

Gerard D Schellenberg

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7714836/publications.pdf>

Version: 2024-02-01

176
papers

21,872
citations

30047

54
h-index

15249

126
g-index

195
all docs

195
docs citations

195
times ranked

25029
citing authors

#	ARTICLE	IF	CITATIONS
1	Meta-analysis of 74,046 individuals identifies 11 new susceptibility loci for Alzheimer's disease. <i>Nature Genetics</i> , 2013, 45, 1452-1458.	9.4	3,741
2	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.	9.4	1,962
3	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.	9.4	1,676
4	Clinical diagnosis of progressive supranuclear palsy: The movement disorder society criteria. <i>Movement Disorders</i> , 2017, 32, 853-864.	2.2	1,402
5	Tau is a candidate gene for chromosome 17 frontotemporal dementia. <i>Annals of Neurology</i> , 1998, 43, 815-825.	2.8	1,257
6	A framework for the interpretation of de novo mutation in human disease. <i>Nature Genetics</i> , 2014, 46, 944-950.	9.4	943
7	Convergence of Genes and Cellular Pathways Dysregulated in Autism Spectrum Disorders. <i>American Journal of Human Genetics</i> , 2014, 94, 677-694.	2.6	819
8	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.	9.4	783
9	New insights into the genetic etiology of Alzheimer's disease and related dementias. <i>Nature Genetics</i> , 2022, 54, 412-436.	9.4	700
10	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.	4.9	376
11	Common polygenic variation enhances risk prediction for Alzheimer's disease. <i>Brain</i> , 2015, 138, 3673-3684.	3.7	359
12	Neurodegeneration and defective neurotransmission in a <i>Caenorhabditis elegans</i> model of tauopathy. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003, 100, 9980-9985.	3.3	336
13	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.	7.1	330
14	Genetic assessment of age-associated Alzheimer disease risk: Development and validation of a polygenic hazard score. <i>PLoS Medicine</i> , 2017, 14, e1002258.	3.9	311
15	Genome-Wide Association Meta-analysis of Neuropathologic Features of Alzheimer's Disease and Related Dementias. <i>PLoS Genetics</i> , 2014, 10, e1004606.	1.5	305
16	Exceptionally low likelihood of Alzheimer's dementia in APOE2 homozygotes from a 5,000-person neuropathological study. <i>Nature Communications</i> , 2020, 11, 667.	5.8	246
17	Sex-Specific Association of Apolipoprotein E With Cerebrospinal Fluid Levels of Tau. <i>JAMA Neurology</i> , 2018, 75, 989.	4.5	223
18	The genetics and neuropathology of Alzheimer's disease. <i>Acta Neuropathologica</i> , 2012, 124, 305-323.	3.9	203

