José Manuel Jiménez-GÃ3mez

List of Publications by Year in descending order

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| # | Article | IF | CITATIONS |
|----|--|------|-----------|
| 1 | The VIL gene CRAWLING ELEPHANT controls maturation and differentiation in tomato via polycomb silencing. PLoS Genetics, 2022, 18, e1009633. | 3.5 | 2 |
| 2 | Interaction between photoperiod and variation in circadian rhythms in tomato. BMC Plant Biology, 2022, 22, 187. | 3.6 | 12 |
| 3 | Effective Mapping by Sequencing to Isolate Causal Mutations in the Tomato Genome. Methods in Molecular Biology, 2021, 2264, 89-103. | 0.9 | 7 |
| 4 | Natural variation in HsfA2 preâ€nRNA splicing is associated with changes in thermotolerance during tomato domestication. New Phytologist, 2020, 225, 1297-1310. | 7.3 | 55 |
| 5 | The impact of transposable elements on tomato diversity. Nature Communications, 2020, 11, 4058. | 12.8 | 92 |
| 6 | <i>ENO</i> regulates tomato fruit size through the floral meristem development network. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 8187-8195. | 7.1 | 108 |
| 7 | Mild drought in the vegetative stage induces phenotypic, gene expression, and DNA methylation plasticity in Arabidopsis but no transgenerational effects. Journal of Experimental Botany, 2020, 71, 3588-3602. | 4.8 | 48 |
| 8 | Functional analysis of <i>FRIGIDA</i> using naturally occurring variation in <i>Arabidopsis thaliana</i> . Plant Journal, 2020, 103, 154-165. | 5.7 | 23 |
| 9 | Sl <scp>HAK</scp> 20: a new player in plant salt tolerance. EMBO Journal, 2020, 39, e104997. | 7.8 | 8 |
| 10 | Arabidopsis SME1 Regulates Plant Development and Response to Abiotic Stress by Determining Spliceosome Activity Specificity. Plant Cell, 2019, 31, 537-554. | 6.6 | 42 |
| 11 | Duplication of a domestication locus neutralized a cryptic variant that caused a breeding barrier in tomato. Nature Plants, 2019, 5, 471-479. | 9.3 | 66 |
| 12 | Pivotal Roles of Cryptochromes 1a and 2 in Tomato Development and Physiology. Plant Physiology, 2019, 179, 732-748. | 4.8 | 40 |
| 13 | Prominent alterations of wild barley leaf transcriptome in response to individual and combined drought acclimation and heat shock conditions. Physiologia Plantarum, 2018, 163, 18-29. | 5.2 | 16 |
| 14 | NPR1 mediates a novel regulatory pathway in cold acclimation by interacting with HSFA1 factors. Nature Plants, 2018, 4, 811-823. | 9.3 | 80 |
| 15 | Mutations in <i>EID1</i> and <i>LNK2</i> caused light-conditional clock deceleration during tomato domestication. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 7135-7140. | 7.1 | 48 |
| 16 | Alleleâ€specific expression and genetic determinants of transcriptomic variations in response to mild water deficit in tomato. Plant Journal, 2018, 96, 635-650. | 5.7 | 28 |
| 17 | New Strategies and Tools in Quantitative Genetics: How to Go from the Phenotype to the Genotype. Annual Review of Plant Biology, 2017, 68, 435-455. | 18.7 | 100 |
| 18 | Multi-Omics of Tomato Glandular Trichomes Reveals Distinct Features of Central Carbon Metabolism Supporting High Productivity of Specialized Metabolites. Plant Cell, 2017, 29, 960-983. | 6.6 | 143 |

