

Teresa J Crease

List of Publications by Year in descending order

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67

papers

4,910

citations

126907

33

h-index

102487

66

g-index

68

all docs

68

docs citations

68

times ranked

5351

citing authors

#	ARTICLE	IF	CITATIONS
1	The Ecoresponsive Genome of <i>Daphnia pulex</i> . <i>Science</i> , 2011, 331, 555-561.	12.6	1,086
2	The correlation between rDNA copy number and genome size in eukaryotes. <i>Genome</i> , 2003, 46, 48-50.	2.0	401
3	Phylogenetic relationships between parthenogens and their sexual relatives: the possible routes to parthenogenesis in animals. <i>Biological Journal of the Linnean Society</i> , 0, 79, 151-163.	1.6	369
4	Clonal Coexistence in <i>Daphnia pulex</i> (Leydig): Another Planktonic Paradox. <i>Science</i> , 1980, 207, 1363-1365.	12.6	237
5	Development of primers for the mitochondrial cytochrome c oxidase I gene in digenetic trematodes (Platyhelminthes) illustrates the challenge of barcoding parasitic helminths. <i>Molecular Ecology Resources</i> , 2009, 9, 75-82.	4.8	210
6	The complete sequence of the mitochondrial genome of <i>Daphnia pulex</i> (Cladocera: Crustacea). <i>Gene</i> , 1999, 233, 89-99.	2.2	193
7	Identification of two QTL influencing upper temperature tolerance in three rainbow trout (<i>Oncorhynchus mykiss</i>) half-sib families. <i>Heredity</i> , 1998, 80, 143-151.	2.6	160
8	The Functional Significance of Ribosomal (r)DNA Variation: Impacts on the Evolutionary Ecology of Organisms. <i>Annual Review of Ecology, Evolution, and Systematics</i> , 2005, 36, 219-242.	8.3	137
9	Clonal diversity in populations of <i>Daphnia pulex</i> reproducing by obligate parthenogenesis. <i>Heredity</i> , 1983, 51, 353-369.	2.6	121
10	Divergence thresholds and divergent biodiversity estimates: can metabarcoding reliably describe zooplankton communities?. <i>Ecology and Evolution</i> , 2015, 5, 2234-2251.	1.9	117
11	Phylogenetics and evolution of a circumarctic species complex (Cladocera: <i>Daphnia pulex</i>). <i>Biological Journal of the Linnean Society</i> , 1998, 65, 347-365.	1.6	111
12	Biogeography and systematics of Bangia (Bangiales, Rhodophyta) based on the Rubisco spacer, rbcL gene and 18S rRNA gene sequences and morphometric analyses. 1. North America. <i>Phycologia</i> , 1998, 37, 195-207.	1.4	92
13	Functional and ecological significance of rDNA intergenic spacer variation in a clonal organism under divergent selection for production rate. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2002, 269, 2373-2379.	2.6	86
14	Mitochondrial DNA variation in North American populations of <i>Daphnia obtusa</i> : continentalism or cryptic endemism?. <i>Molecular Ecology</i> , 2004, 13, 97-107.	3.9	79
15	Holarctic phylogeography of an asexual species complex – II. Allozymic variation and clonal structure in Arctic <i>Daphnia</i> . <i>Molecular Ecology</i> , 1999, 8, 1-13.	3.9	77
16	Probing the relationships of the brachiopod crustaceans. <i>Molecular Phylogenetics and Evolution</i> , 2006, 39, 491-502.	2.7	75
17	THE DISTRIBUTION OF LIFE-HISTORY VARIATION IN THE <i>DAPHNIA PULEX</i> COMPLEX. <i>Evolution; International Journal of Organic Evolution</i> , 1989, 43, 1724-1736.	2.3	71
18	POLYPHYLETIC ORIGINS OF ASEXUALITY IN <i>DAPHNIA PULEX</i> . II. MITOCHONDRIAL-DNA VARIATION. <i>Evolution; International Journal of Organic Evolution</i> , 1989, 43, 1016-1026.	2.3	70