#	ARTICLE	IF	CITATIONS
19	Genome-wide association study identifies four novel loci associated with Alzheimer's endophenotypes and disease modifiers. <i>Acta Neuropathologica</i> , 2017, 133, 839-856.	3.9	199
20	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.	4.1	191
21	Assessment of the genetic variance of late-onset Alzheimer's disease. <i>Neurobiology of Aging</i> , 2016, 41, 200.e13-200.e20.	1.5	174
22	Convergent genetic and expression data implicate immunity in Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2015, 11, 658-671.	0.4	173
23	Genome-wide association study of corticobasal degeneration identifies risk variants shared with progressive supranuclear palsy. <i>Nature Communications</i> , 2015, 6, 7247.	5.8	170
24	Effects of Multiple Genetic Loci on Age at Onset in Late-Onset Alzheimer Disease. <i>JAMA Neurology</i> , 2014, 71, 1394.	4.5	166
25	Gene-Wide Analysis Detects Two New Susceptibility Genes for Alzheimer's Disease. <i>PLoS ONE</i> , 2014, 9, e94661.	1.1	155
26	Association Between Genetic Traits for Immune-Mediated Diseases and Alzheimer Disease. <i>JAMA Neurology</i> , 2016, 73, 691.	4.5	151
27	Genomic variants, genes, and pathways of Alzheimer's disease: An overview. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017, 174, 5-26.	1.1	147
28	Polygenic Overlap Between C-Reactive Protein, Plasma Lipids, and Alzheimer Disease. <i>Circulation</i> , 2015, 131, 2061-2069.	1.6	145
29	Novel Alzheimer Disease Risk Loci and Pathways in African American Individuals Using the African Genome Resources Panel. <i>JAMA Neurology</i> , 2021, 78, 102.	4.5	144
30	Common variants in Alzheimer's disease and risk stratification by polygenic risk scores. <i>Nature Communications</i> , 2021, 12, 3417.	5.8	140
31	Transmission of tauopathy strains is independent of their isoform composition. <i>Nature Communications</i> , 2020, 11, 7.	5.8	121
32	Ancestral origin of ApoE ϵ 4 Alzheimer disease risk in Puerto Rican and African American populations. <i>PLoS Genetics</i> , 2018, 14, e1007791.	1.5	117
33	Immune-related genetic enrichment in frontotemporal dementia: An analysis of genome-wide association studies. <i>PLoS Medicine</i> , 2018, 15, e1002487.	3.9	111
34	Missense variant in TREML2 protects against Alzheimer's disease. <i>Neurobiology of Aging</i> , 2014, 35, 1510.e19-1510.e26.	1.5	110
35	A novel mutation at position +12 in the intron following Exon 10 of the tau gene in familial frontotemporal dementia (FTD-Kumamoto). <i>Annals of Neurology</i> , 2000, 47, 422-429.	2.8	109
36	Genetic architecture of sporadic frontotemporal dementia and overlap with Alzheimer's and Parkinson's diseases. <i>Journal of Neurology, Neurosurgery and Psychiatry</i> , 2017, 88, 152-164.	0.9	107

#	ARTICLE	IF	CITATIONS
37	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.	3.0	106
38	Dissecting the genetic relationship between cardiovascular risk factors and Alzheimer's disease. <i>Acta Neuropathologica</i> , 2019, 137, 209-226.	3.9	100
39	Two rare <i>AKAP9</i> variants are associated with Alzheimer's disease in African Americans. <i>Alzheimer's and Dementia</i> , 2014, 10, 609.	0.4	94
40	Genetic variants and functional pathways associated with resilience to Alzheimer's disease. <i>Brain</i> , 2020, 143, 2561-2575.	3.7	93
41	Shared genetic risk between corticobasal degeneration, progressive supranuclear palsy, and frontotemporal dementia. <i>Acta Neuropathologica</i> , 2017, 133, 825-837.	3.9	90
42	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.	0.4	87
43	Sex-specific genetic predictors of Alzheimer's disease biomarkers. <i>Acta Neuropathologica</i> , 2018, 136, 857-872.	3.9	87
44	SUT-2 potentiates tau-induced neurotoxicity in <i>Caenorhabditis elegans</i> . <i>Human Molecular Genetics</i> , 2009, 18, 1825-1838.	1.4	86
45	Apolipoprotein E genotypes in Parkinson's disease with and without dementia. <i>Annals of Neurology</i> , 1995, 37, 242-245.	2.8	82
46	Polygenic hazard score: an enrichment marker for Alzheimer's associated amyloid and tau deposition. <i>Acta Neuropathologica</i> , 2018, 135, 85-93.	3.9	80
47	Selective Genetic Overlap Between Amyotrophic Lateral Sclerosis and Diseases of the Frontotemporal Dementia Spectrum. <i>JAMA Neurology</i> , 2018, 75, 860.	4.5	79
48	Examination of genetic linkage of chromosome 15 to schizophrenia in a large Veterans Affairs Cooperative Study sample. <i>American Journal of Medical Genetics Part A</i> , 2001, 105, 662-668.	2.4	75
49	Association of an apolipoprotein CII allele with familial dementia of the Alzheimer type. <i>Journal of Neurogenetics</i> , 1987, 4, 97-108.	0.6	73
50	CXCR4 involvement in neurodegenerative diseases. <i>Translational Psychiatry</i> , 2018, 8, 73.	2.4	66
51	Sex differences in the genetic predictors of Alzheimer's pathology. <i>Brain</i> , 2019, 142, 2581-2589.	3.7	65
52	Polygenic hazard score, amyloid deposition and Alzheimer's neurodegeneration. <i>Brain</i> , 2019, 142, 460-470.	3.7	63
53	SUT-1 enables tau-induced neurotoxicity in <i>C. elegans</i> . <i>Human Molecular Genetics</i> , 2007, 16, 1959-1971.	1.4	62
54	<i>APOE</i> mRNA and protein expression in postmortem brain are modulated by an extended haplotype structure. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2010, 153B, 409-417.	1.1	62

