Lindsay A Farrer

List of Publications by Year
in descending order

376	43,251	3933 88	2828 191
papers	citations	h-index	g-index

413	413	413	39928
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Midlife lipid and glucose levels are associated with Alzheimer's disease. Alzheimer's and Dementia, 2023, 19, 181-193.	0.8	23
2	Genomeâ€wide association and multiâ€omics studies identify MGMT as a novel risk gene for Alzheimer's disease among women. Alzheimer's and Dementia, 2023, 19, 896-908.	0.8	19
3	Enhancing Discovery of Genetic Variants for Posttraumatic Stress Disorder Through Integration of Quantitative Phenotypes and Trauma Exposure Information. Biological Psychiatry, 2022, 91, 626-636.	1.3	21
4	Protein phosphatase 2A and complement component 4 are linked to the protective effect of APOE É2 for Alzheimer's disease. Alzheimer's and Dementia, 2022, 18, 2042-2054.	0.8	18
5	Blood and brain transcriptome analysis reveals APOE genotype-mediated and immune-related pathways involved in Alzheimer disease. Alzheimer's Research and Therapy, 2022, 14, 30.	6.2	16
6	An association test of the spatial distribution of rare missense variants within protein structures identifies Alzheimer's diseaseâ€"related patterns. Genome Research, 2022, 32, 778-790.	5.5	5
7	Genome-wide association study of brain arteriolosclerosis. Journal of Cerebral Blood Flow and Metabolism, 2022, 42, 1437-1450.	4.3	2
8	New insights into the genetic etiology of Alzheimerâ€™s disease and related dementias. Nature Genetics, 2022, 54, 412-436.	21.4	700
9	Alzheimerâ€™s disease associated AKAP9 I2558M mutation alters posttranslational modification and interactome of tau and cellular functions in CRISPRâ€edited human neuronal cells. Aging Cell, 2022, 21, e13617.	6.7	7
10	Sex differences in the genetic architecture of cognitive resilience to Alzheimerâ€™s disease. Brain, 2022, 145, 2541-2554.	7.6	26
11	Molecular Quantitative Trait Locus Mapping in Human Complex Diseases. Current Protocols, 2022, 2, e426.	2.9	3
12	Analysis of telomere length variation and Shelterin complex subunit gene expression changes in ethanol-exposed human embryonic stem cells. Journal of Psychiatric Research, 2021, 143, 543-549.	3.1	4
13	Causal Associations Between Modifiable Risk Factors and the Alzheimer's Phenome. Annals of Neurology, 2021, 89, 54-65.	5.3	82
14	Novel Alzheimer Disease Risk Loci and Pathways in African American Individuals Using the African Genome Resources Panel. JAMA Neurology, 2021, 78, 102.	9.0	144
15	Genome-wide association study of phenotypes measuring progression from first cocaine or opioid use to dependence reveals novel risk genes. Exploration of Medicine, 2021, 2, 60-73.	1.5	6
16	Set-Based Rare Variant Expression Quantitative Trait Loci in Blood and Brain from Alzheimer Disease Study Participants. Genes, 2021, 12, 419.	2.4	6
17	Cell-type-specific expression quantitative trait loci associated with Alzheimer disease in blood and brain tissue. Translational Psychiatry, 2021, 11, 250.	4.8	29
18	Common variants in Alzheimerâ€™s disease and risk stratification by polygenic risk scores. Nature Communications, 2021, 12, 3417.	12.8	140

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19	Genome-wide association study of stimulant dependence. Translational Psychiatry, 2021, 11, 363.	4.8	4
20	Association of mitochondrial variants and haplogroups identified by whole exome sequencing with Alzheimer's disease. Alzheimer's and Dementia, 2021, , .	0.8	9
21	Integrative brain transcriptome analysis links complement component 4 and HSPA2 to the APOE Îµ2 protective effect in Alzheimer disease. Molecular Psychiatry, 2021, 26, 6054-6064.	7.9	27
22	Exploration of alcohol use disorder-associated brain miRNAâ€“mRNA regulatory networks. Translational Psychiatry, 2021, 11, 504.	4.8	23
23	A missense variant in SHARPIN mediates Alzheimerâ€™s disease-specific brain damages. Translational Psychiatry, 2021, 11, 590.	4.8	10
