Maarten Koornneef

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178 148 22,091 77 h-index g-index citations papers 181 8.2 6.54 24,647 L-index avg, IF ext. citations ext. papers

#	Paper	IF	Citations
178	A genetic and physiological analysis of late flowering mutants in Arabidopsis thaliana. <i>Molecular Genetics and Genomics</i> , 1991 , 229, 57-66		79 ²
177	The isolation and characterization of abscisic acid-insensitive mutants of Arabidopsis thaliana. <i>Physiologia Plantarum</i> , 1984 , 61, 377-383	4.6	789
176	Seed dormancy and germination. Current Opinion in Plant Biology, 2002, 5, 33-6	9.9	639
175	Influence of the testa on seed dormancy, germination, and longevity in Arabidopsis. <i>Plant Physiology</i> , 2000 , 122, 403-14	6.6	603
174	Induction and analysis of gibberellin sensitive mutants in Arabidopsis thaliana (L.) heynh. <i>Theoretical and Applied Genetics</i> , 1980 , 58, 257-63	6	541
173	Naturally occurring genetic variation in Arabidopsis thaliana. <i>Annual Review of Plant Biology</i> , 2004 , 55, 141-72	30.7	529
172	Induction of dormancy during seed development by endogenous abscisic acid: studies on abscisic acid deficient genotypes of Arabidopsis thaliana (L.) Heynh. <i>Planta</i> , 1983 , 157, 158-65	4.7	468
171	Sucrose-specific induction of anthocyanin biosynthesis in Arabidopsis requires the MYB75/PAP1 gene. <i>Plant Physiology</i> , 2005 , 139, 1840-52	6.6	464
170	The late flowering phenotype of fwa mutants is caused by gain-of-function epigenetic alleles of a homeodomain gene. <i>Molecular Cell</i> , 2000 , 6, 791-802	17.6	460
169	Analysis of Arabidopsis mutants deficient in flavonoid biosynthesis. <i>Plant Journal</i> , 1995 , 8, 659-71	6.9	457
168	The isolation of abscisic acid (ABA) deficient mutants by selection of induced revertants in non-germinating gibberellin sensitive lines of Arabidopsis thaliana (L.) heynh. <i>Theoretical and Applied Genetics</i> , 1982 , 61, 385-93	6	457
167	The TRANSPARENT TESTA12 gene of Arabidopsis encodes a multidrug secondary transporter-like protein required for flavonoid sequestration in vacuoles of the seed coat endothelium. <i>Plant Cell</i> , 2001 , 13, 853-71	11.6	424
166	Cloning of DOG1, a quantitative trait locus controlling seed dormancy in Arabidopsis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006 , 103, 17042-7	11.5	413
165	Large expression differences in genes for iron and zinc homeostasis, stress response, and lignin biosynthesis distinguish roots of Arabidopsis thaliana and the related metal hyperaccumulator Thlaspi caerulescens. <i>Plant Physiology</i> , 2006 , 142, 1127-47	6.6	407
164	The genetics of plant metabolism. <i>Nature Genetics</i> , 2006 , 38, 842-9	36.3	401
163	Isolation and characterization of abscisic acid-deficient Arabidopsis mutants at two new loci. <i>Plant Journal</i> , 1996 , 10, 655-61	6.9	393
162	GENETIC CONTROL OF FLOWERING TIME IN ARABIDOPSIS. <i>Annual Review of Plant Biology</i> , 1998 , 49, 345-370		382

(2008-2000)

161	Naturally occurring variation in Arabidopsis: an underexploited resource for plant genetics. <i>Trends in Plant Science</i> , 2000 , 5, 22-9	13.1	356
160	What has natural variation taught us about plant development, physiology, and adaptation?. <i>Plant Cell</i> , 2009 , 21, 1877-96	11.6	346
159	A QTL for flowering time in Arabidopsis reveals a novel allele of CRY2. <i>Nature Genetics</i> , 2001 , 29, 435-40)36.3	335
158	Gibberellin requirement for Arabidopsis seed germination is determined both by testa characteristics and embryonic abscisic acid. <i>Plant Physiology</i> , 2000 , 122, 415-24	6.6	310
