## Sergio Casas-Flores

List of Publications by Year in descending order

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| #  | Article   | IF  | CITATIONS |
|----|---|-----|-----------|
| 1  | Comparative genome sequence analysis underscores mycoparasitism as the ancestral life style of<br>Trichoderma. Genome Biology, 2011, 12, R40.   | 3.8 | 594       |
| 2  | Colonization of Arabidopsis roots by Trichoderma atroviride promotes growth and enhances<br>systemic disease resistance through jasmonic acid/ethylene and salicylic acid pathways. European<br>Journal of Plant Pathology, 2011, 131, 15-26.   | 0.8 | 231       |
| 3  | The Genomes of Three Uneven Siblings: Footprints of the Lifestyles of Three Trichoderma Species.<br>Microbiology and Molecular Biology Reviews, 2016, 80, 205-327.  | 2.9 | 194       |
| 4  | BLR-1 and BLR-2, key regulatory elements of photoconidiation and mycelial growth in Trichoderma atroviride. Microbiology (United Kingdom), 2004, 150, 3561-3569.  | 0.7 | 163       |
| 5  | Cross Talk between a Fungal Blue-Light Perception System and the Cyclic AMP Signaling Pathway.<br>Eukaryotic Cell, 2006, 5, 499-506.  | 3.4 | 108       |
| 6  | The Epl1 and Sm1 proteins from Trichoderma atroviride and Trichoderma virens differentially<br>modulate systemic disease resistance against different life style pathogens in Solanum lycopersicum.<br>Frontiers in Plant Science, 2015, 6, 77. | 1.7 | 93        |
| 7  | Role of the 4-Phosphopantetheinyl Transferase of <i>Trichoderma virens</i> in Secondary Metabolism<br>and Induction of Plant Defense Responses. Molecular Plant-Microbe Interactions, 2011, 24, 1459-1471.                                      | 1.4 | 89        |
| 8  | Trichoderma as a Model to Study Effector-Like Molecules. Frontiers in Microbiology, 2019, 10, 1030.   | 1.5 | 86        |
| 9  | Mango (Mangifera indica L.) cv. Kent fruit mesocarp de novo transcriptome assembly identifies gene<br>families important for ripening. Frontiers in Plant Science, 2015, 6, 62.   | 1.7 | 76        |
| 10 | Novel light-regulated genes in Trichoderma atroviride: a dissection by cDNA microarrays.<br>Microbiology (United Kingdom), 2006, 152, 3305-3317.  | 0.7 | 74        |
| 11 | The Plant Growth-Promoting Fungus Aspergillus ustus Promotes Growth and Induces Resistance<br>Against Different Lifestyle Pathogens in Arabidopsis thaliana. Journal of Microbiology and<br>Biotechnology, 2011, 21, 686-696.                   | 0.9 | 64        |
| 12 | <i>Trichoderma</i> Histone Deacetylase HDA-2 Modulates Multiple Responses in Arabidopsis. Plant<br>Physiology, 2019, 179, 1343-1361.  | 2.3 | 50        |
| 13 | Humic Substances Mediate Anaerobic Methane Oxidation Linked to Nitrous Oxide Reduction in<br>Wetland Sediments. Frontiers in Microbiology, 2020, 11, 587.   | 1.5 | 50        |
| 14 | Antifungal Nanocomposites Inspired by Titanate Nanotubes for Complete Inactivation of <i>Botrytis<br/>cinerea</i> Isolated from Tomato Infection. ACS Applied Materials & Interfaces, 2016, 8, 31625-31637.                                     | 4.0 | 41        |
| 15 | Molecular Mechanisms of Biocontrol in Trichoderma spp. and Their Applications in Agriculture. , 2014, , 429-453.  |     | 38        |
| 16 | Genome-Wide Identification of Mango (Mangifera indica L.) Polygalacturonases: Expression Analysis of<br>Family Members and Total Enzyme Activity During Fruit Ripening. Frontiers in Plant Science, 2019, 10,<br>969.                           | 1.7 | 34        |
| 17 | Performance of innovative PU-foam and natural fiber-based composites for the biofiltration of a mixture of volatile organic compounds by a fungal biofilm. Journal of Hazardous Materials, 2012, 201-202, 202-208.                              | 6.5 | 32        |
| 18 | New roles for CDC25 in growth control, galactose regulation and cellular differentiation in Saccharomyces cerevisiae. Microbiology (United Kingdom), 2004, 150, 2865-2879.  | 0.7 | 30        |

