

Uell Grossniklaus

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279
papers

22,441
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83
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145
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309
ext. papers

25,993
ext. citations

10.2
avg, IF

6.91
L-index

#	Paper	IF	Citations
279	A gateway cloning vector set for high-throughput functional analysis of genes in planta. <i>Plant Physiology</i> , 2003 , 133, 462-9	6.6	1966
278	Maternal control of embryogenesis by MEDEA, a polycomb group gene in Arabidopsis. <i>Science</i> , 1998 , 280, 446-50	33.3	720
277	P-element-mediated enhancer detection: a versatile method to study development in Drosophila. <i>Genes and Development</i> , 1989 , 3, 1288-300	12.6	567
276	Insights into Land Plant Evolution Garnered from the Marchantia polymorpha Genome. <i>Cell</i> , 2017 , 171, 287-304.e15	56.2	538
275	The Arabidopsis Somatic Embryogenesis Receptor Kinase 1 Gene Is Expressed in Developing Ovules and Embryos and Enhances Embryogenic Competence in Culture. <i>Plant Physiology</i> , 2001 , 127, 803-816	6.6	502
274	The FERONIA receptor-like kinase mediates male-female interactions during pollen tube reception. <i>Science</i> , 2007 , 317, 656-60	33.3	464
273	Cellular efflux of auxin catalyzed by the Arabidopsis MDR/PGP transporter AtPGP1. <i>Plant Journal</i> , 2005 , 44, 179-94	6.9	429
272	Genome-scale proteomics reveals Arabidopsis thaliana gene models and proteome dynamics. <i>Science</i> , 2008 , 320, 938-41	33.3	419
271	P-element-mediated enhancer detection: an efficient method for isolating and characterizing developmentally regulated genes in Drosophila. <i>Genes and Development</i> , 1989 , 3, 1301-13	12.6	345
270	Arabidopsis MSI1 is a component of the MEA/FIE Polycomb group complex and required for seed development. <i>EMBO Journal</i> , 2003 , 22, 4804-14	13	322
269	The Polycomb-group protein MEDEA regulates seed development by controlling expression of the MADS-box gene PHERES1. <i>Genes and Development</i> , 2003 , 17, 1540-53	12.6	316
268	Regulation of Arabidopsis tapetum development and function by DYSFUNCTIONAL TAPETUM1 (DYT1) encoding a putative bHLH transcription factor. <i>Development (Cambridge)</i> , 2006 , 133, 3085-95	6.6	314
267	Apomixis: a developmental perspective. <i>Annual Review of Plant Biology</i> , 2003 , 54, 547-74	30.7	314
266	The Arabidopsis mutant feronia disrupts the female gametophytic control of pollen tube reception. <i>Development (Cambridge)</i> , 2003 , 130, 2149-59	6.6	302
265	Conserved molecular components for pollen tube reception and fungal invasion. <i>Science</i> , 2010 , 330, 968-973	33.3	290
264	Maintenance of genomic imprinting at the Arabidopsis medea locus requires zygotic DDM1 activity. <i>Genes and Development</i> , 1999 , 13, 2971-82	12.6	282
263	Delayed activation of the paternal genome during seed development. <i>Nature</i> , 2000 , 404, 91-4	50.4	265

