Michael W Nachman

List of Publications by Year in descending order

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49 papers

6,594 citations

30 h-index 49 g-index

53 all docs 53 docs citations

53 times ranked 8864 citing authors

#	Article	IF	CITATIONS
1	Estimate of the Mutation Rate per Nucleotide in Humans. Genetics, 2000, 156, 297-304.	2.9	1,023
2	Genomics and the origin of species. Nature Reviews Genetics, 2014, 15, 176-192.	16.3	850
3	The genetic basis of adaptive melanism in pocket mice. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 5268-5273.	7.1	462
4	Subspecific origin and haplotype diversity in the laboratory mouse. Nature Genetics, 2011, 43, 648-655.	21.4	439
5	Recombination rate variation and speciation: theoretical predictions and empirical results from rabbits and mice. Philosophical Transactions of the Royal Society B: Biological Sciences, 2012, 367, 409-421.	4.0	339
6	Adaptive Introgression of Anticoagulant Rodent Poison Resistance by Hybridization between Old World Mice. Current Biology, 2011, 21, 1296-1301.	3.9	282
7	Genome-wide patterns of gene flow across a house mouse hybrid zone. Genome Research, 2008, 18, 67-76.	5.5	235
8	Variation in recombination rate across the genome: evidence and implications. Current Opinion in Genetics and Development, 2002, 12, 657-663.	3.3	214
9	Natural history collections as windows on evolutionary processes. Molecular Ecology, 2016, 25, 864-881.	3.9	199
10	DNA Variability and Recombination Rates at X-Linked Loci in Humans. Genetics, 1998, 150, 1133-1141.	2.9	194
11	Transmission modes of the mammalian gut microbiota. Science, 2018, 362, 453-457.	12.6	189
12	Gene Regulation and Speciation. Trends in Genetics, 2017, 33, 68-80.	6.7	149
13	Insights into mammalian biology from the wild house mouse Mus musculus. ELife, 2015, 4, .	6.0	134
14	Dispersal limitation promotes the diversification of the mammalian gut microbiota. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 13768-13773.	7.1	121
15	Morphological and population genomic evidence that human faces have evolved to signal individual identity. Nature Communications, 2014, 5, 4800.	12.8	117
16	Microsatellite Variation and Recombination Rate in the Human Genome. Genetics, 2000, 156, 1285-1298.	2.9	116
17	Linkage Disequilibrium in Wild Mice. PLoS Genetics, 2007, 3, e144.	3.5	108
18	Gene regulation and speciation in house mice. Genome Research, 2016, 26, 451-461.	5. 5	104

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19	Deleterious mutations in animal mitochondrial DNA. Genetica, 1998, 102/103, 61-69.	1.1	102
20	Dichotomy of single-nucleotide polymorphism haplotypes in olfactory receptor genes and pseudogenes. Nature Genetics, 2000, 26, 221-224.	21.4	92
21	Recombination and Speciation: Loci Near Centromeres Are More Differentiated Than Loci Near Telomeres Between Subspecies of the European Rabbit (<i>Oryctolagus cuniculus</i>). Genetics, 2009, 181, 593-606.	2.9	92
22	Spatial Heterogeneity of Gut Microbial Composition along the Gastrointestinal Tract in Natural Populations of House Mice. PLoS ONE, 2016, 11, e0163720.	2.5	84
23	The Genomic Architecture of Population Divergence between Subspecies of the European Rabbit. PLoS Genetics, 2014, 10, e1003519.	3.5	82
24	Altitudinal variation of the gut microbiota in wild house mice. Molecular Ecology, 2019, 28, 2378-2390.	3.9	77
25	Host genetic determinants of the gut microbiota of wild mice. Molecular Ecology, 2019, 28, 3197-3207.	3.9	76
26	Contrasting Evolutionary Histories of Two Introns of the Duchenne Muscular Dystrophy Gene, Dmd, in Humans. Genetics, 2000, 155, 1855-1864.	2.9	73
27	The genomic basis of environmental adaptation in house mice. PLoS Genetics, 2018, 14, e1007672.	3.5	65
28	Chitinase genes (<i>CHIA</i> s) provide genomic footprints of a post-Cretaceous dietary radiation in placental mammals. Science Advances, 2018, 4, eaar6478.	10.3	55
29	Gene regulation underlies environmental adaptation in house mice. Genome Research, 2018, 28, 1636-1645.	5.5	51
30	Selection on Coding and Regulatory Variation Maintains Individuality in Major Urinary Protein Scent Marks in Wild Mice. PLoS Genetics, 2016, 12, e1005891.	3.5	46
31	The genetic basis of adaptation: lessons from concealing coloration in pocket mice. Genetica, 2005, 123, 125-136.	1.1	33
32	Genome-Wide Patterns of Differentiation Among House Mouse Subspecies. Genetics, 2014, 198, 283-297.	2.9	33
33	Genomics and museum specimens. Molecular Ecology, 2013, 22, 5966-5968.	3.9	29
34	The gut microbiota and Bergmann's rule in wild house mice. Molecular Ecology, 2020, 29, 2300-2311.	3.9	28
35	Haldane and the first estimates of the human mutation rate. Journal of Genetics, 2004, 83, 231-233.	0.7	27
36	Experimental Evidence for Adaptation to Species-Specific Gut Microbiota in House Mice. MSphere, 2019, 4, .	2.9	27

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37	Isolation and characterization of Neisseria musculi sp. nov., from the wild house mouse. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 3585-3593.	1.7	27
38	The genomics of rapid climatic adaptation and parallel evolution in North American house mice. PLoS Genetics, 2021, 17, e1009495.	3.5	26
39	Genomic islands of differentiation in two songbird species reveal candidate genes for hybrid female sterility. Molecular Ecology, 2018, 27, 949-958.	3.9	25
40	The population genetics of crypsis in vertebrates: recent insights from mice, hares, and lizards. Heredity, 2020, 124, 1-14.	2.6	24
41	Gene expression plasticity and desert adaptation in house mice*. Evolution; International Journal of Organic Evolution, 2021, 75, 1477-1491.	2.3	23
42	Deleterious Mutations at the Mitochondrial ND3 Gene in South American Marsh Rats (Holochilus). Genetics, 1998, 150, 359-368.	2.9	23
43	The Contribution of Genetic and Environmental Effects to Bergmann's Rule and Allen's Rule in House Mice. American Naturalist, 2022, 199, 691-704.	2.1	20
44	The Genetics of Adaptive Coat Color in Gophers: Coding Variation at Mc1r Is Not Responsible for Dorsal Color Differences. Journal of Heredity, 2007, 98, 567-574.	2.4	17
45	Nucleotide Variation at Msn and Alas2, Two Genes Flanking the Centromere of the X Chromosome in Humans. Genetics, 2004, 167, 423-437.	2.9	15
46	Speciation and reduced hybrid female fertility in house mice. Evolution; International Journal of Organic Evolution, 2015, 69, 2468-2481.	2.3	15
47	Gene Expression Networks Across Multiple Tissues Are Associated with Rates of Molecular Evolution in Wild House Mice. Genes, 2019, 10, 225.	2.4	12
48	Bidirectional Introgression between <i>Mus musculus domesticus</i> and <i>Mus spretus</i> . Genome Biology and Evolution, 2022, 14, .	2.5	11
49	The genomic basis of high-elevation adaptation in wild house mice (<i>Mus musculus domesticus</i>) from South America. Genetics, 2022, 220, .	2.9	7