157	The Arabidopsis aldehyde oxidase 3 (AAO3) gene product catalyzes the final step in abscisic acid biosynthesis in leaves. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000 , 97, 12908-13	11.5	306
156	In Vivo Inhibition of Seed Development and Reserve Protein Accumulation in Recombinants of Abscisic Acid Biosynthesis and Responsiveness Mutants in Arabidopsis thaliana. <i>Plant Physiology</i> , 1989 , 90, 463-9	6.6	298
155	Arabidopsis thaliana: a model plant for genome analysis. <i>Science</i> , 1998 , 282, 662, 679-82	33.3	297
154	Development of an AFLP based linkage map of Ler, Col and Cvi Arabidopsis thaliana ecotypes and construction of a Ler/Cvi recombinant inbred line population. <i>Plant Journal</i> , 1998 , 14, 259-71	6.9	293
153	Analysis of natural allelic variation at seed dormancy loci of Arabidopsis thaliana. <i>Genetics</i> , 2003 , 164, 711-29	4	284
152	The development of Arabidopsis as a model plant. <i>Plant Journal</i> , 2010 , 61, 909-21	6.9	270
151	Regulatory network construction in Arabidopsis by using genome-wide gene expression quantitative trait loci. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007 , 104, 1708-13	11.5	269
150	Natural allelic variation at seed size loci in relation to other life history traits of Arabidopsis thaliana. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1999 , 96, 4710	- 7 1.5	229
149	The absence of histone H2B monoubiquitination in the Arabidopsis hub1 (rdo4) mutant reveals a role for chromatin remodeling in seed dormancy. <i>Plant Cell</i> , 2007 , 19, 433-44	11.6	225
148	The phenotype of some late-flowering mutants is enhanced by a locus on chromosome 5 that is not effective in the Landsberg erecta wild-type. <i>Plant Journal</i> , 1994 , 6, 911-919	6.9	207
147	System-wide molecular evidence for phenotypic buffering in Arabidopsis. <i>Nature Genetics</i> , 2009 , 41, 166	53 76.3	205
146	Regulation of Arabidopsis thaliana Em genes: role of ABI5. <i>Plant Journal</i> , 2002 , 30, 373-83	6.9	201
145	The BANYULS gene encodes a DFR-like protein and is a marker of early seed coat development. <i>Plant Journal</i> , 1999 , 19, 387-98	6.9	200
144	Seed dormancy and germination. <i>The Arabidopsis Book</i> , 2008 , 6, e0119	3	192

143	Analysis of natural allelic variation at flowering time loci in the Landsberg erecta and Cape Verde Islands ecotypes of Arabidopsis thaliana. <i>Genetics</i> , 1998 , 149, 749-64	4	187
142	Control of FWA gene silencing in Arabidopsis thaliana by SINE-related direct repeats. <i>Plant Journal</i> , 2007 , 49, 38-45	6.9	186
141	ANTHOCYANINLESS2, a homeobox gene affecting anthocyanin distribution and root development in Arabidopsis. <i>Plant Cell</i> , 1999 , 11, 1217-26	11.6	185
140	Analysis of natural allelic variation of Arabidopsis seed germination and seed longevity traits between the accessions Landsberg erecta and Shakdara, using a new recombinant inbred line population. <i>Plant Physiology</i> , 2004 , 135, 432-43	6.6	182
139	Development of a near-isogenic line population of Arabidopsis thaliana and comparison of mapping power with a recombinant inbred line population. <i>Genetics</i> , 2007 , 175, 891-905	4	179
138	Flowering responses to light-breaks in photomorphogenic mutants of Arabidopsis thaliana, a long-day plant. <i>Physiologia Plantarum</i> , 1991 , 83, 209-215	4.6	175
137	A fortunate choice: the history of Arabidopsis as a model plant. <i>Nature Reviews Genetics</i> , 2002 , 3, 883-9	30.1	171
