Lindsay A Farrer

List of Publications by Year
in descending order

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376 papers	43,251 citations	3919 88 h-index	2812 191 g-index
413 all docs	413 docs citations	413 times ranked	39928 citing authors

#	Article	IF	Citations
1	Meta-analysis of 74,046 individuals identifies 11 new susceptibility loci for Alzheimer's disease. Nature Genetics, 2013, 45, 1452-1458.	9.4	3,741
2	Effects of Age, Sex, and Ethnicity on the Association Between Apolipoprotein E Genotype and Alzheimer Disease. JAMA - Journal of the American Medical Association, 1997, 278, 1349.	3.8	3,321
3	Complement Factor H Polymorphism and Age-Related Macular Degeneration. Science, 2005, 308, 421-424.	6.0	2,281
4	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.	9.4	1,962
5	Common variants at MS4A4/MS4A6E, CD2AP, CD33 and EPHA1 are associated with late-onset Alzheimer's disease. Nature Genetics, 2011, 43, 436-441.	9.4	1,676
6	A large genome-wide association study of age-related macular degeneration highlights contributions of rare and common variants. Nature Genetics, 2016, 48, 134-143.	9.4	1,167
7	Analysis of shared heritability in common disorders of the brain. Science, 2018, 360, .	6.0	1,085
8	Genome-wide Analysis of Genetic Loci Associated With Alzheimer Disease. JAMA - Journal of the American Medical Association, 2010, 303, 1832.	3.8	1,064
9	The neuronal sortilin-related receptor SORL1 is genetically associated with Alzheimer disease. Nature Genetics, 2007, 39, 168-177.	9.4	1,045
10	Rare coding variants in PLCG2, ABI3, and TREM2 implicate microglial-mediated innate immunity in Alzheimer's disease. Nature Genetics, 2017, 49, 1373-1384.	9.4	783
11	New insights into the genetic etiology of Alzheimerâ€™s disease and related dementias. Nature Genetics, 2022, 54, 412-436.	9.4	700
12	Seven new loci associated with age-related macular degeneration. Nature Genetics, 2013, 45, 433-439.	9.4	687
13	FUS Phase Separation Is Modulated by a Molecular Chaperone and Methylation of Arginine Cation-Ï€ Interactions. Cell, 2018, 173, 720-734.e15.	13.5	662
14	The PhenX Toolkit: Get the Most From Your Measures. American Journal of Epidemiology, 2011, 174, 253-260.	1.6	610
15	Disclosure ofAPOEGenotype for Risk of Alzheimer's Disease. New England Journal of Medicine, 2009, 361, 245-254.	13.9	490
16	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. Nature Neuroscience, 2018, 21, 1656-1669.	7.1	490
17	Genetic variants near TIMP3 and high-density lipoproteinâ€“associated loci influence susceptibility to age-related macular degeneration. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 7401-7406.	3.3	475
18	Differential modulation of endotoxin responsiveness by human caspase-12 polymorphisms. Nature, 2004, 429, 75-79.	13.7	395

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19	Alzheimer's Disease Neuroimaging Initiative biomarkers as quantitative phenotypes: Genetics core aims, progress, and plans. Alzheimer's and Dementia, 2010, 6, 265-273.	0.4	378
20	Meta-analysis Confirms CR1, CLU, and PICALM as Alzheimer Disease Risk Loci and Reveals Interactions With APOE Genotypes. Archives of Neurology, 2010, 67, 1473.	4.9	376
21	International meta-analysis of PTSD genome-wide association studies identifies sex- and ancestry-specific genetic risk loci. Nature Communications, 2019, 10, 4558.	5.8	363
22	Variants in the ATP-Binding Cassette Transporter (ABCA7), Apolipoprotein E Ïµ4, and the Risk of Late-Onset Alzheimer Disease in African Americans. JAMA - Journal of the American Medical Association, 2013, 309, 1483.	3.8	360
23	GWAS of Cerebrospinal Fluid Tau Levels Identifies Risk Variants for Alzheimerâ€™s Disease. Neuron, 2013, 78, 256-268.	3.8	344