262	Genome-wide high-resolution mapping of exosome substrates reveals hidden features in the Arabidopsis transcriptome. <i>Cell</i> , 2007 , 131, 1340-53	56.2	258
261	Arabidopsis female gametophyte gene expression map reveals similarities between plant and animal gametes. <i>Current Biology</i> , 2010 , 20, 506-12	6.3	255
260	FIDDLEHEAD, a gene required to suppress epidermal cell interactions in Arabidopsis, encodes a putative lipid biosynthetic enzyme. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000 , 97, 1311-6	11.5	233
259	The Arabidopsis thaliana MEDEA Polycomb group protein controls expression of PHERES1 by parental imprinting. <i>Nature Genetics</i> , 2005 , 37, 28-30	36.3	226
258	PAMP (pathogen-associated molecular pattern)-induced changes in plasma membrane compartmentalization reveal novel components of plant immunity. <i>Journal of Biological Chemistry</i> , 2010 , 285, 39140-9	5.4	220
257	Disruption of the pollen-expressed FERONIA homologs ANXUR1 and ANXUR2 triggers pollen tube discharge. <i>Development (Cambridge)</i> , 2009 , 136, 3279-88	6.6	220
256	ATX-1, an Arabidopsis homolog of trithorax, activates flower homeotic genes. <i>Current Biology</i> , 2003 , 13, 627-37	6.3	220
255	Different Polycomb group complexes regulate common target genes in Arabidopsis. <i>EMBO Reports</i> , 2006 , 7, 947-52	6.5	218
254	Transgenerational epigenetic inheritance: how important is it?. <i>Nature Reviews Genetics</i> , 2013 , 14, 228-35	30.1	216
253	Egg cell-secreted EC1 triggers sperm cell activation during double fertilization. <i>Science</i> , 2012 , 338, 1093-3	33.3	216
252	Natural enemies drive geographic variation in plant defenses. <i>Science</i> , 2012 , 338, 116-9	33.3	207
251	The Drosophila sloppy paired locus encodes two proteins involved in segmentation that show homology to mammalian transcription factors. <i>Genes and Development</i> , 1992 , 6, 1030-51	12.6	202
250	ARABIDOPSIS TRITHORAX1 dynamically regulates FLOWERING LOCUS C activation via histone 3 lysine 4 trimethylation. <i>Plant Cell</i> , 2008 , 20, 580-8	11.6	199
249	Hi-C analysis in Arabidopsis identifies the KNOT, a structure with similarities to the flamenco locus of Drosophila. <i>Molecular Cell</i> , 2014 , 55, 678-93	17.6	190
248	ANXUR receptor-like kinases coordinate cell wall integrity with growth at the pollen tube tip via NADPH oxidases. <i>PLoS Biology</i> , 2013 , 11, e1001719	9.7	181
247	Selection of T-DNA-tagged male and female gametophytic mutants by segregation distortion in Arabidopsis. <i>Genetics</i> , 1998 , 149, 621-31	4	172
246	Identification of new members of Fertilisation Independent Seed Polycomb Group pathway involved in the control of seed development in Arabidopsis thaliana. <i>Development (Cambridge)</i> , 2004 , 131, 2971-81	6.6	169
245	The molecular and genetic basis of ovule and megagametophyte development. <i>Seminars in Cell and Developmental Biology</i> , 1998 , 9, 227-38	7.5	166

244	Transcriptional silencing by polycomb-group proteins. <i>Cold Spring Harbor Perspectives in Biology</i> , 2014 , 6, a019331	10.2	163
243	Maternal epigenetic pathways control parental contributions to Arabidopsis early embryogenesis. <i>Cell</i> , 2011 , 145, 707-19	56.2	161
242	The art and design of genetic screens: Arabidopsis thaliana. <i>Nature Reviews Genetics</i> , 2002 , 3, 124-36	30.1	159
241	SHORT INTEGUMENTS1/SUSPENSOR1/CARPEL FACTORY, a Dicer homolog, is a maternal effect gene required for embryo development in Arabidopsis. <i>Plant Physiology</i> , 2002 , 130, 808-22	6.6	155
240	Activation of the U2 snRNA promoter by the octamer motif defines a new class of RNA polymerase II enhancer elements. <i>Genes and Development</i> , 1988 , 2, 1764-78	12.6	155
239	SETH1 and SETH2, two components of the glycosylphosphatidylinositol anchor biosynthetic pathway, are required for pollen germination and tube growth in Arabidopsis. <i>Plant Cell</i> , 2004 , 16, 229-40	11.6	150
238	Selected aspects of transgenerational epigenetic inheritance and resetting in plants. <i>Current Opinion in Plant Biology</i> , 2011 , 14, 195-203	9.9	148
237	Polycomb group and trithorax group proteins in Arabidopsis. <i>Biochimica Et Biophysica Acta Gene Regulatory Mechanisms</i> , 2007 , 1769, 375-82		146
236	CrRLK1L receptor-like kinases: not just another brick in the wall. <i>Current Opinion in Plant Biology</i> , 2012 , 15, 659-69	9.9	145
235	LACHESIS restricts gametic cell fate in the female gametophyte of Arabidopsis. <i>PLoS Biology</i> , 2007 , 5, e47	9.7	142
234	RALF4/19 peptides interact with LRX proteins to control pollen tube growth in. <i>Science</i> , 2017 , 358, 1600-1603	33.9	138
233	Evolutionary ecology of the prezygotic stage. <i>Science</i> , 2004 , 303, 971-5	33.3	138
232	The central cell plays a critical role in pollen tube guidance in Arabidopsis. <i>Plant Cell</i> , 2007 , 19, 3563-77	11.6	136
231	Deterministic protein inference for shotgun proteomics data provides new insights into Arabidopsis pollen development and function. <i>Genome Research</i> , 2009 , 19, 1786-800	9.7	135
230	Developmental genetics of gametophytic apomixis. <i>Trends in Genetics</i> , 2001 , 17, 597-604	8.5	134
229	A calcium dialog mediated by the FERONIA signal transduction pathway controls plant sperm delivery. <i>Developmental Cell</i> , 2014 , 29, 491-500	10.2	133
228	Embryo and endosperm inherit distinct chromatin and transcriptional states from the female gametes in Arabidopsis. <i>Plant Cell</i> , 2010 , 22, 307-20	11.6	133
227	LAF1, a MYB transcription activator for phytochrome A signaling. <i>Genes and Development</i> , 2001 , 15, 2613-25	12.5	128

