

Richard B Flavell

List of Publications by Year in descending order

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17
papers

1,982
citations

516561

16
h-index

887953

17
g-index

17
all docs

17
docs citations

17
times ranked

1767
citing authors

#	ARTICLE	IF	CITATIONS
1	Perspective: 50 years of plant chromosome biology. <i>Plant Physiology</i> , 2021, 185, 731-753.	2.3	1
2	High Resolution Genetic Mapping by Genome Sequencing Reveals Genome Duplication and Tetraploid Genetic Structure of the Diploid <i>Miscanthus sinensis</i> . <i>PLoS ONE</i> , 2012, 7, e33821.	1.1	103
3	Genome-Wide Discovery of <i>cis</i> -Elements in Promoter Sequences Using Gene Expression. <i>OMICS A Journal of Integrative Biology</i> , 2009, 13, 139-151.	1.0	32
4	Insights into corn genes derived from large-scale cDNA sequencing. <i>Plant Molecular Biology</i> , 2009, 69, 179-194.	2.0	212
5	Features of Arabidopsis Genes and Genome Discovered using Full-length cDNAs. <i>Plant Molecular Biology</i> , 2006, 60, 69-85.	2.0	145
6	<i>Agrobacterium</i> T-DNA integration in Arabidopsis is correlated with DNA sequence compositions that occur frequently in gene promoter regions. <i>Functional and Integrative Genomics</i> , 2005, 5, 240-253.	1.4	43
7	Full-length messenger RNA sequences greatly improve genome annotation. <i>Genome Biology</i> , 2002, 3, research0029.1.	13.9	147
8	Developmentally and transgene regulated nuclear processing of primary transcripts of chalcone synthase A in petunia. <i>Plant Journal</i> , 2000, 23, 63-72.	2.8	25
9	Post-transcriptional gene silencing of chalcone synthase in transgenic petunias, cytosine methylation and epigenetic variation. <i>Plant Journal</i> , 1999, 18, 33-42.	2.8	19
10	Details of T-DNA structural organization from a transgenic <i>Petunia</i> population exhibiting co-suppression. <i>Plant Molecular Biology</i> , 1996, 32, 1197-1203.	2.0	103
11	Correlation between the size of the intergenic regulatory region, the status of cytosine methylation of rRNA genes and nucleolar expression in wheat. <i>Molecular Genetics and Genomics</i> , 1993, 236-236, 155-162.	2.4	88
12	Key Features of Cereal Genome Organization as Revealed by the Use of Cytosine Methylation-Sensitive Restriction Endonucleases. <i>Genomics</i> , 1993, 15, 472-482.	1.3	84
13	Protein-binding to reiterated motifs within the wheat rRNA gene promoter and upstream repeats. <i>Plant Molecular Biology</i> , 1992, 20, 911-919.	2.0	19
14	Molecular coevolution: DNA divergence and the maintenance of function. <i>Cell</i> , 1984, 38, 622-623.	13.5	226
15	A chimaeric antibiotic resistance gene as a selectable marker for plant cell transformation. <i>Nature</i> , 1983, 304, 184-187.	13.7	555
16	Sequence organisation analysis of the wheat and rye genomes by interspecies DNA/DNA hybridisation. <i>Journal of Molecular Biology</i> , 1978, 123, 327-359.	2.0	85
17	Nucleotide sequence organisation in the wheat genome. <i>Heredity</i> , 1976, 37, 231-252.	1.2	95