136	Genetic relationships within Brassica rapa as inferred from AFLP fingerprints. <i>Theoretical and Applied Genetics</i> , 2005 , 110, 1301-14	6	167
135	Genotype Invironment interaction QTL mapping in plants: lessons from Arabidopsis. <i>Trends in Plant Science</i> , 2014 , 19, 390-8	13.1	164
134	Genetic analysis of seed-soluble oligosaccharides in relation to seed storability of Arabidopsis. <i>Plant Physiology</i> , 2000 , 124, 1595-604	6.6	163
133	The genetic and molecular dissection of abscisic acid biosynthesis and signal transduction in Arabidopsis. <i>Plant Physiology and Biochemistry</i> , 1998 , 36, 83-89	5.4	159
132	Natural allelic variation identifies new genes in the Arabidopsis circadian system. <i>Plant Journal</i> , 1999 , 20, 67-77	6.9	158
131	Natural variation for seed dormancy in Arabidopsis is regulated by additive genetic and molecular pathways. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010 , 107, 4264-9	11.5	152
130	The genomic landscape of meiotic crossovers and gene conversions in Arabidopsis thaliana. <i>ELife</i> , 2013 , 2, e01426	8.9	149
129	Vacuolar invertase regulates elongation of Arabidopsis thaliana roots as revealed by QTL and mutant analysis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006 , 103, 2994-9	11.5	149
128	Next generation of elevated [CO2] experiments with crops: a critical investment for feeding the future world. <i>Plant, Cell and Environment</i> , 2008 , 31, 1317-24	8.4	145
127	A Seed Shape Mutant of Arabidopsis That Is Affected in Integument Development. <i>Plant Cell</i> , 1994 , 6, 385-392	11.6	142
126	Natural variation and QTL analysis for cationic mineral content in seeds of Arabidopsis thaliana. <i>Plant, Cell and Environment,</i> 2004 , 27, 828-839	8.4	137

125	A genetic analysis of cell culture traits in tomato. Theoretical and Applied Genetics, 1987, 74, 633-41	6	132
124	The earliest stages of adaptation in an experimental plant population: strong selection on QTLS for seed dormancy. <i>Molecular Ecology</i> , 2010 , 19, 1335-51	5.7	127
123	Analysis of natural allelic variation in Arabidopsis using a multiparent recombinant inbred line population. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011 , 108, 4488-93	11.5	122
122	Genetic and molecular analyses of natural variation indicate CBF2 as a candidate gene for underlying a freezing tolerance quantitative trait locus in Arabidopsis. <i>Plant Physiology</i> , 2005 , 139, 1304	4-4-2	120
121	Quantitative trait loci for flowering time and morphological traits in multiple populations of Brassica rapa. <i>Journal of Experimental Botany</i> , 2007 , 58, 4005-16	7	118
120	Chromosome-level assembly of Arabidopsis thaliana Ler reveals the extent of translocation and inversion polymorphisms. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, E4052-60	11.5	118
119	DOG1 expression is predicted by the seed-maturation environment and contributes to geographical variation in germination in Arabidopsis thaliana. <i>Molecular Ecology</i> , 2011 , 20, 3336-49	5.7	112
118	Quantitative trait locus analysis of growth-related traits in a new Arabidopsis recombinant inbred population. <i>Plant Physiology</i> , 2004 , 135, 444-58	6.6	110
117	Seed maturation in Arabidopsis thaliana is characterized by nuclear size reduction and increased chromatin condensation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011 , 108, 20219-24	11.5	107
116	QTL analysis of seed dormancy in Arabidopsis using recombinant inbred lines and MQM mapping. <i>Heredity</i> , 1997 , 79 (Pt 2), 190-200	3.6	106