226	Improved reference genome by single-molecule sequencing and chromosome conformation capture technologies. <i>Horticulture Research</i> , 2018 , 5, 50	7.7	125
225	Interaction of the Arabidopsis polycomb group proteins FIE and MEA mediates their common phenotypes. <i>Current Biology</i> , 2000 , 10, 1535-8	6.3	125
224	Positive darwinian selection at the imprinted MEDEA locus in plants. <i>Nature</i> , 2007 , 448, 349-52	50.4	124
223	Apomixis technology development-virgin births in farmers' fields?. <i>Nature Biotechnology</i> , 2004 , 22, 687-91	14.5	123
222	Tackling drought stress: receptor-like kinases present new approaches. <i>Plant Cell</i> , 2012 , 24, 2262-78	11.6	118
221	Be more specific! Laser-assisted microdissection of plant cells. <i>Trends in Plant Science</i> , 2005 , 10, 397-406	13.1	117
220	Dynamic regulatory interactions of Polycomb group genes: MEDEA autoregulation is required for imprinted gene expression in Arabidopsis. <i>Genes and Development</i> , 2006 , 20, 1081-6	12.6	117
219	The protein expression landscape of the Arabidopsis root. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, 6811-8	11.5	115
218	Regulation and flexibility of genomic imprinting during seed development. <i>Plant Cell</i> , 2011 , 23, 16-26	11.6	114
217	The walls have ears: the role of plant CrRLK1Ls in sensing and transducing extracellular signals. <i>Journal of Experimental Botany</i> , 2011 , 62, 1581-91	7	114
216	Developmentally regulated Drosophila gene family encoding the fork head domain. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1992 , 89, 8754-8	11.5	114
215	Genetic subtraction profiling identifies genes essential for Arabidopsis reproduction and reveals interaction between the female gametophyte and the maternal sporophyte. <i>Genome Biology</i> , 2007 , 8, R204	18.3	113
214	Localized expression of sloppy paired protein maintains the polarity of Drosophila parasegments. <i>Genes and Development</i> , 1994 , 8, 899-913	12.6	111
213	Transcriptional programs of early reproductive stages in Arabidopsis. <i>Plant Physiology</i> , 2004 , 135, 1765-75	15.6	110
212	Genomic imprinting and seed development: endosperm formation with and without sex. <i>Current Opinion in Plant Biology</i> , 2001 , 4, 21-7	9.9	110
211	Arabidopsis genes essential for seedling viability: isolation of insertional mutants and molecular cloning. <i>Genetics</i> , 2001 , 159, 1765-78	4	110
210	Pattern formation during early ovule development in Arabidopsis thaliana. <i>Developmental Biology</i> , 2004 , 273, 321-34	3.1	109
209	Plant germline formation: common concepts and developmental flexibility in sexual and asexual reproduction. <i>Development (Cambridge)</i> , 2015 , 142, 229-41	6.6	108

