

# Mary J O'connell

## List of Publications by Year in descending order

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Version: 2024-02-01

37  
papers

2,501  
citations

471509

17  
h-index

302126

39  
g-index

40  
all docs

40  
docs citations

40  
times ranked

4378  
citing authors

#	ARTICLE	IF	CITATIONS
1	Population Genomics Reveal Recent Speciation and Rapid Evolutionary Adaptation in Polar Bears. <i>Cell</i> , 2014, 157, 785-794.	28.9	363
2	Why prokaryotes have pangenomes. <i>Nature Microbiology</i> , 2017, 2, 17040.	13.3	327
3	Insights into the Evolution of Longevity from the Bowhead Whale Genome. <i>Cell Reports</i> , 2015, 10, 112-122.	6.4	280
4	The Interrelationships of Placental Mammals and the Limits of Phylogenetic Inference. <i>Genome Biology and Evolution</i> , 2016, 8, 330-344.	2.5	195
5	Evolution of sweet taste perception in hummingbirds by transformation of the ancestral umami receptor. <i>Science</i> , 2014, 345, 929-933.	12.6	169
6	The hybrid nature of the Eukaryota and a consilient view of life on Earth. <i>Nature Reviews Microbiology</i> , 2014, 12, 449-455.	28.6	124
7	Growing old, yet staying young: The role of telomeres in bats' exceptional longevity. <i>Science Advances</i> , 2018, 4, eaao0926.	10.3	120
8	Heterogeneous Models Place the Root of the Placental Mammal Phylogeny. <i>Molecular Biology and Evolution</i> , 2013, 30, 2145-2156.	8.9	115
9	Does a tree-like phylogeny only exist at the tips in the prokaryotes?. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2004, 271, 2551-2558.	2.6	114
10	Strigolactone synthesis is ancestral in land plants, but canonical strigolactone signalling is a flowering plant innovation. <i>BMC Biology</i> , 2019, 17, 70.	3.8	92
11	Return to the Sea, Get Huge, Beat Cancer: An Analysis of Cetacean Genomes Including an Assembly for the Humpback Whale ( <i>Megaptera novaeangliae</i> ). <i>Molecular Biology and Evolution</i> , 2019, 36, 1746-1763.	8.9	75
12	The public goods hypothesis for the evolution of life on Earth. <i>Biology Direct</i> , 2011, 6, 41.	4.6	74
13	Inadvertent Paralog Inclusion Drives Artifactual Topologies and Timetree Estimates in Phylogenomics. <i>Molecular Biology and Evolution</i> , 2019, 36, 1344-1356.	8.9	56
14	Insights into Kinesin-1 Activation from the Crystal Structure of KLC2 Bound to JIP3. <i>Structure</i> , 2018, 26, 1486-1498.e6.	3.3	47
15	Adaptive Evolution as a Predictor of Species-Specific Innate Immune Response. <i>Molecular Biology and Evolution</i> , 2015, 32, 1717-1729.	8.9	39
16	The phylogeny of the mammalian heme peroxidases and the evolution of their diverse functions. <i>BMC Evolutionary Biology</i> , 2008, 8, 101.	3.2	31
17	Response to <i>Teladorsagia circumcincta</i> infection in Scottish Blackface lambs with divergent phenotypes for nematode resistance. <i>Veterinary Parasitology</i> , 2014, 206, 200-207.	1.8	28
18	Ribosome heterogeneity in <i>Drosophila melanogaster</i> gonads through paralog-switching. <i>Nucleic Acids Research</i> , 2022, 50, 2240-2257.	14.5	28

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19	The ring of life hypothesis for eukaryote origins is supported by multiple kinds of data. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2015, 370, 20140323.	4.0	19
20	Transcriptional profiling of the ovine abomasal lymph node reveals a role for timing of the immune response in gastrointestinal nematode resistance. <i>Veterinary Parasitology</i> , 2016, 224, 96-108.	1.8	19
21	Surface layer proteins from virulent <i>Clostridium difficile</i> ribotypes exhibit signatures of positive selection with consequences for innate immune response. <i>BMC Evolutionary Biology</i> , 2017, 17, 90.	3.2	19
22	Paternally Expressed Imprinted Genes under Positive Darwinian Selection in <i>Arabidopsis thaliana</i> . <i>Molecular Biology and Evolution</i> , 2019, 36, 1239-1253.	8.9	18
23	New missense variants in <i>REL1A</i> causing hypomineralised amelogenesis imperfecta. <i>Clinical Genetics</i> , 2020, 97, 688-695.	2.0	18
24	Cytoplasmic long noncoding RNAs are differentially regulated and translated during human neuronal differentiation. <i>Rna</i> , 2021, 27, 1082-1101.	3.5	17
25	Selection and the Cell Cycle: Positive Darwinian Selection in a Well-Known DNA Damage Response Pathway. <i>Journal of Molecular Evolution</i> , 2010, 71, 444-457.	1.8	16
26	The role of CAPG in molecular communication between the embryo and the uterine endometrium: Is its function conserved in species with different implantation strategies?. <i>FASEB Journal</i> , 2020, 34, 11015-11029.	0.5	15
27	Gamma Chain Receptor Interleukins: Evidence for Positive Selection Driving the Evolution of Cell-to-Cell Communicators in the Mammalian Immune System. <i>Journal of Molecular Evolution</i> , 2005, 61, 608-619.	1.8	14
28	Functional Consequence of Positive Selection Revealed through Rational Mutagenesis of Human Myeloperoxidase. <i>Molecular Biology and Evolution</i> , 2012, 29, 2039-2046.	8.9	14
29	Reply to "The population genetics of pangenomes". <i>Nature Microbiology</i> , 2017, 2, 1575-1575.	13.3	11
30	Mind the gaps in cellular evolution. <i>Nature</i> , 2017, 541, 297-299.	27.8	8
31	Gene Fusions Derived by Transcriptional Readthrough are Driven by Segmental Duplication in Human. <i>Genome Biology and Evolution</i> , 2019, 11, 2678-2690.	2.5	7
32	Adaptive evolution of the human fatty acid synthase gene: Support for the cancer selection and fat utilization hypotheses?. <i>Gene</i> , 2005, 360, 151-159.	2.2	6
33	An active second dihydrofolate reductase enzyme is not a feature of rat and mouse, but they do have activity in their mitochondria. <i>FEBS Letters</i> , 2015, 589, 1855-1862.	2.8	5
34	In <i>Arabidopsis thaliana</i> codon volatility scores reflect GC3 composition rather than selective pressure. <i>BMC Research Notes</i> , 2012, 5, 359.	1.4	4
35	Spectrum of pathogenic variants and founder effects in amelogenesis imperfecta associated with <i>MMP20</i> . <i>Human Mutation</i> , 2021, 42, 567-576.	2.5	4
36	Adaptive Evolution in TRIF Leads to Discordance between Human and Mouse Innate Immune Signaling. <i>Genome Biology and Evolution</i> , 2021, 13, .	2.5	3

#	ARTICLE	IF	CITATIONS
37	Ghost locus appears. Nature, 2014, 514, 570-571.	27.8	2