#	ARTICLE	IF	CITATIONS
55	Tau-Mediated NMDA Receptor Impairment Underlies Dysfunction of a Selectively Vulnerable Network in a Mouse Model of Frontotemporal Dementia. <i>Journal of Neuroscience</i> , 2014, 34, 16482-16495.	1.7	60
56	Genome-wide, high-content siRNA screening identifies the Alzheimer's genetic risk factor FERMT2 as a major modulator of APP metabolism. <i>Acta Neuropathologica</i> , 2017, 133, 955-966.	3.9	60
57	Presenilin-1 mutation alters NGF-induced neurite outgrowth, calcium homeostasis, and transcription factor (AP-1) activation in PC12 cells. <i>Journal of Neuroscience Research</i> , 1998, 52, 618-624.	1.3	59
58	Association of Rare Coding Mutations With Alzheimer Disease and Other Dementias Among Adults of European Ancestry. <i>JAMA Network Open</i> , 2019, 2, e191350.	2.8	58
59	Missense mutations in the chromosome 14 familial Alzheimer's disease presenilin 1 gene. <i>Human Mutation</i> , 1998, 11, 216-221.	1.1	54
60	Antisense-mediated Exon Skipping Decreases Tau Protein Expression: A Potential Therapy For Tauopathies. <i>Molecular Therapy - Nucleic Acids</i> , 2014, 3, e180.	2.3	54
61	Replication of progressive supranuclear palsy genome-wide association study identifies SLCO1A2 and DUSP10 as new susceptibility loci. <i>Molecular Neurodegeneration</i> , 2018, 13, 37.	4.4	54
62	Modulation of the age at onset of Parkinson's disease by apolipoprotein E genotypes. <i>Annals of Neurology</i> , 1997, 42, 655-658.	2.8	52
63	C9orf72 intermediate repeats are associated with corticobasal degeneration, increased C9orf72 expression and disruption of autophagy. <i>Acta Neuropathologica</i> , 2019, 138, 795-811.	3.9	50
64	Gene expression, methylation and neuropathology correlations at progressive supranuclear palsy risk loci. <i>Acta Neuropathologica</i> , 2016, 132, 197-211.	3.9	49
65	Polygenic hazard scores in preclinical Alzheimer disease. <i>Annals of Neurology</i> , 2017, 82, 484-488.	2.8	49
66	<i>PARK10</i> is a major locus for sporadic neuropathologically confirmed Parkinson disease. <i>Neurology</i> , 2015, 84, 972-980.	1.5	48
67	Rare Functional Variant in TM2D3 is Associated with Late-Onset Alzheimer's Disease. <i>PLoS Genetics</i> , 2016, 12, e1006327.	1.5	47
68	INFERNO: inferring the molecular mechanisms of noncoding genetic variants. <i>Nucleic Acids Research</i> , 2018, 46, 8740-8753.	6.5	46
69	Sex-dependent autosomal effects on clinical progression of Alzheimer's disease. <i>Brain</i> , 2020, 143, 2272-2280.	3.7	46
70	Association of an apolipoprotein CII allele with familial dementia of the Alzheimer type. <i>Journal of Neurogenetics</i> , 1987, 4, 97-108.	0.6	44
71	Genetic and neuroanatomic associations in sporadic frontotemporal lobar degeneration. <i>Neurobiology of Aging</i> , 2014, 35, 1473-1482.	1.5	43
72	Global and local ancestry in African-Americans: Implications for Alzheimer's disease risk. <i>Alzheimer's and Dementia</i> , 2016, 12, 233-243.	0.4	42

