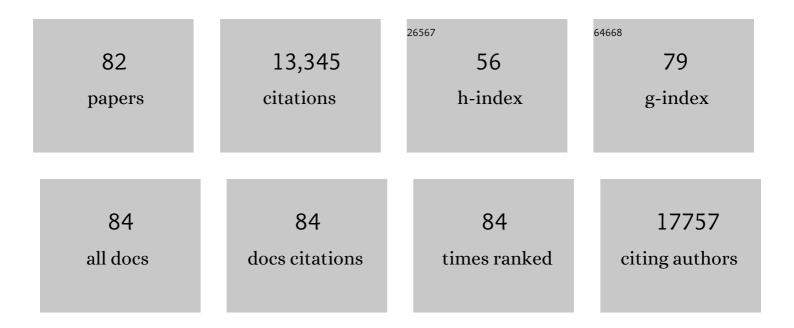
Robert W. Williams

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1880418/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	GeneCup: mining PubMed and GWAS catalog for gene–keyword relationships. G3: Genes, Genomes, Genetics, 2022, 12, .	0.8	8
2	Systems genetics in the rat HXB/BXH family identifies Tti2 as a pleiotropic quantitative trait gene for adult hippocampal neurogenesis and serum glucose. PLoS Genetics, 2022, 18, e1009638.	1.5	3
3	Highlights from the Era of Open Source Web-Based Tools. Journal of Neuroscience, 2021, 41, 927-936.	1.7	19
4	A platform for experimental precision medicine: The extended BXD mouse family. Cell Systems, 2021, 12, 235-247.e9.	2.9	115
5	The genome sequence of the Norway rat, Rattus norvegicus Berkenhout 1769. Wellcome Open Research, 2021, 6, 118.	0.9	16
6	Speeding up eQTL scans in the BXD population using GPUs. G3: Genes, Genomes, Genetics, 2021, 11, .	0.8	2
7	Gene-by-environment modulation of lifespan and weight gain in the murine BXD family. Nature Metabolism, 2021, 3, 1217-1227.	5.1	27
8	Variability and heritability of mouse brain structure: Microscopic MRI atlases and connectomes for diverse strains. Neurolmage, 2020, 222, 117274.	2.1	33
9	Identifying the molecular systems that influence cognitive resilience to Alzheimer's disease in genetically diverse mice. Learning and Memory, 2020, 27, 355-371.	0.5	15
10	Genetic cartography of longevity in humans and mice: Current landscape and horizons. Biochimica Et Biophysica Acta - Molecular Basis of Disease, 2018, 1864, 2718-2732.	1.8	27
11	Postâ€genomic behavioral genetics: From revolution to routine. Genes, Brain and Behavior, 2018, 17, e12441.	1.1	17
12	Reproducibility and replicability of rodent phenotyping in preclinical studies. Neuroscience and Biobehavioral Reviews, 2018, 87, 218-232.	2.9	153
13	Preface to a special issue on genetic models of alcoholism and alcohol-stress interactions. Alcohol, 2017, 58, 23-24.	0.8	1
14	Multi-omics analysis identifies ATF4 as a key regulator of the mitochondrial stress response in mammals. Journal of Cell Biology, 2017, 216, 2027-2045.	2.3	590
15	Orbitofrontal Neuroadaptations and Cross-Species Synaptic Biomarkers in Heavy-Drinking Macaques. Journal of Neuroscience, 2017, 37, 3646-3660.	1.7	43
16	Resources for Systems Genetics. Methods in Molecular Biology, 2017, 1488, 3-29.	0.4	42
17	Genetic Variation in the Social Environment Contributes to Health and Disease. PLoS Genetics, 2017, 13, e1006498.	1.5	110
18	GeneNetwork: framework for web-based genetics. Journal of Open Source Software, 2016, 1, 25.	2.0	51

