Peter L Tiffin

List of Publications by Year
in descending order

101 papers	8,504 citations	53939 47 h-index	56606 87 g-index
114 all docs	114 docs citations	114 times ranked	12317 citing authors

#	Article	IF	Citations
1	Unraveling coevolutionary dynamics using ecological genomics. Trends in Genetics, 2022, 38, 1003-1012.	2.9	4
2	Comparative genomics reveals high rates of horizontal transfer and strong purifying selection on rhizobial symbiosis genes. Proceedings of the Royal Society B: Biological Sciences, 2021, 288, 20201804.	1.2	13
3	Genomic structural variants constrain and facilitate adaptation in natural populations of <i>Theobroma cacao</i> , the chocolate tree. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	50
4	Individual-based eco-evolutionary models for understanding adaptation in changing seas. Proceedings of the Royal Society B: Biological Sciences, 2021, 288, 20212006.	1.2	4
5	Gene Expression Modularity Reveals Footprints of Polygenic Adaptation in Theobroma cacao. Molecular Biology and Evolution, 2020, 37, 110-123.	3.5	22
6	Context Dependence of Local Adaptation to Abiotic and Biotic Environments: A Quantitative and Qualitative Synthesis. American Naturalist, 2020, 195, 412-431.	1.0	55
7	Widely distributed variation in tolerance to Phytophthora palmivora in four genetic groups of cacao. Tree Genetics and Genomes, 2020, 16, 1.	0.6	15
8	Biased Gene Conversion Constrains Adaptation in <i>Arabidopsis thaliana</i>. Genetics, 2020, 215, 831-846.	1.2	15
9	Pleiotropy facilitates local adaptation to distant optima in common ragweed (Ambrosia) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 4	1.5	30
10	A Select and Resequence Approach Reveals Strain-Specific Effects of <i>Medicago</i> Nodule-Specific PLAT-Domain Genes. Plant Physiology, 2020, 182, 463-471.	2.3	13
11	Does adaptation to historical climate shape plant responses to future rainfall patterns? A rainfall manipulation experiment with common ragweed. Oecologia, 2019, 190, 941-953.	0.9	11
12	Climate change is predicted to disrupt patterns of local adaptation in wild and cultivated maize. Proceedings of the Royal Society B: Biological Sciences, 2019, 286, 20190486.	1.2	32
13	Legacy of prior host and soil selection on rhizobial fitness in <i>planta</i>. Evolution; International Journal of Organic Evolution, 2019, 73, 2013-2023.	1.1	19
14	Species distribution models throughout the invasion history of Palmer amaranth predict regions at risk of future invasion and reveal challenges with modeling rapidly shifting geographic ranges. Scientific Reports, 2019, 9, 2426.	1.6	60
15	Sexâ€biased gene expression in flowers, but not leaves, reveals secondary sexual dimorphism in <i>Populus balsamifera</i>. New Phytologist, 2019, 221, 527-539.	3.5	38
16	Select and resequence reveals relative fitness of bacteria in symbiotic and free-living environments. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 2425-2430.	3.3	88
17	Effects of Gene Action, Marker Density, and Timing of Selection on the Performance of Landscape Genomic Scans of Local Adaptation. Journal of Heredity, 2018, 109, 16-28.	1.0	17
18	Genome-Wide Association Analyses in the Model Rhizobium <i>Ensifer meliloti</i>. MSphere, 2018, 3, .	1.3	26

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19	Little plant, big city: a test of adaptation to urban environments in common ragweed (<i>Ambrosia) Tj ETQq1 1 0.784314 rgBT /Overlo	1.2	50