24	Risk of Dementia Among White and African American Relatives of Patients With Alzheimer Disease. JAMA - Journal of the American Medical Association, 2002, 287, 329.	3.8	330
25	A common haplotype lowers PU.1 expression in myeloid cells and delays onset of Alzheimer's disease. Nature Neuroscience, 2017, 20, 1052-1061.	7.1	330
26	Genetic assessment of age-associated Alzheimer disease risk: Development and validation of a polygenic hazard score. PLoS Medicine, 2017, 14, e1002258.	3.9	311
27	Genome-Wide Association Meta-analysis of Neuropathologic Features of Alzheimer's Disease and Related Dementias. PLoS Genetics, 2014, 10, e1004606.	1.5	305
28	Common genetic variants in the CLDN2 and PRSS1-PRSS2 loci alter risk for alcohol-related and sporadic pancreatitis. Nature Genetics, 2012, 44, 1349-1354.	9.4	303
29	Interaction of FKBP5 with Childhood Adversity on Risk for Post-Traumatic Stress Disorder. Neuropsychopharmacology, 2010, 35, 1684-1692.	2.8	299
30	Exceptionally low likelihood of Alzheimerâ€™s dementia in APOE2 homozygotes from a 5,000-person neuropathological study. Nature Communications, 2020, 11, 667.	5.8	246
31	Diabetes Mellitus and Risk of Developing Alzheimer Disease. Archives of Neurology, 2006, 63, 1551.	4.9	245
32	Suicide and attempted suicide in Huntington disease: Implications for preclinical testing of persons at risk. American Journal of Medical Genetics Part A, 1986, 24, 305-311.	2.4	231
33	Brain Expression Genome-Wide Association Study (eGWAS) Identifies Human Disease-Associated Variants. PLoS Genetics, 2012, 8, e1002707.	1.5	225
34	A Comprehensive Genetic Association Study of Alzheimer Disease in African Americans. Archives of Neurology, 2011, 68, 1569.	4.9	221
35	A large-scale genome-wide association study meta-analysis of cannabis use disorder. Lancet Psychiatry,the, 2020, 7, 1032-1045.	3.7	200
36	Genome-wide association study identifies four novel loci associated with Alzheimerâ€™s endophenotypes and disease modifiers. Acta Neuropathologica, 2017, 133, 839-856.	3.9	199

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37	Evidence for a role of the rare p.A152T variant in MAPT in increasing the risk for FTD-spectrum and Alzheimer's diseases. Human Molecular Genetics, 2012, 21, 3500-3512.	1.4	198
38	Genome-Wide Association Study of Opioid Dependence: Multiple Associations Mapped to Calcium and Potassium Pathways. Biological Psychiatry, 2014, 76, 66-74.	0.7	192
39	A systems biology approach towards understanding and treating non-neovascular age-related macular degeneration. Nature Communications, 2019, 10, 3347.	5.8	192
40	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.	4.1	191
41	Correlates of co-occurring ADHD in drug-dependent subjects: Prevalence and features of substance dependence and psychiatric disorders. Addictive Behaviors, 2008, 33, 1199-1207.	1.7	187
42	Hemolysis-associated priapism in sickle cell disease. Blood, 2005, 106, 3264-3267.	0.6	183
43	Assessment of the genetic variance of late-onset Alzheimer's disease. Neurobiology of Aging, 2016, 41, 200.e13-200.e20.	1.5	174
44	Genome-Wide Association Studies of a Broad Spectrum of Antisocial Behavior. JAMA Psychiatry, 2017, 74, 1242.	6.0	174
45	Convergent genetic and expression data implicate immunity in Alzheimer's disease. Alzheimer's and Dementia, 2015, 11, 658-671.	0.4	173
46	Statement on Use of Apolipoprotein E Testing for Alzheimer Disease. JAMA - Journal of the American Medical Association, 1995, 274, 1627.	3.8	172
47	_{Predictive Testing for Huntingtons Disease with Use of a Linked DNA Marker}. New England Journal of Medicine, 1988, 318, 535-542.	13.9	167
48	Effects of Multiple Genetic Loci on Age at Onset in Late-Onset Alzheimer Disease. JAMA Neurology, 2014, 71, 1394.	4.5	166
49	Transethnic genomeâ€wide scan identifies novel Alzheimer's disease loci. Alzheimer's and Dementia, 2017, 13, 727-738.	0.4	166