#	ARTICLE	IF	CITATIONS
73	Rarity of the Alzheimer Diseaseâ€“Protective <i>APP</i> A673T Variant in the United States. <i>JAMA Neurology</i> , 2015, 72, 209.	4.5	41
74	Early-Onset Alzheimer Disease and Candidate Risk Genes Involved in Endolysosomal Transport. <i>JAMA Neurology</i> , 2017, 74, 1113.	4.5	41
75	Association of Long Runs of Homozygosity With Alzheimer Disease Among African American Individuals. <i>JAMA Neurology</i> , 2015, 72, 1313.	4.5	39
76	Discovery of gene-gene interactions across multiple independent data sets of late onset Alzheimer disease from the Alzheimer Disease Genetics Consortium. <i>Neurobiology of Aging</i> , 2016, 38, 141-150.	1.5	39
77	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.	0.4	39
78	Caspase-8, association with Alzheimerâ€™s Disease and functional analysis of rare variants. <i>PLoS ONE</i> , 2017, 12, e0185777.	1.1	38
79	Analysis of Whole-Exome Sequencing Data for Alzheimer Disease Stratified by <i>APOE</i> Genotype. <i>JAMA Neurology</i> , 2019, 76, 1099.	4.5	32
80	Identification of seven novel loci associated with amino acid levels using single-variant and gene-based tests in 8545 Finnish men from the METSIM study. <i>Human Molecular Genetics</i> , 2018, 27, 1664-1674.	1.4	30
81	Activity of the poly(A) binding protein MSUT2 determines susceptibility to pathological tau in the mammalian brain. <i>Science Translational Medicine</i> , 2019, 11, .	5.8	30
82	In vitro amplification of pathogenic tau conserves disease-specific bioactive characteristics. <i>Acta Neuropathologica</i> , 2021, 141, 193-215.	3.9	30
83	Genome-wide pleiotropy analysis of neuropathological traits related to Alzheimerâ€™s disease. <i>Alzheimer's Research and Therapy</i> , 2018, 10, 22.	3.0	27
84	TSC1 loss increases risk for tauopathy by inducing tau acetylation and preventing tau clearance via chaperone-mediated autophagy. <i>Science Advances</i> , 2021, 7, eabg3897.	4.7	27
85	Quality control and integration of genotypes from two calling pipelines for whole genome sequence data in the Alzheimer's disease sequencing project. <i>Genomics</i> , 2019, 111, 808-818.	1.3	26
86	Sex differences in the genetic architecture of cognitive resilience to Alzheimerâ€™s disease. <i>Brain</i> , 2022, 145, 2541-2554.	3.7	26
87	Genome-wide linkage analyses of non-Hispanic white families identify novel loci for familial late-onset Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2016, 12, 2-10.	0.4	24
88	NIAGADS: The NIA Genetics of Alzheimer's Disease Data Storage Site. <i>Alzheimer's and Dementia</i> , 2016, 12, 1200-1203.	0.4	24
89	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.	1.5	24
90	Gene-based analysis in HRC imputed genome wide association data identifies three novel genes for Alzheimerâ€™s disease. <i>PLoS ONE</i> , 2019, 14, e0218111.	1.1	23

