

Jonathan L Haines

List of Publications by Year in descending order

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Version: 2024-02-01

549
papers

82,612
citations

701

121
h-index

517

267
g-index

613
all docs

613
docs citations

613
times ranked

69301
citing authors

#	ARTICLE	IF	CITATIONS
1	Finding the missing heritability of complex diseases. <i>Nature</i> , 2009, 461, 747-753.	27.8	7,490
2	Mutations in Cu/Zn superoxide dismutase gene are associated with familial amyotrophic lateral sclerosis. <i>Nature</i> , 1993, 362, 59-62.	27.8	6,331
3	Meta-analysis of 74,046 individuals identifies 11 new susceptibility loci for Alzheimer's disease. <i>Nature Genetics</i> , 2013, 45, 1452-1458.	21.4	3,741
4	Genetic risk and a primary role for cell-mediated immune mechanisms in multiple sclerosis. <i>Nature</i> , 2011, 476, 214-219.	27.8	2,400
5	Complement Factor H Variant Increases the Risk of Age-Related Macular Degeneration. <i>Science</i> , 2005, 308, 419-421.	12.6	2,232
6	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. <i>Nature Genetics</i> , 2013, 45, 984-994.	21.4	2,067
7	Genetic meta-analysis of diagnosed Alzheimer's disease identifies new risk loci and implicates A β , tau, immunity and lipid processing. <i>Nature Genetics</i> , 2019, 51, 414-430.	21.4	1,962
8	Functional impact of global rare copy number variation in autism spectrum disorders. <i>Nature</i> , 2010, 466, 368-372.	27.8	1,803
9	Common variants at MS4A4/MS4A6E, CD2AP, CD33 and EPHA1 are associated with late-onset Alzheimer's disease. <i>Nature Genetics</i> , 2011, 43, 436-441.	21.4	1,676
10	Risk Alleles for Multiple Sclerosis Identified by a Genomewide Study. <i>New England Journal of Medicine</i> , 2007, 357, 851-862.	27.0	1,529
11	Mapping autism risk loci using genetic linkage and chromosomal rearrangements. <i>Nature Genetics</i> , 2007, 39, 319-328.	21.4	1,272
12	Analysis of immune-related loci identifies 48 new susceptibility variants for multiple sclerosis. <i>Nature Genetics</i> , 2013, 45, 1353-1360.	21.4	1,213
13	A large genome-wide association study of age-related macular degeneration highlights contributions of rare and common variants. <i>Nature Genetics</i> , 2016, 48, 134-143.	21.4	1,167
14	Mutations in UBQLN2 cause dominant X-linked juvenile and adult-onset ALS and ALS/dementia. <i>Nature</i> , 2011, 477, 211-215.	27.8	1,016
15	Common genetic variants on 5p14.1 associate with autism spectrum disorders. <i>Nature</i> , 2009, 459, 528-533.	27.8	912
16	Systematic comparison of phenome-wide association study of electronic medical record data and genome-wide association study data. <i>Nature Biotechnology</i> , 2013, 31, 1102-1111.	17.5	846
17	Convergence of Genes and Cellular Pathways Dysregulated in Autism Spectrum Disorders. <i>American Journal of Human Genetics</i> , 2014, 94, 677-694.	6.2	819
18	Rare coding variants in PLCG2, ABI3, and TREM2 implicate microglial-mediated innate immunity in Alzheimer's disease. <i>Nature Genetics</i> , 2017, 49, 1373-1384.	21.4	783

