

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	Genetic variants in the <i>SHISA6</i> gene are associated with delayed cognitive impairment in two family datasets. <i>Alzheimer's and Dementia</i> , 2023, 19, 611-620.	0.8	4
2	Genome-wide association and multi-omics studies identify <i>MGMT</i> as a novel risk gene for Alzheimer's disease among women. <i>Alzheimer's and Dementia</i> , 2023, 19, 896-908.	0.8	19
3	The National Institute on Aging Late-Onset Alzheimer's Disease Family Based Study: A resource for genetic discovery. <i>Alzheimer's and Dementia</i> , 2022, 18, 1889-1897.	0.8	9
4	Protein phosphatase 2A and complement component 4 are linked to the protective effect of <i>APOE</i> ϵ 2 for Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2022, 18, 2042-2054.	0.8	18
5	Progranulin mutations in clinical and neuropathological Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2022, 18, 2458-2467.	0.8	12
6	An association test of the spatial distribution of rare missense variants within protein structures identifies Alzheimer's disease-related patterns. <i>Genome Research</i> , 2022, 32, 778-790.	5.5	5
7	Genome-wide association study of brain arteriolosclerosis. <i>Journal of Cerebral Blood Flow and Metabolism</i> , 2022, 42, 1437-1450.	4.3	2
8	Admixture Mapping of Alzheimer's disease in Caribbean Hispanics identifies a new locus on 22q13.1. <i>Molecular Psychiatry</i> , 2022, 27, 2813-2820.	7.9	12
9	Genetic architecture of RNA editing regulation in Alzheimer's disease across diverse ancestral populations. <i>Human Molecular Genetics</i> , 2022, 31, 2876-2886.	2.9	2
10	New insights into the genetic etiology of Alzheimer's disease and related dementias. <i>Nature Genetics</i> , 2022, 54, 412-436.	21.4	700
11	Manifestations of Alzheimer's disease genetic risk in the blood are evident in a multiomic analysis in healthy adults aged 18 to 90. <i>Scientific Reports</i> , 2022, 12, 6117.	3.3	12
12	Artificial intelligence-based strategies to identify patient populations and advance analysis in age-related macular degeneration clinical trials. <i>Experimental Eye Research</i> , 2022, 220, 109092.	2.6	2
13	The genetic architecture of Alzheimer disease risk in the Ohio and Indiana Amish. <i>Human Genetics and Genomics Advances</i> , 2022, 3, 100114.	1.7	1
14	Sex differences in the genetic architecture of cognitive resilience to Alzheimer's disease. <i>Brain</i> , 2022, 145, 2541-2554.	7.6	26
15	Effect of OCT B-Scan Density on Sensitivity for Detection of Intraretinal Hyperreflective Foci in Eyes with Age-Related Macular Degeneration. <i>Current Eye Research</i> , 2022, 47, 1294-1299.	1.5	3
16	A locus at 19q13.31 significantly reduces the ApoE ϵ 4 risk for Alzheimer's Disease in African Ancestry. <i>PLoS Genetics</i> , 2022, 18, e1009977.	3.5	19
17	Reproducibility of qualitative assessment of drusen volume in eyes with age related macular degeneration. <i>Eye</i> , 2021, 35, 2594-2600.	2.1	13
18	Causal Associations Between Modifiable Risk Factors and the Alzheimer's Phenome. <i>Annals of Neurology</i> , 2021, 89, 54-65.	5.3	82

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19	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
20	Lower Levels of Education Are Associated with Cognitive Impairment in the Old Order Amish. <i>Journal of Alzheimer's Disease</i> , 2021, 79, 451-458.	2.6	8
21	The GGLEAM Study: Understanding Glaucoma in the Ohio Amish. <i>International Journal of Environmental Research and Public Health</i> , 2021, 18, 1551.	2.6	0
22	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
23	Association of Rare <i>CYP39A1</i> Variants With Exfoliation Syndrome Involving the Anterior Chamber of the Eye. <i>JAMA - Journal of the American Medical Association</i> , 2021, 325, 753.	7.4	16
24	Increased <i>APOE</i> ϵ 4 expression is associated with the difference in Alzheimer's disease risk from diverse ancestral backgrounds. <i>Alzheimer's and Dementia</i> , 2021, 17, 1179-1188.	0.8	33