24	Alzheimerâ€™s disease associated AKAP 9 I2558M mutation alters posttranslational modification and interactome of tau and cellular functions in CRISPRâ€edited human neuronal cells. Alzheimer's and Dementia, 2021, 17, e058592.	0.8	0
25	APOEâ€stratified genomeâ€wide association analysis identifies novel Alzheimer disease candidate risk loci for African Americans. Alzheimer's and Dementia, 2021, 17, e056383.	0.8	2
26	Multivariate analysis of blood and brain transcriptome in Alzheimer's reveals unique APOE Îµ4-related immune pathways.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e054237.	0.8	0
27	Domain specific cognitive functions predict neuropathological traits in the Framingham Heart Study.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e054249.	0.8	0
28	Genome-wide association and multi-omics studies identify MGMT as a novel risk gene for Alzheimer disease among women.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e054483.	0.8	0
29	Multiple viruses detected in human DNA are associated with Alzheimer disease risk.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e054585.	0.8	0
30	Sex differences in the genetic architecture underlying resilience in AD.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e055010.	0.8	0
31	Sex-specific genetic predictors of memory performance.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e056083.	0.8	0
32	Admixture mapping identifies novel regions influencing Alzheimer disease in African Americans.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e056443.	0.8	0
33	Whole exome sequencing study identifies novel rare and common Alzheimerâ€™s-Associated variants involved in immune response and transcriptional regulation. Molecular Psychiatry, 2020, 25, 1859-1875.	7.9	191
34	A large-scale genome-wide association study meta-analysis of cannabis use disorder. Lancet Psychiatry,the, 2020, 7, 1032-1045.	7.4	200
35	Analysis of brain region-specific co-expression networks reveals clustering of established and novel genes associated with Alzheimer disease. Alzheimer's Research and Therapy, 2020, 12, 103.	6.2	9
36	Genetic variants and functional pathways associated with resilience to Alzheimerâ€™s disease. Brain, 2020, 143, 2561-2575.	7.6	93

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37	Novel mechanism underlying the APOE Îµ2 protective effect for Alzheimer disease implicated by integrative genome and transcriptome analysis. Alzheimer's and Dementia, 2020, 16, e040065.	0.8	0
38	Cellâ€type specific eQTLs (ctâ€eQTLs) associated with Alzheimer disease in blood and brain tissue. Alzheimer's and Dementia, 2020, 16, e044149.	0.8	0
39	Genomeâ€wide metaâ€analysis of lateâ€onset Alzheimerâ€™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 Alzheimerâ€™s project (IGAP). Alzheimer's and Dementia, 2020, 16, e044193.	0.8	1
40	Differential effects of apolipoprotein E on the molecular and cellular phenotypes associated with Alzheimerâ€™s disease in isogenic human iPSCâ€derived neurons. Alzheimer's and Dementia, 2020, 16, e044579.	0.8	0
41	Mechanism for the protective effect of APOE Îµ2 against Alzheimer disease is linked to tau and the classical complement pathway. Alzheimer's and Dementia, 2020, 16, e044881.	0.8	0
42	Alzheimerâ€™s disease risk factor mutations in patients diagnosed with Creutzfeltâ€Jakob disease. Alzheimer's and Dementia, 2020, 16, e045035.	0.8	0
43	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.	0.8	0
44	Mapping Alzheimer diseaseâ€“associated regions in the African American population. Alzheimer's and Dementia, 2020, 16, e046072.	0.8	0
45	Earlyâ€mid adulthood measures of HDL, triglycerides and fasting glucose are associated with lateâ€onset Alzheimer disease. Alzheimer's and Dementia, 2020, 16, e046125.	0.8	2
46	Genomeâ€wide interaction study of smoking in Alzheimerâ€™s disease. Alzheimer's and Dementia, 2020, 16, e046149.	0.8	0
