

Susana M Coelho

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

61
papers

2,138
citations

23
h-index

45
g-index

72
ext. papers

2,655
ext. citations

8.2
avg, IF

4.46
L-index

| # | Paper | IF | Citations |
|----|---|------|-----------|
| 61 | The Ectocarpus genome and the independent evolution of multicellularity in brown algae. <i>Nature</i> , 2010 , 465, 617-21 | 50.4 | 645 |
| 60 | Spatiotemporal patterning of reactive oxygen production and Ca(2+) wave propagation in fucus rhizoid cells. <i>Plant Cell</i> , 2002 , 14, 2369-81 | 11.6 | 140 |
| 59 | Development and physiology of the brown alga Ectocarpus siliculosus: two centuries of research. <i>New Phytologist</i> , 2008 , 177, 319-332 | 9.8 | 103 |
| 58 | A haploid system of sex determination in the brown alga Ectocarpus sp. <i>Current Biology</i> , 2014 , 24, 1945-57 | 5.7 | 93 |
| 57 | Complex life cycles of multicellular eukaryotes: new approaches based on the use of model organisms. <i>Gene</i> , 2007 , 406, 152-70 | 3.8 | 92 |
| 56 | Life-cycle-generation-specific developmental processes are modified in the immediate upright mutant of the brown alga Ectocarpus siliculosus. <i>Development (Cambridge)</i> , 2008 , 135, 1503-12 | 6.6 | 76 |
| 55 | OUROBOROS is a master regulator of the gametophyte to sporophyte life cycle transition in the brown alga Ectocarpus. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011 , 108, 11518-23 | 11.5 | 69 |
| 54 | Sexual dimorphism and the evolution of sex-biased gene expression in the brown alga ectocarpus. <i>Molecular Biology and Evolution</i> , 2015 , 32, 1581-97 | 8.3 | 68 |
| 53 | A sequence-tagged genetic map for the brown alga Ectocarpus siliculosus provides large-scale assembly of the genome sequence. <i>New Phytologist</i> , 2010 , 188, 42-51 | 9.8 | 51 |
| 52 | The Algal Revolution. <i>Trends in Plant Science</i> , 2017 , 22, 726-738 | 13.1 | 48 |
| 51 | Re-annotation, improved large-scale assembly and establishment of a catalogue of noncoding loci for the genome of the model brown alga Ectocarpus. <i>New Phytologist</i> , 2017 , 214, 219-232 | 9.8 | 46 |
| 50 | Evolution and regulation of complex life cycles: a brown algal perspective. <i>Current Opinion in Plant Biology</i> , 2014 , 17, 1-6 | 9.9 | 41 |
| 49 | Role of endoreduplication and apomeiosis during parthenogenetic reproduction in the model brown alga Ectocarpus. <i>New Phytologist</i> , 2010 , 188, 111-21 | 9.8 | 40 |
| 48 | How to cultivate Ectocarpus. <i>Cold Spring Harbor Protocols</i> , 2012 , 2012, 258-61 | 1.2 | 39 |
| 47 | Evolution and maintenance of haploid-diploid life cycles in natural populations: The case of the marine brown alga Ectocarpus. <i>Evolution; International Journal of Organic Evolution</i> , 2015 , 69, 1808-22 | 3.8 | 36 |
| 46 | Ectocarpus: a model organism for the brown algae. <i>Cold Spring Harbor Protocols</i> , 2012 , 2012, 193-8 | 1.2 | 33 |
| 45 | Convergent recruitment of TALE homeodomain life cycle regulators to direct sporophyte development in land plants and brown algae. <i>ELife</i> , 2019 , 8, | 8.9 | 33 |

