John K Colbourne

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5729681/publications.pdf

Version: 2024-02-01

117 papers

9,959 citations

44069 48 h-index 95 g-index

126 all docs

126 docs citations

126 times ranked 10845 citing authors

#	Article	IF	CITATIONS
1	Refining the evolutionary time machine: An assessment of whole genome amplification using single historical <i>Daphnia</i> eggs. Molecular Ecology Resources, 2022, 22, 946-961.	4.8	3
2	Multigenerational Exposure to Nanoâ€TiO ₂ Induces Ageing as a Stress Response Mitigated by Environmental Interactions. Advanced NanoBiomed Research, 2021, 1, 2000083.	3.6	3
3	Extensive standing genetic variation from a small number of founders enables rapid adaptation in Daphnia. Nature Communications, 2021, 12, 4306.	12.8	27
4	Progress towards an OECD reporting framework for transcriptomics and metabolomics in regulatory toxicology. Regulatory Toxicology and Pharmacology, 2021, 125, 105020.	2.7	46
5	Vision of a near future: Bridging the human health–environment divide. Toward an integrated strategy to understand mechanisms across species for chemical safety assessment. Toxicology in Vitro, 2020, 62, 104692.	2.4	33
6	Centennial clonal stability of asexual <i>Daphnia</i> in Greenland lakes despite climate variability. Ecology and Evolution, 2020, 10, 14178-14188.	1.9	4
7	Spatio-temporal processes drive fine-scale genetic structure in an otherwise panmictic seabird population. Scientific Reports, 2020, 10, 20725.	3.3	2
8	Sex biased expression and co-expression networks in development, using the hymenopteran Nasonia vitripennis. PLoS Genetics, 2020, 16, e1008518.	3.5	11
9	A comprehensive epigenomic analysis of phenotypically distinguishable, genetically identical female and male Daphnia pulex. BMC Genomics, 2020, 21, 17.	2.8	25
10	Multigenerational Exposures of <i>Daphnia Magna</i> to Pristine and Aged Silver Nanoparticles: Epigenetic Changes and Phenotypical Ageing Related Effects. Small, 2020, 16, e2000301.	10.0	31
11	Improved Algal Toxicity Test System for Robust Omics-Driven Mode-of-Action Discovery in Chlamydomonas reinhardtii. Metabolites, 2019, 9, 94.	2.9	4
12	Transgenerational response to early spring warming in Daphnia. Scientific Reports, 2019, 9, 4449.	3.3	21
13	Dynamics of Cadmium Acclimation in <i>Daphnia pulex</i> : Linking Fitness Costs, Cross-Tolerance, and Hyper-Induction of Metallothionein. Environmental Science & Environmental	10.0	16
14	Review of and Recommendations for Monitoring Contaminants and their Effects in the San Francisco Bayâ^'Delta. San Francisco Estuary and Watershed Science, 2019, 17, .	0.4	3
15	Mixtures of Aluminum and Indium Induce More than Additive Phenotypic and Toxicogenomic Responses in <i>Daphnia magna</i> . Environmental Science & Envi	10.0	19
16	Invertebrate Model Species in AOP Development. , 2018, , 75-106.		3
17	The Toxicogenome of <i>Hyalella azteca</i> : A Model for Sediment Ecotoxicology and Evolutionary Toxicology. Environmental Science & Eamp; Technology, 2018, 52, 6009-6022.	10.0	79
18	Evolutionary transition from blood feeding to obligate nonbiting in a mosquito. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 1009-1014.	7.1	23

