

Jeffrey M. Good

List of Publications by Year in descending order

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Version: 2024-02-01

66
papers

12,626
citations

109321

35
h-index

106344

65
g-index

78
all docs

78
docs citations

78
times ranked

13721
citing authors

#	ARTICLE	IF	CITATIONS
1	A Draft Sequence of the Neandertal Genome. <i>Science</i> , 2010, 328, 710-722.	12.6	3,588
2	Genetic history of an archaic hominin group from Denisova Cave in Siberia. <i>Nature</i> , 2010, 468, 1053-1060.	27.8	1,537
3	Harnessing the power of RADseq for ecological and evolutionary genomics. <i>Nature Reviews Genetics</i> , 2016, 17, 81-92.	16.3	1,169
4	The complete mitochondrial DNA genome of an unknown hominin from southern Siberia. <i>Nature</i> , 2010, 464, 894-897.	27.8	659
5	A Complete Neandertal Mitochondrial Genome Sequence Determined by High-Throughput Sequencing. <i>Cell</i> , 2008, 134, 416-426.	28.9	503
6	Targeted Retrieval and Analysis of Five Neandertal mtDNA Genomes. <i>Science</i> , 2009, 325, 318-321.	12.6	456
7	The bonobo genome compared with the chimpanzee and human genomes. <i>Nature</i> , 2012, 486, 527-531.	27.8	445
8	Rabbit genome analysis reveals a polygenic basis for phenotypic change during domestication. <i>Science</i> , 2014, 345, 1074-1079.	12.6	343
9	Unlocking the vault: next-generation museum population genomics. <i>Molecular Ecology</i> , 2013, 22, 6018-6032.	3.9	329
10	Targeted capture in evolutionary and ecological genomics. <i>Molecular Ecology</i> , 2016, 25, 185-202.	3.9	295
11	Targeted Investigation of the Neandertal Genome by Array-Based Sequence Capture. <i>Science</i> , 2010, 328, 723-725.	12.6	255
12	Transcriptome-based exon capture enables highly cost-effective comparative genomic data collection at moderate evolutionary scales. <i>BMC Genomics</i> , 2012, 13, 403.	2.8	253
13	Adaptive introgression underlies polymorphic seasonal camouflage in snowshoe hares. <i>Science</i> , 2018, 360, 1355-1358.	12.6	234
14	INVESTIGATING THE EVOLUTIONARY HISTORY OF THE PACIFIC NORTHWEST MESIC FOREST ECOSYSTEM: HYPOTHESIS TESTING WITHIN A COMPARATIVE PHYLOGEOGRAPHIC FRAMEWORK. <i>Evolution; International Journal of Organic Evolution</i> , 2005, 59, 1639-1652.	2.3	183
15	Ancient hybridization and mitochondrial capture between two species of chipmunks. <i>Molecular Ecology</i> , 2008, 17, 1313-1327.	3.9	162
16	A Complex Genetic Basis to X-Linked Hybrid Male Sterility Between Two Species of House Mice. <i>Genetics</i> , 2008, 179, 2213-2228.	2.9	143
17	ASYMMETRY AND POLYMORPHISM OF HYBRID MALE STERILITY DURING THE EARLY STAGES OF SPECIATION IN HOUSE MICE. <i>Evolution; International Journal of Organic Evolution</i> , 2007, 62, 071115145922007-???	2.3	139
18	Widespread Over-Expression of the X Chromosome in Sterile F1 Hybrid Mice. <i>PLoS Genetics</i> , 2010, 6, e1001148.	3.5	111