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|----|---|------|----|
| 44 | Multiple gene movements into and out of haploid sex chromosomes. <i>Genome Biology</i> , 2017 , 18, 104 | 18.3 | 31 |
| 43 | Phylogeny and Evolution of the Brown Algae. <i>Critical Reviews in Plant Sciences</i> , 2020 , 39, 281-321 | 5.6 | 31 |
| 42 | microRNAs and the evolution of complex multicellularity: identification of a large, diverse complement of microRNAs in the brown alga <i>Ectocarpus</i> . <i>Nucleic Acids Research</i> , 2015 , 43, 6384-98 | 20.1 | 28 |
| 41 | High-density genetic map and identification of QTLs for responses to temperature and salinity stresses in the model brown alga <i>Ectocarpus</i> . <i>Scientific Reports</i> , 2017 , 7, 43241 | 4.9 | 25 |
| 40 | Algal models in plant biology. <i>Journal of Experimental Botany</i> , 2011 , 62, 2425-30 | 7 | 25 |
| 39 | Development of PCR-Based Markers to Determine the Sex of Kelps. <i>PLoS ONE</i> , 2015 , 10, e0140535 | 3.7 | 23 |
| 38 | Algal Sex Determination and the Evolution of Anisogamy. <i>Annual Review of Microbiology</i> , 2019 , 73, 267-291 | 15 | 22 |
| 37 | The cell-wall active mannuronan C5-epimerases in the model brown alga <i>Ectocarpus</i> : From gene context to recombinant protein. <i>Glycobiology</i> , 2016 , 26, 973-983 | 5.8 | 22 |
| 36 | Gene silencing in <i>Fucus</i> embryos: developmental consequences of RNAi-mediated cytoskeletal disruption. <i>Journal of Phycology</i> , 2013 , 49, 819-29 | 3 | 20 |
| 35 | The <i>Ectocarpus</i> IMMEDIATE UPRIGHT gene encodes a member of a novel family of cysteine-rich proteins with an unusual distribution across the eukaryotes. <i>Development (Cambridge)</i> , 2017 , 144, 409-418 | 6.6 | 18 |
| 34 | Rapid turnover of life-cycle-related genes in the brown algae. <i>Genome Biology</i> , 2019 , 20, 35 | 18.3 | 17 |
| 33 | Non-cell autonomous regulation of life cycle transitions in the model brown alga <i>Ectocarpus</i> . <i>New Phytologist</i> , 2013 , 197, 503-510 | 9.8 | 17 |
| 32 | The Pseudoautosomal Regions of the U/V Sex Chromosomes of the Brown Alga <i>Ectocarpus</i> Exhibit Unusual Features. <i>Molecular Biology and Evolution</i> , 2015 , 32, 2973-85 | 8.3 | 16 |
| 31 | Rapid Evolution of microRNA Loci in the Brown Algae. <i>Genome Biology and Evolution</i> , 2017 , 9, 740-749 | 3.9 | 15 |
| 30 | Genetic crosses between <i>Ectocarpus</i> strains. <i>Cold Spring Harbor Protocols</i> , 2012 , 2012, 262-5 | 1.2 | 13 |
| 29 | DISTAG/TBCCd1 Is Required for Basal Cell Fate Determination in. <i>Plant Cell</i> , 2017 , 29, 3102-3122 | 11.6 | 12 |
| 28 | A key role for sex chromosomes in the regulation of parthenogenesis in the brown alga <i>Ectocarpus</i> . <i>PLoS Genetics</i> , 2019 , 15, e1008211 | 6 | 11 |
| 27 | instaGRAAL: chromosome-level quality scaffolding of genomes using a proximity ligation-based scaffolder. <i>Genome Biology</i> , 2020 , 21, 148 | 18.3 | 11 |