#	Article	IF	Citations
19	Early transcriptional response pathways in <i>Daphnia magna</i> are coordinated in networks of crustaceanâ€specific genes. Molecular Ecology, 2018, 27, 886-897.	3.9	38
20	Mitotic activity patterns and cytoskeletal changes throughout the progression of diapause developmental program in Daphnia. BMC Cell Biology, 2018, 19, 30.	3.0	33
21	How omics technologies can enhance chemical safety regulation: perspectives from academia, government, and industry. Environmental Toxicology and Chemistry, 2018, 37, 1252-1259.	4.3	12
22	Pattern of DNA Methylation in Daphnia: Evolutionary Perspective. Genome Biology and Evolution, 2018, 10, 1988-2007.	2.5	47
23	The transcriptomic and proteomic responses of Daphnia pulex to changes in temperature and food supply comprise environment-specific and clone-specific elements. BMC Genomics, 2018, 19, 376.	2.8	19
24	Arsenic Reduces Gene Expression Response to Changing Salinity in Killifish. Environmental Science & Expression Response to Changing Salinity in Killifish. Environmental Science & Expression Response to Changing Salinity in Killifish.	10.0	5
25	Scan, extract, wrap, compute—a 3D method to analyse morphological shape differences. PeerJ, 2018, 6, e4861.	2.0	12
26	The Role of Omics in the Application of Adverse Outcome Pathways for Chemical Risk Assessment. Toxicological Sciences, 2017, 158, 252-262.	3.1	161
27	Conserved Transcription Factors Steer Growth-Related Genomic Programs in Daphnia. Genome Biology and Evolution, 2017, 9, 1821-1842.	2.5	13
28	The Landscape of Extreme Genomic Variation in the Highly Adaptable Atlantic Killifish. Genome Biology and Evolution, 2017, 9, 659-676.	2.5	43
29	OGS2: genome re-annotation of the jewel wasp Nasonia vitripennis. BMC Genomics, 2016, 17, 678.	2.8	35
30	The genomic landscape of rapid repeated evolutionary adaptation to toxic pollution in wild fish. Science, 2016, 354, 1305-1308.	12.6	348
31	Thermal variation and factors influencing vertical migration behavior in Daphnia populations. Journal of Thermal Biology, 2016, 60, 70-78.	2.5	17
32	Daphnia magna transcriptome by RNA-Seq across 12 environmental stressors. Scientific Data, 2016, 3, 160030.	5.3	89
33	Male meiosis in Crustacea: synapsis, recombination, epigenetics and fertility in Daphnia magna. Chromosoma, 2016, 125, 769-787.	2.2	17
34	Evolutionary divergence of core and post-translational circadian clock genes in the pitcher-plant mosquito, Wyeomyia smithii. BMC Genomics, 2015, 16, 754.	2.8	12
35	Ionotropic Glutamate Receptors Mediate Inducible Defense in the Water Flea Daphnia pulex. PLoS ONE, 2015, 10, e0121324.	2.5	23
36	Novel Cadmium Responsive MicroRNAs in <i>Daphnia pulex</i> . Environmental Science & Environmental Scie	10.0	34

#	Article	IF	Citations
37	Synergistic interactions of biotic and abiotic environmental stressors on gene expression. Genome, 2015, 58, 99-109.	2.0	17
38	Differential transcriptomic responses of ancient and modern <i>Daphnia</i> genotypes to phosphorus supply. Molecular Ecology, 2015, 24, 123-135.	3.9	38
39	Potential for sexual conflict assessed via testosterone-mediated transcriptional changes in liver and muscle of a songbird. Journal of Experimental Biology, 2014, 217, 507-17.	1.7	28
40	Functional genomics of acclimation and adaptation in response to thermal stress in Daphnia. BMC Genomics, 2014, 15, 859.	2.8	68
41	Genomes of the rice pest brown planthopper and its endosymbionts reveal complex complementary contributions for host adaptation. Genome Biology, 2014, 15, 521.	8.8	404
42	Functional genomics of intraspecific variation in carbon and phosphorus kinetics in <i>Daphnia</i> Journal of Experimental Zoology, 2014, 321, 387-398.	1.2	24
43	A millennialâ€scale chronicle of evolutionary responses to cultural eutrophication in <i>Daphnia</i> . Ecology Letters, 2014, 17, 360-368.	6.4	178
44	An SNP-based second-generation genetic map of Daphnia magna and its application to QTL analysis of phenotypic traits. BMC Genomics, 2014, 15, 1033.	2.8	49
45	The Genomics of Cladoceran Physiology. , 2014, , 235-258.		2
46	Natural Selection Canalizes Expression Variation of Environmentally Induced Plasticity-Enabling Genes. Molecular Biology and Evolution, 2014, 31, 3002-3015.	8.9	48
47	Genomic data integration for ecological and evolutionary traits in non-model organisms. BMC Genomics, 2014, 15, 490.	2.8	36
48	Genome-Wide Transcription Profiles Reveal Genotype-Dependent Responses of Biological Pathways and Gene-Families in Daphnia Exposed to Single and Mixed Stressors. Environmental Science & Emp; Technology, 2014, 48, 3513-3522.	10.0	51
49	Transcriptome assembly and microarray construction for Enchytraeus crypticus, a model oligochaete to assess stress response mechanisms derived from soil conditions. BMC Genomics, 2014, 15, 302.	2.8	35
50	Molecular cloning of doublesex genes of four cladocera (water flea) species. BMC Genomics, 2013, 14, 239.	2.8	53
51	Why so many unknown genes? Partitioning orphans from a representative transcriptome of the lone star tick Amblyomma americanum. BMC Genomics, 2013, 14, 135.	2.8	42
52	Multiple origins of pyrethroid insecticide resistance across the species complex of a nontarget aquatic crustacean, <i>Hyalella azteca</i> List Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 16532-16537.	7.1	131
53	Transcription patterns of genes encoding four metallothionein homologs in Daphnia pulex exposed to copper and cadmium are time- and homolog-dependent. Aquatic Toxicology, 2013, 142-143, 422-430.	4.0	26
54	The evolutionary time machine: using dormant propagules to forecast how populations can adapt to changing environments. Trends in Ecology and Evolution, 2013, 28, 274-282.	8.7	123