208	CLO/GFA1 and ATO are novel regulators of gametic cell fate in plants. <i>Plant Journal</i> , 2008 , 56, 913-21	6.9	100
207	The MADS domain protein DIANA acts together with AGAMOUS-LIKE80 to specify the central cell in <i>Arabidopsis</i> ovules. <i>Plant Cell</i> , 2008 , 20, 2088-101	11.6	97
206	The pollen tube: a soft shell with a hard core. <i>Plant Journal</i> , 2013 , 73, 617-27	6.9	93
205	Transcriptome analysis of the <i>Arabidopsis</i> megaspore mother cell uncovers the importance of RNA helicases for plant germline development. <i>PLoS Biology</i> , 2011 , 9, e1001155	9.7	93
204	A versatile and reliable two-component system for tissue-specific gene induction in <i>Arabidopsis</i> . <i>Plant Physiology</i> , 2006 , 141, 1194-204	6.6	92
203	A Bsister MADS-box gene involved in ovule and seed development in petunia and <i>Arabidopsis</i> . <i>Plant Journal</i> , 2006 , 47, 934-46	6.9	91
202	A powerful method for transcriptional profiling of specific cell types in eukaryotes: laser-assisted microdissection and RNA sequencing. <i>PLoS ONE</i> , 2012 , 7, e29685	3.7	88
201	A dynamic reciprocal RBR-PRC2 regulatory circuit controls <i>Arabidopsis</i> gametophyte development. <i>Current Biology</i> , 2008 , 18, 1680-6	6.3	87
200	How to avoid sex: the genetic control of gametophytic apomixis. <i>Plant Cell</i> , 2001 , 13, 1491-8	11.6	87
199	Genomic imprinting, methylation and molecular evolution of maize Enhancer of zeste (Mez) homologs. <i>Plant Journal</i> , 2007 , 49, 325-37	6.9	86
198	The maternal to zygotic transition in animals and plants. <i>Cold Spring Harbor Symposia on Quantitative Biology</i> , 2008 , 73, 89-100	3.9	85
197	Evolutionary origins of the endosperm in flowering plants. <i>Genome Biology</i> , 2002 , 3, reviews1026	18.3	85
196	Members of the RKD transcription factor family induce an egg cell-like gene expression program. <i>Plant Journal</i> , 2011 , 67, 280-91	6.9	79
195	Analysis of transposon insertion mutants highlights the diversity of mechanisms underlying male progamic development in <i>Arabidopsis</i> . <i>Genetics</i> , 2004 , 167, 1975-86	4	78
194	Apomixis in agriculture: the quest for clonal seeds. <i>Sexual Plant Reproduction</i> , 2001 , 14, 179-87		77
193	She's the boss: signaling in pollen tube reception. <i>Current Opinion in Plant Biology</i> , 2011 , 14, 622-7	9.9	76
192	CHR11, a chromatin-remodeling factor essential for nuclear proliferation during female gametogenesis in <i>Arabidopsis thaliana</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005 , 102, 17231-6	11.5	76
191	VERDANDI is a direct target of the MADS domain ovule identity complex and affects embryo sac differentiation in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2010 , 22, 1702-15	11.6	75