115	Co-variation between seed dormancy, growth rate and flowering time changes with latitude in Arabidopsis thaliana. <i>PLoS ONE</i> , 2013 , 8, e61075	3.7	102
114	Natural variation at Strubbelig Receptor Kinase 3 drives immune-triggered incompatibilities between Arabidopsis thaliana accessions. <i>Nature Genetics</i> , 2010 , 42, 1135-9	36.3	99
113	Accumulation of C19-gibberellins in the gibberellin-insensitive dwarf mutantgai of Arabidopsis thaliana (L.) Heynh. <i>Planta</i> , 1990 , 182, 501-5	4.7	99
112	The conserved splicing factor SUA controls alternative splicing of the developmental regulator ABI3 in Arabidopsis. <i>Plant Cell</i> , 2010 , 22, 1936-46	11.6	98
111	The isolation and characterization of gibberellin-deficient mutants in tomato. <i>Theoretical and Applied Genetics</i> , 1990 , 80, 852-7	6	95
110	Far-red light-insensitive, phytochrome A-deficient mutants of tomato. <i>Molecular Genetics and Genomics</i> , 1995 , 246, 133-41		94
109	Gene function beyond the single trait: natural variation, gene effects, and evolutionary ecology in Arabidopsis thaliana. <i>Plant, Cell and Environment</i> , 2005 , 28, 2-20	8.4	92
108	Genetic differences in seed longevity of various Arabidopsis mutants. <i>Physiologia Plantarum</i> , 2004 , 121, 448-461	4.6	91

107	An integrated genetic/RFLP map of the Arabidopsis thaliana genome. <i>Plant Journal</i> , 1993 , 3, 745-754	6.9	90
106	RFLP markers linked to the root knot nematode resistance gene Mi in tomato. <i>Theoretical and Applied Genetics</i> , 1991 , 81, 661-7	6	89
105	Integrative analyses of genetic variation in enzyme activities of primary carbohydrate metabolism reveal distinct modes of regulation in Arabidopsis thaliana. <i>Genome Biology</i> , 2008 , 9, R129	18.3	82
104	New Arabidopsis recombinant inbred line populations genotyped using SNPWave and their use for mapping flowering-time quantitative trait loci. <i>Genetics</i> , 2006 , 172, 1867-76	4	80
103	Paths to selection on life history loci in different natural environments across the native range of Arabidopsis thaliana. <i>Molecular Ecology</i> , 2013 , 22, 3552-66	5.7	78
102	Association mapping of leaf traits, flowering time, and phytate content in Brassica rapa. <i>Genome</i> , 2007 , 50, 963-73	2.4	78
101	The aurea mutant of tomato is deficient in spectrophotometrically and immunochemically detectable phytochrome. <i>Plant Molecular Biology</i> , 1987 , 9, 97-107	4.6	76
100	Somaclonal variation in tomato: effect of explant source and a comparison with chemical mutagenesis. <i>Theoretical and Applied Genetics</i> , 1990 , 80, 817-25	6	75
99	Molecular analysis of the phytochrome deficiency in an aurea mutant of tomato. <i>Molecular Genetics and Genomics</i> , 1988 , 213, 9-14		7 ²
98	Breeding of a tomato genotype readily accessible to genetic manipulation. <i>Plant Science</i> , 1986 , 45, 201-	2508	7 <u>2</u>
97	Physiological interactions of phytochromes A, B1 and B2 in the control of development in tomato. <i>Plant Journal</i> , 2000 , 24, 345-56	6.9	71
96	Combined genetic and modeling approaches reveal that epidermal cell area and number in leaves are controlled by leaf and plant developmental processes in Arabidopsis. <i>Plant Physiology</i> , 2008 , 148, 1117-27	6.6	70
95	From phenotypic to molecular polymorphisms involved in naturally occurring variation of plant development. <i>International Journal of Developmental Biology</i> , 2005 , 49, 717-32	1.9	70
94	Characterization and mapping of a gene controlling shoot regeneration in tomato. <i>Plant Journal</i> , 1993 , 3, 131-141	6.9	70