#	Article	IF	CITATIONS
19	Correlation Trait Loci (CTL) mapping: phenotype network inference subject to genotype. Journal of Open Source Software, 2016, 1, 87.	2.0	3
20	Common genetic variants influence human subcortical brain structures. Nature, 2015, 520, 224-229.	13.7	772
21	Functionally Enigmatic Genes: A Case Study of the Brain Ignorome. PLoS ONE, 2014, 9, e88889.	1.1	77
22	Genetics of Gene Expression in CNS. International Review of Neurobiology, 2014, 116, 195-231.	0.9	26
23	Mitonuclear protein imbalance as a conserved longevity mechanism. Nature, 2013, 497, 451-457.	13.7	846
24	Metabolic Stress Modulates Alzheimer's β-Secretase Gene Transcription via SIRT1-PPARγ-PGC-1 in Neurons. Cell Metabolism, 2013, 17, 685-694.	7.2	170
25	Genetic variation of the cutaneous HPA axis: An analysis of UVB-induced differential responses. Gene, 2013, 530, 1-7.	1.0	36
26	A promoter polymorphism in the Per3 gene is associated with alcohol and stress response. Translational Psychiatry, 2012, 2, e73-e73.	2.4	63
27	Fine-Scale Maps of Recombination Rates and Hotspots in the Mouse Genome. Genetics, 2012, 191, 757-764.	1.2	82
28	Genetic and Molecular Network Analysis of Behavior. International Review of Neurobiology, 2012, 104, 135-157.	0.9	17
29	Systems Genetics of Metabolism: The Use of the BXD Murine Reference Panel for Multiscalar Integration of Traits. Cell, 2012, 150, 1287-1299.	13.5	212
30	Murine Gut Microbiota Is Defined by Host Genetics and Modulates Variation of Metabolic Traits. PLoS ONE, 2012, 7, e39191.	1.1	198
31	Genetic Dissection of Behavioral Flexibility: Reversal Learning in Mice. Biological Psychiatry, 2011, 69, 1109-1116.	0.7	97
32	Detection, Validation, and Downstream Analysis of Allelic Variation in Gene Expression. Genetics, 2010, 184, 119-128.	1.2	60
33	Strain Differences in Stress Responsivity Are Associated with Divergent Amygdala Gene Expression and Glutamate-Mediated Neuronal Excitability. Journal of Neuroscience, 2010, 30, 5357-5367.	1.7	224
34	Metabolic Networks of Longevity. Cell, 2010, 142, 9-14.	13.5	190
35	Host Genetic Variation Affects Resistance to Infection with a Highly Pathogenic H5N1 Influenza A Virus in Mice. Journal of Virology, 2009, 83, 10417-10426.	1.5	169
36	Antisense transcription: A critical look in both directions. Cellular and Molecular Life Sciences, 2009, 66, 94-112.	2.4	104

#	Article	IF	CITATIONS
37	The Diasporin Pathway: a tumor progression-related transcriptional network that predicts breast cancer survival. Clinical and Experimental Metastasis, 2008, 25, 357-369.	1.7	70
38	The Collaborative Cross at Oak Ridge National Laboratory: developing a powerful resource for systems genetics. Mammalian Genome, 2008, 19, 382-389.	1.0	245
39	Exploiting regulatory variation to identify genes underlying quantitative resistance to the wheat stem rust pathogen Puccinia graminis f. sp. tritici in barley. Theoretical and Applied Genetics, 2008, 117, 261-272.	1.8	43
40	The Neuroscience Information Framework: A Data and Knowledge Environment for Neuroscience. Neuroinformatics, 2008, 6, 149-160.	1.5	189
41	Alcohol trait and transcriptional genomic analysis of C57BL/6 substrains. Genes, Brain and Behavior, 2008, 7, 677-689.	1.1	81
42	Interleukin 17–producing T helper cells and interleukin 17 orchestrate autoreactive germinal center development in autoimmune BXD2 mice. Nature Immunology, 2008, 9, 166-175.	7.0	639
43	Dissection of a QTL Hotspot on Mouse Distal Chromosome 1 that Modulates Neurobehavioral Phenotypes and Gene Expression. PLoS Genetics, 2008, 4, e1000260.	1.5	98
44	Variation in Mouse Basolateral Amygdala Volume is Associated With Differences in Stress Reactivity and Fear Learning. Neuropsychopharmacology, 2008, 33, 2595-2604.	2.8	123
45	High Susceptibility to Experimental Myopia in a Mouse Model with a Retinal ON Pathway Defect. , 2008, 49, 706.		106
46	The p47 GTPases <i>ligp2</i> and <i>lrgb10</i> Regulate Innate Immunity and Inflammation to Murine <i>Chlamydia psittaci</i> Infection. Journal of Immunology, 2007, 179, 1814-1824.	0.4	79
47	Genome-level analysis of genetic regulation of liver gene expression networks. Hepatology, 2007, 46, 548-557.	3.6	49
48	How replicable are mRNA expression QTL?. Mammalian Genome, 2006, 17, 643-656.	1.0	56
49	Combined Expression Trait Correlations and Expression Quantitative Trait Locus Mapping. PLoS Genetics, 2006, 2, e6.	1.5	97
50	A High-Resolution Single Nucleotide Polymorphism Genetic Map of the Mouse Genome. PLoS Biology, 2006, 4, e395.	2.6	243
51	Natural variation and genetic covariance in adult hippocampal neurogenesis. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 780-785.	3.3	181
52	Uncovering regulatory pathways that affect hematopoietic stem cell function using 'genetical genomics'. Nature Genetics, 2005, 37, 225-232.	9.4	366
53	Complex trait analysis of gene expression uncovers polygenic and pleiotropic networks that modulate nervous system function. Nature Genetics, 2005, 37, 233-242.	9.4	695
54	Genetic Segregation of Spontaneous Erosive Arthritis and Generalized Autoimmune Disease in the BXD2 Recombinant Inbred Strain of Mice. Scandinavian Journal of Immunology, 2005, 61, 128-138.	1.3	59

