial leaf surface of salt secreting halophytic <i>Oryza coarctata</i> Roxb. show distinct morphotypes: Isolation for molecular and functional analysis. <i>Plant Science</i> , 2019, 285, 248-257.                                | 3.6 | 16        |
| 13 | A quick, easy and cost-effective in planta method to develop direct transformants in wheat. <i>3 Biotech</i> , 2019, 9, 180.  | 2.2 | 6         |
| 14 | CRISPR for Crop Improvement: An Update Review. <i>Frontiers in Plant Science</i> , 2018, 9, 985.  | 3.6 | 425       |
| 15 | Molecular Evolution of the Negative Regulatory Gene (NIFL) from <i>Azotobacter Chroococcum</i> and its Nitrogenase Activity. <i>Biosciences, Biotechnology Research Asia</i> , 2018, 15, 397-406.   | 0.5 | 0         |
| 16 | An Environmentally Friendly Engineered <i>Azotobacter</i> Strain That Replaces a Substantial Amount of Urea Fertilizer while Sustaining the Same Wheat Yield. <i>Applied and Environmental Microbiology</i> , 2017, 83, .                             | 3.1 | 41        |
| 17 | Evaluation of entomopathogenic fungus against <i>Alternaria porri</i> (Ellis) causing purple blotch disease of onion. <i>Archives of Phytopathology and Plant Protection</i> , 2015, 48, 135-144.   | 1.3 | 12        |
| 18 | GhDRIN1, a novel drought-induced gene of upland cotton ( <i>Gossypium hirsutum</i> L.) confers abiotic and biotic stress tolerance in transgenic tobacco. <i>Biotechnology Letters</i> , 2015, 37, 907-919.   | 2.2 | 8         |