

Ramin Yadegari

List of Publications by Year in descending order

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35
papers

4,336
citations

236925

25
h-index

377865

34
g-index

38
all docs

38
docs citations

38
times ranked

3912
citing authors

#	ARTICLE	IF	CITATIONS
1	Broadening the impact of plant science through innovative, integrative, and inclusive outreach. <i>Plant Direct</i> , 2021, 5, e00316.	1.9	14
2	FERTILIZATION-INDEPENDENT SEED-Polycomb Repressive Complex 2 Plays a Dual Role in Regulating Type I MADS-Box Genes in Early Endosperm Development. <i>Plant Physiology</i> , 2018, 177, 285-299.	4.8	60
3	Laser-Capture Microdissection of Maize Kernel Compartments for RNA-Seq-Based Expression Analysis. <i>Methods in Molecular Biology</i> , 2018, 1676, 153-163.	0.9	9
4	Opaque-2 Regulates a Complex Gene Network Associated with Cell Differentiation and Storage Functions of Maize Endosperm. <i>Plant Cell</i> , 2018, 30, 2425-2446.	6.6	83
5	Maize opaque mutants are no longer so opaque. <i>Plant Reproduction</i> , 2018, 31, 319-326.	2.2	12
6	Arabidopsis CALCINEURIN B-LIKE10 Functions Independently of the SOS Pathway during Reproductive Development in Saline Conditions. <i>Plant Physiology</i> , 2016, 171, 369-379.	4.8	31
7	ACTIN-RELATED PROTEIN 6 regulates DISRUPTED MEIOTIC cDNA 1 gene expression in <i>Arabidopsis thaliana</i> ovules. <i>Molecular Reproduction and Development</i> , 2015, 82, 499-499.	2.0	0
8	RNA Sequencing of Laser-Capture Microdissected Compartments of the Maize Kernel Identifies Regulatory Modules Associated with Endosperm Cell Differentiation. <i>Plant Cell</i> , 2015, 27, 513-531.	6.6	206
9	Temporal patterns of gene expression in developing maize endosperm identified through transcriptome sequencing. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 7582-7587.	7.1	146
10	RNA-Seq analysis of laser-capture microdissected cells of the developing central starchy endosperm of maize. <i>Genomics Data</i> , 2014, 2, 242-245.	1.3	13
11	Maize early endosperm growth and development: From fertilization through cell type differentiation. <i>American Journal of Botany</i> , 2014, 101, 1259-1274.	1.7	80
12	ACTIN-RELATED PROTEIN6 Regulates Female Meiosis by Modulating Meiotic Gene Expression in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2014, 26, 1612-1628.	6.6	68
13	Dynamic Expression of Imprinted Genes Associates with Maternally Controlled Nutrient Allocation during Maize Endosperm Development. <i>Plant Cell</i> , 2013, 25, 3212-3227.	6.6	97
14	Identification of genes expressed in the angiosperm female gametophyte. <i>Journal of Experimental Botany</i> , 2011, 62, 1593-1599.	4.8	17
15	Identification of transcription-factor genes expressed in the Arabidopsis female gametophyte. <i>BMC Plant Biology</i> , 2010, 10, 110.	3.6	60
16	Plant SMU-1 and SMU-2 Homologues Regulate Pre-mRNA Splicing and Multiple Aspects of Development. <i>Plant Physiology</i> , 2009, 151, 1498-1512.	4.8	37
17	RPK1 and TOAD2 Are Two Receptor-like Kinases Redundantly Required for Arabidopsis Embryonic Pattern Formation. <i>Developmental Cell</i> , 2007, 12, 943-956.	7.0	137
18	Segregation distortion in Arabidopsis gametophytic factor 1 (<i>gfa1</i>) mutants is caused by a deficiency of an essential RNA splicing factor. <i>Sexual Plant Reproduction</i> , 2007, 20, 87-97.	2.2	23

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19	Partially redundant functions of two SET-domain polycomb-group proteins in controlling initiation of seed development in Arabidopsis. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 13244-13249.	7.1	123
20	The role of JAGGED in shaping lateral organs. Development (Cambridge), 2004, 131, 1101-1110.	2.5	277
21	Female Gametophyte Development. Plant Cell, 2004, 16, S133-S141.	6.6	370
22	Development and Function of the Angiosperm Female Gametophyte. Annual Review of Genetics, 2002, 36, 99-124.	7.6	197
23	RASPBERRY3 Gene Encodes a Novel Protein Important for Embryo Development. Plant Physiology, 2002, 129, 691-705.	4.8	35
24	The Arabidopsis Embryo Mutant schlepperless Has a Defect in the Chaperonin-60± Gene. Plant Physiology, 2001, 126, 717-730.	4.8	124
25	Mutations in the FIE and MEA Genes That Encode Interacting Polycomb Proteins Cause Parent-of-Origin Effects on Seed Development by Distinct Mechanisms. Plant Cell, 2000, 12, 2367.	6.6	2
26	Mutations in the <i>FIE</i> and <i>MEA</i> Genes That Encode Interacting Polycomb Proteins Cause Parent-of-Origin Effects on Seed Development by Distinct Mechanisms. Plant Cell, 2000, 12, 2367-2381.	6.6	231
27	Imprinting of the <i>MEDEA</i> Polycomb Gene in the Arabidopsis Endosperm. Plant Cell, 1999, 11, 1945-1952.	6.6	313
28	Mutations in <i>FIE</i>, a WD Polycomb Group Gene, Allow Endosperm Development without Fertilization. Plant Cell, 1999, 11, 407-415.	6.6	407
29	Control of fertilization-independent endosperm development by the <i>MEDEA</i> polycomb gene in <i>Arabidopsis</i>. Proceedings of the National Academy of Sciences of the United States of America, 1999, 96, 4186-4191.	7.1	331
30	Imprinting of the MEDEA Polycomb Gene in the Arabidopsis Endosperm. Plant Cell, 1999, 11, 1945.	6.6	31
31	Embryogenesis in Dicotyledonous Plants. Advances in Cellular and Molecular Biology of Plants, 1997, , 3-52.	0.2	19
32	Plant Embryogenesis: Zygote to Seed. Science, 1994, 266, 605-614.	12.6	534
33	Cell Differentiation and Morphogenesis Are Uncoupled in Arabidopsis raspberry Embryos. Plant Cell, 1994, 6, 1713.	6.6	64
34	Structure and nucleotide sequence of a Drosophila melanogaster protein kinase C gene.. EMBO Journal, 1987, 6, 433-441.	7.8	128
35	The isolation, characterization and sequence of two divergent ?-tubulin genes from soybean (Glycine) Tj ETQq1 1 0,784314 rsgBT /Ov	3.9	57