93	Genetic dissection of blue-light sensing in tomato using mutants deficient in cryptochrome 1 and phytochromes A, B1 and B2. <i>Plant Journal</i> , 2001 , 25, 427-40	6.9	68
92	Cell division activity during apical hook development. <i>Plant Physiology</i> , 2001 , 125, 219-26	6.6	67
91	A strong effect of growth medium and organ type on the identification of QTLs for phytate and mineral concentrations in three Arabidopsis thaliana RIL populations. <i>Journal of Experimental Botany</i> , 2009 , 60, 1409-25	7	66
90	Community standards for Arabidopsis genetics. <i>Plant Journal</i> , 1997 , 12, 247-253	6.9	66

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89	The role of cryptochrome 2 in flowering in Arabidopsis. <i>Plant Physiology</i> , 2003 , 133, 1504-16	6.6	63
88	Quantitative trait loci for glucosinolate accumulation in Brassica rapa leaves. <i>New Phytologist</i> , 2008 , 179, 1017-1032	9.8	62
87	Quantitative genetics in the age of omics. Current Opinion in Plant Biology, 2008, 11, 123-8	9.9	62
86	Genetic basis for natural variation in seed vitamin E levels in Arabidopsis thaliana. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006 , 103, 18834-41	11.5	61
85	The root-knot nematode resistance gene (Mi) in tomato: construction of a molecular linkage map and identification of dominant cDNA markers in resistant genotypes. <i>Plant Journal</i> , 1992 , 2, 971-982	6.9	61
84	PHOTOMORPHOGENETIC MUTANTS OF HIGHER PLANTS*. <i>Photochemistry and Photobiology</i> , 1988 , 48, 833-841	3.6	61
83	Arabidopsis semidwarfs evolved from independent mutations in GA20ox1, ortholog to green revolution dwarf alleles in rice and barley. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 15818-23	11.5	59
82	Mapping QTLs for mineral accumulation and shoot dry biomass under different Zn nutritional conditions in Chinese cabbage (Brassica rapa L. ssp. pekinensis). <i>Plant and Soil</i> , 2008 , 310, 25-40	4.2	56
81	Importance of the B2 domain of the Arabidopsis ABI3 protein for Em and 2S albumin gene regulation. <i>Plant Molecular Biology</i> , 1999 , 40, 1045-54	4.6	56
80	Photophysiology and phytochrome content of long-hypocotyl mutant and wild-type cucumber seedlings. <i>Plant Physiology</i> , 1988 , 87, 264-8	6.6	56
79	Properties of proteins and the glassy matrix in maturation-defective mutant seeds of Arabidopsis thaliana. <i>Plant Journal</i> , 1998 , 16, 133-43	6.9	53
78	PHOTOCONTROL OF ANTHOCYANIN SYNTHESIS IN TOMATO SEEDLINGS: A GENETIC APPROACH*. <i>Photochemistry and Photobiology</i> , 1989 , 50, 107-111	3.6	53
77	Tomato chromosome 6: effect of alien chromosomal segments on recombinant frequencies. <i>Genome</i> , 1996 , 39, 485-91	2.4	50
76	Characterization of green seed, an enhancer of abi3-1 in Arabidopsis that affects seed longevity. <i>Plant Physiology</i> , 2003 , 132, 1077-84	6.6	49
75	Cytogenetic tools for Arabidopsis thaliana. <i>Chromosome Research</i> , 2003 , 11, 183-94	4.4	49
74	Molecular, genetic and evolutionary analysis of a paracentric inversion in Arabidopsis thaliana. <i>Plant Journal</i> , 2016 , 88, 159-178	6.9	47
73	Characterization of mutants with reduced seed dormancy at two novel rdo loci and a further characterization of rdo1 and rdo2 in Arabidopsis. <i>Physiologia Plantarum</i> , 2002 , 115, 604-612	4.6	47
72	Natural modifiers of seed longevity in the Arabidopsis mutants abscisic acid insensitive3-5 (abi3-5) and leafy cotyledon1-3 (lec1-3). <i>New Phytologist</i> , 2009 , 184, 898-908	9.8	46

71	Changing paradigms in plant breeding. <i>Plant Physiology</i> , 2001 , 125, 156-9	6.6	44
