

# Thomas Peterson

## List of Publications by Year in descending order

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49  
papers

2,889  
citations

236925

25  
h-index

206112

48  
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54  
all docs

54  
docs citations

54  
times ranked

2692  
citing authors

#	ARTICLE	IF	CITATIONS
1	Excision and reinsertion of <i>Ac</i> macrotransposons in maize. <i>Genetics</i> , 2022, 221, .	2.9	4
2	A Tutorial of EDTA: Extensive De Novo TE Annotator. <i>Methods in Molecular Biology</i> , 2021, 2250, 55-67.	0.9	22
3	Rapid Detection of Transposon-Induced Genome Rearrangements. <i>Methods in Molecular Biology</i> , 2021, 2250, 131-139.	0.9	3
4	Transposon-induced inversions activate gene expression in the maize pericarp. <i>Genetics</i> , 2021, 218, .	2.9	7
5	Ectopic Expression of a Maize Gene Is Induced by Composite Insertions Generated Through Alternative Transposition. <i>Genetics</i> , 2020, 216, 1039-1049.	2.9	9
6	Small RNA-Mediated <i>De Novo</i> Silencing of <i>Ac/Ds</i> Transposons Is Initiated by Alternative Transposition in Maize. <i>Genetics</i> , 2020, 215, 393-406.	2.9	11
7	TIR-Learner, a New Ensemble Method for TIR Transposable Element Annotation, Provides Evidence for Abundant New Transposable Elements in the Maize Genome. <i>Molecular Plant</i> , 2019, 12, 447-460.	8.3	90
8	Benchmarking transposable element annotation methods for creation of a streamlined, comprehensive pipeline. <i>Genome Biology</i> , 2019, 20, 275.	8.8	579
9	Evolutionary Impacts of Alternative Transposition. , 2018, , 113-130.		6
10	Maize Transposon Storm Kicks up a <i>White Cap</i> . <i>Genetics</i> , 2017, 206, 87-89.	2.9	1
11	Generation and Analysis of Transposon <i>Ac/Ds</i> -Induced Chromosomal Rearrangements in Rice Plants. <i>Methods in Molecular Biology</i> , 2016, 1469, 49-61.	0.9	4
12	Genes and Small RNA Transcripts Exhibit Dosage-Dependent Expression Pattern in Maize Copy-Number Alterations. <i>Genetics</i> , 2016, 203, 1133-1147.	2.9	12
13	Alternative Transposition Generates New Chimeric Genes and Segmental Duplications at the Maize <i>p1</i> Locus. <i>Genetics</i> , 2015, 201, 925-935.	2.9	13
14	Transposition-mediated DNA re-replication in maize. <i>ELife</i> , 2014, 3, e03724.	6.0	21
15	Generation of Tandem Direct Duplications by Reversed-Ends Transposition of Maize <i>Ac</i> Elements. <i>PLoS Genetics</i> , 2013, 9, e1003691.	3.5	43
16	Isolation of Sequences Flanking <i>Ac</i> Insertion Sites by <i>Ac</i> Casting. <i>Methods in Molecular Biology</i> , 2013, 1057, 117-122.	0.9	8
17	Survey of Natural and Transgenic Gene Markers Used to Monitor Transposon Activity. <i>Methods in Molecular Biology</i> , 2013, 1057, 43-58.	0.9	0
18	A transgenic system for generation of transposon <i>Ac/Ds</i> -induced chromosome rearrangements in rice. <i>Theoretical and Applied Genetics</i> , 2012, 125, 1449-1462.	3.6	20