#	Article	IF	CITATIONS
55	Quantitative Trait Locus Analysis Using Recombinant Inbred Intercrosses. Genetics, 2005, 170, 1299-1311.	1.2	75
56	Inferring gene transcriptional modulatory relations: a genetical genomics approach. Human Molecular Genetics, 2005, 14, 1119-1125.	1.4	76
57	Ethanol-Responsive Brain Region Expression Networks: Implications for Behavioral Responses to Acute Ethanol in DBA/2J versus C57BL/6J Mice. Journal of Neuroscience, 2005, 25, 2255-2266.	1.7	251
58	Measurement of Refractive State and Deprivation Myopia in Two Strains of Mice. Optometry and Vision Science, 2004, 81, 99-110.	0.6	164
59	The Collaborative Cross, a community resource for the genetic analysis of complex traits. Nature Genetics, 2004, 36, 1133-1137.	9.4	1,034
60	WebQTL: rapid exploratory analysis of gene expression and genetic networks for brain and behavior. Nature Neuroscience, 2004, 7, 485-486.	7.1	176
61	Genetic structure of the LXS panel of recombinant inbred mouse strains: a powerful resource for complex trait analysis. Mammalian Genome, 2004, 15, 637-647.	1.0	95
62	Towards Effective and Rewarding Data Sharing. Neuroinformatics, 2003, 1, 289-296.	1.5	78
63	WebQTL: Web-Based Complex Trait Analysis. Neuroinformatics, 2003, 1, 299-308.	1.5	249
64	Genetic Correlates of Gene Expression in Recombinant Inbred Strains: A Relational Model System to Explore Neurobehavioral Phenotypes. Neuroinformatics, 2003, 1, 343-358.	1.5	118
65	A strategy for the integration of QTL, gene expression, and sequence analyses. Mammalian Genome, 2003, 14, 733-747.	1.0	69
66	Genetic architecture of the mouse hippocampus: identification of gene loci with selective regional effects. Genes, Brain and Behavior, 2003, 2, 238-252.	1.1	72
67	The nature and identification of quantitative trait loci: a community's view. Nature Reviews Genetics, 2003, 4, 911-916.	7.7	390
68	Increased brain size and glial cell number in CD81-null mice. Journal of Comparative Neurology, 2002, 453, 22-32.	0.9	87
69	Genetic dissection of complex and quantitative traits: from fantasy to reality via a community effort. Mammalian Genome, 2002, 13, 175-178.	1.0	191
70	QTL analysis and genomewide mutagenesis in mice: complementary genetic approaches to the dissection of complex traits. Behavior Genetics, 2001, 31, 5-15.	1.4	78
71	Genetic control of retinal projections in inbred strains of albino mice. Journal of Comparative Neurology, 1995, 354, 459-469.	0.9	53
72	Target recognition and visual maps in the thalamus of achiasmatic dogs. Nature, 1994, 367, 637-639.	13.7	84

#	ARTICLE	IF	CITATIONS
73	Structure of clonal and polyclonal cell arrays in chimeric mouse retina Proceedings of the National Academy of Sciences of the United States of America, 1992, 89, 1184-1188.	3.3	52
74	Lineage versus environment in embryonic retina: a revisionist perspective. Trends in Neurosciences, 1992, 15, 368-373.	4.2	48
75	A novel cytoarchitectonic area induced experimentally within the primate visual cortex Proceedings of the National Academy of Sciences of the United States of America, 1991, 88, 2083-2087.	3.3	242
76	Photoreceptor mosaic: Number and distribution of rods and cones in the rhesus monkey retina. Journal of Comparative Neurology, 1990, 297, 499-508.	0.9	145
77	Elimination of neurons from the rhesus monkey's lateral geniculate nucleus during development. Journal of Comparative Neurology, 1988, 272, 424-436.	0.9	125
78	Three-dimensional counting: An accurate and direct method to estimate numbers of cells in sectioned material. Journal of Comparative Neurology, 1988, 278, 344-352.	0.9	423
79	The Control of Neuron Number. Annual Review of Neuroscience, 1988, 11, 423-453.	5.0	527
80	Formation of retinal ganglion cell topography during prenatal development. Science, 1987, 236, 848-851.	6.0	99
81	Growth cones, dying axons, and developmental fluctuations in the fiber population of the cat's optic nerve. Journal of Comparative Neurology, 1986, 246, 32-69.	0.9	201
82	Dispersion of growing axons within the optic nerve of the embryonic monkey Proceedings of the National Academy of Sciences of the United States of America, 1985, 82, 3906-3910.	3.3	73