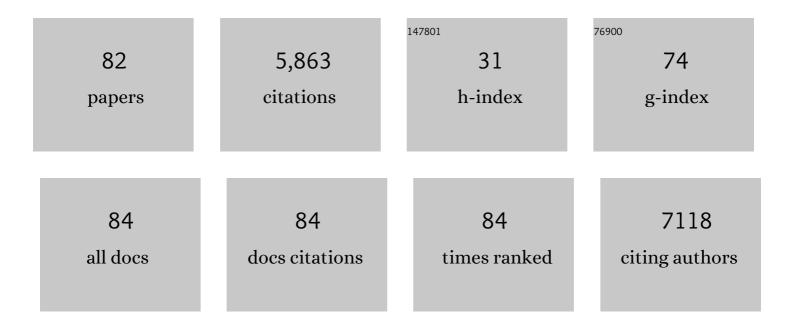
List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Comparison of Demographic and Clinical Features of Bipolar Disorder in Persons of African and European Ancestry. Journal of Racial and Ethnic Health Disparities, 2023, 10, 367-372.	3.2	1
2	Motivating and Discouraging Factors for Bipolar Patient Participation in Genomic Research. Public Health Genomics, 2021, 24, 89-98.	1.0	2
3	Current practices in nutrition management and disease incidence of common marmosets ( <i>Callithrix jacchus</i> ). Journal of Medical Primatology, 2021, 50, 164-175.	0.6	8
4	The roles of borderline personality disorder symptoms and dispositional capability for suicide in suicidal ideation and suicide attempts: Examination of the COMT Val158Met polymorphism. Psychiatry Research, 2021, 302, 114011.	3.3	3
5	Pedigree reconstruction and distant pairwise relatedness estimation from genome sequence data: A demonstration in a population of rhesus macaques ( <i>Macaca mulatta</i> ). Molecular Ecology Resources, 2021, 21, 1333-1346.	4.8	3
6	Decreased core symptoms of mania and utilization of lithium/mood stabilizing anticonvulsants in U.S. bipolar I patients of African vs European ancestry. Journal of Affective Disorders, 2020, 260, 361-365.	4.1	7
7	The BDNF Val66Met Polymorphism Moderates the Relationship Between Posttraumatic Stress Disorder and Trauma Script-evoked Attentional Bias to Cocaine Cues Among Patients with Cocaine Dependence. Journal of Anxiety Disorders, 2020, 72, 102223.	3.2	4
8	In Vitro Effects of Ligand Bias on Primate Mu Opioid Receptor Downstream Signaling. International Journal of Molecular Sciences, 2020, 21, 3999.	4.1	5
9	Genome-wide DNA methylomic differences between dorsolateral prefrontal and temporal pole cortices of bipolar disorder. Journal of Psychiatric Research, 2019, 117, 45-54.	3.1	24
10	Comparative genomics of Bifidobacterium species isolated from marmosets and humans. American Journal of Primatology, 2019, 81, e983.	1.7	12
11	Alcohol-induced changes in the gut microbiome and metabolome of rhesus macaques. Psychopharmacology, 2019, 236, 1531-1544.	3.1	16
12	Genetics of human brain evolution. Progress in Brain Research, 2019, 250, 3-39.	1.4	4
13	The Genome of the Common Marmoset. , 2019, , 313-333.		1
14	MicrobiomeR: An R Package for Simplified and Standardized Microbiome Analysis Workflows. Journal of Open Source Software, 2019, 4, 1299.	4.6	6
15	Altered neuro-inflammatory gene expression in hippocampus in major depressive disorder. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2018, 82, 177-186.	4.8	108
16	Risk-taking behaviors and stressors differentially predict suicidal preparation, non-fatal suicide attempts, and suicide deaths. Psychiatry Research, 2018, 270, 160-167.	3.3	27
17	Convergent Balancing Selection on the Mu-Opioid Receptor in Primates. Molecular Biology and Evolution, 2017, 34, 1629-1643.	8.9	12
18	Transcriptomic profiling of the ventral tegmental area and nucleus accumbens in rhesus macaques following long-term cocaine self-administration. Drug and Alcohol Dependence, 2017, 175, 9-23.	3.2	23