#	Article	IF	CITATIONS
55	Evolutionary history of alpine and subalpine <i><scp>D</scp>aphnia</i> in western <scp>N</scp> orth <scp>A</scp> merica. Freshwater Biology, 2013, 58, 1512-1522.	2.4	13
56	A mutation in the receptor Methoprene-tolerant alters juvenile hormone response in insects and crustaceans. Nature Communications, 2013, 4, 1856.	12.8	100
57	Function and Evolution of DNA Methylation in Nasonia vitripennis. PLoS Genetics, 2013, 9, e1003872.	3.5	162
58	Fine-Scale Mapping of the Nasonia Genome to Chromosomes Using a High-Density Genotyping Microarray. G3: Genes, Genomes, Genetics, 2013, 3, 205-215.	1.8	33
59	Testosterone Affects Neural Gene Expression Differently in Male and Female Juncos: A Role for Hormones in Mediating Sexual Dimorphism and Conflict. PLoS ONE, 2013, 8, e61784.	2.5	52
60	Gut Pathology and Responses to the Microsporidium Nosema ceranae in the Honey Bee Apis mellifera. PLoS ONE, 2012, 7, e37017.	2.5	204
61	Functional characterization of four metallothionein genes in Daphnia pulex exposed to environmental stressors. Aquatic Toxicology, 2012, 110-111, 54-65.	4.0	35
62	De novo transcriptome sequencing in a songbird, the dark-eyed junco (Junco hyemalis): genomic tools for an ecological model system. BMC Genomics, 2012, 13, 305.	2.8	35
63	Identification of maternally-loaded RNA transcripts in unfertilized eggs of Tribolium castaneum. BMC Genomics, 2012, 13, 671.	2.8	15
64	Identification of Pathways, Gene Networks, and Paralogous Gene Families in Daphnia pulex Responding to Exposure to the Toxic Cyanobacterium Microcystis aeruginosa. Environmental Science & Eamp; Technology, 2012, 46, 8448-8457.	10.0	52
65	The evolution of salinity tolerance in <i>Daphnia</i> : a functional genomics approach. Ecology Letters, 2012, 15, 794-802.	6.4	88
66	The genome of the green anole lizard and a comparative analysis with birds and mammals. Nature, 2011, 477, 587-591.	27.8	575
67	How do consumers deal with stoichiometric constraints? Lessons from functional genomics using Daphnia pulex. Molecular Ecology, 2011, 20, 2341-2352.	3.9	51
68	The Ecoresponsive Genome of <i>Daphnia pulex</i> . Science, 2011, 331, 555-561.	12.6	1,086
69	Will genetic adaptation of natural populations to chemical pollution result in lower or higher tolerance to future climate change?. Integrated Environmental Assessment and Management, 2011, 7, 141-143.	2.9	7
70	Genomics of Environmentally Induced Phenotypes in 2 Extremely Plastic Arthropods. Journal of Heredity, 2011, 102, 512-525.	2.4	41
71	Targeted bisulfite sequencing by solution hybrid selection and massively parallel sequencing. Nucleic Acids Research, 2011, 39, e127-e127.	14.5	61
72	Dual Labeled Expression-Tiling Microarray Protocol for Empirical Annotation of Genome Sequences. CGB Technical Report, 2011, 2011, .	1.5	3