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|----|--|------|-----------|
| 19 | Variation in the flowering gene SELF PRUNING 5G promotes day-neutrality and early yield in tomato. Nature Genetics, 2017, 49, 162-168. | 21.4 | 344 |
| 20 | A footprint of desiccation tolerance in the genome of Xerophyta viscosa. Nature Plants, 2017, 3, 17038. | 9.3 | 123 |
| 21 | Construction of a High-Density Genetic Map from RNA-Seq Data for an Arabidopsis Bay-0 × Shahdara RIL Population. Frontiers in Genetics, 2017, 8, 201. | 2.3 | 15 |
| 22 | CLAUSA is a MYB Transcription Factor that Promotes Leaf Differentiation by Attenuating Cytokinin Signaling. Plant Cell, 2016, 28, tpc.00211.2016. | 6.6 | 40 |
| 23 | Alternative splicing enhances transcriptome complexity in desiccating seeds. Journal of Integrative Plant Biology, 2016, 58, 947-958. | 8.5 | 26 |
| 24 | The LSM1-7 Complex Differentially Regulates Arabidopsis Tolerance to Abiotic Stress Conditions by Promoting Selective mRNA Decapping. Plant Cell, 2016, 28, 505-520. | 6.6 | 60 |
| 25 | Domestication selected for deceleration of the circadian clock in cultivated tomato. Nature Genetics, 2016, 48, 89-93. | 21.4 | 165 |
| 26 | Analysis of Circadian Leaf Movements. Methods in Molecular Biology, 2016, 1398, 71-79. | 0.9 | 7 |
| 27 | Targeted and Untargeted Approaches Unravel Novel Candidate Genes and Diagnostic SNPs for Quantitative Resistance of the Potato (Solanum tuberosum L.) to Phytophthora infestans Causing the Late Blight Disease. PLoS ONE, 2016, 11, e0156254. | 2.5 | 51 |
| 28 | The relationship between flowering time and growth responses to drought in the Arabidopsis Landsberg erecta x Antwerp-1 population. Frontiers in Plant Science, 2014, 5, 609. | 3.6 | 44 |
| 29 | Network types and their application in natural variation studies in plants. Current Opinion in Plant Biology, 2014, 18, 80-86. | 7.1 | 9 |
| 30 | The genome of the stress-tolerant wild tomato species Solanum pennellii. Nature Genetics, 2014, 46, 1034-1038. | 21.4 | 391 |
| 31 | Nonsense-Mediated mRNA Decay Modulates Immune Receptor Levels to Regulate Plant Antibacterial Defense. Cell Host and Microbe, 2014, 16, 376-390. | 11.0 | 126 |
| 32 | Functional analysis of the Landsberg erecta allele of FRIGIDA. BMC Plant Biology, 2014, 14, 218. | 3.6 | 20 |
| 33 | Comparative transcriptomics reveals patterns of selection in domesticated and wild tomato. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, E2655-62. | 7.1 | 325 |
| 34 | Fine genetic mapping of RXopJ4, a bacterial spot disease resistance locus from Solanum pennellii LA716. Theoretical and Applied Genetics, 2013, 126, 601-609. | 3.6 | 51 |
| 35 | A Quantitative Genetic Basis for Leaf Morphology in a Set of Precisely Defined Tomato Introgression Lines. Plant Cell, 2013, 25, 2465-2481. | 6.6 | 209 |
| 36 | Native Environment Modulates Leaf Size and Response to Simulated Foliar Shade across Wild Tomato Species. PLoS ONE, 2012, 7, e29570. | 2.5 | 54 |

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|----|--|-----|-----------|
| 37 | Next generation quantitative genetics in plants. Frontiers in Plant Science, 2011, 2, 77. | 3.6 | 11 |
| 38 | Network Quantitative Trait Loci Mapping of Circadian Clock Outputs Identifies Metabolic Pathway-to-Clock Linkages in <i>Arabidopsis</i> Â Â. Plant Cell, 2011, 23, 471-485. | 6.6 | 139 |
| 39 | Genomic Analysis of QTLs and Genes Altering Natural Variation in Stochastic Noise. PLoS Genetics, 2011, 7, e1002295. | 3.5 | 107 |
| 40 | Network Analysis Identifies ELF3 as a QTL for the Shade Avoidance Response in Arabidopsis. PLoS Genetics, 2010, 6, e1001100. | 3.5 | 120 |
| 41 | Sequence diversity in three tomato species: SNPs, markers, and molecular evolution. BMC Plant Biology, 2009, 9, 85. | 3.6 | 44 |