190	Functional redundancy: the respective roles of the two sloppy paired genes in <i>Drosophila</i> segmentation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1994 , 91, 6324-8	11.5	73
189	Stearoyl-acyl carrier protein desaturases are associated with floral isolation in sexually deceptive orchids. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011 , 108, 5696-701	11.5	72
188	Dynamic regulation of Polycomb group activity during plant development. <i>Current Opinion in Plant Biology</i> , 2012 , 15, 523-9	9.9	70
187	Cytoplasmic Ca ²⁺ changes dynamically during the interaction of the pollen tube with synergid cells. <i>Development (Cambridge)</i> , 2012 , 139, 4202-9	6.6	68
186	The triploid endosperm genome of <i>Arabidopsis</i> adopts a peculiar, parental-dosage-dependent chromatin organization. <i>Plant Cell</i> , 2007 , 19, 1782-94	11.6	68
185	Receptor-like cytoplasmic kinase MARIS functions downstream of CrRLK1L-dependent signaling during tip growth. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015 , 112, 12211-6	11.5	67
184	Contribution of epigenetic variation to adaptation in <i>Arabidopsis</i> . <i>Nature Communications</i> , 2018 , 9, 4446	17.4	67
183	Characterization of the phosphoproteome of mature <i>Arabidopsis</i> pollen. <i>Plant Journal</i> , 2012 , 72, 89-101	6.9	65
182	Characterization of chromosomal architecture in <i>Arabidopsis</i> by chromosome conformation capture. <i>Genome Biology</i> , 2013 , 14, R129	18.3	64
181	Epigenetic variation, inheritance, and selection in plant populations. <i>Cold Spring Harbor Symposia on Quantitative Biology</i> , 2012 , 77, 97-104	3.9	63
180	<i>Marchantia</i> MpRKD Regulates the Gametophyte-Sporophyte Transition by Keeping Egg Cells Quiescent in the Absence of Fertilization. <i>Current Biology</i> , 2016 , 26, 1782-1789	6.3	62
179	The RPN1 subunit of the 26S proteasome in <i>Arabidopsis</i> is essential for embryogenesis. <i>Plant Cell</i> , 2005 , 17, 2723-37	11.6	62
178	Intronic regulatory elements determine the divergent expression patterns of AGAMOUS-LIKE6 subfamily members in <i>Arabidopsis</i> . <i>Plant Journal</i> , 2009 , 59, 987-1000	6.9	61
177	Functional analysis of related CrRLK1L receptor-like kinases in pollen tube reception. <i>EMBO Reports</i> , 2015 , 16, 107-15	6.5	60
176	Genomic imprinting during seed development. <i>Advances in Genetics</i> , 2002 , 46, 165-214	3.3	60
175	The <i>Arabidopsis</i> CUL4-DDB1 complex interacts with MSI1 and is required to maintain MEDEA parental imprinting. <i>EMBO Journal</i> , 2011 , 30, 731-43	13	59
174	RETINOBLASTOMA RELATED1 mediates germline entry in. <i>Science</i> , 2017 , 356,	33.3	55
173	Epigenetic inheritance of expression states in plant development: the role of Polycomb group proteins. <i>Current Opinion in Cell Biology</i> , 2002 , 14, 773-9	9	55

172	Genomic imprinting in the Arabidopsis embryo is partly regulated by PRC2. <i>PLoS Genetics</i> , 2013 , 9, e1003862	3.8	54
171	Confocal microscopy of whole ovules for analysis of reproductive development: the elongate1 mutant affects meiosis II. <i>Plant Journal</i> , 2005 , 43, 309-20	6.9	54
170	Model organisms--A historical perspective. <i>Journal of Proteomics</i> , 2010 , 73, 2054-63	3.9	53
169	Molecular control of autonomous embryo and endosperm development. <i>Sexual Plant Reproduction</i> , 2008 , 21, 79-88		53
168	Theoretical and experimental evidence indicates that there is no detectable auxin gradient in the angiosperm female gametophyte. <i>Development (Cambridge)</i> , 2013 , 140, 4544-53	6.6	52
167	Apomictic and sexual germline development differ with respect to cell cycle, transcriptional, hormonal and epigenetic regulation. <i>PLoS Genetics</i> , 2014 , 10, e1004476	6	49
166	Epigenetic control of plant development: new layers of complexity. <i>Current Opinion in Plant Biology</i> , 2004 , 7, 11-9	9.9	49
165	Arabidopsis CUL3A and CUL3B genes are essential for normal embryogenesis. <i>Plant Journal</i> , 2005 , 43, 437-48	6.9	49
164	Epigenetic regulation and reprogramming during gamete formation in plants. <i>Current Opinion in Genetics and Development</i> , 2011 , 21, 124-33	4.9	48
163	The Arabidopsis Somatic Embryogenesis Receptor Kinase 1 Gene Is Expressed in Developing Ovules and Embryos and Enhances Embryogenic Competence in Culture. <i>Plant Physiology</i> , 2001 , 127, 803-816	6.6	48
162	Genomic Imprinting in the Endosperm Is Systematically Perturbed in Abortive Hybrid Tomato Seeds. <i>Molecular Biology and Evolution</i> , 2016 , 33, 2935-2946	8.3	47
161	A bright future for apomixis. <i>Trends in Plant Science</i> , 1998 , 3, 415-416	13.1	46
160	Chromatin modification and remodeling during early seed development. <i>Current Opinion in Genetics and Development</i> , 2007 , 17, 473-9	4.9	46
159	Genetic interaction of an origin recognition complex subunit and the Polycomb group gene MEDEA during seed development. <i>Plant Cell</i> , 2004 , 16, 1035-46	11.6	46
158	Developmental regulation of expression and activity of multiple forms of the Drosophila RAC protein kinase. <i>Journal of Biological Chemistry</i> , 1995 , 270, 4066-75	5.4	45
157	TURAN and EVAN mediate pollen tube reception in Arabidopsis Synergids through protein glycosylation. <i>PLoS Biology</i> , 2015 , 13, e1002139	9.7	43
156	LRX Proteins Play a Crucial Role in Pollen Grain and Pollen Tube Cell Wall Development. <i>Plant Physiology</i> , 2018 , 176, 1981-1992	6.6	43
155	Identification of imprinted genes subject to parent-of-origin specific expression in Arabidopsis thaliana seeds. <i>BMC Plant Biology</i> , 2011 , 11, 113	5.3	42