20	The complete replicons of 16 Ensifer meliloti strains offer insights into intra- and inter-replicon gene transfer, transposon-associated loci, and repeat elements. Microbial Genomics, 2018, 4, .	1.0	21
21	Validating Genome-Wide Association Candidates Controlling Quantitative Variation in Nodulation. Plant Physiology, 2017, 173, 921-931.	2.3	71
22	A Guide to Genomeâ€Wide Association Mapping in Plants. Current Protocols in Plant Biology, 2017, 2, 22-38.	2.8	75
23	Exploring structural variation and gene family architecture with De Novo assemblies of 15 Medicago genomes. BMC Genomics, 2017, 18, 261.	1.2	87
24	Sanctions, Partner Recognition, and Variation in Mutualism. American Naturalist, 2017, 190, 491-505.	1.0	10
25	Transcriptomic basis of genome by genome variation in a legumeâ€rhizobia mutualism. Molecular Ecology, 2017, 26, 6122-6135.	2.0	40
26	Living in the city: urban environments shape the evolution of a native annual plant. Global Change Biology, 2017, 23, 2082-2089.	4.2	52
27	Strategies for optimizing BioNano and Dovetail explored through a second reference quality assembly for the legume model, Medicago truncatula. BMC Genomics, 2017, 18, 578.	1.2	54
28	Hybrid assembly with long and short reads improves discovery of gene family expansions. BMC Genomics, 2017, 18, 541.	1.2	51
29	ODG: Omics database generator - a tool for generating, querying, and analyzing multi-omics comparative databases to facilitate biological understanding. BMC Bioinformatics, 2017, 18, 367.	1.2	13
30	The Role of Deleterious Substitutions in Crop Genomes. Molecular Biology and Evolution, 2016, 33, 2307-2317.	3.5	83
31	Adaptation to climate through flowering phenology: a case study in <i>Medicago truncatula</i>. Molecular Ecology, 2016, 25, 3397-3415.	2.0	36
32	Relocation, highâ€latitude warming and host genetic identity shape the foliar fungal microbiome of poplars. Molecular Ecology, 2015, 24, 235-248.	2.0	125
33	Potential of Association Mapping and Genomic Selection to Explore PI 88788 Derived Soybean Cyst Nematode Resistance. Plant Genome, 2014, 7, plantgenome2013.11.0039.	1.6	63
34	Selection on Horizontally Transferred and Duplicated Genes in Sinorhizobium (Ensifer), the Root-Nodule Symbionts of Medicago. Genome Biology and Evolution, 2014, 6, 1199-1209.	1.1	17
35	Local adaptation and range boundary formation in response to complex environmental gradients across the geographical range of <i>Clarkia xantiana</i> ssp<i>. xantiana</i>. Journal of Ecology, 2014, 102, 95-107.	1.9	49
36	Highâ€density genomeâ€wide association mapping implicates an <scp>F</scp>â€box encoding gene in <i><scp>M</scp>edicago truncatula</i> resistance to <i><scp>A</scp>phanomyces euteiches</i>. New Phytologist, 2014, 201, 1328-1342.	3.5	86

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37	Advances and limits of using population genetics to understand local adaptation. Trends in Ecology and Evolution, 2014, 29, 673-680.	4.2	329
38	Indirect effects drive evolutionary responses to global change. New Phytologist, 2014, 201, 335-343.	3.5	31
39	Timing for success: expression phenotype and local adaptation related to latitude in the boreal forest tree, Populus balsamifera. Tree Genetics and Genomes, 2014, 10, 911-922.	0.6	7
40	Genomic Signature of Adaptation to Climate in <i>Medicago truncatula</i>. Genetics, 2014, 196, 1263-1275.	1.2	160
41	Comparative genomics of the core and accessory genomes of 48 Sinorhizobium strains comprising five genospecies. Genome Biology, 2013, 14, R17.	13.9	164