#	Article	IF	Citations
73	Dual Labeled Expression Microarray Protocol for High-Throughput Genomic Investigations. CGB Technical Report, 2011, 2011, .	1.5	8
74	Conservation, loss, and redeployment of Wnt ligands in protostomes: implications for understanding the evolution of segment formation. BMC Evolutionary Biology, 2010, 10, 374.	3.2	153
75	Predator-induced defences in Daphnia pulex: Selection and evaluation of internal reference genes for gene expression studies with real-time PCR. BMC Molecular Biology, 2010, 11, 50.	3.0	52
76	AphidBase: a centralized bioinformatic resource for annotation of the pea aphid genome. Insect Molecular Biology, 2010, 19, 5-12.	2.0	108
77	Characteristics of the genome of <i>Arsenophonus nasoniae</i> , sonâ€killer bacterium of the wasp <i>Nasonia</i> . Insect Molecular Biology, 2010, 19, 75-89.	2.0	94
78	The draft genome sequence of <i>Arsenophonus nasoniae</i> , sonâ€killer bacterium of <i>Nasonia vitripennis</i> , reveals genes associated with virulence and symbiosis. Insect Molecular Biology, 2010, 19, 59-73.	2.0	46
79	Functional and Evolutionary Insights from the Genomes of Three Parasitoid <i>Nasonia</i> Species. Science, 2010, 327, 343-348.	12.6	808
80	ESTPiper – a web-based analysis pipeline for expressed sequence tags. BMC Genomics, 2009, 10, 174.	2.8	17
81	The components of the Daphnia pulex immune system as revealed by complete genome sequencing. BMC Genomics, 2009, 10, 175.	2.8	93
82	Sequencing and de novo analysis of a coral larval transcriptome using 454 GSFlx. BMC Genomics, 2009, 10, 219.	2.8	405
83	The scale of divergence: A phylogenetic appraisal of intercontinental allopatric speciation in a passively dispersed freshwater zooplankton genus. Molecular Phylogenetics and Evolution, 2009, 50, 423-436.	2.7	146
84	Ecological genomics in Daphnia: stress responses and environmental sex determination. Heredity, 2008, 100, 184-190.	2.6	95
85	Daphnia as an emerging model for toxicological genomics. Advances in Experimental Biology, 2008, 2, 165-328.	0.1	91
86	A machine-learning approach to combined evidence validation of genome assemblies. Bioinformatics, 2008, 24, 744-750.	4.1	23
87	Recent advances in crustacean genomics. Integrative and Comparative Biology, 2008, 48, 852-868.	2.0	54
88	Sampling Daphnia's expressed genes: preservation, expansion and invention of crustacean genes with reference to insect genomes. BMC Genomics, 2007, 8, 217.	2.8	25
89	Profiling sex-biased gene expression during parthenogenetic reproduction in Daphnia pulex. BMC Genomics, 2007, 8, 464.	2.8	51
90	Gene response profiles for Daphnia pulex exposed to the environmental stressor cadmium reveals novel crustacean metallothioneins. BMC Genomics, 2007, 8, 477.	2.8	108