25	Automated identification of clinical features from sparsely annotated 3-dimensional medical imaging. <i>Npj Digital Medicine</i> , 2021, 4, 44.	10.9	16
26	Using the PhenX Toolkit to Select Standard Measurement Protocols for Your Research Study. <i>Current Protocols</i> , 2021, 1, e149.	2.9	16
27	Polygenic Risk Score for Alzheimer's Disease in Caribbean Hispanics. <i>Annals of Neurology</i> , 2021, 90, 366-376.	5.3	15
28	Common variants in Alzheimer's disease and risk stratification by polygenic risk scores. <i>Nature Communications</i> , 2021, 12, 3417.	12.8	140
29	Association of mitochondrial variants and haplogroups identified by whole exome sequencing with Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2021, . .	0.8	9
30	Linkage of Alzheimer disease families with Puerto Rican ancestry identifies a chromosome 9 locus. <i>Neurobiology of Aging</i> , 2021, 104, 115.e1-115.e7.	3.1	4
31	COMPARISON OF SPECTRALIS AND CIRRUS OPTICAL COHERENCE TOMOGRAPHY FOR THE DETECTION OF INCOMPLETE AND COMPLETE RETINAL PIGMENT EPITHELIUM AND OUTER RETINAL ATROPHY. <i>Retina</i> , 2021, 41, 1851-1857.	1.7	5
32	Association of Smoking, Alcohol Consumption, Blood Pressure, Body Mass Index, and Glycemic Risk Factors With Age-Related Macular Degeneration. <i>JAMA Ophthalmology</i> , 2021, 139, 1299.	2.5	29
33	Plasma Metabolomics of Intermediate and Neovascular Age-Related Macular Degeneration Patients. <i>Cells</i> , 2021, 10, 3141.	4.1	13
34	Large-scale sequencing studies expand the known genetic architecture of Alzheimer's disease. <i>Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring</i> , 2021, 13, e12255.	2.4	4
35	Assessment of AD-related plasma biomarkers in diverse ancestral populations. <i>Alzheimer's and Dementia</i> , 2021, 17, .	0.8	0
36	Does higher educational attainment influence functional capabilities among African Americans with Alzheimer's disease?. <i>Alzheimer's and Dementia</i> , 2021, 17, .	0.8	0

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37	Association of a locus on chromosome 17 with earlier age at onset of cognitive impairment in a familial Amish dataset. <i>Alzheimer's and Dementia</i> , 2021, 17, e056288.	0.8	0
38	Genome-wide association for protective variants in Alzheimer's disease in the Midwestern Amish. <i>Alzheimer's and Dementia</i> , 2021, 17, e056363.	0.8	0
39	Preferential preservation of constructional praxis delayed recall compared to word list delayed recall in the Amish. <i>Alzheimer's and Dementia</i> , 2021, 17, e056386.	0.8	0
40	APOE ϵ -stratified genome-wide association analysis identifies novel Alzheimer disease candidate risk loci for African Americans. <i>Alzheimer's and Dementia</i> , 2021, 17, e056383.	0.8	2
41	Genetic risk score for Alzheimer's disease in the Amish highlights differences in the genetic architecture compared to other European ancestry populations.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e053304.	0.8	0
42	Genome-wide association and multi-omics studies identify MGMT as a novel risk gene for Alzheimer disease among women.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e054483.	0.8	0
43	Multiple viruses detected in human DNA are associated with Alzheimer disease risk.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e054585.	0.8	0
44	Sex differences in the genetic architecture underlying resilience in AD.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e055010.	0.8	0
45	Exome sequencing identifies rare damaging variants in the ATB8B4 and ABCA1 genes as novel risk factors for Alzheimer's disease.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e055982.	0.8	1
46	Sex-specific genetic predictors of memory performance.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e056083.	0.8	0
47	Expression quantitative trait loci (eQTL) analysis in a diverse Alzheimer disease cohort reveals ancestry-specific regulatory architectures.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e056211.	0.8	0
48	Suggestive linkage and association of preserved cognition to chromosome 18 in genetically at-risk Amish.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e056306.	0.8	0