42	Selection, genomeâ€wide fitness effects and evolutionary rates in the model legume <i><scp>M</scp>edicago truncatula</i>. Molecular Ecology, 2013, 22, 3525-3538.	2.0	54
43	Phylogenetic Signal Variation in the Genomes of Medicago (Fabaceae). Systematic Biology, 2013, 62, 424-438.	2.7	51
44	The adaptive potential of <i><scp>P</scp>opulus balsamifera </i><scp>L</scp>. to phenology requirements in a warmer global climate. Molecular Ecology, 2013, 22, 1214-1230.	2.0	91
45	Insights from population genetics for range limits of a widely distributed native plant. American Journal of Botany, 2013, 100, 744-753.	0.8	12
46	Estimating heritability using genomic data. Methods in Ecology and Evolution, 2013, 4, 1151-1158.	2.2	54
47	Epigenetic and Genetic Influences on DNA Methylation Variation in Maize Populations. Plant Cell, 2013, 25, 2783-2797.	3.1	227
48	Host Genotype Shapes the Foliar Fungal Microbiome of Balsam Poplar (Populus balsamifera). PLoS ONE, 2013, 8, e53987.	1.1	213
49	Candidate Genes and Genetic Architecture of Symbiotic and Agronomic Traits Revealed by Whole-Genome, Sequence-Based Association Genetics in Medicago truncatula. PLoS ONE, 2013, 8, e65688.	1.1	156
50	Reshaping of the maize transcriptome by domestication. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 11878-11883.	3.3	154
51	Fine-Scale Population Recombination Rates, Hotspots, and Correlates of Recombination in the Medicago truncatula Genome. Genome Biology and Evolution, 2012, 4, 726-737.	1.1	62
52	Reduced pollinator service and elevated pollen limitation at the geographic range limit of an annual plant. Ecology, 2012, 93, 1036-1048.	1.5	119
53	Pleistocene Speciation in the Genus Populus (Salicaceae). Systematic Biology, 2012, 61, 401.	2.7	100
54	Role of climate and competitors in limiting fitness across range edges of an annual plant. Ecology, 2012, 93, 1604-1613.	1.5	81

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55	Local Adaptation in the Flowering-Time Gene Network of Balsam Poplar, Populus balsamifera L.. Molecular Biology and Evolution, 2012, 29, 3143-3152.	3.5	106
56	Comparative population genomics of maize domestication and improvement. Nature Genetics, 2012, 44, 808-811.	9.4	816
57	Population Genomics of the Facultatively Mutualistic Bacteria Sinorhizobium meliloti and S. medicae. PLoS Genetics, 2012, 8, e1002868.	1.5	69
58	Interactions between Soil Habitat and Geographic Range Location Affect Plant Fitness. PLoS ONE, 2012, 7, e36015.	1.1	28
59	Whole-genome nucleotide diversity, recombination, and linkage disequilibrium in the model legume <i>Medicago truncatula</i>. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, E864-70.	3.3	220
60	Population Genetics and the Evolution of Geographic Range Limits in an Annual Plant. American Naturalist, 2011, 178, S44-S57.	1.0	44
61	Climateâ€driven local adaptation of ecophysiology and phenology in balsam poplar, <i>Populus balsamifera</i> L. (Salicaceae). American Journal of Botany, 2011, 98, 99-108.	0.8	103
62	Local Selection Across a Latitudinal Gradient Shapes Nucleotide Diversity in Balsam Poplar, <i>Populus balsamifera</i> L. Genetics, 2011, 188, 941-952.	1.2	47
63	Nucleotide diversity and linkage disequilibrium in balsam poplar (<i>Populus balsamifera</i>). New Phytologist, 2010, 186, 526-536.	3.5	70
64	Plant functional type classifications in tropical dry forests in Costa Rica: leaf habit versus taxonomic approaches. Functional Ecology, 2010, 24, 927-936.	1.7	112