47	Structural characterization of rare missense variants within known neurodegenerative disease proteins. Alzheimer's and Dementia, 2020, 16, e046405.	0.8	0
48	Defining Alzheimerâ€™s disease subtypes using polygenic risk scores integrated with genomic and brain transcriptomic profiles. Alzheimer's and Dementia, 2020, 16, e046449.	0.8	3
49	Genome wide association study of chronic traumatic encephalopathy. Alzheimer's and Dementia, 2020, 16, e046505.	0.8	0
50	Expanding the genetic architecture of nicotine dependence and its shared genetics with multiple traits. Nature Communications, 2020, 11, 5562.	12.8	80
51	Genomeâ€wide association study of rate of cognitive decline in Alzheimer's disease patients identifies novel genes and pathways. Alzheimer's and Dementia, 2020, 16, 1134-1145.	0.8	28
52	Association of OPRM1 Functional Coding Variant With Opioid Use Disorder. JAMA Psychiatry, 2020, 77, 1072.	11.0	135
53	Prefrontal cortex eQTLs/mQTLs enriched in genetic variants associated with alcohol use disorder and other diseases. Epigenomics, 2020, 12, 789-800.	2.1	15
54	Sex-dependent autosomal effects on clinical progression of Alzheimerâ€™s disease. Brain, 2020, 143, 2272-2280.	7.6	46

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55	An analysis of the effect of mu-opioid receptor gene (OPRM1) promoter region DNA methylation on the response of naltrexone treatment of alcohol dependence. Pharmacogenomics Journal, 2020, 20, 672-680.	2.0	9
56	Genomic influences on self-reported childhood maltreatment. Translational Psychiatry, 2020, 10, 38.	4.8	47
57	Leveraging genome-wide data to investigate differences between opioid use vs. opioid dependence in 41,176 individuals from the Psychiatric Genomics Consortium. Molecular Psychiatry, 2020, 25, 1673-1687.	7.9	82
58	Genome-Wide Association Study of Opioid Cessation. Journal of Clinical Medicine, 2020, 9, 180.	2.4	17
59	Exceptionally low likelihood of Alzheimerâ€™s dementia in APOE2 homozygotes from a 5,000-person neuropathological study. Nature Communications, 2020, 11, 667.	12.8	246
60	Welcome message from the Editor-in-Chief. , 2020, 1, 1-3.		0
61	Identifying factors associated with opioid cessation in a biracial sample using machine learning. , 2020, 1, 27-41.		1
62	APOE Promoter Polymorphism-219T/G is an Effect Modifier of the Influence of APOE Îµ4 on Alzheimerâ€™s Disease Risk in a Multiracial Sample. Journal of Clinical Medicine, 2019, 8, 1236.	2.4	40
63	A systems biology approach towards understanding and treating non-neovascular age-related macular degeneration. Nature Communications, 2019, 10, 3347.	12.8	192
64	International meta-analysis of PTSD genome-wide association studies identifies sex- and ancestry-specific genetic risk loci. Nature Communications, 2019, 10, 4558.	12.8	363
65	Comparison of methods for multivariate gene-based association tests for complex diseases using common variants. European Journal of Human Genetics, 2019, 27, 811-823.	2.8	24
66	The genetics and epigenetics of Neonatal Abstinence Syndrome. Seminars in Fetal and Neonatal Medicine, 2019, 24, 105-110.	2.3	36
67	Analysis of Whole-Exome Sequencing Data for Alzheimer Disease Stratified by APOE Genotype. JAMA Neurology, 2019, 76, 1099.	9.0	32
68	Salivary microRNAs identified by small RNA sequencing and machine learning as potential biomarkers of alcohol dependence. Epigenomics, 2019, 11, 739-749.	2.1	19
69	CpGâ€related SNPs in the MS4A region have a doseâ€dependent effect on risk of lateâ€“onset Alzheimer disease. Aging Cell, 2019, 18, e12964.	6.7	8
70	The Utah Protocol for Postmortem Eye Phenotyping and Molecular Biochemical Analysis. , 2019, 60, 1204.		25
71	Association of Rare Coding Mutations With Alzheimer Disease and Other Dementias Among Adults of European Ancestry. JAMA Network Open, 2019, 2, e191350.	5.9	58
72	Genetic meta-analysis of diagnosed Alzheimerâ€™s disease identifies new risk loci and implicates AÎ², tau, immunity and lipid processing. Nature Genetics, 2019, 51, 414-430.	21.4	1,962