154	Adaptation and extinction in experimentally fragmented landscapes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010 , 107, 19120-5	11.5	41
153	Dosage-sensitive function of retinoblastoma related and convergent epigenetic control are required during the Arabidopsis life cycle. <i>PLoS Genetics</i> , 2010 , 6, e1000988	6	41
152	Selection-driven evolution of sex-biased genes is consistent with sexual selection in Arabidopsis thaliana. <i>Molecular Biology and Evolution</i> , 2014 , 31, 574-83	8.3	40
151	Measuring the Mechanical Properties of Plant Cell Walls. <i>Plants</i> , 2015 , 4, 167-82	4.5	40
150	SNP-Ratio Mapping (SRM): identifying lethal alleles and mutations in complex genetic backgrounds by next-generation sequencing. <i>Genetics</i> , 2012 , 191, 1381-6	4	39
149	The Polycomb group protein MEDEA and the DNA methyltransferase MET1 interact to repress autonomous endosperm development in Arabidopsis. <i>Plant Journal</i> , 2013 , 73, 776-87	6.9	38
148	Extensive epigenetic reprogramming during the life cycle of Marchantia polymorpha. <i>Genome Biology</i> , 2018 , 19, 9	18.3	37
147	Transcriptome and proteome data reveal candidate genes for pollinator attraction in sexually deceptive orchids. <i>PLoS ONE</i> , 2013 , 8, e64621	3.7	37
146	Diverse functions of Polycomb group proteins during plant development. <i>Seminars in Cell and Developmental Biology</i> , 2003 , 14, 77-84	7.5	37
145	Analysis of plant germline development by high-throughput RNA profiling: technical advances and new insights. <i>Plant Journal</i> , 2012 , 70, 18-29	6.9	36
144	HiCdat: a fast and easy-to-use Hi-C data analysis tool. <i>BMC Bioinformatics</i> , 2015 , 16, 277	3.6	36
143	Identification of a DNA methylation-independent imprinting control region at the Arabidopsis MEDEA locus. <i>Genes and Development</i> , 2012 , 26, 1837-50	12.6	36
142	The genetic basis of pollinator adaptation in a sexually deceptive orchid. <i>PLoS Genetics</i> , 2012 , 8, e1002889		36
141	Arabidopsis GLAUCE promotes fertilization-independent endosperm development and expression of paternally inherited alleles. <i>Development (Cambridge)</i> , 2007 , 134, 4107-17	6.6	35
140	Structural basis for recognition of RALF peptides by LRX proteins during pollen tube growth. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 7494-7503	11.5	34
139	Characterization of the three Arabidopsis thaliana RAD21 cohesins reveals differential responses to ionizing radiation. <i>Journal of Experimental Botany</i> , 2006 , 57, 971-83	7	34
138	Intrachromosomal excision of a hybrid Ds element induces large genomic deletions in Arabidopsis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004 , 101, 2969-74	11.5	34
137	Different yet similar: evolution of imprinting in flowering plants and mammals. <i>F1000prime Reports</i> , 2014 , 6, 63		33