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19	Transcontinental Phylogeography of the <i>Daphnia pulex</i> Species Complex. PLoS ONE, 2012, 7, e46620.	2.5	69
20	A Hierarchical Molecular Phylogeny within the Genus <i>Daphnia</i> . Molecular Phylogenetics and Evolution, 1995, 4, 395-407.	2.7	68
21	HOLARCTIC PHYLOGEOGRAPHY OF AN ASEXUAL SPECIES COMPLEX I. MITOCHONDRIAL DNA VARIATION IN ARCTIC <i>DAPHNIA</i> . Evolution; International Journal of Organic Evolution, 1999, 53, 777-792.	2.3	66
22	Speciation with gene flow and the genetics of habitat transitions. Molecular Ecology, 2012, 21, 1411-1422.	3.9	61
23	Allozyme and mtDNA variation in populations of the <i>Daphnia pulex</i> complex from both sides of the Rocky Mountains. Heredity, 1997, 79, 242-251.	2.6	53
24	The Unusually Long Small-Subunit Ribosomal RNA of the Crustacean, <i>Daphnia pulex</i> : Sequence and Predicted Secondary Structure. Journal of Molecular Evolution, 1998, 46, 307-313.	1.8	52
25	Phylogenetic evidence for a single long-lived clade of crustacean cyclic parthenogens and its implications for the evolution of sex. Proceedings of the Royal Society B: Biological Sciences, 1999, 266, 791-797.	2.6	51
26	Molecular Evolution of <i>Daphnia</i> Immunity Genes: Polymorphism in a Gram-Negative Binding Protein Gene and an α -2-Macroglobulin Gene. Journal of Molecular Evolution, 2004, 59, 498-506.	1.8	45
27	Molecular characterization of clonal population structure and biogeography of arctic apomictic <i>Daphnia</i> from Greenland and Iceland. Molecular Ecology, 1996, 5, 107-118.	3.9	40
28	Pokey, a New DNA Transposon in <i>Daphnia</i> (Cladocera: Crustacea). Journal of Molecular Evolution, 2002, 55, 664-673.	1.8	40
29	D- and L-lactate dehydrogenases during invertebrate evolution. BMC Evolutionary Biology, 2008, 8, 268.	3.2	40
30	THE MOLECULAR THROUGH ECOLOGICAL GENETICS OF ABNORMAL ABDOMEN IN DROSOPHILA MERCATORUM. I. BASIC GENETICS. Genetics, 1985, 111, 805-818.	2.9	40
31	Polyphyletic Origins of Asexuality in <i>Daphnia pulex</i> . II. Mitochondrial-DNA Variation. Evolution; International Journal of Organic Evolution, 1989, 43, 1016.	2.3	37
32	Holarctic Phylogeography of an Asexual Species Complex I. Mitochondrial DNA Variation in Arctic <i>Daphnia</i> . Evolution; International Journal of Organic Evolution, 1999, 53, 777.	2.3	35
33	Rates of Recombination in the Ribosomal DNA of Apomictically Propagated <i>Daphnia obtusa</i> Lines. Genetics, 2007, 175, 311-320.	2.9	35
34	Mitochondrial DNA diversity in the pea aphid <i>Acyrtosiphon pisum</i> . Genome, 1994, 37, 858-865.	2.0	34
35	Partial mitochondrial DNA sequence of the crustacean <i>Daphnia pulex</i> . Current Genetics, 1994, 25, 66-72.	1.7	32
36	Evolution of the Transposable Element Pokey in the Ribosomal DNA of Species in the Subgenus <i>Daphnia</i> (Crustacea: Cladocera). Molecular Biology and Evolution, 2004, 21, 1727-1739.	8.9	32