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19	Function and underlying mechanisms of seasonal colour moulting in mammals and birds: what keeps them changing in a warming world?. <i>Biological Reviews</i> , 2018, 93, 1478-1498.	10.4	109
20	Rates of Protein Evolution Are Positively Correlated with Developmental Timing of Expression During Mouse Spermatogenesis. <i>Molecular Biology and Evolution</i> , 2005, 22, 1044-1052.	8.9	94
21	Winter color polymorphisms identify global hot spots for evolutionary rescue from climate change. <i>Science</i> , 2018, 359, 1033-1036.	12.6	91
22	Negligible nuclear introgression despite complete mitochondrial capture between two species of chipmunks. <i>Evolution; International Journal of Organic Evolution</i> , 2015, 69, 1961-1972.	2.3	88
23	Meiotic Sex Chromosome Inactivation Is Disrupted in Sterile Hybrid Male House Mice. <i>Genetics</i> , 2013, 193, 819-828.	2.9	86
24	The Genomic Architecture of Population Divergence between Subspecies of the European Rabbit. <i>PLoS Genetics</i> , 2014, 10, e1003519.	3.5	82
25	Temporal genomic contrasts reveal rapid evolutionary responses in an alpine mammal during recent climate change. <i>PLoS Genetics</i> , 2019, 15, e1008119.	3.5	70
26	The genomic basis of environmental adaptation in house mice. <i>PLoS Genetics</i> , 2018, 14, e1007672.	3.5	65
27	The composite regulatory basis of the large X-effect in mouse speciation. <i>Molecular Biology and Evolution</i> , 2017, 34, msw243.	8.9	59
28	Rapid neo-sex chromosome evolution and incipient speciation in a major forest pest. <i>Nature Communications</i> , 2017, 8, 1593.	12.8	59
29	Phylogeography of the red-tailed chipmunk (<i>Tamias ruficaudus</i>), a northern Rocky Mountain endemic. <i>Molecular Ecology</i> , 2001, 10, 2683-2695.	3.9	57
30	Bonobos Fall within the Genomic Variation of Chimpanzees. <i>PLoS ONE</i> , 2011, 6, e21605.	2.5	57
31	The Evolution of Polymorphic Hybrid Incompatibilities in House Mice. <i>Genetics</i> , 2018, 209, 845-859.	2.9	50
32	Adaptive Evolution of Proteins Secreted during Sperm Maturation: An Analysis of the Mouse Epididymal Transcriptome. <i>Molecular Biology and Evolution</i> , 2008, 25, 383-392.	8.9	49
33	Phylogenomic Insights into Mouse Evolution Using a Pseudoreference Approach. <i>Genome Biology and Evolution</i> , 2017, 9, 726-739.	2.5	47
34	Spermatogenesis and the Evolution of Mammalian Sex Chromosomes. <i>Trends in Genetics</i> , 2018, 34, 722-732.	6.7	47
35	The Legacy of Recurrent Introgression during the Radiation of Hares. <i>Systematic Biology</i> , 2021, 70, 593-607.	5.6	47
36	The Contribution of the Y Chromosome to Hybrid Male Sterility in House Mice. <i>Genetics</i> , 2012, 191, 1271-1281.	2.9	45