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| 26 | Origin and evolution of sex-determination systems in the brown algae. <i>New Phytologist</i> , 2019 , 222, 1751-1756 | 17.56 | 10 |
| 25 | Genetic Diversity in the UV Sex Chromosomes of the Brown Alga. <i>Genes</i> , 2018 , 9, | 4.2 | 10 |
| 24 | Isolation and regeneration of protoplasts from Ectocarpus. <i>Cold Spring Harbor Protocols</i> , 2012 , 2012, 361-4 | 1.2 | 10 |
| 23 | Genome-wide comparison of ultraviolet and ethyl methanesulphonate mutagenesis methods for the brown alga Ectocarpus. <i>Marine Genomics</i> , 2015 , 24 Pt 1, 109-13 | 1.9 | 9 |
| 22 | The origin and evolution of the sexes: Novel insights from a distant eukaryotic lineage. <i>Comptes Rendus - Biologies</i> , 2016 , 339, 252-7 | 1.4 | 9 |
| 21 | Histone modifications during the life cycle of the brown alga Ectocarpus. <i>Genome Biology</i> , 2021 , 22, 12 | 18.3 | 9 |
| 20 | Genetic regulation of life cycle transitions in the brown alga Ectocarpus. <i>Plant Signaling and Behavior</i> , 2011 , 6, 1858-60 | 2.5 | 8 |
| 19 | Brown Algal Model Organisms. <i>Annual Review of Genetics</i> , 2020 , 54, 71-92 | 14.5 | 7 |
| 18 | Targeted CRISPR-Cas9-based gene knockouts in the model brown alga Ectocarpus. <i>New Phytologist</i> , 2021 , 231, 2077-2091 | 9.8 | 7 |
| 17 | Cell cycles and endocycles in the model brown seaweed, Ectocarpus siliculosus. <i>Plant Signaling and Behavior</i> , 2010 , 5, 1473-5 | 2.5 | 5 |
| 16 | Immunostaining of Ectocarpus cells. <i>Cold Spring Harbor Protocols</i> , 2012 , 2012, 369-72 | 1.2 | 5 |
| 15 | Evolution of life cycles and reproductive traits: insights from the brown algae | | 5 |
| 14 | : an evo-devo model for the brown algae. <i>EvoDevo</i> , 2020 , 11, 19 | 3.2 | 5 |
| 13 | Evolution of life cycles and reproductive traits: Insights from the brown algae. <i>Journal of Evolutionary Biology</i> , 2021 , 34, 992-1009 | 2.3 | 5 |
| 12 | Emergence of Ectocarpus as a Model System to Study the Evolution of Complex Multicellularity in the Brown Algae. <i>Advances in Marine Genomics</i> , 2015 , 153-162 | | 4 |
| 11 | Unusual Patterns of Mitochondrial Inheritance in the Brown Alga Ectocarpus. <i>Molecular Biology and Evolution</i> , 2019 , 36, 2778-2789 | 8.3 | 3 |
| 10 | Histone modifications during the life cycle of the brown alga Ectocarpus | | 3 |
| 9 | Chromosome-level quality scaffolding of brown algal genomes using InstaGRAAL, a proximity ligation-based scaffolder | | 2 |

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| 8 | Rapid Turnover of Life-Cycle-Related Genes in the Brown Algae | | 2 |
| 7 | A partially sex-reversed giant kelp sheds light into the mechanisms of sexual differentiation in a UV sexual system. <i>New Phytologist</i> , 2021 , 232, 252-263 | 9.8 | 2 |
| 6 | Chromatin landscape associated with sexual differentiation in a UV sex determination system.. <i>Nucleic Acids Research</i> , 2022 , | 20.1 | 2 |
| 5 | Production and Bioassay of a Diffusible Factor That Induces Gametophyte-to-Sporophyte Developmental Reprogramming in the Brown Alga. <i>Bio-protocol</i> , 2020 , 10, e3753 | 0.9 | 1 |
| 4 | Biochemical characteristics of a diffusible factor that induces gametophyte to sporophyte switching in the brown alga Ectocarpus. <i>Journal of Phycology</i> , 2021 , 57, 742-753 | 3 | 1 |
| 3 | Switching it up: algal insights into sexual transitions. <i>Plant Reproduction</i> , 2021 , 34, 287-296 | 3.9 | 1 |
| 2 | A partially sex-reversed giant kelp sheds light into the mechanisms of sexual differentiation in a UV sexual system | | 1 |
| 1 | An Efficient Chromatin Immunoprecipitation Protocol for the Analysis of Histone Modification Distributions in the Brown Alga Ectocarpus. <i>Methods and Protocols</i> , 2022 , 5, 36 | 2.5 | 0 |