65	Genomic diversity, population structure, and migration following rapid range expansion in the Balsam Poplar,<i>Populus balsamifera</i>. Molecular Ecology, 2010, 19, 1212-1226.	2.0	101
66	Pervasive gene content variation and copy number variation in maize and its undomesticated progenitor. Genome Research, 2010, 20, 1689-1699.	2.4	309
67	Heterosis Is Prevalent for Multiple Traits in Diverse Maize Germplasm. PLoS ONE, 2009, 4, e7433.	1.1	173
68	Elevated carbon dioxide concentrations indirectly affect plant fitness by altering plant tolerance to herbivory. Oecologia, 2009, 161, 401-410.	0.9	24
69	STABILIZING MECHANISMS IN A LEGUME-RHIZOBIUM MUTUALISM. Evolution; International Journal of Organic Evolution, 2009, 63, 652-662.	1.1	174
70	Selective histories of poplar protease inhibitors: elevated polymorphism, purifying selection, and positive selection driving divergence of recent duplicates. New Phytologist, 2009, 183, 740-750.	3.5	29
71	Transgenerational effects of global environmental change: long-term CO2 and nitrogen treatments influence offspring growth response to elevated CO2. Oecologia, 2008, 158, 141-150.	0.9	41
72	GEOGRAPHIC VARIATION IN ADAPTATION AT THE MOLECULAR LEVEL: A CASE STUDY OF PLANT IMMUNITY GENES. Evolution; International Journal of Organic Evolution, 2008, 62, 3069-3081.	1.1	37

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73	Evolution in ecological field experiments: implications for effect size. Ecology Letters, 2008, 11, 199-207.	3.0	66
74	The quest for adaptive evolution: a theoretical challenge in a maze of data. Current Opinion in Plant Biology, 2008, 11, 110-115.	3.5	16
75	DIRECT AND INDIRECT EFFECTS OF CO₂, NITROGEN, AND COMMUNITY DIVERSITY ON PLANTâ€"ENEMY INTERACTIONS. Ecology, 2008, 89, 226-236.	1.5	28
76	Context dependence in the coevolution of plant and rhizobial mutualists. Proceedings of the Royal Society B: Biological Sciences, 2007, 274, 1905-1912.	1.2	209
77	Population Structure and Its Effects on Patterns of Nucleotide Polymorphism in Teosinte (Zea mays) Tj ETQq1 1 0.784314 rgBT /Over	1.2	55
78	Sequence diversity and haplotype associations with phenotypic responses to crowding: GIGANTEA affects fruit set in Arabidopsis thaliana. Molecular Ecology, 2007, 16, 3050-3062.	2.0	21
79	Strong ecological but weak evolutionary effects of elevated CO 2 on a recombinant inbred population of Arabidopsis thaliana. New Phytologist, 2007, 175, 351-362.	3.5	37
80	An integrated approach for the comparative analysis of a multigene family: The nicotianamine synthase genes of barley. Functional and Integrative Genomics, 2007, 7, 169-179.	1.4	10
81	Molecular evolution of plant immune system genes. Trends in Genetics, 2006, 22, 662-670.	2.9	111
82	Genetic Diversity and the Evolutionary History of Plant Immunity Genes in Two Species of Zea. Molecular Biology and Evolution, 2005, 22, 2480-2490.	3.5	31
83	Comparative Evolutionary Histories of Chitinase Genes in the Genus Zea and Family PoaceaeSequence data from this article have been deposited with the EMBL/GenBank Data Libraries under accession nos. AY532721, AY532722, AY532723, AY532724, AY532725, AY532726, AY532727, AY532728, AY532729, AY532730, AY532731, AY532732, AY532733, AY532734, AY532735, AY532736, AY532737, AY532738, AY532739, AY532740, AY532741, AY532742, AY53274?, AY532744, AY532745, AY532746, AY532747, AY532748, AY532749, AY532750,	1.2	44