70	Analysis of a plant complex resistance gene locus underlying immune-related hybrid incompatibility and its occurrence in nature. <i>PLoS Genetics</i> , 2014 , 10, e1004848	6	43
69	Genetic analysis identifies quantitative trait loci controlling rosette mineral concentrations in Arabidopsis thaliana under drought. <i>New Phytologist</i> , 2009 , 184, 180-192	9.8	42
68	Quantitative trait loci analysis of phytate and phosphate concentrations in seeds and leaves of Brassica rapa. <i>Plant, Cell and Environment</i> , 2008 , 31, 887-900	8.4	42
67	The effect of daylength on the transition to flowering in phytochrome-deficient, late-flowering and double mutants of Arabidopsis thaliana. <i>Physiologia Plantarum</i> , 1995 , 95, 260-266	4.6	42
66	Quantitative trait loci and candidate genes underlying genotype by environment interaction in the response of Arabidopsis thaliana to drought. <i>Plant, Cell and Environment</i> , 2015 , 38, 585-99	8.4	41
65	Six-Rowed Spike3 (VRS3) Is a Histone Demethylase That Controls Lateral Spikelet Development in Barley. <i>Plant Physiology</i> , 2017 , 174, 2397-2408	6.6	41
64	Genetic analysis 1992 , 83-99		41
63	Variations in constitutive and inducible UV-B tolerance; dissecting photosystem II protection in Arabidopsis thaliana accessions. <i>Physiologia Plantarum</i> , 2010 , 138, 22-34	4.6	39
62	Tomato: a crop species amenable to improvement by cellular and molecular methods. <i>Euphytica</i> , 1989 , 42, 1-23	2.1	38
61	Tomato protoplast DNA transformation: physical linkage and recombination of exogenous DNA sequences. <i>Plant Molecular Biology</i> , 1987 , 8, 383-94	4.6	37
60	Analysis of phytochrome-deficient yellow-green-2 and aurea mutants of tomato. <i>Plant Journal</i> , 1996 , 9, 173-182	6.9	36
59	The Footprint of Polygenic Adaptation on Stress-Responsive Cis-Regulatory Divergence in the Arabidopsis Genus. <i>Molecular Biology and Evolution</i> , 2016 , 33, 2088-101	8.3	34
58	The ELONGATED gene of Arabidopsis acts independently of light and gibberellins in the control of elongation growth. <i>Plant Journal</i> , 1996 , 9, 305-12	6.9	34
57	Mutations in Barley Row Type Genes Have Pleiotropic Effects on Shoot Branching. <i>PLoS ONE</i> , 2015 , 10, e0140246	3.7	32
56	Epistatic natural allelic variation reveals a function of AGAMOUS-LIKE6 in axillary bud formation in Arabidopsis. <i>Plant Cell</i> , 2012 , 24, 2364-79	11.6	31
55	NLR locus-mediated trade-off between abiotic and biotic stress adaptation in Arabidopsis. <i>Nature Plants</i> , 2017 , 3, 17072	11.5	30
54	Fine mapping of a major QTL for awn length in barley using a multiparent mapping population. <i>Theoretical and Applied Genetics</i> , 2017 , 130, 269-281	6	30

(2015-1997)

53	QTL analysis of seed dormancy in Arabidopsis using recombinant inbred lines and MQM mapping. <i>Heredity</i> , 1997 , 79, 190-200	3.6	30
52	Altered photosynthetic performance of a natural Arabidopsis accession is associated with atrazine resistance. <i>Journal of Experimental Botany</i> , 2005 , 56, 1625-34	7	30
51	A multi-marker association method for genome-wide association studies without the need for population structure correction. <i>Nature Communications</i> , 2016 , 7, 13299	17.4	28
50	Characterization of natural variation for zinc, iron and manganese accumulation and zinc exposure response in Brassica rapa L <i>Plant and Soil</i> , 2007 , 291, 167-180	4.2	28
49	Mutations in and 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	11.5	28