#	ARTICLE	IF	CITATIONS
91	VCPA: genomic variant calling pipeline and data management tool for Alzheimer's Disease Sequencing Project. <i>Bioinformatics</i> , 2019, 35, 1768-1770.	1.8	23
92	One for all and all for One: Improving replication of genetic studies through network diffusion. <i>PLoS Genetics</i> , 2018, 14, e1007306.	1.5	22
93	Inferring the Molecular Mechanisms of Noncoding Alzheimer's Disease-Associated Genetic Variants. <i>Journal of Alzheimer's Disease</i> , 2019, 72, 301-318.	1.2	19
94	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.	0.4	19
95	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.	1.5	19
96	High copy wildtype human 1N4R tau expression promotes early pathological tauopathy accompanied by cognitive deficits without progressive neurofibrillary degeneration. <i>Acta Neuropathologica Communications</i> , 2015, 3, 33.	2.4	18
97	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.	0.4	18
98	Progranulin mutations in clinical and neuropathological Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2022, 18, 2458-2467.	0.4	12
99	A scoring strategy combining statistics and functional genomics supports a possible role for common polygenic variation in autism. <i>Frontiers in Genetics</i> , 2014, 5, 33.	1.1	10
100	Association of mitochondrial variants and haplogroups identified by whole exome sequencing with Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2021, , .	0.4	9
101	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.	3.0	8
102	Evidence against DNA polymerase γ as a candidate gene for Werner syndrome. <i>Human Genetics</i> , 1994, 93, 507-12.	1.8	7
103	Genetic influences on cognition in progressive supranuclear palsy. <i>Movement Disorders</i> , 2017, 32, 1764-1771.	2.2	6
104	Impact of apolipoprotein E genotypes on vitamin E and memantine treatment outcomes in Alzheimer's disease. <i>Alzheimer's and Dementia: Translational Research and Clinical Interventions</i> , 2018, 4, 344-349.	1.8	6
105	Insoluble Tau From Human FTDP-17 Cases Exhibit Unique Transmission Properties In Vivo. <i>Journal of Neuropathology and Experimental Neurology</i> , 2020, 79, 941-949.	0.9	6
106	A novel mutation at position +12 in the intron following Exon 10 of the tau gene in familial frontotemporal dementia (FTD-Kumamoto). <i>Annals of Neurology</i> , 2000, 47, 422-429.	2.8	6
107	LRP10 variants in progressive supranuclear palsy. <i>Neurobiology of Aging</i> , 2020, 94, 311.e5-311.e10.	1.5	6
108	Latent trait modeling of tau neuropathology in progressive supranuclear palsy. <i>Acta Neuropathologica</i> , 2021, 141, 667-680.	3.9	5