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19	Meta-analysis of genome scans and replication identify CD6, IRF8 and TNFRSF1A as new multiple sclerosis susceptibility loci. <i>Nature Genetics</i> , 2009, 41, 776-782.	21.4	729
20	Multiple sclerosis genomic map implicates peripheral immune cells and microglia in susceptibility. <i>Science</i> , 2019, 365, .	12.6	710
21	New insights into the genetic etiology of Alzheimer's disease and related dementias. <i>Nature Genetics</i> , 2022, 54, 412-436.	21.4	700
22	Seven new loci associated with age-related macular degeneration. <i>Nature Genetics</i> , 2013, 45, 433-439.	21.4	687
23	The Electronic Medical Records and Genomics (eMERGE) Network: past, present, and future. <i>Genetics in Medicine</i> , 2013, 15, 761-771.	2.4	611
24	The PhenX Toolkit: Get the Most From Your Measures. <i>American Journal of Epidemiology</i> , 2011, 174, 253-260.	3.4	610
25	Heterozygosity for a Surfactant Protein C Gene Mutation Associated with Usual Interstitial Pneumonitis and Cellular Nonspecific Interstitial Pneumonitis in One Kindred. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2002, 165, 1322-1328.	5.6	597
26	Interleukin 7 receptor α chain (IL7R) shows allelic and functional association with multiple sclerosis. <i>Nature Genetics</i> , 2007, 39, 1083-1091.	21.4	578
27	Isolation of a novel gene underlying batten disease, CLN3. <i>Cell</i> , 1995, 82, 949-957.	28.9	554
28	A genome-wide scan for common alleles affecting risk for autism. <i>Human Molecular Genetics</i> , 2010, 19, 4072-4082.	2.9	538
29	Mitochondrial Polymorphisms Significantly Reduce the Risk of Parkinson Disease. <i>American Journal of Human Genetics</i> , 2003, 72, 804-811.	6.2	507
30	Genome-wide association study identifies a new breast cancer susceptibility locus at 6q25.1. <i>Nature Genetics</i> , 2009, 41, 324-328.	21.4	481
31	Genetic linkage of bilateral acoustic neurofibromatosis to a DNA marker on chromosome 22. <i>Nature</i> , 1987, 329, 246-248.	27.8	478
32	Genetic variants near <i>TIMP3</i> and high-density lipoprotein-associated loci influence susceptibility to age-related macular degeneration. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 7401-7406.	7.1	475
33	Genome-Wide Association Study Confirms SNPs in <i>SNCA</i> and the <i>MAPT</i> Region as Common Risk Factors for Parkinson Disease. <i>Annals of Human Genetics</i> , 2010, 74, 97-109.	0.8	417
34	Linkage of a Gene Causing Familial Amyotrophic Lateral Sclerosis to Chromosome 21 and Evidence of Genetic-Locus Heterogeneity. <i>New England Journal of Medicine</i> , 1991, 324, 1381-1384.	27.0	407
35	Meta-analysis Confirms CR1, CLU, and PICALM as Alzheimer Disease Risk Loci and Reveals Interactions With APOE Genotypes. <i>Archives of Neurology</i> , 2010, 67, 1473.	4.5	376
36	Variants in the ATP-Binding Cassette Transporter (ABCA7), Apolipoprotein E ϵ 4, and the Risk of Late-Onset Alzheimer Disease in African Americans. <i>JAMA - Journal of the American Medical Association</i> , 2013, 309, 1483.	7.4	360