50	Genetic analysis of quantitative phenotypes in AD and MCI: imaging, cognition and biomarkers. Brain Imaging and Behavior, 2014, 8, 183-207.	1.1	161
51	The Genetics of Adult-Onset Neuropsychiatric Disease: Complexities and Conundra?. Science, 2003, 302, 822-826.	6.0	160
52	A network model to predict the risk of death in sickle cell disease. Blood, 2007, 110, 2727-2735.	0.6	159
53	BCL11A is a major HbF quantitative trait locus in three different populations with Î²-hemoglobinopathies. Blood Cells, Molecules, and Diseases, 2008, 41, 255-258.	0.6	158
54	Identification of a rare coding variant in complement 3 associated with age-related macular degeneration. Nature Genetics, 2013, 45, 1375-1379.	9.4	158

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55	Sickle cell leg ulcers: associations with haemolysis and SNPs in Klotho, TEK and genes of the TGF-beta/BMP pathway. British Journal of Haematology, 2006, 133, 570-578.	1.2	155
56	Gene-Wide Analysis Detects Two New Susceptibility Genes for Alzheimer's Disease. PLoS ONE, 2014, 9, e94661.	1.1	155
57	Estimating risk curves for first-degree relatives of patients with Alzheimerâ€™s disease: The REVEAL study. Genetics in Medicine, 2004, 6, 192-196.	1.1	153
58	Meta-analysis of the Association Between Variants in SORL1 and Alzheimer Disease. Archives of Neurology, 2011, 68, 99.	4.9	153
59	Genome-wide Association Study Identifies New Susceptibility Loci for Posttraumatic Stress Disorder. Biological Psychiatry, 2013, 74, 656-663.	0.7	150
60	SORL1 Is Genetically Associated with Late-Onset Alzheimerâ€™s Disease in Japanese, Koreans and Caucasians. PLoS ONE, 2013, 8, e58618.	1.1	149
61	A genome-wide association study of plasma total IgE concentrations in the Framingham Heart Study. Journal of Allergy and Clinical Immunology, 2012, 129, 840-845.e21.	1.5	148
62	Genome-wide Association Study of Cannabis Dependence Severity, Novel Risk Variants, and Shared Genetic Risks. JAMA Psychiatry, 2016, 73, 472.	6.0	148
63	Genomeâ€wide association study of the rate of cognitive decline in Alzheimer's disease. Alzheimer's and Dementia, 2014, 10, 45-52.	0.4	147
64	Fetal hemoglobin in sickle cell anemia: genome-wide association studies suggest a regulatory region in the 5â€² olfactory receptor gene cluster. Blood, 2010, 115, 1815-1822.	0.6	146
65	Assessment of genetic risk for alzheimer's disease among first-degree relatives. Annals of Neurology, 1989, 25, 485-493.	2.8	145
66	Polygenic Overlap Between C-Reactive Protein, Plasma Lipids, and Alzheimer Disease. Circulation, 2015, 131, 2061-2069.	1.6	145
67	Diabetes mellitus in Huntington disease. Clinical Genetics, 1985, 27, 62-67.	1.0	144
68	Novel late-onset Alzheimer disease loci variants associate with brain gene expression. Neurology, 2012, 79, 221-228.	1.5	144
69	Novel Alzheimer Disease Risk Loci and Pathways in African American Individuals Using the African Genome Resources Panel. JAMA Neurology, 2021, 78, 102.	4.5	144
70	Common variants in Alzheimerâ€™s disease and risk stratification by polygenic risk scores. Nature Communications, 2021, 12, 3417.	5.8	140
71	Association of Granulomatosis With Polyangiitis (Wegener's) With HLAâ€“DPB1*04 and SEMA6A Gene Variants: Evidence From Genomeâ€Wide Analysis. Arthritis and Rheumatism, 2013, 65, 2457-2468.	6.7	138
72	Identification of Alzheimer disease-associated variants in genes that regulate retromer function. Neurobiology of Aging, 2012, 33, 2231.e15-2231.e30.	1.5	135

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73	Association of OPRM1 Functional Coding Variant With Opioid Use Disorder. JAMA Psychiatry, 2020, 77, 1072.	6.0	135
74	Identification of multiple loci for Alzheimer disease in a consanguineous Israeli-Arab community. Human Molecular Genetics, 2003, 12, 415-422.	1.4	117