48	Multiple loci and genetic interactions involving flowering time genes regulate stem branching among natural variants of Arabidopsis. <i>New Phytologist</i> , 2013 , 199, 843-57	9.8	27
47	Relation among plant growth, carbohydrates and flowering time in the Arabidopsis Landsberg erecta x Kondara recombinant inbred line population. <i>Plant, Cell and Environment</i> , 2010 , 33, 1369-82	8.4	27
46	Photomorphogenic mutants of higher plants 1994 , 601-628		27
45	Genotype-environment interactions affecting preflowering physiological and morphological traits of Brassica rapa grown in two watering regimes. <i>Journal of Experimental Botany</i> , 2014 , 65, 697-708	7	26
44	Somatic hybridization as a tool for tomato breeding. <i>Euphytica</i> , 1994 , 79, 265-277	2.1	26
43	Histochemical analysis reveals organ-specific quantitative trait loci for enzyme activities in Arabidopsis. <i>Plant Physiology</i> , 2004 , 134, 237-45	6.6	25
42	Natural variation for anthocyanin accumulation under high-light and low-temperature stress is attributable to the ENHANCER OF AG-4 2 (HUA2) locus in combination with PRODUCTION OF ANTHOCYANIN PIGMENT1 (PAP1) and PAP2. <i>New Phytologist</i> , 2015 , 206, 422-435	9.8	23
41	Madeiran Arabidopsis thaliana Reveals Ancient Long-Range Colonization and Clarifies Demography in Eurasia. <i>Molecular Biology and Evolution</i> , 2018 , 35, 564-574	8.3	22
40	Genetic analysis of morphological traits in a new, versatile, rapid-cycling Brassica rapa recombinant inbred line population. <i>Frontiers in Plant Science</i> , 2012 , 3, 183	6.2	22
39	A mixed model QTL analysis for a complex cross population consisting of a half diallel of two-way hybrids in Arabidopsis thaliana: analysis of simulated data. <i>Euphytica</i> , 2008 , 161, 107-114	2.1	22
38	Genetic and molecular organization of the short arm and pericentromeric region of tomato chromsome 6. <i>Euphytica</i> , 1994 , 79, 169-174	2.1	22
37	Conserved histidine of metal transporter AtNRAMP1 is crucial for optimal plant growth under manganese deficiency at chilling temperatures. <i>New Phytologist</i> , 2014 , 202, 1173-1183	9.8	21
36	phenoVein-A Tool for Leaf Vein Segmentation and Analysis. <i>Plant Physiology</i> , 2015 , 169, 2359-70	6.6	21

35	The genetic architecture of freezing tolerance varies across the range of Arabidopsis thaliana. <i>Plant, Cell and Environment</i> , 2016 , 39, 2570-2579	8.4	20
34	QTL analysis. <i>Methods in Molecular Biology</i> , 2006 , 323, 79-99	1.4	18
33	Reciprocal cybrids reveal how organellar genomes affect plant phenotypes. <i>Nature Plants</i> , 2020 , 6, 13-	2111.5	17
32	Plants regenerated from auxin-auxotrophic variants are inviable. <i>Molecular Genetics and Genomics</i> , 1988 , 215, 58-64		16
31	A comparison of population types used for QTL mapping in Arabidopsis thaliana. <i>Plant Genetic Resources: Characterisation and Utilisation</i> , 2011 , 9, 185-188	1	15
30	Genetic variation in flowering time inArabidopsis thaliana. <i>Seminars in Cell and Developmental Biology</i> , 1996 , 7, 381-389	7.5	14
29	Isolation and characterization of nitrate reductase-deficient mutants in tomato (Lycopersicon esculentum Mill.). <i>Molecular Genetics and Genomics</i> , 1991 , 227, 458-64		14
28	Genetic Aspects of Seed Dormancy113-132		13
27	Variation in seed dormancy quantitative trait loci in Arabidopsis thaliana originating from one site. <i>PLoS ONE</i> , 2011 , 6, e20886	3.7	13
26	Phenotype of Arabidopsis thaliana semi-dwarfs with deep roots and high growth rates under water-limiting conditions is independent of the GA5 loss-of-function alleles. <i>Annals of Botany</i> , 2015 , 116, 321-31	4.1	12