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37	CWAS of Cerebrospinal Fluid Tau Levels Identifies Risk Variants for Alzheimer's Disease. <i>Neuron</i> , 2013, 78, 256-268.	8.1	344
38	Individual common variants exert weak effects on the risk for autism spectrum disorders. <i>Human Molecular Genetics</i> , 2012, 21, 4781-4792.	2.9	334
39	A common haplotype lowers PU.1 expression in myeloid cells and delays onset of Alzheimer's disease. <i>Nature Neuroscience</i> , 2017, 20, 1052-1061.	14.8	330
40	Cigarette Smoking Strongly Modifies the Association of LOC387715 and Age-Related Macular Degeneration. <i>American Journal of Human Genetics</i> , 2006, 78, 852-864.	6.2	316
41	Genome-wide meta-analysis identifies novel multiple sclerosis susceptibility loci. <i>Annals of Neurology</i> , 2011, 70, 897-912.	5.3	314
42	Class II HLA interactions modulate genetic risk for multiple sclerosis. <i>Nature Genetics</i> , 2015, 47, 1107-1113.	21.4	312
43	Mapping Multiple Sclerosis Susceptibility to the HLA-DR Locus in African Americans. <i>American Journal of Human Genetics</i> , 2004, 74, 160-167.	6.2	311
44	Genetic assessment of age-associated Alzheimer disease risk: Development and validation of a polygenic hazard score. <i>PLoS Medicine</i> , 2017, 14, e1002258.	8.4	311
45	Genome-Wide Association Meta-analysis of Neuropathologic Features of Alzheimer's Disease and Related Dementias. <i>PLoS Genetics</i> , 2014, 10, e1004606.	3.5	305
46	Common genetic variants in the CLDN2 and PRSS1-PRSS2 loci alter risk for alcohol-related and sporadic pancreatitis. <i>Nature Genetics</i> , 2012, 44, 1349-1354.	21.4	303
47	Robust Replication of Genotype-Phenotype Associations across Multiple Diseases in an Electronic Medical Record. <i>American Journal of Human Genetics</i> , 2010, 86, 560-572.	6.2	302
48	Cardiac Sodium Channel (<i>SCN5A</i>) Variants Associated with Atrial Fibrillation. <i>Circulation</i> , 2008, 117, 1927-1935.	1.6	292
49	Age at Onset in Two Common Neurodegenerative Diseases Is Genetically Controlled. <i>American Journal of Human Genetics</i> , 2002, 70, 985-993.	6.2	291
50	PheKB: a catalog and workflow for creating electronic phenotype algorithms for transportability. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2016, 23, 1046-1052.	4.4	284
51	Heterogeneity at the HLA-DRB1 locus and risk for multiple sclerosis. <i>Human Molecular Genetics</i> , 2006, 15, 2813-2824.	2.9	279
52	Common Variants at 9p21 and 8q22 Are Associated with Increased Susceptibility to Optic Nerve Degeneration in Glaucoma. <i>PLoS Genetics</i> , 2012, 8, e1002654.	3.5	276
53	The genetic defect in familial Alzheimer's disease is not tightly linked to the amyloid β -protein gene. <i>Nature</i> , 1987, 329, 156-157.	27.8	275
54	Genome-wide association analyses identify multiple loci associated with central corneal thickness and keratoconus. <i>Nature Genetics</i> , 2013, 45, 155-163.	21.4	269

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55	A High-Density Screen for Linkage in Multiple Sclerosis. <i>American Journal of Human Genetics</i> , 2005, 77, 454-467.	6.2	268
56	Analysis of European mitochondrial haplogroups with Alzheimer disease risk. <i>Neuroscience Letters</i> , 2004, 365, 28-32.	2.1	264
57	Quality Control Procedures for Genome-Wide Association Studies. <i>Current Protocols in Human Genetics</i> , 2011, 68, Unit1.19.	3.5	259
58	Complete Genomic Screen in Parkinson Disease. <i>JAMA - Journal of the American Medical Association</i> , 2001, 286, 2239.	7.4	257
59	Temperature-sensitive mutations in the III ^{IV} cytoplasmic loop region of the skeletal muscle sodium channel gene in paramyotonia congenita. <i>Cell</i> , 1992, 68, 769-774.	28.9	249
60	Human gene for torsion dystonia located on chromosome 9q32-q34. <i>Neuron</i> , 1989, 2, 1427-1434.	8.1	246
61	Exome sequencing of extended families with autism reveals genes shared across neurodevelopmental and neuropsychiatric disorders. <i>Molecular Autism</i> , 2014, 5, 1.	4.9	246
62	Exceptionally low likelihood of Alzheimer's dementia in APOE2 homozygotes from a 5,000-person neuropathological study. <i>Nature Communications</i> , 2020, 11, 667.	12.8	246
63	Genome-wide Association Study Implicates a Chromosome 12 Risk Locus for Late-Onset Alzheimer Disease. <i>American Journal of Human Genetics</i> , 2009, 84, 35-43.	6.2	242
64	Genetics, statistics and human disease: analytical retooling for complexity. <i>Trends in Genetics</i> , 2004, 20, 640-647.	6.7	230
65	Brain Expression Genome-Wide Association Study (eGWAS) Identifies Human Disease-Associated Variants. <i>PLoS Genetics</i> , 2012, 8, e1002707.	3.5	225
66	Meta-analysis of genome scans of age-related macular degeneration. <i>Human Molecular Genetics</i> , 2005, 14, 2257-2264.	2.9	224
67	Sex-Specific Association of Apolipoprotein E With Cerebrospinal Fluid Levels of Tau. <i>JAMA Neurology</i> , 2018, 75, 989.	9.0	223
68	Linkage of recessive familial amyotrophic lateral sclerosis to chromosome 2q33-q35. <i>Nature Genetics</i> , 1994, 7, 425-428.	21.4	221
69	Genome-wide analyses identify 68 new loci associated with intraocular pressure and improve risk prediction for primary open-angle glaucoma. <i>Nature Genetics</i> , 2018, 50, 778-782.	21.4	214
70	Genome-wide analysis of multi-ancestry cohorts identifies new loci influencing intraocular pressure and susceptibility to glaucoma. <i>Nature Genetics</i> , 2014, 46, 1126-1130.	21.4	212
71	Genome-wide association analysis identifies TXNRD2, ATXN2 and FOXC1 as susceptibility loci for primary open-angle glaucoma. <i>Nature Genetics</i> , 2016, 48, 189-194.	21.4	211
72	Glutathione S-transferase omega-1 modifies age-at-onset of Alzheimer disease and Parkinson disease. <i>Human Molecular Genetics</i> , 2003, 12, 3259-3267.	2.9	208