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19	Persistent negative effects of alcohol drinking on aspects of novelty-directed behavior in male rhesus macaques. Alcohol, 2017, 63, 19-26.	1.7	5
20	Spontaneous endometriosis in rhesus macaques: evidence for a genetic association with specific Mamu-A1 alleles. Primate Biology, 2017, 4, 117-125.	1.0	1
21	The population genomics of rhesus macaques ( <i>Macaca mulatta</i> ) based on whole-genome sequences. Genome Research, 2016, 26, 1651-1662.	5.5	101
22	Biogeography of the Intestinal Mucosal and Lumenal Microbiome in the Rhesus Macaque. Cell Host and Microbe, 2015, 17, 385-391.	11.0	273
23	Evolutionary conservation in genes underlying human psychiatric disorders. Frontiers in Human Neuroscience, 2014, 8, 283.	2.0	27
24	Genetic substructure in cynomolgus macaques (Macaca fascicularis) on the island of Mauritius. BMC Genomics, 2014, 15, 748.	2.8	17
25	Systematic mapping of occluded genes by cell fusion reveals prevalence and stability of <i>cis</i> -mediated silencing in somatic cells. Genome Research, 2014, 24, 267-280.	5.5	12
26	Bringing non-human primate research into the post-genomic era: how monkeys are teaching us about elite controllers of HIV/AIDS. Genome Biology, 2014, 15, 507.	8.8	2
27	Development and validation of a SNPâ€based assay for inferring the genetic ancestry of rhesus macaques ( <i>Macaca mulatta</i> ). American Journal of Primatology, 2014, 76, 1105-1113.	1.7	23
28	Twinning and survivorship of captive common marmosets (Callithrix jacchus) and cotton-top tamarins (Saguinus oedipus). Journal of the American Association for Laboratory Animal Science, 2014, 53, 7-11.	1.2	8
29	Large-scale polymorphism discovery in macaque G-protein coupled receptors. BMC Genomics, 2013, 14, 703.	2.8	6
30	Nonhuman Primate Models in the Genomic Era: A Paradigm Shift. ILAR Journal, 2013, 54, 154-165.	1.8	50
31	How Brains Are Built: Genetics and Evolution. Brain, Behavior and Evolution, 2013, 81, 71-73.	1.7	2
32	Trace Amine Associated Receptor 1 Modulates Behavioral Effects of Ethanol. Substance Abuse: Research and Treatment, 2013, 7, SART.S12110.	0.9	32
33	Naturally occurring, physiologically normal, primate chimeras. Chimerism, 2012, 3, 43-44.	0.7	11
34	Genetic correlates of the evolving primate brain. Progress in Brain Research, 2012, 195, 27-44.	1.4	5
35	The resurgence and genetic implications of New World primates in biomedical research. Trends in Genetics, 2012, 28, 586-591.	6.7	29
36	Trace Amine Associated Receptor 1 Signaling in Activated Lymphocytes. Journal of NeuroImmune Pharmacology, 2012, 7, 866-876.	4.1	64

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37	Augmentation of methamphetamine-induced behaviors in transgenic mice lacking the trace amine-associated receptor 1. Pharmacology Biochemistry and Behavior, 2012, 101, 201-207.	2.9	77
38	Quantitative molecular assessment of chimerism across tissues in marmosets and tamarins. BMC Genomics, 2012, 13, 98.	2.8	46
39	Refinement of primate copy number variationhotspots identifies candidate genomic regions evolving under positive selection. Genome Biology, 2011, 12, R52.	8.8	58
40	Expanding whole exome resequencing into non-human primates. Genome Biology, 2011, 12, R87.	9.6	68
41	Growth-associated protein-43 and ephrin B3 induction in the brain of adult SIV-infected rhesus macaques. Journal of NeuroVirology, 2011, 17, 455-468.	2.1	2
42	Comparative genetic approaches to the evolution of human brain and behavior. American Journal of Human Biology, 2011, 23, 53-64.	1.6	4
43	Functional evolution of the trace amine associated receptors in mammals and the loss of TAAR1 in dogs. BMC Evolutionary Biology, 2010, 10, 51.	3.2	31
44	Normal thermoregulatory responses to 3â€iodothyronamine, trace amines and amphetamineâ€like psychostimulants in trace amine associated receptor 1 knockout mice. Journal of Neuroscience Research, 2010, 88, 1962-1969.	2.9	77
45	<i>TPH2</i> 5′―and 3′―egulatory polymorphisms are differentially associated with HPA axis function and self―njurious behavior in rhesus monkeys. Genes, Brain and Behavior, 2010, 9, 335-347.	2.2	27
46	A pharmacogenetic model of naltrexone-induced attenuation of alcohol consumption in rhesus monkeysâ~†. Drug and Alcohol Dependence, 2010, 109, 252-256.	3.2	48
47	The effect of rearing experience and TPH2 genotype on HPA axis function and aggression in rhesus monkeys: A retrospective analysis. Hormones and Behavior, 2010, 57, 184-191.	2.1	29
48	Polymorphisms in the 3′ UTR of the serotonin transporter are associated with cognitive flexibility in rhesus macaques. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2009, 150B, 467-475.	1.7	23
49	Resources for genetic management and genomics research on nonâ€human primates at the National Primate Research Centers (NPRCs). Journal of Medical Primatology, 2009, 38, 17-23.	0.6	23
50	Bioinformatic approaches to identifying orthologs and assessing evolutionary relationships. Methods, 2009, 49, 50-55.	3.8	12
51	Resource brief: The National Non-Human Primate DNA Bank. Methods, 2009, 49, 3-4.	3.8	3
52	Human Expression Variation in the Mu-Opioid Receptor is Paralleled in Rhesus Macaque. Behavior Genetics, 2008, 38, 390-395.	2.1	21
53	Functional characterization of the human TPH2 5′ regulatory region: untranslated region and polymorphisms modulate gene expression in vitro. Human Genetics, 2008, 122, 645-657.	3.8	106
54	Cloning, expression, and functional analysis of rhesus monkey trace amineâ€associated receptor 6: Evidence for lack of monoaminergic association. Journal of Neuroscience Research, 2008, 86, 3435-3446.	2.9	25