75	Ancestral origin of ApoE Îµ4 Alzheimer disease risk in Puerto Rican and African American populations. PLoS Genetics, 2018, 14, e1007791.	1.5	117
76	A 3-bp deletion in the HBS1L-MYB intergenic region on chromosome 6q23 is associated with HbF expression. Blood, 2011, 117, 4935-4945.	0.6	116
77	A rare mutation in UNC5C predisposes to late-onset Alzheimer's disease and increases neuronal cell death. Nature Medicine, 2014, 20, 1452-1457.	15.2	116
78	An Î±-2-macroglobulin insertion-deletion polymorphism in Alzheimer disease. Nature Genetics, 1999, 22, 19-21.	9.4	115
79	Missense variant in TREML2 protects against Alzheimer's disease. Neurobiology of Aging, 2014, 35, 1510.e19-1510.e26.	1.5	110
80	Genetics of age-related macular degeneration (AMD). Human Molecular Genetics, 2017, 26, R45-R50.	1.4	109
81	Analysis of the glucocerebrosidase gene in Parkinson's disease. Movement Disorders, 2005, 20, 367-370.	2.2	107
82	Association of MAPT haplotypes with Alzheimerâ€™s disease risk and MAPT brain gene expression levels. Alzheimer's Research and Therapy, 2014, 6, 39.	3.0	106
83	SORCS1 alters amyloid precursor protein processing and variants may increase Alzheimer's disease risk. Annals of Neurology, 2011, 69, 47-64.	2.8	104
84	Variation in Nicotinic Acetylcholine Receptor Genes is Associated with Multiple Substance Dependence Phenotypes. Neuropsychopharmacology, 2010, 35, 1921-1931.	2.8	103
85	Who seeks genetic susceptibility testing for Alzheimerâ€™s disease? Findings from a multisite, randomized clinical trial. Genetics in Medicine, 2004, 6, 197-203.	1.1	101
86	Dissecting the genetic relationship between cardiovascular risk factors and Alzheimerâ€™s disease. Acta Neuropathologica, 2019, 137, 209-226.	3.9	100
87	Association Between Apolipoprotein E Genotype and Alzheimer Disease in African American Subjects. Archives of Neurology, 2002, 59, 594.	4.9	98
88	Comprehensive Search for Alzheimer Disease Susceptibility Loci in the APOE Region. Archives of Neurology, 2012, 69, 1270.	4.9	97
89	Linkage of congenital, recessive deafness (DFNB4) to chromosome 7q31 and evidence for genetic heterogeneity in the Middle Eastern Druze population. Human Molecular Genetics, 1995, 4, 1637-1642.	1.4	96
90	Association Between Angiotensin-Converting Enzyme and Alzheimer Disease. Archives of Neurology, 2000, 57, 210.	4.9	96

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91	Two rare AKAP9 variants are associated with Alzheimer's disease in African Americans. Alzheimer's and Dementia, 2014, 10, 609.	0.4	94
92	Genetic variants and functional pathways associated with resilience to Alzheimerâ€™s disease. Brain, 2020, 143, 2561-2575.	3.7	93
93	Apolipoprotein E genotype in patients with alzheimer's disease: Implications for the risk of dementia among relatives. Annals of Neurology, 1995, 38, 797-808.	2.8	87
94	Polymorphisms in the PON gene cluster are associated with Alzheimer disease. Human Molecular Genetics, 2006, 15, 77-85.	1.4	87
95	Two novel loci, COBL and SLC10A2, for Alzheimer's disease in African Americans. Alzheimer's and Dementia, 2017, 13, 119-129.	0.4	87
96	Association between SORL1 and Alzheimer's disease in a genome-wide study. NeuroReport, 2007, 18, 1761-1764.	0.6	83
97	Reasons for Seeking Genetic Susceptibility Testing Among First-Degree Relatives of People With Alzheimer Disease. Alzheimer Disease and Associated Disorders, 2003, 17, 86-93.	0.6	82
98	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.	4.1	82
99	Causal Associations Between Modifiable Risk Factors and the Alzheimer's Phenome. Annals of Neurology, 2021, 89, 54-65.	2.8	82
100	A family history study of male sexual orientation using three independent samples. Behavior Genetics, 1999, 29, 79-86.	1.4	80
101	Expanding the genetic architecture of nicotine dependence and its shared genetics with multiple traits. Nature Communications, 2020, 11, 5562.	5.8	80