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73	A Genome-wide Association Study of Autism Reveals a Common Novel Risk Locus at 5p14.1. <i>Annals of Human Genetics</i> , 2009, 73, 263-273.	0.8	207
74	Neonatal Pulmonary Hypertension. <i>New England Journal of Medicine</i> , 2001, 344, 1832-1838.	27.0	202
75	Parkin mutations and susceptibility alleles in late-onset Parkinson's disease. <i>Annals of Neurology</i> , 2003, 53, 624-629.	5.3	201
76	Mutations in a Novel CLN6-Encoded Transmembrane Protein Cause Variant Neuronal Ceroid Lipofuscinosis in Man and Mouse. <i>American Journal of Human Genetics</i> , 2002, 70, 324-335.	6.2	199
77	Genome-wide association study identifies four novel loci associated with Alzheimer's endophenotypes and disease modifiers. <i>Acta Neuropathologica</i> , 2017, 133, 839-856.	7.7	199
78	Evidence for a role of the rare p.A152T variant in MAPT in increasing the risk for FTD-spectrum and Alzheimer's diseases. <i>Human Molecular Genetics</i> , 2012, 21, 3500-3512.	2.9	198
79	Prevalence of Mutations in TIGR/Myocilin in Patients with Adult and Juvenile Primary Open-Angle Glaucoma. <i>American Journal of Human Genetics</i> , 1998, 63, 1549-1552.	6.2	197
80	Genome-wide meta-analysis identifies 127 open-angle glaucoma loci with consistent effect across ancestries. <i>Nature Communications</i> , 2021, 12, 1258.	12.8	196
81	Clustering of autoimmune diseases in families with a high-risk for multiple sclerosis: a descriptive study. <i>Lancet Neurology</i> , The, 2006, 5, 924-931.	10.2	194
82	Incorporating language phenotypes strengthens evidence of linkage to autism. <i>American Journal of Medical Genetics Part A</i> , 2001, 105, 539-547.	2.4	192
83	Multitrait analysis of glaucoma identifies new risk loci and enables polygenic prediction of disease susceptibility and progression. <i>Nature Genetics</i> , 2020, 52, 160-166.	21.4	192
84	A noise-reduction GWAS analysis implicates altered regulation of neurite outgrowth and guidance in autism. <i>Molecular Autism</i> , 2011, 2, 1.	4.9	191
85	Whole exome sequencing study identifies novel rare and common Alzheimer's-Associated variants involved in immune response and transcriptional regulation. <i>Molecular Psychiatry</i> , 2020, 25, 1859-1875.	7.9	191
86	Common variants near ABCA1, AFAP1 and GMDS confer risk of primary open-angle glaucoma. <i>Nature Genetics</i> , 2014, 46, 1120-1125.	21.4	186
87	The role of the <i>CD58</i> locus in multiple sclerosis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 5264-5269.	7.1	185
88	Construction of a GT polymorphism map of human 9q. <i>Genomics</i> , 1992, 12, 229-240.	2.9	181
89	A novel approach of homozygous haplotype sharing identifies candidate genes in autism spectrum disorder. <i>Human Genetics</i> , 2012, 131, 565-579.	3.8	180
90	Functional Candidate Genes in Age-Related Macular Degeneration: Significant Association with <i>VEGF</i> , <i>VLDLR</i> , and <i>LRP6</i> ., 2006, 47, 329.		178