49	Admixture mapping identifies novel regions influencing Alzheimer disease in African Americans.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e056443.	0.8	0
50	Genome-wide association study of cognitive status and decline in the Amish.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e056525.	0.8	0
51	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
52	Segregation, linkage, GWAS, and sequencing. , 2020, , 7-23.		0
53	CHOROIDAL VASCULARITY INDEX AND CHOROIDAL THICKNESS IN EYES WITH RETICULAR PSEUDODRUSEN. <i>Retina</i> , 2020, 40, 612-617.	1.7	40
54	Statistical driver genes as a means to uncover missing heritability for age-related macular degeneration. <i>BMC Medical Genomics</i> , 2020, 13, 95.	1.5	0

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55	Analysis of brain region-specific co-expression networks reveals clustering of established and novel genes associated with Alzheimer disease. <i>Alzheimer's Research and Therapy</i> , 2020, 12, 103.	6.2	9
56	Genetic variants and functional pathways associated with resilience to Alzheimer's disease. <i>Brain</i> , 2020, 143, 2561-2575.	7.6	93
57	Comparative trans-ethnic meta-analysis of whole exome sequencing variation for Alzheimer's disease (AD) in 18,402 individuals of the Alzheimer's Disease Sequencing Project (ADSP). <i>Alzheimer's and Dementia</i> , 2020, 16, e041583.	0.8	0
58	Sex differences in genetic predictors of resilience to Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2020, 16, e043259.	0.8	0
59	Using linkage analysis to identify novel gene-gene interactions in Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2020, 16, e043435.	0.8	1
60	Longitudinal assessment of cognitive decline in the Amish. <i>Alzheimer's and Dementia</i> , 2020, 16, e043440.	0.8	0
61	Genome-wide meta-analysis of late-onset Alzheimer's disease using rare variant imputation in 65,602 subjects identifies risk loci with roles in memory, neurodevelopment, and cardiometabolic traits: The international genomics of Alzheimer's project (IGAP). <i>Alzheimer's and Dementia</i> , 2020, 16, e044193.	0.8	1
62	Mechanism for the protective effect of APOE ϵ 2 against Alzheimer disease is linked to tau and the classical complement pathway. <i>Alzheimer's and Dementia</i> , 2020, 16, e044881.	0.8	0
63	Search for protective genetic variants in Alzheimer disease in the U.S. Midwestern Amish. <i>Alzheimer's and Dementia</i> , 2020, 16, e045350.	0.8	0
64	A multiancestry analysis of Alzheimer's disease coexpressed gene networks identifies a common immune signaling pathway regulated by granulocyte-colony stimulating factor (G-CSF). <i>Alzheimer's and Dementia</i> , 2020, 16, e045361.	0.8	0
65	Increased <i>APOE4</i> expression is associated with reactive A1 astrocytes and may confer the difference in Alzheimer disease risk from different ancestral backgrounds. <i>Alzheimer's and Dementia</i> , 2020, 16, e045415.	0.8	0
66	Assessing whole genome sequencing variation for Alzheimer's disease in 4707 individuals from the Alzheimer's Disease Sequencing Project (ADSP). <i>Alzheimer's and Dementia</i> , 2020, 16, e045548.	0.8	0
67	Transcriptomic characterization of a Puerto Rican Alzheimer disease cohort implicates convergent immune-related pathways. <i>Alzheimer's and Dementia</i> , 2020, 16, e045890.	0.8	0
68	Mapping Alzheimer disease-associated regions in the African American population. <i>Alzheimer's and Dementia</i> , 2020, 16, e046072.	0.8	0
69	Education and its effect on risk and age at onset in Alzheimer disease (AD) in African Americans. <i>Alzheimer's and Dementia</i> , 2020, 16, e046078.	0.8	0
70	iPSC-derived neurons and microglia with an African-specific ABCA7 frameshift deletion have impaired function. <i>Alzheimer's and Dementia</i> , 2020, 16, e046109.	0.8	1
71	Genome-wide interaction study of smoking in Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2020, 16, e046149.	0.8	0
72	Structural characterization of rare missense variants within known neurodegenerative disease proteins. <i>Alzheimer's and Dementia</i> , 2020, 16, e046405.	0.8	0

