## Peter J Prentis

List of Publications by Year in descending order

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73 papers

3,545 citations

218592 26 h-index 149623 56 g-index

77 all docs

77 docs citations

times ranked

77

5763 citing authors

#	Article	IF	Citations
1	Venoms for all occasions: The functional toxin profiles of different anatomical regions in sea anemones are related to their ecological function. Molecular Ecology, 2022, 31, 866-883.	2.0	21
2	Horticultural innovation by viral-induced gene regulation of carotenogenesis. Horticulture Research, 2022, 9, .	2.9	4
3	Fruit Fly Larval Survival in Picked and Unpicked Tomato Fruit of Differing Ripeness and Associated Gene Expression Patterns. Insects, 2022, 13, 451.	1.0	2
4	A disulfide-stabilised helical hairpin fold in acrorhagin I: An emerging structural motif in peptide toxins. Journal of Structural Biology, 2021, 213, 107692.	1.3	10
5	Tentacle Morphological Variation Coincides with Differential Expression of Toxins in Sea Anemones. Toxins, 2021, 13, 452.	1.5	12
6	The Tentacular Spectacular: Evolution of Regeneration in Sea Anemones. Genes, 2021, 12, 1072.	1.0	5
7	Gene selection for studying frugivore-plant interactions: a review and an example using Queensland fruit fly in tomato. PeerJ, 2021, 9, e11762.	0.9	2
8	Effect of tomato fruit cultivar and ripening stage on Bactrocera tryoni (Froggatt) egg and larval survival. Journal of Applied Entomology, 2020, 144, 797-805.	0.8	4
9	Quantitative Genetic Assessment of Female Reproductive Traits in a Domesticated Pacific White Shrimp (Penaeus vannamei) Line in China. Scientific Reports, 2020, 10, 7840.	1.6	6
10	The effect of diet change and insulin dysregulation on the fecal microbiome of ponies. Journal of Experimental Biology, 2020, 223, .	0.8	4
11	Genetic parameters for growth and survival traits in a base population of Pacific white shrimp (Litopenaeus vannamei) developed from domesticated strains in China. Aquaculture, 2020, 523, 735148.	1.7	15
12	The genome of the sea anemone Actinia equina (L.): Meiotic toolkit genes and the question of sexual reproduction. Marine Genomics, 2020, 53, 100753.	0.4	17
13	Modulation of Placental Gene Expression in Small-for-Gestational-Age Infants. Genes, 2020, 11, 80.	1.0	10
14	The Rapid Regenerative Response of a Model Sea Anemone Species Exaiptasia pallida Is Characterised by Tissue Plasticity and Highly Coordinated Cell Communication. Marine Biotechnology, 2020, 22, 285-307.	1.1	12
15	Characterising Functional Venom Profiles of Anthozoans and Medusozoans within Their Ecological Context. Marine Drugs, 2020, 18, 202.	2.2	28
16	Structural and functional characterisation of a novel peptide from the Australian sea anemone Actinia tenebrosa. Toxicon, 2019, 168, 104-112.	0.8	11
17	The draft genome of <i>Actinia tenebrosa</i> reveals insights into toxin evolution. Ecology and Evolution, 2019, 9, 11314-11328.	0.8	28
18	Tools and Strategies for Long-Read Sequencing and De Novo Assembly of Plant Genomes. Trends in Plant Science, 2019, 24, 700-724.	4.3	80