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73	A regulatory variant of CHRM3 is associated with cannabis-induced hallucinations in European Americans. Translational Psychiatry, 2019, 9, 309.	4.8	3
74	Dissecting the genetic relationship between cardiovascular risk factors and Alzheimerâ€™s disease. Acta Neuropathologica, 2019, 137, 209-226.	7.7	100
75	A rare missense variant of CASP7 is associated with familial lateâ€onset Alzheimer's disease. Alzheimer's and Dementia, 2019, 15, 441-452.	0.8	39
76	Genome-wide Association Study Identifies a Regulatory Variant of RGMA Associated With Opioid Dependence in European Americans. Biological Psychiatry, 2018, 84, 762-770.	1.3	64
77	Genetic Variation in Genes Underlying Diverse Dementias May Explain a Small Proportion of Cases in the Alzheimerâ€™s Disease Sequencing Project. Dementia and Geriatric Cognitive Disorders, 2018, 45, 1-17.	1.5	22
78	FUS Phase Separation Is Modulated by a Molecular Chaperone and Methylation of Arginine Cation-Ï€ Interactions. Cell, 2018, 173, 720-734.e15.	28.9	662
79	Genomeâ€wide association study of Alzheimer's disease endophenotypes at prediagnosis stages. Alzheimer's and Dementia, 2018, 14, 623-633.	0.8	64
80	Tau Phosphorylation is Impacted by Rare AKAP9 Mutations Associated with Alzheimer Disease in African Americans. Journal of NeuroImmune Pharmacology, 2018, 13, 254-264.	4.1	19
81	Rare genetic variation implicated in non-Hispanic white families with Alzheimer disease. Neurology: Genetics, 2018, 4, e286.	1.9	27
82	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. Nature Neuroscience, 2018, 21, 1656-1669.	14.8	490
83	Ancestral origin of ApoE Îµ4 Alzheimer disease risk in Puerto Rican and African American populations. PLoS Genetics, 2018, 14, e1007791.	3.5	117
84	Genome-wide association study identifies glutamate ionotropic receptor GRIA4 as a risk gene for comorbid nicotine dependence and major depression. Translational Psychiatry, 2018, 8, 208.	4.8	14
85	Risk Locus Identification Ties Alcohol Withdrawal Symptoms to <scp>SORCS</scp>2. Alcoholism: Clinical and Experimental Research, 2018, 42, 2337-2348.	2.4	14
86	Variation in TMEM106B in chronic traumatic encephalopathy. Acta Neuropathologica Communications, 2018, 6, 115.	5.2	38
87	One for all and all for One: Improving replication of genetic studies through network diffusion. PLoS Genetics, 2018, 14, e1007306.	3.5	22
88	Analysis of shared heritability in common disorders of the brain. Science, 2018, 360, .	12.6	1,085
89	Genome-wide pleiotropy analysis of neuropathological traits related to Alzheimerâ€™s disease. Alzheimer's Research and Therapy, 2018, 10, 22.	6.2	27
90	Genomeâ€wide association study of cognitive flexibility assessed by the Wisconsin Card Sorting Test. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2018, 177, 511-519.	1.7	4

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91	Male-specific epistasis between WWC1 and TLN2 genes is associated with Alzheimer's disease. Neurobiology of Aging, 2018, 72, 188.e3-188.e12.	3.1	24
92	Targeted Sequencing of Alzheimer Disease Genes in African Americans Implicates Novel Risk Variants. Frontiers in Neuroscience, 2018, 12, 592.	2.8	24
93	Translational studies support a role for serotonin 2B receptor (HTR2B) gene in aggression-related cannabis response. Molecular Psychiatry, 2018, 23, 2277-2286.	7.9	20
94	Genomeâ€wide association study of body mass index in subjects with alcohol dependence. Addiction Biology, 2017, 22, 535-549.	2.6	21
95	Association of maternal and infant variants inPNOCandCOMTgenes with neonatal abstinence syndrome severity. American Journal on Addictions, 2017, 26, 42-49.	1.4	39
96	A Common Variant of IL-6R is Associated with Elevated IL-6 Pathway Activity in Alzheimerâ€™s Disease Brains. Journal of Alzheimer's Disease, 2017, 56, 1037-1054.	2.6	44