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73	Joint linkage and association mapping of preserved cognition in the old-order Amish. <i>Alzheimer's and Dementia</i> , 2020, 16, e046416.	0.8	0
74	Use of local genetic ancestry to assess <i>TOMM40</i> -523 and risk for Alzheimer disease. <i>Neurology: Genetics</i> , 2020, 6, e404.	1.9	12
75	Association of <i>APOE</i> With Primary Open-Angle Glaucoma Suggests a Protective Effect for <i>APOE</i> ϵ 4. <i>Investigative Ophthalmology and Visual Science</i> , 2020, 61, 3.		23
76	Immune and Inflammatory Pathways Implicated by Whole Blood Transcriptomic Analysis in a Diverse Ancestry Alzheimer's Disease Cohort. <i>Journal of Alzheimer's Disease</i> , 2020, 76, 1047-1060.	2.6	6
77	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
78	Exceptionally low likelihood of Alzheimer's dementia in <i>APOE2</i> homozygotes from a 5,000-person neuropathological study. <i>Nature Communications</i> , 2020, 11, 667.	12.8	246
79	Hadoop and PySpark for reproducibility and scalability of genomic sequencing studies. <i>Pacific Symposium on Biocomputing</i> , 2020, 25, 523-534.	0.7	0
80	AMISH EYE STUDY. <i>Retina</i> , 2019, 39, 1540-1550.	1.7	17
81	Rare variants and loci for age-related macular degeneration in the Ohio and Indiana Amish. <i>Human Genetics</i> , 2019, 138, 1171-1182.	3.8	7
82	<i>APOE</i> Promoter Polymorphism-219T/G is an Effect Modifier of the Influence of <i>APOE</i> ϵ 4 on Alzheimer's Disease Risk in a Multiracial Sample. <i>Journal of Clinical Medicine</i> , 2019, 8, 1236.	2.4	40
83	Association of a Primary Open-Angle Glaucoma Genetic Risk Score With Earlier Age at Diagnosis. <i>JAMA Ophthalmology</i> , 2019, 137, 1190.	2.5	32
84	Sex differences in the genetic predictors of Alzheimer's pathology. <i>Brain</i> , 2019, 142, 2581-2589.	7.6	65
85	Association of Genetic Variants With Primary Open-Angle Glaucoma Among Individuals With African Ancestry. <i>JAMA - Journal of the American Medical Association</i> , 2019, 322, 1682.	7.4	50
86	Education Moderates the Relation Between <i>APOE</i> ϵ 4 and Memory in Nondemented Non-Hispanic Black Older Adults. <i>Journal of Alzheimer's Disease</i> , 2019, 72, 495-506.	2.6	14
87	Pathway Analysis Integrating Genome-Wide and Functional Data Identifies <i>PLCG2</i> as a Candidate Gene for Age-Related Macular Degeneration. <i>Investigative Ophthalmology and Visual Science</i> , 2019, 60, 4041.		10
88	Multiple sclerosis genomic map implicates peripheral immune cells and microglia in susceptibility. <i>Science</i> , 2019, 365, .	12.6	710
89	Genetic Correlations Between Diabetes and Glaucoma: An Analysis of Continuous and Dichotomous Phenotypes. <i>American Journal of Ophthalmology</i> , 2019, 206, 245-255.	3.3	12
90	Analysis of Whole-Exome Sequencing Data for Alzheimer Disease Stratified by <i>APOE</i> Genotype. <i>JAMA Neurology</i> , 2019, 76, 1099.	9.0	32

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91	CpG-related SNPs in the MS4A region have a dose-dependent effect on risk of late-onset Alzheimer disease. <i>Aging Cell</i> , 2019, 18, e12964.	6.7	8
92	RNA editing alterations in a multi-ethnic Alzheimer disease cohort converge on immune and endocytic molecular pathways. <i>Human Molecular Genetics</i> , 2019, 28, 3053-3061.	2.9	19
93	Association of Rare Coding Mutations With Alzheimer Disease and Other Dementias Among Adults of European Ancestry. <i>JAMA Network Open</i> , 2019, 2, e191350.	5.9	58
94	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
95	O3131: HIGHLY PENETRANT LATE-ONSET ALZHEIMER DISEASE VARIANTS IN NOTCH3 IN ASHKENAZI JEWS. <i>Alzheimer's and Dementia</i> , 2019, 15, P918.	0.8	0
96	Dissecting the genetic relationship between cardiovascular risk factors and Alzheimer's disease. <i>Acta Neuropathologica</i> , 2019, 137, 209-226.	7.7	100
97	A rare missense variant of <i>CASP7</i> is associated with familial late-onset Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2019, 15, 441-452.	0.8	39