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19	A process of convergent amplification and tissueâ€specific expression dominates the evolution of toxin and toxinâ€like genes in sea anemones. Molecular Ecology, 2019, 28, 2272-2289.	2.0	48
20	A molecular and morphometric assessment of the systematics of the Macropus complex clarifies the tempo and mode of kangaroo evolution. Zoological Journal of the Linnean Society, 2019, 186, 793-812.	1.0	23
21	A Versatile and Robust Serine Protease Inhibitor Scaffold from Actinia tenebrosa. Marine Drugs, 2019, 17, 701.	2.2	9
22	A non-disruptive method for obtaining DNA samples from sea anemones (Cnidaria: Anthozoa:) Tj ETQq0 0 0 rgB	T /Overloc 0.4	k 10 Tf 50 62
23	Plant-Mediated Female Transcriptomic Changes Post-Mating in a Tephritid Fruit Fly, Bactrocera tryoni. Genome Biology and Evolution, 2018, 10, 94-107.	1.1	12
24	Evidence for a Large Expansion and Subfunctionalization of Globin Genes in Sea Anemones. Genome Biology and Evolution, 2018, 10, 1892-1901.	1.1	8
25	A transcriptome-wide assessment of differentially expressed genes among two highly divergent, yet sympatric, lineages of the freshwater Atyid shrimp, Paratya australiensis. Hydrobiologia, 2018, 825, 189-196.	1.0	10
26	Sea Anemones: Quiet Achievers in the Field of Peptide Toxins. Toxins, 2018, 10, 36.	1.5	87
27	Insights into the phylogenetic and molecular evolutionary histories of <i>Fad</i> and <i>Elovl</i> gene families in Actiniaria. Ecology and Evolution, 2018, 8, 5323-5335.	0.8	17
28	Transcriptomic investigation of wound healing and regeneration in the cnidarian Calliactis polypus. Scientific Reports, 2017, 7, 41458.	1.6	35
29	Expression patterns of two carbonic anhydrase genes, Na+/K+-ATPase and V-type H+-ATPase, in the freshwater crayfish, Cherax quadricarinatus, exposed to low pH and high pH. Australian Journal of Zoology, 2017, 65, 50.	0.6	8
30	Occasional hybridization between a native and invasive <i>Senecio </i> species in Australia is unlikely to contribute to invasive success. PeerJ, 2017, 5, e3630.	0.9	1
31	Insights into the innate immunome of actiniarians using a comparative genomic approach. BMC Genomics, 2016, 17, 850.	1.2	42
32	Domestication and the storage starch biosynthesis pathway: signatures of selection from a whole sorghum genome sequencing strategy. Plant Biotechnology Journal, 2016, 14, 2240-2253.	4.1	38
33	Gene coevolution and regulation lock cyclic plant defence peptides to their targets. New Phytologist, 2016, 210, 717-730.	3.5	58
34	Comparative Analysis and Distribution of Omega-3 lcPUFA Biosynthesis Genes in Marine Molluscs. PLoS ONE, 2015, 10, e0136301.	1.1	29
35	Transcriptome analysis and characterisation of gill-expressed carbonic anhydrase and other key osmoregulatory genes in freshwater crayfish Cherax quadricarinatus. Data in Brief, 2015, 5, 187-193.	0.5	16
36	Expression and characterization of digestive enzyme genes from hepatopancreatic transcripts from redclaw crayfish ( <i>Cherax quadricarinatus</i> ). Aquaculture Nutrition, 2015, 21, 868-880.	1.1	14

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37	Comparative analysis of gill transcriptomes of two freshwater crayfish, Cherax cainii and C. destructor. Marine Genomics, 2015, 22, 11-13.	0.4	15
38	Analysis, characterisation and expression of gill-expressed carbonic anhydrase genes in the freshwater crayfish Cherax quadricarinatus. Gene, 2015, 564, 176-187.	1.0	26
39	Genetic Bottlenecks in Time and Space: Reconstructing Invasions from Contemporary and Historical Collections. PLoS ONE, 2014, 9, e106874.	1.1	16
40	The plasticity of NBS resistance genes in sorghum is driven by multiple evolutionary processes. BMC Plant Biology, 2014, 14, 253.	1.6	49
41	Genomic Resources Notes accepted 1 December 2013 - 31 January 2014. Molecular Ecology Resources, 2014, 14, 664-665.	2.2	5
42	The Anadara trapezia transcriptome: A resource for molluscan physiological genomics. Marine Genomics, 2014, 18, 113-115.	0.4	18
43	A transcriptome resource for the koala (Phascolarctos cinereus): insights into koala retrovirus transcription and sequence diversity. BMC Genomics, 2014, 15, 786.	1.2	49
44	Sexual selection in true fruit flies ( <scp>D</scp> iptera: <scp>T</scp> ephritidae): transcriptome and experimental evidences for phytochemicals increasing male competitive ability. Molecular Ecology, 2014, 23, 4645-4657.	2.0	35
45	Assembly and annotation of a non-model gastropod (Nerita melanotragus) transcriptome: a comparison of De novo assemblers. BMC Research Notes, 2014, 7, 488.	0.6	27
46	The koala immunological toolkit: sequence identification and comparison of key markers of the koala (Phascolarctos cinereus) immune response. Australian Journal of Zoology, 2014, 62, 195.	0.6	15
47	Design and development of a sampling platform to study long distance seed dispersal. , 2014, , .		0
48	Molecular characterisation and expression analysis of Interferon gamma in response to natural Chlamydia infection in the koala, Phascolarctos cinereus. Gene, 2013, 527, 570-577.	1.0	28
49	Characterisation of candidate nuclear genes for species delineation in the genus Cherax. Conservation Genetics Resources, 2013, 5, 331-333.	0.4	0
50	Whole-genome sequencing reveals untapped genetic potential in Africa's indigenous cereal crop sorghum. Nature Communications, 2013, 4, 2320.	5.8	405
51	Allelic variation at a single gene increases food value in a drought-tolerant staple cereal. Nature Communications, 2013, 4, 1483.	5.8	41
52	CONVERGENCE AND DIVERGENCE DURING THE ADAPTATION TO SIMILAR ENVIRONMENTS BY AN AUSTRALIAN GROUNDSEL. Evolution; International Journal of Organic Evolution, 2013, 67, 2515-2529.	1.1	66
53	Genomic evidence for the parallel evolution of coastal forms in the <i>Senecio lautus</i> complex. Molecular Ecology, 2013, 22, 2941-2952.	2.0	109
54	Understanding the genetic basis of invasiveness. Molecular Ecology, 2013, 22, 2366-2368.	2.0	15