#	ARTICLE	IF	CITATIONS
109	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.	2.4	5
110	Integration of GWAS and brain transcriptomic analyses in a multiethnic sample of 35,245 older adults identifies <i>DCDC2</i> gene as predictor of episodic memory maintenance. <i>Alzheimer's and Dementia</i> , 2022, 18, 1797-1811.	0.4	5
111	THE FREQUENCY OF C4B VARIANTS OF COMPLEMENT IN FAMILIAL AND SPORADIC ALZHEIMER DISEASE. <i>Alzheimer Disease and Associated Disorders</i> , 1987, 1, 251-255.	0.6	4
112	[O1-03-01]: GENOME-WIDE RARE VARIANT IMPUTATION AND TISSUE-SPECIFIC TRANSCRIPTOMIC ANALYSIS IDENTIFY NOVEL RARE VARIANT CANDIDATE LOCI IN LATE-ONSET ALZHEIMER'S DISEASE: THE ALZHEIMER'S DISEASE GENETICS CONSORTIUM. <i>Alzheimer's and Dementia</i> , 2017, 13, P189.	0.4	4
113	Fibrillation and molecular characteristics are coherent with clinical and pathological features of 4-repeat tauopathy caused by MAPT variant G273R. <i>Neurobiology of Disease</i> , 2020, 146, 105079.	2.1	4
114	Copy Number Variation Identification on 3,800 Alzheimer's Disease Whole Genome Sequencing Data from the Alzheimer's Disease Sequencing Project. <i>Frontiers in Genetics</i> , 2021, 12, 752390.	1.1	4
115	Alzheimer's Disease Variant Portal: A Catalog of Genetic Findings for Alzheimer's Disease. <i>Journal of Alzheimer's Disease</i> , 2022, 86, 461-477.	1.2	4
116	Neuropathological lesions and their contribution to dementia and cognitive impairment in a heterogeneous clinical population. <i>Alzheimer's and Dementia</i> , 2022, 18, 2403-2412.	0.4	4
117	The Alzheimer's disease sequencing project follow up study (ADSP-FUS): Increasing ethnic diversity in Alzheimer's genetics research with addition of potential new cohorts. <i>Alzheimer's and Dementia</i> , 2020, 16, e046400.	0.4	3
118	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
119	Genome-wide association study of brain arteriolosclerosis. <i>Journal of Cerebral Blood Flow and Metabolism</i> , 2022, 42, 1437-1450.	2.4	2
120	APOE-stratified genome-wide association analysis identifies novel Alzheimer disease candidate risk loci for African Americans. <i>Alzheimer's and Dementia</i> , 2021, 17, e056383.	0.4	2
121	P1-045: EXOME ARRAY ANALYSIS IDENTIFIES NOVEL RISK VARIANTS FOR ALZHEIMER'S DISEASE WITH ONSET BEFORE 65 YEARS. , 2014, 10, P319-P319.		1
122	P1-018: Rare Deleterious And Loss-of-Function Variants in <i>OPRL1</i> and <i>GAS2L2</i> Contribute to the Risk of Late-Onset Alzheimer's Disease: Alzheimer's Disease Sequencing Project Case-Control Study. <i>Alzheimer's and Dementia</i> , 2016, 12, P406.	0.4	1
123	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.	0.4	1
124	NIA genetics of Alzheimer's disease data storage site (NIAGADS): Update 2020. <i>Alzheimer's and Dementia</i> , 2020, 16, e044284.	0.4	1
125	Gene-Environment Interactions in Progressive Supranuclear Palsy. <i>Frontiers in Neurology</i> , 2021, 12, 664796.	1.1	1
126	The Seattle Alzheimer's disease data set. <i>Genetic Epidemiology</i> , 1993, 10, 365-369.	0.6	0

#	ARTICLE	IF	CITATIONS
127	Regulation of Four-Repeat tau Expression: Interactions between Exon and Intron Splicing Regulatory Sequences. , 0, , 87-95.		0
128	F1â€01â€02: Alzheimer's Disease Sequencing Project: Search for Alzheimer's Disease Resilience Genes That May Modify Disease Susceptibility in Specific Apoe Genotype Backgrounds. Alzheimer's and Dementia, 2016, 12, P162.	0.4	0
129	P2â€097: The Alzheimerâ€™s Disease Sequencing Project (ADSP): Data Production, Management, and Availability. Alzheimer's and Dementia, 2016, 12, P648.	0.4	0
130	P2â€077: Alzheimer's Disease Sequencing Project: Search for Alzheimer's Disease Resilience Genes That May Modify Disease Susceptibility in Specific <i>Apoe</i> Genotype Backgrounds. Alzheimer's and Dementia, 2016, 12, P638.	0.4	0
131	P2-083: Computational Identification of Regulatory Mechanisms Affected By Noncoding Variants Associated with Late-Onset Alzheimer's Disease. , 2016, 12, P640-P641.		0
132	P2â€085: Further Stratification of <i>APOE</i> E4â€Negative Subjects Identifies Novel Genes for Alzheimer's Disease. Alzheimer's and Dementia, 2016, 12, P641.	0.4	0
133	P3â€082: Assessment of the Genetic Variance of Lateâ€Onset Alzheimerâ€™s Disease. Alzheimer's and Dementia, 2016, 12, P849.	0.4	0
134	P3-093: NIA Genetics of Alzheimerâ€™s Disease Data Storage Site (NIAGADS): 2016 Update. , 2016, 12, P855-P856.		0
135	P3â€096: Secondary Analyses of International Genomics of Alzheimer's Project Stage I GWAS Summary Data Identifies Additional Variants Associated With Lateâ€Onset Alzheimer's Disease. Alzheimer's and Dementia, 2016, 12, P856.	0.4	0
136	P4â€048: Convergent Analysis of Endophenotypes in Progressive Supranuclear Palsy. Alzheimer's and Dementia, 2016, 12, P1032.	0.4	0
137	F1-01-03: Rare Deleterious and Loss-of-Function Variants in OPRL1 and GAS2L2 Contribute to the Risk of Late-Onset Alzheimerâ€™s Disease: Alzheimerâ€™s Disease Sequencing Project Case-Control Study. , 2016, 12, P163-P163.		0
138	O1â€03â€02: <i>ABCA7</i> Frameshift Deletion Associated with Alzheimerâ€™s Disease in African Americans. Alzheimer's and Dementia, 2016, 12, P177.	0.4	0
139	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
140	O1â€09â€03: Whole Genome Sequencing in Familial Lateâ€Onset Alzheimerâ€™s Disease Identifies Variations in TTC3 and FSIP2. Alzheimer's and Dementia, 2016, 12, P197.	0.4	0
141	O2â€06â€03: Tissueâ€Specific Genomeâ€Wide Predictions of Genetically Regulated Expression in Alzheimer's Disease. Alzheimer's and Dementia, 2016, 12, P239.	0.4	0
142	O2â€10â€06: A Common Allele in <i>SPI1</i> Lowers Risk and Delays Age at Onset for Alzheimer's Disease. Alzheimer's and Dementia, 2016, 12, P253.	0.4	0
143	P1â€122: Multivariate Phenotypes Association Study of Neuropathological Features of Alzheimer's Disease and Related Dementias. Alzheimer's and Dementia, 2016, 12, P450.	0.4	0
144	P1-117: Blood Gene Expression Changes Implicated in Alzheimerâ€™s Disease. , 2016, 12, P448-P448.		0