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55	Functional variation in the 3′ untranslated region of the serotonin transporter in human and rhesus macaque. Genes, Brain and Behavior, 2008, 7, 690-697.	2.2	31
56	Genetic basis of human brain evolution. Trends in Neurosciences, 2008, 31, 637-644.	8.6	88
57	Exploring the Origins of the Human Brain through Molecular Evolution. Brain, Behavior and Evolution, 2008, 72, 168-177.	1.7	14
58	Extensive contribution of embryonic stem cells to the development of an evolutionarily divergent host. Human Molecular Genetics, 2008, 17, 27-37.	2.9	29
59	Analysis of copy number variation in the rhesus macaque genome identifies candidate loci for evolutionary and human disease studies. Human Molecular Genetics, 2008, 17, 1127-1136.	2.9	101
60	Rhesus Monkey Trace Amine-Associated Receptor 1 Signaling: Enhancement by Monoamine Transporters and Attenuation by the D2 Autoreceptor in Vitro. Journal of Pharmacology and Experimental Therapeutics, 2007, 321, 116-127.	2.5	103
61	Evolutionary and Biomedical Insights from the Rhesus Macaque Genome. Science, 2007, 316, 222-234.	12.6	1,283
62	Uncovering the mutation-fixation correlation in short lineages. BMC Evolutionary Biology, 2007, 7, 168.	3.2	9
63	Response to Comment on "Ongoing Adaptive Evolution of ASPM, a Brain Size Determinant in Homo sapiens" and "Microcephalin, a Gene Regulating Brain Size, Continues to Evolve Adaptively in Humans". Science, 2006, 313, 172b-172b.	12.6	51
64	Molecular evolution of the brain size regulator genes CDK5RAP2 and CENPJ. Gene, 2006, 375, 75-79.	2.2	77
65	SPEED: a molecular-evolution-based database of mammalian orthologous groups. Bioinformatics, 2006, 22, 2835-2837.	4.1	9
66	Systematically Assessing the Influence of 3-Dimensional Structural Context on the Molecular Evolution of Mammalian Proteomes. Molecular Biology and Evolution, 2006, 23, 2131-2133.	8.9	35
67	A primate-specific acceleration in the evolution of the caspase-dependent apoptosis pathway. Human Molecular Genetics, 2006, 15, 3034-3040.	2.9	20
68	Multiple independent origins of sex chromosomes in amniotes. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 18031-18032.	7.1	20
69	Evidence that the adaptive allele of the brain size gene microcephalin introgressed into Homo sapiens from an archaic Homo lineage. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 18178-18183.	7.1	127
70	Sonic Hedgehog, a key development gene, experienced intensified molecular evolution in primates. Human Molecular Genetics, 2006, 15, 2031-2037.	2.9	37
71	Ongoing Adaptive Evolution of <i>ASPM</i> , a Brain Size Determinant in <i>Homo sapiens</i> . Science, 2005, 309, 1720-1722.	12.6	445
72	The X chromosome: not just her brother's keeper. Nature Genetics, 2005, 37, 343-345.	21.4	16

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73	A highly unexpected strong correlation between fixation probability of nonsynonymous mutations and mutation rate. Trends in Genetics, 2005, 21, 381-385.	6.7	45
74	<i>Microcephalin</i> , a Gene Regulating Brain Size, Continues to Evolve Adaptively in Humans. Science, 2005, 309, 1717-1720.	12.6	447
75	Positive selection on the human genome. Human Molecular Genetics, 2004, 13, R245-R254.	2.9	215
76	Adaptive evolution of ASPM, a major determinant of cerebral cortical size in humans. Human Molecular Genetics, 2004, 13, 489-494.	2.9	232
77	How mammalian sex chromosomes acquired their peculiar gene content. BioEssays, 2004, 26, 159-169.	2.5	103
78	Reconstructing the evolutionary history of microcephalin, a gene controlling human brain size. Human Molecular Genetics, 2004, 13, 1139-1145.	2.9	191
79	Accelerated Evolution of Nervous System Genes in the Origin of Homo sapiens. Cell, 2004, 119, 1027-1040.	28.9	404
80	Diverse fates of paralogs following segmental duplication of telomeric genesâ~†â~†â~†. Genomics, 2004, 84, 239-247.	2.9	19
81	Effects of chromosomal rearrangements on human-chimpanzee molecular evolution. Genomics, 2004, 84, 757-761.	2.9	23
82	Genomic Divergence Between Human and Chimpanzee Estimated from Large-Scale Alignments of	2.4	64

Genomic Sequences. Journal of Heredity, 2001, 92, 481-489.