97	Genome-wide association study identifies four novel loci associated with Alzheimerâ€™s endophenotypes and disease modifiers. Acta Neuropathologica, 2017, 133, 839-856.	7.7	199
98	Transethnic genomeâ€wide scan identifies novel Alzheimer's disease loci. Alzheimer's and Dementia, 2017, 13, 727-738.	0.8	166
99	A common haplotype lowers PU.1 expression in myeloid cells and delays onset of Alzheimer's disease. Nature Neuroscience, 2017, 20, 1052-1061.	14.8	330
100	Genetic Risk Variants Associated With Comorbid Alcohol Dependence and Major Depression. JAMA Psychiatry, 2017, 74, 1234.	11.0	74
101	Genome-Wide Association Studies of a Broad Spectrum of Antisocial Behavior. JAMA Psychiatry, 2017, 74, 1242.	11.0	174
102	Oxytocin receptor gene polymorphisms, attachment, and PTSD: Results from the National Health and Resilience in Veterans Study. Journal of Psychiatric Research, 2017, 94, 139-147.	3.1	46
103	Rare coding variants in PLCG2, ABI3, and TREM2 implicate microglial-mediated innate immunity in Alzheimer's disease. Nature Genetics, 2017, 49, 1373-1384.	21.4	783
104	Early-Onset Alzheimer Disease and Candidate Risk Genes Involved in Endolysosomal Transport. JAMA Neurology, 2017, 74, 1113.	9.0	41
105	[P3â€“431]: DEEP LEARNING APPLICATION IN IDENTIFYING PROTEOMIC RISK MARKERS FOR ALZHEIMER's DISEASE. Alzheimer's and Dementia, 2017, 13, P1133.	0.8	3
106	[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. Alzheimer's and Dementia, 2017, 13, P189.	0.8	4
107	[O2â€“08â€“04]: NOVEL GENETIC VARIANTS ASSOCIATED WITH FAMILIAL LATEâ€ONSET ALZHEIMER DISEASE IN THE ALZHEIMER's DISEASE SEQUENCING PROJECT. Alzheimer's and Dementia, 2017, 13, P572.	0.8	0
108	Genetic factor common to schizophrenia and HIV infection is associated with risky sexual behavior: antagonistic vs. synergistic pleiotropic SNPs enriched for distinctly different biological functions. Human Genetics, 2017, 136, 75-83.	3.8	17

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109	Genetics of age-related macular degeneration (AMD). Human Molecular Genetics, 2017, 26, R45-R50.	2.9	109
110	Two novel loci, COBL and SLC10A2, for Alzheimer's disease in African Americans. Alzheimer's and Dementia, 2017, 13, 119-129.	0.8	87
111	The Interplay Between Risky Sexual Behaviors and Alcohol Dependence: Genome-Wide Association and Neuroimaging Support for LHPP as a Risk Gene. Neuropsychopharmacology, 2017, 42, 598-605.	5.4	40
112	[P1â€“242]: GENOMEâ€WIDE ASSOCIATION STUDY OF ALZHEIMER DISEASE ENDOPHENOTYPES AT PRECLINICAL AND MCI STAGES. Alzheimer's and Dementia, 2017, 13, P337.	0.8	0
113	[P3â€“092]: TAU PHOSPHORYLATION IS IMPACTED BY RARE ADâ€ASSOCIATED AKAP9 MUTATIONS SPECIFIC TO AFRICAN AMERICANS. Alzheimer's and Dementia, 2017, 13, P969.	0.8	0
114	[O2â€“08â€“02]: SEXâ€SPECIFIC ANALYSIS OF THE ADSP CASEâ€CONTROL WHOLEâ€EXOME SEQUENCING DATASET. Alzheimer's and Dementia, 2017, 13, P571.	0.8	0
115	A phased SNP-based classification of sickle cell anemia HBB haplotypes. BMC Genomics, 2017, 18, 608.	2.8	31
116	Caspase-8, association with Alzheimerâ€™s Disease and functional analysis of rare variants. PLoS ONE, 2017, 12, e0185777.	2.5	38
117	Genetic assessment of age-associated Alzheimer disease risk: Development and validation of a polygenic hazard score. PLoS Medicine, 2017, 14, e1002258.	8.4	311
118	A putative causal relationship between genetically determined female body shape and posttraumatic stress disorder. Genome Medicine, 2017, 9, 99.	8.2	31
119	S100A10 identified in a genome-wide gene Ã— cannabis dependence interaction analysis of risky sexual behaviours. Journal of Psychiatry and Neuroscience, 2017, 42, 252-261.	2.4	9
120	Human induced pluripotent stem cells illuminate pathways and novel treatment targets for age-related macular degeneration. Stem Cell Investigation, 2017, 4, 92-92.	3.0	1
121	Genome-wide Association Study of Cannabis Dependence Severity, Novel Risk Variants, and Shared Genetic Risks. JAMA Psychiatry, 2016, 73, 472.	11.0	148