#	Article	IF	CITATIONS
91	A microsatellite-based genetic linkage map of the waterflea, Daphnia pulex: On the prospect of crustacean genomics. Genomics, 2006, 88, 415-430.	2.9	76
92	The systematics of AustralianDaphniaandDaphniopsis(Crustacea: Cladocera): a shared phylogenetic history transformed by habitat-specific rates of evolution. Biological Journal of the Linnean Society, 2006, 89, 469-488.	1.6	41
93	Selection of Surrogate Animal Species for Comparative Toxicogenomics. , 2006, , 33-75.		5
94	EVOLUTIONARY HISTORY OF CONTAGIOUS ASEXUALITY IN DAPHNIA PULEX. Evolution; International Journal of Organic Evolution, 2005, 59, 800-813.	2.3	102
95	wFleaBase: the Daphnia genome database. BMC Bioinformatics, 2005, 6, 45.	2.6	125
96	EVOLUTIONARY HISTORY OF CONTAGIOUS ASEXUALITY IN DAPHNIA PULEX. Evolution; International Journal of Organic Evolution, 2005, 59, 800.	2.3	15
97	Evolutionary history of contagious asexuality in Daphnia pulex. Evolution; International Journal of Organic Evolution, 2005, 59, 800-13.	2.3	43
98	Five hundred and twenty-eight microsatellite markers for ecological genomic investigations using Daphnia. Molecular Ecology Notes, 2004, 4, 485-490.	1.7	58
99	Molecular Evolution of Daphnia Immunity Genes: Polymorphism in a Gram-Negative Binding Protein Gene and an ?-2-Macroglobulin Gene. Journal of Molecular Evolution, 2004, 59, 498-506.	1.8	45
100	Reconstruction of Centuries-old Daphnia Communities in a Lake Recovering from Acidification and Metal Contamination. Ambio, 2003, 32, 214-218.	5. 5	34
101	Reconstruction of Centuries-old Daphnia Communities in a Lake Recovering from Acidification and Metal Contamination. Ambio, 2003, 32, 214.	5.5	5
102	ACCELERATED MOLECULAR EVOLUTION IN HALOPHILIC CRUSTACEANS. Evolution; International Journal of Organic Evolution, 2002, 56, 909-926.	2.3	101
103	Rapid, local adaptation of zooplankton behavior to changes in predation pressure in the absence of neutral genetic changes. Proceedings of the National Academy of Sciences of the United States of America, 2001, 98, 6256-6260.	7.1	373
104	Reconstructing the history of intercontinental dispersal in <i>Daphnia lumholtzi</i> by use of genetic markers. Limnology and Oceanography, 2000, 45, 1414-1419.	3.1	49
105	The Quantitative and Molecular Genetic Architecture of a Subdivided Species. Evolution; International Journal of Organic Evolution, 1999, 53, 100.	2.3	102
106	Holarctic Phylogeography of an Asexual Species Complex I. Mitochondrial DNA Variation in Arctic Daphnia. Evolution; International Journal of Organic Evolution, 1999, 53, 777.	2.3	35
107	THE QUANTITATIVE AND MOLECULAR GENETIC ARCHITECTURE OF A SUBDIVIDED SPECIES. Evolution; International Journal of Organic Evolution, 1999, 53, 100-110.	2.3	192
108	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

#	Article	IF	CITATIONS
109	The Unusually Long Small-Subunit Ribosomal RNA of the Crustacean, Daphnia pulex: Sequence and Predicted Secondary Structure. Journal of Molecular Evolution, 1998, 46, 307-313.	1.8	52
110	Phylogenetics and evolution of a circumarctic species complex (Cladocera: Daphnia pulex). Biological Journal of the Linnean Society, 1998, 65, 347-365.	1.6	111
111	Evolution of Enterohemorrhagic <i>Escherichia coli</i> Hemolysin Plasmids and the Locus for Enterocyte Effacement in Shiga Toxin-Producing <i>E. coli</i> Infection and Immunity, 1998, 66, 2553-2561.	2.2	88
112	Long–term genetic shifts in a microcrustacean egg bank associated with anthropogenic changes in the Lake Constance ecosystem. Proceedings of the Royal Society B: Biological Sciences, 1997, 264, 1613-1618.	2.6	129
113	The systematics of North American Daphnia (Crustacea: Anomopoda): a molecular phylogenetic approach. Philosophical Transactions of the Royal Society B: Biological Sciences, 1996, 351, 349-360.	4.0	185
114	DNA fingerprinting of bluegill sunfish (Lepomis macrochirus) using (GT)n microsatellites and its potential for assessment of mating success. Canadian Journal of Fisheries and Aquatic Sciences, 1996, 53, 342-349.	1.4	58
115	Phylogenetics and Evolution of theDaphnia longispinaGroup (Crustacea) Based on 12S rDNA Sequence and Allozyme Variation. Molecular Phylogenetics and Evolution, 1996, 5, 495-510.	2.7	144
116	DNA fingerprinting of bluegill sunfish (Lepomis macrochirus) using (GT) _n microsatellites and its potential for assessment of mating success. Canadian Journal of Fisheries and Aquatic Sciences, 1996, 53, 342-349.	1.4	24
117	A Global Grid for Analysis of Arthropod Evolution. , 0, , .		7