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91	Protective effect of complement factor B and complement component 2 variants in age-related macular degeneration. <i>Human Molecular Genetics</i> , 2007, 16, 1986-1992.	2.9	175
92	Age-Related Maculopathy: A Genomewide Scan with Continued Evidence of Susceptibility Loci within the 1q31, 10q26, and 17q25 Regions. <i>American Journal of Human Genetics</i> , 2004, 75, 174-189.	6.2	174
93	Assessment of the genetic variance of late-onset Alzheimer's disease. <i>Neurobiology of Aging</i> , 2016, 41, 200.e13-200.e20.	3.1	174
94	Convergent genetic and expression data implicate immunity in Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2015, 11, 658-671.	0.8	173
95	Genome- and Phenome-Wide Analyses of Cardiac Conduction Identifies Markers of Arrhythmia Risk. <i>Circulation</i> , 2013, 127, 1377-1385.	1.6	167
96	Effects of Multiple Genetic Loci on Age at Onset in Late-Onset Alzheimer Disease. <i>JAMA Neurology</i> , 2014, 71, 1394.	9.0	166
97	Rapidly progressive Alzheimer's disease features distinct structures of amyloid- β . <i>Brain</i> , 2015, 138, 1009-1022.	7.6	166
98	Transethnic genome-wide scan identifies novel Alzheimer's disease loci. <i>Alzheimer's and Dementia</i> , 2017, 13, 727-738.	0.8	166
99	Evidence for Polygenic Susceptibility to Multiple Sclerosis "The Shape of Things to Come. <i>American Journal of Human Genetics</i> , 2010, 86, 621-625.	6.2	162
100	Neurofibromatosis 2: Clinical and DNA Linkage Studies of a Large Kindred. <i>New England Journal of Medicine</i> , 1988, 319, 278-283.	27.0	161
101	The Next PAGE in Understanding Complex Traits: Design for the Analysis of Population Architecture Using Genetics and Epidemiology (PAGE) Study. <i>American Journal of Epidemiology</i> , 2011, 174, 849-859.	3.4	161
102	A highly polymorphic locus very tightly linked to the Huntington's disease gene. <i>Nature</i> , 1988, 332, 734-736.	27.8	159
103	Evaluation of copy number variations reveals novel candidate genes in autism spectrum disorder-associated pathways. <i>Human Molecular Genetics</i> , 2012, 21, 3513-3523.	2.9	158
104	Identification of a rare coding variant in complement 3 associated with age-related macular degeneration. <i>Nature Genetics</i> , 2013, 45, 1375-1379.	21.4	158
105	A second major histocompatibility complex susceptibility locus for multiple sclerosis. <i>Annals of Neurology</i> , 2007, 61, 228-236.	5.3	156
106	Common variants near CAV1 and CAV2 are associated with primary open-angle glaucoma in Caucasians from the USA. <i>Human Molecular Genetics</i> , 2011, 20, 4707-4713.	2.9	156
107	Whole-Exome Sequencing Links a Variant in DHDDS to Retinitis Pigmentosa. <i>American Journal of Human Genetics</i> , 2011, 88, 201-206.	6.2	155
108	Gene-Wide Analysis Detects Two New Susceptibility Genes for Alzheimer's Disease. <i>PLoS ONE</i> , 2014, 9, e94661.	2.5	155