84	Population Genetic Evidence for Rapid Changes in Intraspecific Diversity and Allelic Cycling of a Specialist Defense Gene in ZeaSequence data from this article have been deposited with the EMBL/GenBank Data Libraries under accession nos. AY320258â€"320280, AY52550, AY52551, AY52552, AY52553, AY52554, AY52555, AY52556, AY52557, AY52558, AY52559, and AY549598, AY549599, AY549600, AY549601, AY549602, AY549603, AY549604, AY549605, AY549606, AY549607, AY549608, AY549609, AY549610, AY549611, AY549612, AY549613, AY5. Genetics, 2004, 168, 425-434.	1.2	34
85	PATTERNS OF REPRODUCTIVE ISOLATION IN THREE ANGIOSPERM GENERA. Evolution; International Journal of Organic Evolution, 2004, 58, 1195-1208.	1.1	213
86	MEASURING TOLERANCE TO HERBIVORY WITH NATURAL OR IMPOSED DAMAGE: A REPLY TO LEHTILÃ.... Evolution; International Journal of Organic Evolution, 2003, 57, 681-682.	1.1	5
87	MEASURING TOLERANCE TO HERBIVORY WITH NATURAL OR IMPOSED DAMAGE: A REPLY TO LEHTILÃ„. Evolution; International Journal of Organic Evolution, 2003, 57, 681.	1.1	11
88	Testing for Environmentally Induced Bias in Phenotypic Estimates of Natural Selection: Theory and Practice. American Naturalist, 2002, 160, 511-523.	1.0	219
89	COMPETITION AND TIME OF DAMAGE AFFECT THE PATTERN OF SELECTION ACTING ON PLANT DEFENSE AGAINST HERBIVORES. Ecology, 2002, 83, 1981-1990.	1.5	88
90	Coding Sequence Divergence Between Two Closely Related Plant Species: Arabidopsis thaliana and Brassica rapa ssp. pekinensis. Journal of Molecular Evolution, 2002, 54, 746-753.	0.8	54

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91	Coding Sequence Divergence Between Two Closely Related Plant Species: Arabidopsis thaliana and Brassica rapa ssp. pekinensis. Journal of Molecular Evolution, 2002, 54, 746-753.	0.8	69
92	Molecular Evolution of the Wound-Induced Serine Protease Inhibitor wip1 in Zea and Related Genera. Molecular Biology and Evolution, 2001, 18, 2092-2101.	3.5	61
93	Sequence Diversity in the Tetraploid Zea perennis and the Closely Related Diploid Z. diploperennis: Insights From Four Nuclear Loci. Genetics, 2001, 158, 401-412.	1.2	41
94	MEASURING TOLERANCE TO HERBIVORY: ACCURACY AND PRECISION OF ESTIMATES MADE USING NATURAL VERSUS IMPOSED DAMAGE. Evolution; International Journal of Organic Evolution, 2000, 54, 1024-1029.	1.1	111
95	Mechanisms of tolerance to herbivore damage:what do we know?. Evolutionary Ecology, 2000, 14, 523-536.	0.5	323
96	MEASURING TOLERANCE TO HERBIVORY: ACCURACY AND PRECISION OF ESTIMATES MADE USING NATURAL VERSUS IMPOSED DAMAGE. Evolution; International Journal of Organic Evolution, 2000, 54, 1024.	1.1	11
97	Are Tolerance, Avoidance, and Antibiosis Evolutionarily and Ecologically Equivalent Responses of Plants to Herbivores?. American Naturalist, 2000, 155, 128-138.	1.0	107
98	Genetic Constraints and Selection Acting on Tolerance to Herbivory in the Common Morning Glory Ipomoea purpurea. American Naturalist, 1999, 154, 700-716.	1.0	205
99	Patterns of evolutionary rate variation among genes of the anthocyanin biosynthetic pathway. Molecular Biology and Evolution, 1999, 16, 266-274.	3.5	199
100	Control of expression patterns of anthocyanin structural genes by two loci in the common morning glory.. Genes and Genetic Systems, 1998, 73, 105-110.	0.2	28
101	Response of Corn Grain Yield to Early and Late Killed Red Clover Green Manure and Subirrigation. Journal of Production Agriculture, 1998, 11, 112-121.	0.4	13