102	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.	0.7	78
103	Performance of random forest when SNPs are in linkage disequilibrium. BMC Bioinformatics, 2009, 10, 78.	1.2	76
104	ABCA7 frameshift deletion associated with Alzheimer disease in African Americans. Neurology: Genetics, 2016, 2, e79.	0.9	74
105	Genetic Risk Variants Associated With Comorbid Alcohol Dependence and Major Depression. JAMA Psychiatry, 2017, 74, 1234.	6.0	74
106	Integrating GWASs and Human Protein Interaction Networks Identifies a Gene Subnetwork Underlying Alcohol Dependence. American Journal of Human Genetics, 2013, 93, 1027-1034.	2.6	72
107	Association of single nucleotide polymorphisms in klotho with priapism in sickle cell anaemia. British Journal of Haematology, 2005, 128, 266-272.	1.2	71
108	Systems biology-based analysis implicates a novel role for vitamin D metabolism in the pathogenesis of age-related macular degeneration. Human Genomics, 2011, 5, 538.	1.4	70

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109	ABCC9 gene polymorphism is associated with hippocampal sclerosis of aging pathology. Acta Neuropathologica, 2014, 127, 825-843.	3.9	70
110	Familial Alzheimer's disease: Progress and problems. Neurobiology of Aging, 1989, 10, 417-425.	1.5	69
111	Nonsteroidal anti-inflammatory drug use and Alzheimer's disease risk: the MIRAGE Study. BMC Geriatrics, 2005, 5, 2.	1.1	69
112	Association of Polymorphisms in the Angiotensin-Converting Enzyme Gene with Alzheimer Disease in an Israeli Arab Community. American Journal of Human Genetics, 2006, 78, 871-877.	2.6	69
113	Potential ethnic modifiers in the assessment and treatment of Alzheimer's disease: challenges for the future. International Psychogeriatrics, 2007, 19, 539-558.	0.6	65
114	Genome-wide Association Study Identifies a Regulatory Variant of RGMA Associated With Opioid Dependence in European Americans. Biological Psychiatry, 2018, 84, 762-770.	0.7	64
115	Genomeâ€wide association study of Alzheimer's disease endophenotypes at prediagnosis stages. Alzheimer's and Dementia, 2018, 14, 623-633.	0.4	64
116	Relation between atherogenic dyslipidemia and the Adult Treatment Program-III definition of metabolic syndrome (Genetic Epidemiology of Metabolic Syndrome Project). American Journal of Cardiology, 2005, 95, 194-198.	0.7	63
117	Childhood Adversity Increases Risk for Nicotine Dependence and Interacts with Î±5 Nicotinic Acetylcholine Receptor Genotype Specifically in Males. Neuropsychopharmacology, 2012, 37, 669-676.	2.8	63
118	Serum heat shock protein 70 level as a biomarker of exceptional longevity. Mechanisms of Ageing and Development, 2006, 127, 862-868.	2.2	62
119	Rare Nonsynonymous Variants in Alpha-4 Nicotinic Acetylcholine Receptor Gene Protect Against Nicotine Dependence. Biological Psychiatry, 2011, 70, 528-536.	0.7	62
120	Association of Distinct Variants in SORL1 With Cerebrovascular and Neurodegenerative Changes Related to Alzheimer Disease. Archives of Neurology, 2008, 65, 1640.	4.9	60
121	<scp>PLXNA</scp>4 is associated with <scp>A</scp>lzheimer disease and modulates tau phosphorylation. Annals of Neurology, 2014, 76, 379-392.	2.8	60
122	Multiple loci influencing hippocampal degeneration identified by genome scan. Annals of Neurology, 2012, 72, 65-75.	2.8	59
123	Genomewide Association Study for Maximum Number of Alcoholic Drinks in European Americans and African Americans. Alcoholism: Clinical and Experimental Research, 2015, 39, 1137-1147.	1.4	58
124	Association of Rare Coding Mutations With Alzheimer Disease and Other Dementias Among Adults of European Ancestry. JAMA Network Open, 2019, 2, e191350.	2.8	58
125	Î´-Catenin Is Genetically and Biologically Associated with Cortical Cataract and Future Alzheimer-Related Structural and Functional Brain Changes. PLoS ONE, 2012, 7, e43728.	1.1	58