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55	Isolation and characterisation of novel microsatellite and mitochondrial DNA markers for the Eastern Water Dragon (Physignathus lesueurii). Conservation Genetics Resources, 2012, 4, 113-116.	0.4	5
56	Anthropogenic landscape change promotes asymmetric dispersal and limits regional patch occupancy in a spatially structured bird population. Journal of Animal Ecology, 2012, 81, 940-952.	1.3	44
57	Lack of Low Frequency Variants Masks Patterns of Non-Neutral Evolution following Domestication. PLoS ONE, 2011, 6, e23041.	1.1	17
58	Diversification history and hybridisation of Dacrydium (Podocarpaceae) in remote Oceania. Australian Journal of Botany, 2011, 59, 262.	0.3	27
59	Allelic variation of the $\hat{l}^2$ -, $\hat{l}^3$ - and $\hat{l}'$ -kafirin genes in diverse Sorghum genotypes. Theoretical and Applied Genetics, 2010, 121, 1227-1237.	1.8	39
60	Massively parallel sequencing and analysis of expressed sequence tags in a successful invasive plant. Annals of Botany, 2010, 106, 1009-1017.	1.4	33
61	Characterization and multiplexing of ESTâ€SSR primers in <i>Cynodon</i> (Poaceae) species <sup>1</sup> . American Journal of Botany, 2010, 97, e99-e101.	0.8	21
62	Development and characterization of microsatellite loci for <i>Khaya senegalensis</i> (Meliaceae) <sup>1</sup> . American Journal of Botany, 2010, 97, e111-3.	0.8	15
63	A landscape genetics approach for quantifying the relative influence of historic and contemporary habitat heterogeneity on the genetic connectivity of a rainforest bird. Molecular Ecology, 2009, 18, 2945-2960.	2.0	70
64	Understanding invasion history: genetic structure and diversity of two globally invasive plants and implications for their management. Diversity and Distributions, 2009, 15, 822-830.	1.9	57
65	Something in the way you move: dispersal pathways affect invasion success. Trends in Ecology and Evolution, 2009, 24, 136-144.	4.2	680
66	Biogeographic concepts define invasion biology. Trends in Ecology and Evolution, 2009, 24, 586-586.	4.2	29
67	Fine-scale patterns of genetic variation indicate non-equilibrium gene frequency divergence in the stream lily, Helmholtzia glaberrima. Freshwater Biology, 2008, 53, 973-980.	1.2	10
68	Phylogenetic relationships in the monocot order Commelinales, with a focus on PhilydraceaeThis paper is one of a selection of papers published in the Special Issue on Systematics Research Botany, 2008, 86, 719-731.	0.5	22
69	Adaptive evolution in invasive species. Trends in Plant Science, 2008, 13, 288-294.	4.3	724
70	Micro-geographic landscape features demarcate seedling genetic structure in the stream lily, Helmholtzia glaberrima. Aquatic Botany, 2007, 87, 111-115.	0.8	3
71	Can hybridization cause local extinction: a case for demographic swamping of the Australian native <i>Senecio pinnatifolius</i> by the invasive <i>Senecio madagascariensis</i> New Phytologist, 2007, 176, 902-912.	3.5	68
72	Significance of post-germination buoyancy in Helmholtzia glaberrima and Philydrum lanuginosum (Philydraceae). Australian Journal of Botany, 2006, 54, 11.	0.3	8

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73	Genetic structuring of the stream lily Helmholtzia glaberrima (Philydraceae) within Toolona Creek, south-eastern Queensland. Australian Journal of Botany, 2004, 52, 201.	0.3	21