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37	Human Adaptive Evolution at Myostatin (GDF8), a Regulator of Muscle Growth. <i>American Journal of Human Genetics</i> , 2006, 79, 1089-1097.	6.2	41
38	Comparative Population Genomics of the Ejaculate in Humans and the Great Apes. <i>Molecular Biology and Evolution</i> , 2013, 30, 964-976.	8.9	40
39	A high-throughput method for unbiased quantitation and categorization of nuclear morphology. <i>Biology of Reproduction</i> , 2019, 100, 1250-1260.	2.7	38
40	Parental origin growth effects and the evolution of hybrid inviability in dwarf hamsters. <i>Evolution; International Journal of Organic Evolution</i> , 2014, 68, 3134-3148.	2.3	37
41	Genomic imprinting, disrupted placental expression, and speciation. <i>Evolution; International Journal of Organic Evolution</i> , 2016, 70, 2690-2703.	2.3	32
42	Contrasting Levels of Molecular Evolution on the Mouse X Chromosome. <i>Genetics</i> , 2016, 203, 1841-1857.	2.9	32
43	The Origin and Spread of Locally Adaptive Seasonal Camouflage in Snowshoe Hares. <i>American Naturalist</i> , 2020, 196, 316-332.	2.1	29
44	Molecular Correlates of Genes Exhibiting RNAi Phenotypes in <i>Caenorhabditis elegans</i> . <i>Genome Research</i> , 2003, 13, 2651-2657.	5.5	28
45	The transcriptional landscape of seasonal coat colour moult in the snowshoe hare. <i>Molecular Ecology</i> , 2017, 26, 4173-4185.	3.9	27
46	Whole exome sequencing of wild-derived inbred strains of mice improves power to link phenotype and genotype. <i>Mammalian Genome</i> , 2017, 28, 416-425.	2.2	25
47	The population genetics of crypsis in vertebrates: recent insights from mice, hares, and lizards. <i>Heredity</i> , 2020, 124, 1-14.	2.6	24
48	Adaptive Protein Evolution and Regulatory Divergence in <i>Drosophila</i> . <i>Molecular Biology and Evolution</i> , 2006, 23, 1101-1103.	8.9	23
49	Extraordinary Sequence Divergence at <i>Tsga8</i> , an X-linked Gene Involved in Mouse Spermiogenesis. <i>Molecular Biology and Evolution</i> , 2011, 28, 1675-1686.	8.9	22
50	Diversification, Introgression, and Rampant Cytonuclear Discordance in Rocky Mountains Chipmunks (<i>Sciuridae: Tamias</i>). <i>Systematic Biology</i> , 2021, 70, 908-921.	5.6	20
51	Convergent evolution of seasonal camouflage in response to reduced snow cover across the snowshoe hare range*. <i>Evolution; International Journal of Organic Evolution</i> , 2020, 74, 2033-2045.	2.3	19
52	Transposable Element Orientation Bias in the <i>Drosophila melanogaster</i> Genome. <i>Journal of Molecular Evolution</i> , 2005, 61, 733-741.	1.8	18
53	Molecular Evolution across Mouse Spermatogenesis. <i>Molecular Biology and Evolution</i> , 2022, 39, .	8.9	18
54	Multiple paternity in wild-caught <i>Drosophila mojavensis</i> . <i>Molecular Ecology</i> , 2006, 15, 2253-2260.	3.9	16

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55	Transcriptomic regulation of seasonal coat color change in hares. <i>Ecology and Evolution</i> , 2020, 10, 1180-1192.	1.9	16
56	On Characterizing Adaptive Events Unique to Modern Humans. <i>Genome Biology and Evolution</i> , 2011, 3, 791-798.	2.5	15
57	The Conflict within and the Escalating War between the Sex Chromosomes. <i>PLoS Genetics</i> , 2012, 8, e1002955.	3.5	14
58	Unraveling patterns of disrupted gene expression across a complex tissue. <i>Evolution; International Journal of Organic Evolution</i> , 2022, 76, 275-291.	2.3	14
59	An Annotated Draft Genome of the Mountain Hare (<i>Lepus timidus</i>). <i>Genome Biology and Evolution</i> , 2020, 12, 3656-3662.	2.5	13
60	Comparative Phylogenomic Assessment of Mitochondrial Introgression among Several Species of Chipmunks (<i>TAMIAS</i>). <i>Genome Biology and Evolution</i> , 2016, 9, evw254.	2.5	12
61	X chromosome-dependent disruption of placental regulatory networks in hybrid dwarf hamsters. <i>Genetics</i> , 2021, 218, .	2.9	10
62	Stage-specific disruption of X chromosome expression during spermatogenesis in sterile house mouse hybrids. <i>G3: Genes, Genomes, Genetics</i> , 2022, 12, .	1.8	8
63	Automated Nuclear Cartography Reveals Conserved Sperm Chromosome Territory Localization across 2 Million Years of Mouse Evolution. <i>Genes</i> , 2019, 10, 109.	2.4	7
64	The genomic basis of high-elevation adaptation in wild house mice (<i>Mus musculus domesticus</i>) from South America. <i>Genetics</i> , 2022, 220, .	2.9	7
65	The Evolution of Widespread Recombination Suppression on the Dwarf Hamster (<i>Phodopus</i>) X Chromosome. <i>Genome Biology and Evolution</i> , 2022, 14, .	2.5	2
66	Genomic resolution of cryptic species diversity in chipmunks. <i>Evolution; International Journal of Organic Evolution</i> , 2022, 76, 2004-2019.	2.3	2