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37	The Behavior of a <i>Daphnia pulex</i> Transposable Element in Cyclically and Obligately Parthenogenetic Populations. <i>Journal of Molecular Evolution</i> , 2001, 53, 63-69.	1.8	27
38	The Association Between Breeding System and Transposable Element Dynamics in <i>Daphnia Pulex</i> . <i>Journal of Molecular Evolution</i> , 2008, 66, 643-654.	1.8	27
39	Copy number variation of ribosomal DNA and Pokey transposons in natural populations of <i>Daphnia</i> . <i>Mobile DNA</i> , 2012, 3, 4.	3.6	26
40	Identification of two QTL influencing upper temperature tolerance in three rainbow trout (<i>Oncorhynchus mykiss</i>) half-sib families. <i>Heredity</i> , 1998, 80, 143-151.	2.6	25
41	Sequence of the intergenic spacer between the 28S and 18S rRNA-encoding genes of the crustacean, <i>Daphnia pulex</i> . <i>Gene</i> , 1993, 134, 245-249.	2.2	24
42	Variation in transcriptional responses to copper exposure across <i>Daphnia pulex</i> lineages. <i>Aquatic Toxicology</i> , 2019, 210, 85-97.	4.0	23
43	Evolution of the nuclear ribosomal DNA intergenic spacer in four species of the <i>Daphnia pulex</i> complex. <i>BMC Genetics</i> , 2011, 12, 13.	2.7	22
44	Evolutionary factors affecting Lactate dehydrogenase A and B variation in the <i>Daphnia pulex</i> species complex. <i>BMC Evolutionary Biology</i> , 2011, 11, 212.	3.2	20
45	A test for the production of sexual pheromones by <i>Daphnia magna</i> (Crustacea: Cladocera). <i>Freshwater Biology</i> , 1983, 13, 491-496.	2.4	18
46	Heterosis in <i>Daphnia</i> : A Reassessment. <i>American Naturalist</i> , 1982, 119, 427-434.	2.1	17
47	Selection on the Structural Stability of a Ribosomal RNA Expansion Segment in <i>Daphnia obtusa</i> . <i>Molecular Biology and Evolution</i> , 2005, 22, 1309-1319.	8.9	16
48	Impact of ploidy level on the distribution of Pokey element insertions in the <i>Daphnia pulex</i> complex. <i>Mobile DNA</i> , 2014, 5, 1.	3.6	16
49	Alternative splicing is highly variable among <i>Daphnia pulex</i> lineages in response to acute copper exposure. <i>BMC Genomics</i> , 2020, 21, 433.	2.8	15
50	The Distribution of Life-History Variation in the <i>Daphnia pulex</i> Complex. <i>Evolution; International Journal of Organic Evolution</i> , 1989, 43, 1724.	2.3	13
51	In and out of the rRNA genes: characterization of Pokey elements in the sequenced <i>Daphnia</i> genome. <i>Mobile DNA</i> , 2013, 4, 20.	3.6	13
52	Genetic changes within an aphid clone: homogenization of rDNA intergenic spacers after insecticide selection. <i>Biological Journal of the Linnean Society</i> , 2003, 79, 101-105.	1.6	11
53	The effect of transposon <i>< i>Pokey</i></i> insertions on sequence variation in the 28S rRNA gene of <i>< i>Daphnia pulex</i></i> . <i>Genome</i> , 2008, 51, 988-1000.	2.0	11
54	Allozyme and mtDNA variation in populations of the <i>Daphnia pulex</i> complex from both sides of the Rocky Mountains. <i>Heredity</i> , 1997, 79, 242-251.	2.6	10

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55	Copy Number of the Transposon, Pokey, in rDNA Is Positively Correlated with rDNA Copy Number in <i>Daphnia obtusa</i> . PLoS ONE, 2014, 9, e114773.	2.5	8
56	Cloning and characterization of <i>Daphnia</i> mitochondrial DNA. Journal of Molecular Evolution, 1991, 33, 152-155.	1.8	7
57	Evolution of a transposon in <i>Daphnia</i> hybrid genomes. Mobile DNA, 2013, 4, 7.	3.6	6
58	Partial sequence of the mitochondrial genome of the crustacean <i>Daphnia pulex</i> . Current Genetics, 1997, 31, 48-54.	1.7	5
59	Evolution of Repeated Sequences in the Ribosomal DNA Intergenic Spacer of 32 Arthropod Species. Journal of Molecular Evolution, 2010, 70, 247-259.	1.8	5
60	Metal exposure causes rDNA copy number to fluctuate in mutation accumulation lines of <i>Daphnia pulex</i> . Aquatic Toxicology, 2020, 226, 105556.	4.0	5
61	Length Variation in 18S rRNA Expansion Segment 43/e4 of <i>Daphnia obtusa</i> : Ancient or Recurring Polymorphism?. Journal of Molecular Evolution, 2009, 69, 142-149.	1.8	4
62	Gene Expression Variation in Duplicate Lactate dehydrogenase Genes: Do Ecological Species Show Distinct Responses?. PLoS ONE, 2014, 9, e103964.	2.5	4
63	Detection of "Lost" Plasmids from <i>Escherichia coli</i> Using Excess Ampicillin. Analytical Biochemistry, 1996, 236, 181-182.	2.4	3
64	Bioinformatics for Biomonitoring: Species Detection and Diversity Estimates Across Next-Generation Sequencing Platforms. Advances in Ecological Research, 2018, , 1-32.	2.7	3
65	Distribution of the DNA transposon family, Pokey in the <i>Daphnia pulex</i> species complex. Mobile DNA, 2016, 7, 11.	3.6	2
66	A simple and effective chemiluminescent DNA fingerprinting technique using digoxigenin-labelled minisatellite probes. Canadian Journal of Zoology, 1997, 75, 817-820.	1.0	1
67	Population cycles can maintain foraging polymorphism. Proceedings of the Royal Society B: Biological Sciences, 1999, 266, 1277-1281.	2.6	1