25	Flowering responses to light-breaks in photomorphogenic mutants of Arabidopsis thaliana, a long-day plant. <i>Physiologia Plantarum</i> , 1991 , 83, 209-215	4.6	12
24	Quantitative trait loci controlling leaf venation in Arabidopsis. <i>Plant, Cell and Environment</i> , 2017 , 40, 1429-1441	8.4	10
23	Analysis of nuclear and organellar DNA of somatic hybrid calli and plants between Lycopersicon spp. and Nicotiana spp. <i>Molecular Genetics and Genomics</i> , 1993 , 241, 707-18		10
22	Genes for seed longevity in barley identified by genomic analysis on near isogenic lines. <i>Plant, Cell and Environment</i> , 2018 , 41, 1895-1911	8.4	8
21	The TRANSPARENT TESTA12 Gene of Arabidopsis Encodes a Multidrug Secondary Transporter-Like Protein Required for Flavonoid Sequestration in Vacuoles of the Seed Coat Endothelium. <i>Plant Cell</i> , 2001 , 13, 853	11.6	8
20	Signals of speciation within Arabidopsis thaliana in comparison with its relatives. <i>Current Opinion in Plant Biology</i> , 2012 , 15, 205-11	9.9	6
19	Genetic analysis. <i>Methods in Molecular Biology</i> , 2006 , 323, 65-77	1.4	5
18	Asymmetric fusion between protoplasts of tomato (Lycopersicon esculentum Mill.) and gamma-irradiated protoplasts of potato (Solanum tuberosum L.): the effects of gamma irradiation. Molecular Genetics and Genomics 1994, 242, 313-20		5

LIST OF PUBLICATIONS

17	Use of leaky nitrate reductase-deficient mutants of tomato (Lycopersicon esculentum Mill.) for selection of somatic hybrid cell lines with wild type potato (Solanum tuberosum L.). <i>Plant Cell, Tissue and Organ Culture</i> , 1992 , 31, 151-154	2.7	5
16	QTL and candidate genes associated with leaf anion concentrations in response to phosphate supply in Arabidopsis thaliana. <i>BMC Plant Biology</i> , 2019 , 19, 410	5.3	3
15	Optimization of Direct Gene Transfer in Tomato. <i>Current Plant Science and Biotechnology in Agriculture</i> , 1988 , 357-358		3
14	Natural Variation in Arabidopsis thaliana 2011 , 123-151		3
13	Identification and characterization of quantitative trait loci that control seed dormancy in Arabidopsis. <i>Methods in Molecular Biology</i> , 2011 , 773, 165-84	1.4	2
12	An integrated genetic/RFLP map of the Arabidopsis thaliana genome 1993 , 3, 745		2
11	The effect of daylength on the transition to flowering in phytochrome-deficient, late-flowering and double mutants of Arabidopsis thaliana. <i>Physiologia Plantarum</i> , 1995 , 95, 260-266	4.6	2
10	and Are Key Regulators of Flowering Time in the Biennial/Perennial Species. <i>Frontiers in Plant Science</i> , 2020 , 11, 582577	6.2	2
9	A Central Role for Genetics in Plant Biology. Annual Review of Plant Biology, 2021, 72, 1-16	30.7	2
8	Dormancy in Plants 2007 ,		1
7	Conserved Mechanisms of Dormancy and Germination as Targets for Manipulation of Agricultural Problems 2006 , 11-32		1
6	The evolutionary dynamics of genetic incompatibilities introduced by duplicated genes in Arabidopsis thaliana		1
5	The Evolutionary Dynamics of Genetic Incompatibilities Introduced by Duplicated Genes in Arabidopsis thaliana. <i>Molecular Biology and Evolution</i> , 2021 , 38, 1225-1240	8.3	О
4	Genetic Aspects of Seed Dormancy 2018 , 113-132		O
3	My favourite flowering image. <i>Journal of Experimental Botany</i> , 2013 , 64, 5801-3	7	
2	Interview with Maarten Koornneef. <i>Trends in Plant Science</i> , 2015 , 20, 135-6	13.1	
1	Spotlight on phytochrome nomenclature. <i>Plant Molecular Biology Reporter</i> , 1994 , 12, S50-S56	1.7	