136	Starch Turnover and Metabolism during Flower and Early Embryo Development. <i>Plant Physiology</i> , 2016 , 172, 2388-2402	6.6	31
135	Apomixis Allows the Transgenerational Fixation of Phenotypes in Hybrid Plants. <i>Current Biology</i> , 2016 , 26, 331-7	6.3	31
134	Plant germline development: a tale of cross-talk, signaling, and cellular interactions. <i>Sexual Plant Reproduction</i> , 2011 , 24, 91-5		31
133	Plant genetics: a decade of integration. <i>Nature Genetics</i> , 2003 , 33 Suppl, 294-304	36.3	31
132	An egg apparatus-specific enhancer of Arabidopsis, identified by enhancer detection. <i>Plant Physiology</i> , 2005 , 139, 1421-32	6.6	31
131	The female gametophyte: an emerging model for cell type-specific systems biology in plant development. <i>Frontiers in Plant Science</i> , 2015 , 6, 907	6.2	29
130	Unveiling the gene-expression profile of pollen. <i>Genome Biology</i> , 2003 , 5, 205	18.3	29
129	Genomic imprinting in plants. <i>Results and Problems in Cell Differentiation</i> , 1999 , 25, 23-40	1.4	29
128	Quantifying growth mechanics of living, growing plant cells in situ using microrobotics. <i>Micro and Nano Letters</i> , 2011 , 6, 311	0.9	28
127	Transposons and tandem repeats are not involved in the control of genomic imprinting at the MEDEA locus in Arabidopsis. <i>Cold Spring Harbor Symposia on Quantitative Biology</i> , 2004 , 69, 465-75	3.9	28
126	Massively Parallelized Pollen Tube Guidance and Mechanical Measurements on a Lab-on-a-Chip Platform. <i>PLoS ONE</i> , 2016 , 11, e0168138	3.7	28
125	The first high-resolution DNA "methylome". <i>Cell</i> , 2006 , 126, 1025-8	56.2	27
124	Female gametophytic cell specification and seed development require the function of the putative Arabidopsis INCENP ortholog WYRD. <i>Development (Cambridge)</i> , 2011 , 138, 3409-20	6.6	26
123	Haplotype-resolved genomes of geminivirus-resistant and geminivirus-susceptible African cassava cultivars. <i>BMC Biology</i> , 2019 , 17, 75	7.3	25
122	Characterization of size-dependent mechanical properties of tip-growing cells using a lab-on-chip device. <i>Lab on A Chip</i> , 2016 , 17, 82-90	7.2	25
121	Genomic origin and organization of the allopolyploid <i>Primula egaliksensis</i> investigated by in situ hybridization. <i>Annals of Botany</i> , 2008 , 101, 919-27	4.1	25
120	Chromosome conformation capture-based studies reveal novel features of plant nuclear architecture. <i>Current Opinion in Plant Biology</i> , 2017 , 36, 149-157	9.9	24
119	Real-time automated characterization of 3D morphology and mechanics of developing plant cells. <i>International Journal of Robotics Research</i> , 2015 , 34, 1136-1146	5.7	24

118	Rcount: simple and flexible RNA-Seq read counting. <i>Bioinformatics</i> , 2015 , 31, 436-7	7.2	24
117	Seed Production Affects Maternal Growth and Senescence in Arabidopsis. <i>Plant Physiology</i> , 2016 , 171, 392-404	6.6	24
116	3D Manipulation and Imaging of Plant Cells using Acoustically Activated Microbubbles. <i>Small Methods</i> , 2019 , 3, 1800527	12.8	23
115	Computational analysis and characterization of UCE-like elements (ULEs) in plant genomes. <i>Genome Research</i> , 2012 , 22, 2455-66	9.7	23
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