#	ARTICLE	IF	CITATIONS
145	P1â€129: Structural Variation (SV) in Heterogenous Wholeâ€Genome Sequencing Data from 111 Families at Risk For Alzheimer's Disease: Alzheimer's Disease Sequencing Project SV Study. Alzheimer's and Dementia, 2016, 12, P453.	0.4	0
146	[P3â€097]: NIA GENETICS OF ALZHEIMER'S DISEASE DATA STORAGE SITE (NIAGADS): 2017. Alzheimer's and Dementia, 2017, 13, P971.	0.4	0
147	[P4â€074]: INTEGRATIVE SYSTEMS BIOLOGY APPROACH TO IDENTIFY NOVEL RISK FACTORS FOR PSP. Alzheimer's and Dementia, 2017, 13, P1286.	0.4	0
148	[P3â€090]: THE ALZHEIMER'S DISEASE SEQUENCING PROJECT (ADSP) DATA UPDATE 2017. Alzheimer's and Dementia, 2017, 13, P968.	0.4	0
149	P4â€044: THE GCAD CLOUDâ€BASED WORKFLOW FOR PROCESSING WHOLE EXOME AND WHOLE GENOME DATA FROM THE ALZHEIMER'S DISEASE SEQUENCING PROJECT. Alzheimer's and Dementia, 2018, 14, P1450.	0.4	0
150	P2â€171: THE POLY(A) BINDING PROTEIN MSUT2 MODULATES GLIOSIS IN TAUOPATHY. Alzheimer's and Dementia, 2018, 14, P734.	0.4	0
151	P1â€156: GENEâ€BASED ANALYSES IN WHOLE GENOME SEQUENCING OF FAMILIAL LATEâ€ONSET ALZHEIMER'S DISEASE. Alzheimer's and Dementia, 2018, 14, P336.	0.4	0
152	P2â€279: CSF SMALL RNA BIOMARKERS FOR ALZHEIMER'S DISEASE. Alzheimer's and Dementia, 2018, 14, P785.	0.4	0
153	P2â€106: AFRICAN AMERICAN WHOLE EXOME SEQUENCING SUGGESTS RISK CODING VARIANTS IN IDH1 GENE. Alzheimer's and Dementia, 2018, 14, P709.	0.4	0
154	P1â€139: THE CONTRIBUTION OF SEXâ€SPECIFIC ASSOCIATIONS IN GENETIC STUDIES OF ALZHEIMER'S DISEASE PATHOLOGY. Alzheimer's and Dementia, 2018, 14, P327.	0.4	0
155	P2â€108: WHOLEâ€GENOME SEQUENCING IN NONâ€HISPANIC WHITE FAMILIES IMPLICATES RARE VARIATION IN LATEâ€ONSET ALZHEIMER'S DISEASE RISK. Alzheimer's and Dementia, 2018, 14, P710.	0.4	0
156	O3â€13â€01: HIGHLY PENETRANT LATEâ€ONSET ALZHEIMER DISEASE VARIANTS IN NOTCH3 IN ASHKENAZI JEWS. Alzheimer's and Dementia, 2019, 15, P918.	0.4	0
157	Sex differences in genetic predictors of resilience to Alzheimerâ€™s disease. Alzheimer's and Dementia, 2020, 16, e043259.	0.4	0
158	Alzheimerâ€™s disease variant portal (ADVP): Harmonized genetics data and evidence collection for Alzheimerâ€™s disease. Alzheimer's and Dementia, 2020, 16, e044090.	0.4	0
159	Genomeâ€wide profiling of the noncoding regulatory mechanisms in Alzheimerâ€™s disease. Alzheimer's and Dementia, 2020, 16, e044268.	0.4	0
160	Mechanism for the protective effect of APOE Î¼2 against Alzheimer disease is linked to tau and the classical complement pathway. Alzheimer's and Dementia, 2020, 16, e044881.	0.4	0
161	Assessing whole genome sequencing variation for Alzheimerâ€™s disease in 4707 individuals from the Alzheimerâ€™s Disease Sequencing Project (ADSP). Alzheimer's and Dementia, 2020, 16, e045548.	0.4	0
162	Pleiotropy analyses using TADs identify genomic regions affecting risk of AD and stroke. Alzheimer's and Dementia, 2020, 16, e045975.	0.4	0