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109	Linkage Disequilibrium at the Angelman Syndrome Gene UBE3A in Autism Families. <i>Genomics</i> , 2001, 77, 105-113.	2.9	154
110	Meta-analysis of the Association Between Variants in SORL1 and Alzheimer Disease. <i>Archives of Neurology</i> , 2011, 68, 99.	4.5	153
111	A Genomewide Scan for Early-Onset Coronary Artery Disease in 438 Families: The GENECARD Study. <i>American Journal of Human Genetics</i> , 2004, 75, 436-447.	6.2	152
112	Localization of the gene for familial dysautonomia on chromosome 9 and definition of DNA markers for genetic diagnosis. <i>Nature Genetics</i> , 1993, 4, 160-164.	21.4	149
113	SORL1 Is Genetically Associated with Late-Onset Alzheimer's Disease in Japanese, Koreans and Caucasians. <i>PLoS ONE</i> , 2013, 8, e58618.	2.5	149
114	Polygenic Overlap Between C-Reactive Protein, Plasma Lipids, and Alzheimer Disease. <i>Circulation</i> , 2015, 131, 2061-2069.	1.6	145
115	Novel late-onset Alzheimer disease loci variants associate with brain gene expression. <i>Neurology</i> , 2012, 79, 221-228.	1.1	144
116	Novel Alzheimer Disease Risk Loci and Pathways in African American Individuals Using the African Genome Resources Panel. <i>JAMA Neurology</i> , 2021, 78, 102.	9.0	144
117	Rapid Communication: Cu/Zn Superoxide Dismutase Activity in Familial and Sporadic Amyotrophic Lateral Sclerosis. <i>Journal of Neurochemistry</i> , 1994, 62, 384-387.	3.9	143
118	Investigation of autism and GABA receptor subunit genes in multiple ethnic groups. <i>Neurogenetics</i> , 2006, 7, 167-174.	1.4	141
119	Common variants in Alzheimer's disease and risk stratification by polygenic risk scores. <i>Nature Communications</i> , 2021, 12, 3417.	12.8	140
120	A pooled case-control study of the apolipoprotein E (APOE) gene in age-related maculopathy. <i>Ophthalmic Genetics</i> , 2002, 23, 209-223.	1.2	136
121	A linkage disequilibrium map of the 14cMb 15q12 GABA _A receptor subunit cluster and association to autism. <i>American Journal of Medical Genetics Part A</i> , 2004, 131B, 51-59.	2.4	135
122	Localization of the huntington's disease gene to a small segment of chromosome 4 flanked by D4S10 and the telomere. <i>Cell</i> , 1987, 50, 565-571.	28.9	130
123	Genome-wide and Ordered-Subset linkage analyses provide support for autism loci on 17q and 19p with evidence of phenotypic and interlocus genetic correlates. <i>BMC Medical Genetics</i> , 2005, 6, 1.	2.1	130
124	Dementia Revealed: Novel Chromosome 6 Locus for Late-Onset Alzheimer Disease Provides Genetic Evidence for Folate-Pathway Abnormalities. <i>PLoS Genetics</i> , 2010, 6, e1001130.	3.5	130
125	Recurrent Tissue-Specific mtDNA Mutations Are Common in Humans. <i>PLoS Genetics</i> , 2013, 9, e1003929.	3.5	130
126	Imputation and quality control steps for combining multiple genome-wide datasets. <i>Frontiers in Genetics</i> , 2014, 5, 370.	2.3	130

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127	Mitochondrial haplogroups and peripheral neuropathy during antiretroviral therapy: an adult AIDS clinical trials group study. <i>Aids</i> , 2005, 19, 1341-1349.	2.2	129
128	Association of Polymorphisms in the Apolipoprotein E Region with Susceptibility to and Progression of Multiple Sclerosis. <i>American Journal of Human Genetics</i> , 2002, 70, 708-717.	6.2	125
129	Genetic susceptibility to Alzheimer disease. <i>Trends in Genetics</i> , 1995, 11, 504-508.	6.7	120
130	C3 R102G polymorphism increases risk of age-related macular degeneration. <i>Human Molecular Genetics</i> , 2008, 17, 1821-1824.	2.9	120
131	New insights into the genetics of primary open-angle glaucoma based on meta-analyses of intraocular pressure and optic disc characteristics.. <i>Human Molecular Genetics</i> , 2017, 26, ddw399.	2.9	120
132	Identification of Two Novel Loci for Dominantly Inherited Familial Amyotrophic Lateral Sclerosis. <i>American Journal of Human Genetics</i> , 2003, 73, 397-403.	6.2	119
133	Identification of Genomic Predictors of Atrioventricular Conduction. <i>Circulation</i> , 2010, 122, 2016-2021.	1.6	117
134	Ancestral origin of ApoE ϵ 4 Alzheimer disease risk in Puerto Rican and African American populations. <i>PLoS Genetics</i> , 2018, 14, e1007791.	3.5	117
135	A rare mutation in UNC5C predisposes to late-onset Alzheimer's disease and increases neuronal cell death. <i>Nature Medicine</i> , 2014, 20, 1452-1457.	30.7	116
136	An ϵ -2-macroglobulin insertion-deletion polymorphism in Alzheimer disease. <i>Nature Genetics</i> , 1999, 22, 19-21.	21.4	115
137	Replication analysis identifies TYK2 as a multiple sclerosis susceptibility factor. <i>European Journal of Human Genetics</i> , 2009, 17, 1309-1313.	2.8	115
138	Low-Frequency and Rare-Coding Variation Contributes to Multiple Sclerosis Risk. <i>Cell</i> , 2018, 175, 1679-1687.e7.	28.9	115
139	Distribution ofWDR36DNA Sequence Variants in Patients with Primary Open-Angle Glaucoma. , 2006, 47, 2542.		114
140	Genetic association study of exfoliation syndrome identifies a protective rare variant at LOXL1 and five new susceptibility loci. <i>Nature Genetics</i> , 2017, 49, 993-1004.	21.4	114
141	Exploratory Subsetting of Autism Families Based on Savant Skills Improves Evidence of Genetic Linkage to 15q11-q13. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2003, 42, 856-863.	0.5	112
142	Genome-wide association study identifies seven novel susceptibility loci for primary open-angle glaucoma. <i>Human Molecular Genetics</i> , 2018, 27, 1486-1496.	2.9	111
143	A comparison of neurological, metabolic, structural, and genetic evaluations in persons at risk for Huntington's disease. <i>Annals of Neurology</i> , 1990, 28, 614-621.	5.3	110
144	Peakwide Mapping on Chromosome 3q13 Identifies the Kalirin Gene as a Novel Candidate Gene for Coronary Artery Disease. <i>American Journal of Human Genetics</i> , 2007, 80, 650-663.	6.2	110