126	Structural Interactions between Inhibitor and Substrate Docking Sites Give Insight into Mechanisms of Human PS1 Complexes. Structure, 2014, 22, 125-135.	1.6	56

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127	Shared genetic contribution to ischemic stroke and Alzheimer's disease. Annals of Neurology, 2016, 79, 739-747.	2.8	56
128	Rating the severity and character of transient cocaine-induced delusions and hallucinations with a new instrument, the Scale for Assessment of Positive Symptoms for Cocaine-Induced Psychosis (SAPS-CIP). Drug and Alcohol Dependence, 2005, 80, 23-33.	1.6	55
129	Variations in opioid receptor genes in neonatal abstinence syndrome. Drug and Alcohol Dependence, 2015, 155, 253-259.	1.6	55
130	Association of Polymorphisms ofIGF1Rand Genes in the Transforming Growth Factorâ€“Î²/Bone Morphogenetic Protein Pathway with Bacteremia in Sickle Cell Anemia. Clinical Infectious Diseases, 2006, 43, 593-598.	2.9	54
131	Convergence of linkage, gene expression and association data demonstrates the influence of the RAR-related orphan receptor alpha (RORA) gene on neovascular AMD: A systems biology based approach. Vision Research, 2010, 50, 698-715.	0.7	54
132	A search for age-related macular degeneration risk variants in Alzheimer disease genes and pathways. Neurobiology of Aging, 2014, 35, 1510.e7-1510.e18.	1.5	53
133	An anthropometric assessment of Huntington's disease patients and families. American Journal of Physical Anthropology, 1985, 67, 185-194.	2.1	52
134	Rare and common variants in extracellular matrix gene Fibrillin 2 (FBN2) are associated with macular degeneration. Human Molecular Genetics, 2014, 23, 5827-5837.	1.4	52
135	Allele Ïµ4 of Apolipoprotein E Shows a Dose Effect on Age at Onset of Pick Disease. Experimental Neurology, 1995, 136, 162-170.	2.0	50
136	Fetal hemoglobin in sickle cell anemia: Genetic studies of the Arab-Indian haplotype. Blood Cells, Molecules, and Diseases, 2013, 51, 22-26.	0.6	50
137	Association between bleomycin hydrolase and Alzheimer's disease in caucasians. Annals of Neurology, 1998, 44, 808-811.	2.8	48
138	Statin use and the risk of Alzheimer's disease: The MIRAGE Study. , 2006, 2, 96-103.		48
139	Intronic variants in the dopa decarboxylase (DDC) gene are associated with smoking behavior in European-Americans and African-Americans. Human Molecular Genetics, 2006, 15, 2192-2199.	1.4	48
140	Estimated glomerular filtration rate in sickle cell anemia is associated with polymorphisms of bone morphogenetic protein receptor 1B. American Journal of Hematology, 2007, 82, 179-184.	2.0	48
141	Protective variant for hippocampal atrophy identified by whole exome sequencing. Annals of Neurology, 2015, 77, 547-552.	2.8	48
142	The natural history of Huntington disease: Possible role of â€œaging genesâ€• American Journal of Medical Genetics Part A, 1984, 18, 115-123.	2.4	47
143	High Prevalence of Mild Cognitive Impairment and Alzheimer's Disease in Arabic Villages in Northern Israel: Impact of Gender and Education. Journal of Alzheimer's Disease, 2012, 29, 431-439.	1.2	47
144	Genomic influences on self-reported childhood maltreatment. Translational Psychiatry, 2020, 10, 38.	2.4	47

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145	Novel ATP7B mutations causing Wilson disease in several Israeli ethnic groups. , 1998, 11, 145-151.		46
146	Polymorphisms in the Promoter Region of Catalase Gene and Essential Hypertension. Disease Markers, 2005, 21, 3-7.	0.6	46
147	Comorbid Psychiatric Diagnoses and Their Association with Cocaine-Induced Psychosis in Cocaine-Dependent Subjects. American Journal on Addictions, 2007, 16, 343-351.	1.3	46
148	Identification of Novel Candidate Genes for Alzheimer's Disease by Autozygosity Mapping using Genome Wide SNP Data. Journal of Alzheimer's Disease, 2011, 23, 349-359.	1.2	46