#	ARTICLE	IF	CITATIONS
163	Multimodal genome-wide meta-analysis of brain amyloidosis reveals heterogeneity across CSF, PET, and pathological amyloid measures. <i>Alzheimer's and Dementia</i> , 2020, 16, e046009.	0.4	0
164	Mapping Alzheimer disease-associated regions in the African American population. <i>Alzheimer's and Dementia</i> , 2020, 16, e046072.	0.4	0
165	Genome-wide interaction study of smoking in Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2020, 16, e046149.	0.4	0
166	Heritability analyses show partial genetic overlap between (non-Mendelian) early and late onset Alzheimer disease due to an intriguing APOE effect. <i>Alzheimer's and Dementia</i> , 2021, 17, e056143.	0.4	0
167	NIA genetics of Alzheimer's disease data storage site (NIAGADS): 2021 update.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e052258.	0.4	0
168	Copy number variation (CNV) identification and association study on 3,928 Alzheimer's disease whole genome sequencing data from the Alzheimer's Disease Sequencing Project (ADSP).. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e052721.	0.4	0
169	Characterization of regulatory roles of genetic signals curated from more than 200 GWA studies in the Alzheimer's Disease Variant Portal (ADVP).. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e054255.	0.4	0
170	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.	0.4	0
171	Multiple viruses detected in human DNA are associated with Alzheimer disease risk.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e054585.	0.4	0
172	Sex differences in the genetic architecture underlying resilience in AD.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e055010.	0.4	0
173	Sex-specific genetic predictors of memory performance.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e056083.	0.4	0
174	The Alzheimer's Disease Sequencing Project - Follow Up Study (ADSP-FUS): Increasing ethnic diversity in Alzheimer's genetics research with the addition of potential new cohorts.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e056101.	0.4	0
175	Admixture mapping identifies novel regions influencing Alzheimer disease in African Americans.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e056443.	0.4	0
176	A large-scale, whole genome sequencing study of unexplained early-onset Alzheimer disease.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e056664.	0.4	0