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145	Missense variant in TREML2 protects against Alzheimer's disease. <i>Neurobiology of Aging</i> , 2014, 35, 1510.e19-1510.e26.	3.1	110
146	Genetic Variants Associated with Optic Nerve Vertical Cup-to-Disc Ratio Are Risk Factors for Primary Open Angle Glaucoma in a US Caucasian Population. , 2011, 52, 1788.		109
147	Comprehensive follow-up of the first genome-wide association study of multiple sclerosis identifies KIF21B and TMEM39A as susceptibility loci. <i>Human Molecular Genetics</i> , 2010, 19, 953-962.	2.9	108
148	Endothelial Nitric Oxide Synthase Gene Variants and Primary Open-Angle Glaucoma: Interactions with Sex and Postmenopausal Hormone Use. , 2010, 51, 971.		107
149	Association of MAPT haplotypes with Alzheimer's disease risk and MAPT brain gene expression levels. <i>Alzheimer's Research and Therapy</i> , 2014, 6, 39.	6.2	106
150	DNA sequence variants in the LOXL1 gene are associated with pseudoexfoliation glaucoma in a U.S. clinic-based population with broad ethnic diversity. <i>BMC Medical Genetics</i> , 2008, 9, 5.	2.1	105
151	A common variant near TGFB3 is associated with primary open angle glaucoma. <i>Human Molecular Genetics</i> , 2015, 24, 3880-3892.	2.9	105
152	SORCS1 alters amyloid precursor protein processing and variants may increase Alzheimer's disease risk. <i>Annals of Neurology</i> , 2011, 69, 47-64.	5.3	104
153	Dissecting the genetic relationship between cardiovascular risk factors and Alzheimer's disease. <i>Acta Neuropathologica</i> , 2019, 137, 209-226.	7.7	100
154	Ordered-Subsets Linkage Analysis Detects Novel Alzheimer Disease Loci on Chromosomes 2q34 and 15q22. <i>American Journal of Human Genetics</i> , 2003, 73, 1041-1051.	6.2	99
155	Mitochondrial DNA Polymorphism A4917G Is Independently Associated with Age-Related Macular Degeneration. <i>PLoS ONE</i> , 2008, 3, e2091.	2.5	99
156	Association Between Apolipoprotein E Genotype and Alzheimer Disease in African American Subjects. <i>Archives of Neurology</i> , 2002, 59, 594.	4.5	98
157	Comprehensive Search for Alzheimer Disease Susceptibility Loci in the APOE Region. <i>Archives of Neurology</i> , 2012, 69, 1270.	4.5	97
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