149	Linkage Analysis Followed by Association Show NRG1 Associated with Cannabis Dependence in African Americans. Biological Psychiatry, 2012, 72, 637-644.	0.7	46
150	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.	1.5	46
151	Sex-dependent autosomal effects on clinical progression of Alzheimerâ€™s disease. Brain, 2020, 143, 2272-2280.	3.7	46
152	Autosomal Dominant Orthostatic Hypotensive Disorder Maps to Chromosome 18q. American Journal of Human Genetics, 1998, 63, 1425-1430.	2.6	45
153	Identification of the genetic locus for keratosis palmaris et plantaris on chromosome 17 near the RARA and keratin type I genes. Nature Genetics, 1993, 5, 158-162.	9.4	44
154	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.	1.2	44
155	Anthropometric discrimination among affected, at-risk, and not-at-risk individuals in families with Huntington disease. American Journal of Medical Genetics Part A, 1985, 21, 307-316.	2.4	42
156	The human cationic amino acid transporter (ATRC1): Physical and genetic mapping to 13q12â€“q14. Genomics, 1992, 12, 430-434.	1.3	42
157	Education Attenuates the Effect of Medial Temporal Lobe Atrophy on Cognitive Function in Alzheimer's Disease: The MIRAGE Study. Journal of Alzheimer's Disease, 2009, 17, 855-862.	1.2	42
158	Global and local ancestry in Africanâ€Americans: Implications for Alzheimer's disease risk. Alzheimer's and Dementia, 2016, 12, 233-243.	0.4	42
159	Initial Assessment of the Pathogenic Mechanisms of the Recently Identified Alzheimer Risk Loci. Annals of Human Genetics, 2013, 77, 85-105.	0.3	41
160	Rarity of the Alzheimer Diseaseâ€“ProtectiveAPPA673T Variant in the United States. JAMA Neurology, 2015, 72, 209.	4.5	41
161	Segregation of a rare TTC3 variant in an extended family with late-onset Alzheimer disease. Neurology: Genetics, 2016, 2, e41.	0.9	41
162	Early-Onset Alzheimer Disease and Candidate Risk Genes Involved in Endolysosomal Transport. JAMA Neurology, 2017, 74, 1113.	4.5	41

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163	GENETICS AND THE DEMENTIA PATIENT. Neurologist, 1997, 3, 13-30.	0.4	40
164	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.	2.8	40
165	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.	1.0	40
166	No genetic association between the LRP receptor and sporadic or late-onset familial Alzheimer disease. Neurogenetics, 1998, 1, 179-183.	0.7	39
167	Multiple QTLs influencing triglyceride and HDL and total cholesterol levels identified in families with atherogenic dyslipidemia. Journal of Lipid Research, 2005, 46, 2202-2213.	2.0	39
168	Education effects on cognitive function in a healthy aged Arab population. International Psychogeriatrics, 2007, 19, 593-603.	0.6	39
169	GABRG1 and GABRA2 Variation Associated with Alcohol Dependence in African Americans. Alcoholism: Clinical and Experimental Research, 2012, 36, 588-593.	1.4	39
170	Association of Long Runs of Homozygosity With Alzheimer Disease Among African American Individuals. JAMA Neurology, 2015, 72, 1313.	4.5	39
171	Genome-Wide Association Study of Copy Number Variations (CNVs) with Opioid Dependence. Neuropsychopharmacology, 2015, 40, 1016-1026.	2.8	39
172	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.	1.5	39
173	Association of maternal and infant variants inPNOCandCOMTgenes with neonatal abstinence syndrome severity. American Journal on Addictions, 2017, 26, 42-49.	1.3	39
174	A rare missense variant of CASP7 is associated with familial lateâ€onset Alzheimer's disease. Alzheimer's and Dementia, 2019, 15, 441-452.	0.4	39
175	Caspase-8, association with Alzheimerâ€™s Disease and functional analysis of rare variants. PLoS ONE, 2017, 12, e0185777.	1.1	38
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