122	Genomeâ€wide linkage analyses of nonâ€Hispanic white families identify novel loci for familial lateâ€onset Alzheimer's disease. Alzheimer's and Dementia, 2016, 12, 2-10.	0.8	24
123	Assessment of the genetic variance of late-onset Alzheimer's disease. Neurobiology of Aging, 2016, 41, 200.e13-200.e20.	3.1	174
124	Segregation of a rare TTC3 variant in an extended family with late-onset Alzheimer disease. Neurology: Genetics, 2016, 2, e41.	1.9	41
125	Network-driven plasma proteomics expose molecular changes in the Alzheimerâ€™s brain. Molecular Neurodegeneration, 2016, 11, 31.	10.8	34
126	A candidate transacting modulator of fetal hemoglobin gene expression in the Arabâ€”Indian haplotype of sickle cell anemia. American Journal of Hematology, 2016, 91, 1118-1122.	4.1	16

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127	Shared genetic contribution to ischemic stroke and Alzheimer's disease. Annals of Neurology, 2016, 79, 739-747.	5.3	56
128	Polygenic risk for alcohol dependence associates with alcohol consumption, cognitive function and social deprivation in a populationâ€based cohort. Addiction Biology, 2016, 21, 469-480.	2.6	27
129	The ticking clock of Cayo Santiago macaques and its implications for understanding human circadian rhythm disorders. American Journal of Primatology, 2016, 78, 117-126.	1.7	5
130	ABCA7 frameshift deletion associated with Alzheimer disease in African Americans. Neurology: Genetics, 2016, 2, e79.	1.9	74
131	Homozygosity for a haplotype in the HBG2â€OR51B4 region is exclusive to Arabâ€Indian haplotype sickle cell anemia. American Journal of Hematology, 2016, 91, E308-11.	4.1	13
132	Discovery of gene-gene interactions across multiple independent data sets of late onset Alzheimer disease from the Alzheimer Disease Genetics Consortium. Neurobiology of Aging, 2016, 38, 141-150.	3.1	39
133	A large genome-wide association study of age-related macular degeneration highlights contributions of rare and common variants. Nature Genetics, 2016, 48, 134-143.	21.4	1,167
134	Global and local ancestry in Africanâ€Americans: Implications for Alzheimer's disease risk. Alzheimer's and Dementia, 2016, 12, 233-243.	0.8	42
135	Estimating the Risk for Conversion from Mild Cognitive Impairment to Alzheimer's Disease in an Elderly Arab Community. Journal of Alzheimer's Disease, 2015, 45, 865-871.	2.6	12
136	Genomewide Association Study for Maximum Number of Alcoholic Drinks in European Americans and African Americans. Alcoholism: Clinical and Experimental Research, 2015, 39, 1137-1147.	2.4	58
137	Further analyses support the association between light eye color and alcohol dependence. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2015, 168, 757-760.	1.7	1
138	Reply. Annals of Neurology, 2015, 78, 836-837.	5.3	0
139	Ancestry of the Timorese: age-related macular degeneration associated genotype and allele sharing among human populations from throughout the world. Frontiers in Genetics, 2015, 6, 238.	2.3	9
140	Rarity of the Alzheimer Diseaseâ€“ProtectiveAPPA673T Variant in the United States. JAMA Neurology, 2015, 72, 209.	9.0	41
141	Linkage analyses in Caribbean Hispanic families identify novel loci associated with familial lateâ€onset Alzheimer's disease. Alzheimer's and Dementia, 2015, 11, 1397-1406.	0.8	24
142	Variations in opioid receptor genes in neonatal abstinence syndrome. Drug and Alcohol Dependence, 2015, 155, 253-259.	3.2	55
143	Genome-Wide Association Study of Nicotine Dependence in American Populations: Identification of Novel Risk Loci in Both African-Americans and European-Americans. Biological Psychiatry, 2015, 77, 493-503.	1.3	78
144	Protective variant for hippocampal atrophy identified by whole exome sequencing. Annals of Neurology, 2015, 77, 547-552.	5.3	48

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145	Expanding the genomic roadmap of Alzheimer's disease. Lancet Neurology, The, 2015, 14, 783-785.	10.2	11