98	Genetic Variation in Genes Underlying Diverse Dementias May Explain a Small Proportion of Cases in the Alzheimer's Disease Sequencing Project. <i>Dementia and Geriatric Cognitive Disorders</i> , 2018, 45, 1-17.	1.5	22
99	Whole genome sequencing of Caribbean Hispanic families with late-onset Alzheimer's disease. <i>Annals of Clinical and Translational Neurology</i> , 2018, 5, 406-417.	3.7	42
100	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
101	Analysis combining correlated glaucoma traits identifies five new risk loci for open-angle glaucoma. <i>Scientific Reports</i> , 2018, 8, 3124.	3.3	33
102	Functional annotation of genomic variants in studies of late-onset Alzheimer's disease. <i>Bioinformatics</i> , 2018, 34, 2724-2731.	4.1	30
103	Identification of rare noncoding sequence variants in gamma-aminobutyric acid A receptor, alpha 4 subunit in autism spectrum disorder. <i>Neurogenetics</i> , 2018, 19, 17-26.	1.4	5
104	Rare genetic variation implicated in non-Hispanic white families with Alzheimer disease. <i>Neurology: Genetics</i> , 2018, 4, e286.	1.9	27
105	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
106	The Carnitine Shuttle Pathway is Altered in Patients With Neovascular Age-Related Macular Degeneration. , 2018, 59, 4978.		37
107	Low-Frequency and Rare-Coding Variation Contributes to Multiple Sclerosis Risk. <i>Cell</i> , 2018, 175, 1679-1687.e7.	28.9	115
108	Alzheimer Disease: Perspectives from Epidemiology and Genetics. <i>Journal of Law, Medicine and Ethics</i> , 2018, 46, 694-698.	0.9	33

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109	Pedigree Selection and Information Content. <i>Current Protocols in Human Genetics</i> , 2018, 97, e56.	3.5	3
110	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
111	Cross-ancestry genome-wide association analysis of corneal thickness strengthens link between complex and Mendelian eye diseases. <i>Nature Communications</i> , 2018, 9, 1864.	12.8	63
112	One for all and all for One: Improving replication of genetic studies through network diffusion. <i>PLoS Genetics</i> , 2018, 14, e1007306.	3.5	22
113	Sex-specific genetic predictors of Alzheimer's disease biomarkers. <i>Acta Neuropathologica</i> , 2018, 136, 857-872.	7.7	87
114	Family-Based Genome-Wide Association Study of South Indian Pedigrees Supports <i>WNT7B</i> as a Central Corneal Thickness Locus. , 2018, 59, 2495.		11
115	Genome-wide pleiotropy analysis of neuropathological traits related to Alzheimer's disease. <i>Alzheimer's Research and Therapy</i> , 2018, 10, 22.	6.2	27
116	Sex-Specific Association of Apolipoprotein E With Cerebrospinal Fluid Levels of Tau. <i>JAMA Neurology</i> , 2018, 75, 989.	9.0	223
117	Male-specific epistasis between <i>WWC1</i> and <i>TLN2</i> genes is associated with Alzheimer's disease. <i>Neurobiology of Aging</i> , 2018, 72, 188.e3-188.e12.	3.1	24
118	Testosterone Pathway Genetic Polymorphisms in Relation to Primary Open-Angle Glaucoma: An Analysis in Two Large Datasets. , 2018, 59, 629.		14
119	Assay for Transposase-Accessible Chromatin Using Sequencing (ATAC-seq) Data Analysis. <i>Current Protocols in Human Genetics</i> , 2017, 92, 20.4.1-20.4.13.	3.5	9
120	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
121	A Common Variant of <i>IL-6R</i> is Associated with Elevated <i>IL-6</i> Pathway Activity in Alzheimer's Disease Brains. <i>Journal of Alzheimer's Disease</i> , 2017, 56, 1037-1054.	2.6	44
122	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
123	Transethnic genome-wide scan identifies novel Alzheimer's disease loci. <i>Alzheimer's and Dementia</i> , 2017, 13, 727-738.	0.8	166
124	A common haplotype lowers <i>PU.1</i> expression in myeloid cells and delays onset of Alzheimer's disease. <i>Nature Neuroscience</i> , 2017, 20, 1052-1061.	14.8	330
125	Genetic association study of exfoliation syndrome identifies a protective rare variant at <i>LOXL1</i> and five new susceptibility loci. <i>Nature Genetics</i> , 2017, 49, 993-1004.	21.4	114
