## Javier Gallego-Bartolome

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8246683/publications.pdf

Version: 2024-02-01

25 papers

3,166 citations

331670 21 h-index 25 g-index

26 all docs

26 docs citations

26 times ranked

4034 citing authors

#	Article	IF	CITATIONS
1	Molecular mechanism for the interaction between gibberellin and brassinosteroid signaling pathways in <i>Arabidopsis</i> . Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 13446-13451.	7.1	327
2	Hormonal regulation of temperatureâ€induced growth in Arabidopsis. Plant Journal, 2009, 60, 589-601.	5.7	271
3	A One Precursor One siRNA Model for Pol IV-Dependent siRNA Biogenesis. Cell, 2015, 163, 445-455.	28.9	260
4	Molecular Mechanism of Action of Plant DRM De Novo DNA Methyltransferases. Cell, 2014, 157, 1050-1060.	28.9	245
5	Site-specific manipulation of Arabidopsis loci using CRISPR-Cas9 SunTag systems. Nature Communications, 2019, 10, 729.	12.8	215
6	Targeted DNA demethylation of the <i>Arabidopsis</i> genome using the human TET1 catalytic domain. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E2125-E2134.	7.1	190
7	Mechanism of DNA Methylation-Directed Histone Methylation by KRYPTONITE. Molecular Cell, 2014, 55, 495-504.	9.7	186
8	A DNA methylation reader complex that enhances gene transcription. Science, 2018, 362, 1182-1186.	12.6	181
9	Polarization of PIN3â€dependent auxin transport for hypocotyl gravitropic response in <i>Arabidopsis thaliana</i> . Plant Journal, 2011, 67, 817-826.	5.7	171
10	Gibberellins modulate light signaling pathways to prevent Arabidopsis seedling deâ€etiolation in darkness. Plant Journal, 2008, 53, 324-335.	5.7	160
11	Co-targeting RNA Polymerases IV and V Promotes Efficient De Novo DNA Methylation in Arabidopsis. Cell, 2019, 176, 1068-1082.e19.	28.9	124
12	Transcriptional Diversification and Functional Conservation between DELLA Proteins in Arabidopsis. Molecular Biology and Evolution, 2010, 27, 1247-1256.	8.9	123
13	DNA methylation in plants: mechanisms and tools for targeted manipulation. New Phytologist, 2020, 227, 38-44.	7.3	116
14	RNA-directed DNA methylation involves co-transcriptional small-RNA-guided slicing of polymerase V transcripts in Arabidopsis. Nature Plants, 2018, 4, 181-188.	9.3	106
15	Hierarchy of hormone action controlling apical hook development in Arabidopsis. Plant Journal, 2011, 67, 622-634.	5.7	92
16	DNA methylome of the 20-gigabase Norway spruce genome. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E8106-E8113.	7.1	85
17	DNA methylation-linked chromatin accessibility affects genomic architecture in <i>Arabidopsis</i> Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	70
18	DELLA-Induced Early Transcriptional Changes during Etiolated Development in Arabidopsis thaliana. PLoS ONE, 2011, 6, e23918.	2.5	63

#	Article	IF	CITATIONS
19	A Hormonal Regulatory Module That Provides Flexibility to Tropic Responses  Â. Plant Physiology, 2011, 156, 1819-1825.	4.8	33
20	CryoEM structures of Arabidopsis DDR complexes involved in RNA-directed DNA methylation. Nature Communications, 2019, 10, 3916.	12.8	31
21	Identification of Multiple Proteins Coupling Transcriptional Gene Silencing to Genome Stability in Arabidopsis thaliana. PLoS Genetics, 2016, 12, e1006092.	3.5	30
22	Arabidopsis MORC proteins function in the efficient establishment of RNA directed DNA methylation. Nature Communications, 2021, 12, 4292.	12.8	28
23	The characterization of Mediator 12 and 13 as conditional positive gene regulators in Arabidopsis. Nature Communications, 2020, 11, 2798.	12.8	22
24	Ectopic targeting of CG DNA methylation in Arabidopsis with the bacterial SssI methyltransferase. Nature Communications, 2021, 12, 3130.	12.8	20
25	Comprehensive identification of SWI/SNF complex subunits underpins deep eukaryotic ancestry and reveals new plant components. Communications Biology, 2022, 5, .	4.4	17