146	Eye color: A potential indicator of alcohol dependence risk in European Americans. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2015, 168, 347-353.	1.7	7
147	Polygenic Overlap Between C-Reactive Protein, Plasma Lipids, and Alzheimer Disease. Circulation, 2015, 131, 2061-2069.	1.6	145
148	Association of Long Runs of Homozygosity With Alzheimer Disease Among African American Individuals. JAMA Neurology, 2015, 72, 1313.	9.0	39
149	Convergent genetic and expression data implicate immunity in Alzheimer's disease. Alzheimer's and Dementia, 2015, 11, 658-671.	0.8	173
150	Genome-Wide Association Study of Copy Number Variations (CNVs) with Opioid Dependence. Neuropsychopharmacology, 2015, 40, 1016-1026.	5.4	39
151	Nf1 Regulates Alcohol Dependence-Associated Excessive Drinking and Gamma-Aminobutyric Acid Release in the Central Amygdala in Mice and Is Associated with Alcohol Dependence in Humans. Biological Psychiatry, 2015, 77, 870-879.	1.3	14
152	FLT1Genetic Variation Predisposes to Neovascular AMD in Ethnically Diverse Populations and Alters Systemic FLT1 Expression. , 2014, 55, 3543.		20
153	Genome-Wide Association Meta-analysis of Neuropathologic Features of Alzheimer's Disease and Related Dementias. PLoS Genetics, 2014, 10, e1004606.	3.5	305
154	Evaluating the role of a galanin enhancer genotype on a range of metabolic, depressive and addictive phenotypes. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2014, 165, 654-664.	1.7	4
155	P1-035: BIVARIATE GENOME-WIDE ASSOCIATION STUDY OF ALZHEIMER DISEASE ENDOPHENOTYPES IDENTIFIES NOVEL LOCI. , 2014, 10, P316-P316.		0
156	Reply. Arthritis and Rheumatology, 2014, 66, 1401-1401.	5.6	1
157	A rare mutation in UNC5C predisposes to late-onset Alzheimer's disease and increases neuronal cell death. Nature Medicine, 2014, 20, 1452-1457.	30.7	116
158	Deep resequencing of 17 glutamate system genes identifies rare variants in <scp>DISC1</scp> and <scp>GRIN2B</scp> affecting risk of opioid dependence. Addiction Biology, 2014, 19, 955-964.	2.6	22
159	Association of MAPT haplotypes with Alzheimerâ€™s disease risk and MAPT brain gene expression levels. Alzheimer's Research and Therapy, 2014, 6, 39.	6.2	106
160	Effects of Multiple Genetic Loci on Age at Onset in Late-Onset Alzheimer Disease. JAMA Neurology, 2014, 71, 1394.	9.0	166
161	P2-131: WHOLE-EXOME SEQUENCING OF HISPANIC EARLY-ONSET ALZHEIMER DISEASE FAMILIES IDENTIFIES RARE VARIANTS IN MULTIPLE ALZHEIMER'S-RELATED GENES. , 2014, 10, P518-P519.		0
162	Structural Interactions between Inhibitor and Substrate Docking Sites Give Insight into Mechanisms of Human PS1 Complexes. Structure, 2014, 22, 125-135.	3.3	56

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163	Exploring the genetic architecture of alcohol dependence in African-Americans via analysis of a genomewide set of common variants. Human Genetics, 2014, 133, 617-624.	3.8	15
164	Genetic analysis of quantitative phenotypes in AD and MCI: imaging, cognition and biomarkers. Brain Imaging and Behavior, 2014, 8, 183-207.	2.1	161
165	Genomeâ€wide association study of the rate of cognitive decline in Alzheimer's disease. Alzheimer's and Dementia, 2014, 10, 45-52.	0.8	147
166	<scp>PLXNA</scp>4 is associated with <scp>A</scp>lzheimer disease and modulates tau phosphorylation. Annals of Neurology, 2014, 76, 379-392.	5.3	60
167	ABCC9 gene polymorphism is associated with hippocampal sclerosis of aging pathology. Acta Neuropathologica, 2014, 127, 825-843.	7.7	70
168	A Case of Inappropriate Apolipoprotein E Testing in Alzheimerâ€™s Disease Due to Lack of an Informed Consent Discussion. American Journal of Alzheimer's Disease and Other Dementias, 2014, 29, 590-595.	1.9	0
169	Two rare AKAP9 variants are associated with Alzheimer's disease in African Americans. Alzheimer's and Dementia, 2014, 10, 609.	0.8	94
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