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|----|--|-----|-----------|
| 19 | The separation between the 5′-3′ ends in long RNA molecules is short and nearly constant. Nucleic Acids<br>Research, 2014, 42, 13963-13968.  | 6.5 | 30        |
| 20 | The Small GTPases in Fungal Signaling Conservation and Function. Cells, 2021, 10, 1039.  | 1.8 | 29        |
| 21 | Mesocarp RNA-Seq analysis of mango ( Mangifera indica L.) identify quarantine postharvest treatment<br>effects on gene expression. Scientia Horticulturae, 2018, 227, 146-153.   | 1.7 | 25        |
| 22 | Histone Deacetylase HDA-2 Regulates Trichoderma atroviride Growth, Conidiation, Blue Light<br>Perception, and Oxidative Stress Responses. Applied and Environmental Microbiology, 2017, 83, .  | 1.4 | 22        |
| 23 | Generation of Sporothrix schenckii mutants expressing the green fluorescent protein suitable for the study of host-fungus interactions. Fungal Biology, 2018, 122, 1023-1030.  | 1.1 | 21        |
| 24 | Thermophile mats of microalgae growing on the woody structure of a cooling tower of a<br>thermoelectric power plant in Central Mexico. Revista Mexicana De Biodiversidad, 2016, 87, 277-287.   | 0.4 | 20        |
| 25 | Silencing of <em>OCH1 </em> unveils the role of <em>Sporothrix schenckii</em><br><em>N</em> -linked glycans during the host–fungus interaction. Infection and<br>Drug Resistance, 2019, Volume 12, 67-85.  | 1.1 | 20        |
| 26 | Histone acetyltransferase TGF-1 regulates Trichoderma atroviride secondary metabolism and mycoparasitism. PLoS ONE, 2018, 13, e0193872.  | 1.1 | 19        |
| 27 | TBRG-1 a Ras-like protein in Trichoderma virens involved in conidiation, development, secondary<br>metabolism, mycoparasitism, and biocontrol unveils a new family of Ras-GTPases. Fungal Genetics and<br>Biology, 2020, 136, 103292.  | 0.9 | 19        |
| 28 | Three Decades of Fungal Transformation: Novel Technologies. , 2004, 267, 315-326.  |     | 18        |
| 29 | Performance and Bacterial Population Composition of an n-Hexane Degrading Biofilter Working<br>Under Fluctuating Conditions. Applied Biochemistry and Biotechnology, 2014, 174, 832-844.   | 1.4 | 15        |
| 30 | The small RNAâ€mediated gene silencing machinery is required in Arabidopsis for stimulation of growth,<br>systemic disease resistance, and suppression of the nitrileâ€specifier gene <i>NSP4</i> by <i>Trichoderma<br/>atroviride</i> . Plant Journal, 2022, 109, 873-890.                        | 2.8 | 13        |
| 31 | Community of thermoacidophilic and arsenic resistant microorganisms isolated from a deep profile of mine heaps. AMB Express, 2015, 5, 132.   | 1.4 | 12        |
| 32 | Photo-assisted inactivation of Escherichia coli bacteria by silver functionalized titanate nanotubes,<br>Ag/H2Ti2O5·H2O. Photochemical and Photobiological Sciences, 2017, 16, 854-860.  | 1.6 | 12        |
| 33 | Secretome Analysis of Arabidopsis–Trichoderma atroviride Interaction Unveils New Roles for the<br>Plant Glutamate:Glyoxylate Aminotransferase GGAT1 in Plant Growth Induced by the Fungus and<br>Resistance against Botrytis cinerea. International Journal of Molecular Sciences, 2021, 22, 6804. | 1.8 | 12        |
| 34 | The <i><scp>T</scp>richodermaÂatroviride</i> photolyaseâ€encoding gene is transcriptionally regulated<br>by nonâ€canonical light response elements. FEBS Journal, 2013, 280, 3697-3708.  | 2.2 | 11        |
| 35 | Automated, continuous video microscopy tracking of hyphal growth. Fungal Genetics and Biology, 2019, 123, 25-32.   | 0.9 | 11        |
| 36 | IPA-1 a Putative Chromatin Remodeler/Helicase-Related Protein of <i>Trichoderma virens</i> Plays<br>Important Roles in Antibiosis Against <i>Rhizoctonia solani</i> and Induction of <i>Arabidopsis</i> Systemic Disease Resistance, Molecular Plant-Microbe Interactions, 2020, 33, 808-824       | 1.4 | 10        |

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| 37 | Light induces oxidative damage and protein stability in the fungal photoreceptor Vivid. PLoS ONE, 2018, 13, e0201028.   | 1.1 | 9         |
| 38 | 3 The Bright and Dark Sides of Fungal Life. , 2016, , 41-77.  |     | 8         |
| 39 | Microbial contamination in methanol biofilters inoculated with a pure strain of <i>Pichia pastoris</i> : A potential limitation for waste revalorization. Biotechnology Progress, 2019, 35, e2715.                                | 1.3 | 5         |
| 40 | Actividad de dos hongos entomopatÃ <sup>3</sup> genos, identificados molecularmente, sobre Bactericera<br>cockerelli. Revista Colombiana De Entomologia, 2017, 43, 27.  | 0.1 | 5         |
| 41 | Differential distribution of transcripts from genes involved in polyamine biosynthesis in bean plants.<br>Biologia Plantarum, 2006, 50, 551-558.  | 1.9 | 4         |
| 42 | Unraveling the photoactive annihilation mechanism of nanostructures as effective green tools for<br>inhibiting the proliferation of the phytopathogenic bacteriumPseudomonas syringae. Nanoscale<br>Advances, 2019, 1, 2258-2267. | 2.2 | 4         |
| 43 | Trichoderma in the rhizosphere. , 2020, , 3-38.   |     | 4         |
| 44 | Characterization of the trypsin-III from Monterey sardine (Sardinops caeruleus): Insights on the cold-adaptation from the A236N mutant. International Journal of Biological Macromolecules, 2020, 164, 2701-2710.                 | 3.6 | 2         |
| 45 | Molecular modeling and expression analysis of a MADS-box cDNA from mango (Mangifera indica L.). 3<br>Biotech, 2014, 4, 357-365.   | 1.1 | 1         |
| 46 | 13 Nematophagous Fungi. , 2016, , 247-267.  |     | 1         |
| 47 | Analysis of bacterial communities of infected primary teeth in a Mexican population. Medicina Oral,<br>Patologia Oral Y Cirugia Bucal, 2020, 25, e668-e674.   | 0.7 | 0         |