126	Genetic correlations between intraocular pressure, blood pressure and primary open-angle glaucoma: a multi-cohort analysis. <i>European Journal of Human Genetics</i> , 2017, 25, 1261-1267.	2.8	18

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127	The molecular genetics of eye diseases. <i>Human Molecular Genetics</i> , 2017, 26, R1-R1.	2.9	5
128	Rare coding variants in <i>PLCG2</i> , <i>ABI3</i> , and <i>TREM2</i> implicate microglial-mediated innate immunity in Alzheimer's disease. <i>Nature Genetics</i> , 2017, 49, 1373-1384.	21.4	783
129	A population-specific reference panel empowers genetic studies of Anabaptist populations. <i>Scientific Reports</i> , 2017, 7, 6079.	3.3	16
130	Early-Onset Alzheimer Disease and Candidate Risk Genes Involved in Endolysosomal Transport. <i>JAMA Neurology</i> , 2017, 74, 1113.	9.0	41
131	Age at natural menopause genetic risk score in relation to age at natural menopause and primary open-angle glaucoma in a US-based sample. <i>Menopause</i> , 2017, 24, 150-156.	2.0	6
132	[O1â€“03â€“01]: GENOMEâ€“WIDE RARE VARIANT IMPUTATION AND TISSUEâ€“SPECIFIC TRANSCRIPTOMIC ANALYSIS IDENTIFY NOVEL RARE VARIANT CANDIDATE LOCI IN LATEâ€“ONSET ALZHEIMER'S DISEASE: THE ALZHEIMER'S DISEASE GENETICS CONSORTIUM. <i>Alzheimer's and Dementia</i> , 2017, 13, P189.	0.8	4
133	Two novel loci, <i>COBL</i> and <i>SLC10A2</i> , for Alzheimer's disease in African Americans. <i>Alzheimer's and Dementia</i> , 2017, 13, 119-129.	0.8	87
134	[O2â€“08â€“02]: SEXâ€“SPECIFIC ANALYSIS OF THE ADSP CASEâ€“CONTROL WHOLEâ€“EXOME SEQUENCING DATASET. <i>Alzheimer's and Dementia</i> , 2017, 13, P571.	0.8	0
135	[O2â€“08â€“03]: WHOLEâ€“GENOME SEQUENCING IN FAMILIAL LATEâ€“ONSET ALZHEIMER'S DISEASE IDENTIFIES RARE VARIATION IN AD CANDIDATE GENES. <i>Alzheimer's and Dementia</i> , 2017, 13, P571.	0.8	1
136	Joint Analysis of Nuclear and Mitochondrial Variants in Age-Related Macular Degeneration Identifies Novel Loci <i>TRPM1</i> and <i>ABHD2/RLBP1</i> . , 2017, 58, 4027.		21
137	Caspase-8, association with Alzheimer's Disease and functional analysis of rare variants. <i>PLoS ONE</i> , 2017, 12, e0185777.	2.5	38
138	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
139	Introducing COCOS: codon consequence scanner for annotating reading frame changes induced by stop-lost and frame shift variants. <i>Bioinformatics</i> , 2017, 33, btw820.	4.1	2
140	Progression Rate From Intermediate to Advanced Age-Related Macular Degeneration Is Correlated With the Number of Risk Alleles at the CFH Locus. , 2016, 57, 6107.		18
141	Genetic Association Analysis of Drusen Progression. , 2016, 57, 2225.		12
142	The Application of Genetic Risk Scores in Age-Related Macular Degeneration: A Review. <i>Journal of Clinical Medicine</i> , 2016, 5, 31.	2.4	31
143	A Common Variant in <i>MIR182</i> Is Associated With Primary Open-Angle Glaucoma in the NEIGHBORHOOD Consortium. , 2016, 57, 4528.		42
144	Assessing the Association of Mitochondrial Genetic Variation With Primary Open-Angle Glaucoma Using Gene-Set Analyses. , 2016, 57, 5046.		44

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145	DNA variants in <i>CACNA1C</i> modify Parkinson disease risk only when vitamin D level is deficient. <i>Neurology: Genetics</i> , 2016, 2, e72.	1.9	11
146	Heritability of Choroidal Thickness in the Amish. <i>Ophthalmology</i> , 2016, 123, 2537-2544.	5.2	24
147	P3082: Assessment of the Genetic Variance of Late-Onset Alzheimer's Disease. <i>Alzheimer's and Dementia</i> , 2016, 12, P849.	0.8	0
148	O10302: <i>ABCA7</i> Frameshift Deletion Associated with Alzheimer's Disease in African Americans. <i>Alzheimer's and Dementia</i> , 2016, 12, P177.	0.8	0
149	O1-03-03: Identification of Novel Candidate Genes for Early-Onset Alzheimer's Disease Through Integrated Whole-Exome Sequencing and Exome Chip Array Association Analysis. , 2016, 12, P177-P178.		2
150	O1-03-05: High-Resolution Imputation in Genome-Wide Association Studies of Late-Onset Alzheimer's Disease Identifies Novel Rare Variant Associations. , 2016, 12, P178-P179.		0
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