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19	Transposable Elements as Catalysts for Chromosome Rearrangements. <i>Methods in Molecular Biology</i> , 2011, 701, 315-326.	0.9	41
20	Transposon Ac/Ds -induced chromosomal rearrangements at the rice OsRLG5 locus. <i>Nucleic Acids Research</i> , 2011, 39, e149-e149.	14.5	19
21	Genome Rearrangements in Maize Induced by Alternative Transposition of Reversed Ac/Ds Termini. <i>Genetics</i> , 2011, 188, 59-67.	2.9	47
22	Fusion of reverse-oriented Ds termini following abortive transposition in Arabidopsis: implications for the mechanism of Ac/Ds transposition. <i>Plant Cell Reports</i> , 2010, 29, 413-417.	5.6	9
23	Spatial Configuration of Transposable Element Ac Termini Affects Their Ability to Induce Chromosomal Breakage in Maize. <i>Plant Cell</i> , 2010, 22, 744-754.	6.6	24
24	Alternative Ac/Ds transposition induces major chromosomal rearrangements in maize. <i>Genes and Development</i> , 2009, 23, 755-765.	5.9	61
25	Reversed end Ds element: a novel tool for chromosome engineering in Arabidopsis. <i>Plant Molecular Biology</i> , 2008, 68, 399-411.	3.9	14
26	Epigenetic Modifications of Distinct Sequences of the p1 Regulatory Gene Specify Tissue-Specific Expression Patterns in Maize. <i>Genetics</i> , 2007, 175, 1059-1070.	2.9	35
27	Comparative Structural and Functional Characterization of Sorghum and Maize Duplications Containing Orthologous Myb Transcription Regulators of 3-Deoxyflavonoid Biosynthesis. <i>Plant Molecular Biology</i> , 2006, 60, 185-199.	3.9	47
28	Transposition of Reversed Ac Element Ends Generates Novel Chimeric Genes in Maize. <i>PLoS Genetics</i> , 2006, 2, e164.	3.5	53
29	Gene Conversion Between Direct Noncoding Repeats Promotes Genetic and Phenotypic Diversity at a Regulatory Locus of Zea mays (L.). <i>Genetics</i> , 2006, 174, 753-762.	2.9	6
30	Transformation of maize with the p1 transcription factor directs production of silk maysin, a corn earworm resistance factor, in concordance with a hierarchy of floral organ pigmentation. <i>Plant Biotechnology Journal</i> , 2005, 3, 225-235.	8.3	28
31	Comparisons of Maize pericarp color1 Alleles Reveal Paralogous Gene Recombination and an Organ-Specific Enhancer Region. <i>Plant Cell</i> , 2005, 17, 903-914.	6.6	46
32	A Segmental Deletion Series Generated by Sister-Chromatid Transposition of Ac Transposable Elements in Maize. <i>Genetics</i> , 2005, 171, 333-344.	2.9	44
33	Transposition of Reversed Ac Element Ends Generates Chromosome Rearrangements in Maize. <i>Genetics</i> , 2004, 167, 1929-1937.	2.9	80
34	A maize QTL for silk maysin levels contains duplicated Myb-homologous genes which jointly regulate flavone biosynthesis. <i>Plant Molecular Biology</i> , 2003, 52, 1-15.	3.9	80
35	Functional analysis of two matrix attachment region (MAR) elements in transgenic maize plants. <i>Transgenic Research</i> , 2003, 12, 137-154.	2.4	20
36	The Maize Unstable factor for orange1 Is a Dominant Epigenetic Modifier of a Tissue Specifically Silent Allele of pericarp color1. <i>Genetics</i> , 2003, 163, 1135-1146.	2.9	40

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37	Tissue-specific patterns of a maize Myb transcription factor are epigenetically regulated. <i>Plant Journal</i> , 2001, 27, 467-478.	5.7	56
38	Transgene-Induced Silencing Identifies Sequences Involved in the Establishment of Paramutation of the Maize <i>p1</i> Gene. <i>Plant Cell</i> , 2001, 13, 319.	6.6	2
39	Transgene-Induced Silencing Identifies Sequences Involved in the Establishment of Paramutation of the Maize <i>p1</i> Gene. <i>Plant Cell</i> , 2001, 13, 319-335.	6.6	87
40	Complex structure of a maize Myb gene promoter: functional analysis in transgenic plants. <i>Plant Journal</i> , 2000, 22, 471-482.	5.7	69
41	A Segmental Gene Duplication Generated Differentially Expressed myb-Homologous Genes in Maize. <i>Plant Cell</i> , 2000, 12, 2311.	6.6	4
42	A Segmental Gene Duplication Generated Differentially Expressed myb-Homologous Genes in Maize. <i>Plant Cell</i> , 2000, 12, 2311-2322.	6.6	110
43	Hierarchical Patterns of Transgene Expression Indicate Involvement of Developmental Mechanisms in the Regulation of the Maize <i>P1-rr</i> Promoter. <i>Genetics</i> , 2000, 156, 839-846.	2.9	23
44	Ac Insertion Site Affects the Frequency of Transposon-Induced Homologous Recombination at the Maize <i>p1</i> Locus. <i>Genetics</i> , 2000, 156, 2007-2017.	2.9	30
45	Characterization of the regulatory elements of the maize <i>P-rr</i> gene by transient expression assays. <i>Plant Molecular Biology</i> , 1999, 39, 11-19.	3.9	41
46	Genome Rearrangements by Nonlinear Transposons in Maize. <i>Genetics</i> , 1999, 153, 1403-1410.	2.9	89
47	Isolation and characterization of a maize gene encoding chalcone flavonone isomerase. <i>Molecular Genetics and Genomics</i> , 1994, 242, 1-8.	2.4	70
48	The myb-homologous <i>P</i> gene controls phlobaphene pigmentation in maize floral organs by directly activating a flavonoid biosynthetic gene subset. <i>Cell</i> , 1994, 76, 543-553.	28.9	644
49	Isolation and molecular analysis of the maize <i>P</i> locus. <i>Molecular Genetics and Genomics</i> , 1989, 219, 225-234.	2.4	109