

# Kevin Morgan

## List of Publications by Year in Descending Order

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**Version:** 2024-04-26

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

168  
papers

15,291  
citations

42  
h-index

123  
g-index

183  
ext. papers

19,090  
ext. citations

7.9  
avg, IF

5.14  
L-index

#	Paper	IF	Citations
168	Prion-like $\beta$ -synuclein pathology in the brain of infants with Krabbe disease.. <i>Brain</i> , <b>2022</b> ,	11.2	2
167	New insights into the genetic etiology of Alzheimer's disease and related dementias.. <i>Nature Genetics</i> , <b>2022</b> ,	36.3	27
166	Comment on: Association between serum urate, gout and comorbidities: a case-control study using data from the UK Biobank: Reply. <i>Rheumatology</i> , <b>2021</b> ,	3.9	
165	Association between serum urate, gout and comorbidities: a case-control study using data from the UK Biobank. <i>Rheumatology</i> , <b>2021</b> , 60, 3243-3251	3.9	4
164	Variants in urate transporters, ADH1B, GCKR and MEPE genes associate with transition from asymptomatic hyperuricaemia to gout: results of the first gout versus asymptomatic hyperuricaemia GWAS in Caucasians using data from the UK Biobank. <i>Annals of the Rheumatic Diseases</i> , <b>2021</b> , 80, 1220-1226	2.4	6
163	A meta-analysis of epigenome-wide association studies in Alzheimer's disease highlights novel differentially methylated loci across cortex. <i>Nature Communications</i> , <b>2021</b> , 12, 3517	17.4	8
162	Response to: 'Correspondence on 'Variants in urate transporters, ADH1B, GCKR and MEPE genes associated with transition from asymptomatic hyperuricaemia to gout: results of the first gout versus asymptomatic hyperuricaemia GWAS in Caucasians using data from the UK Biobank'' by Takei. <i>Annals of the Rheumatic Diseases</i> , <b>2021</b>	2.4	1
161	Common variants in Alzheimer's disease and risk stratification by polygenic risk scores. <i>Nature Communications</i> , <b>2021</b> , 12, 3417	17.4	23
160	Genome-wide association findings from the brains for dementia research cohort. <i>Neurobiology of Aging</i> , <b>2021</b> , 107, 159-167	5.6	1
159	Genetic variants in glutamate-, A $\beta$ and tau-related pathways determine polygenic risk for Alzheimer's disease. <i>Neurobiology of Aging</i> , <b>2021</b> , 101, 299.e13-299.e21	5.6	3
158	A genetic link between risk for Alzheimer's disease and severe COVID-19 outcomes via the OAS1 gene. <i>Brain</i> , <b>2021</b> ,	11.2	14
157	Genome sequencing analysis identifies new loci associated with Lewy body dementia and provides insights into its genetic architecture. <i>Nature Genetics</i> , <b>2021</b> , 53, 294-303	36.3	31
156	Recalibrating the epigenetic clock: implications for assessing biological age in the human cortex. <i>Brain</i> , <b>2020</b> , 143, 3763-3775	11.2	32
155	Psychological stress, cognitive decline and the development of dementia in amnesic mild cognitive impairment. <i>Scientific Reports</i> , <b>2020</b> , 10, 3618	4.9	6
154	Analysis of neurodegenerative disease-causing genes in dementia with Lewy bodies. <i>Acta Neuropathologica Communications</i> , <b>2020</b> , 8, 5	7.3	15
153	Genetic risk for Alzheimer's disease influences neuropathology via multiple biological pathways. <i>Brain Communications</i> , <b>2020</b> , 2, fcaa167	4.5	1
152	A rare loss-of-function variant of ADAM17 is associated with late-onset familial Alzheimer disease. <i>Molecular Psychiatry</i> , <b>2020</b> , 25, 629-639	15.1	27

151	Alzheimer's disease polygenic risk score as a predictor of conversion from mild-cognitive impairment. <i>Translational Psychiatry</i> , <b>2019</b> , 9, 154	8.6	31
150	The Epistasis Project: A Multi-Cohort Study of the Effects of BDNF, DBH, and SORT1 Epistasis on Alzheimer's Disease Risk. <i>Journal of Alzheimer's Disease</i> , <b>2019</b> , 68, 1535-1547	4.3	5
149	Heritability and genetic variance of dementia with Lewy bodies. <i>Neurobiology of Disease</i> , <b>2019</b> , 127, 492-501	5.0	15
148	Gene-based analysis in HRC imputed genome wide association data identifies three novel genes for Alzheimer's disease. <i>PLoS ONE</i> , <b>2019</b> , 14, e0218111	3.7	12
147	Cognitive Change in Schizophrenia and Other Psychoses in the Decade Following the First Episode. <i>American Journal of Psychiatry</i> , <b>2019</b> , 176, 811-819	11.9	64
146	Genetic variability in response to amyloid beta deposition influences Alzheimer's disease risk. <i>Brain Communications</i> , <b>2019</b> , 1, fcz022	4.5	29
145	Genetic meta-analysis of diagnosed Alzheimer's disease identifies new risk loci and implicates A $\beta$ tau, immunity and lipid processing. <i>Nature Genetics</i> , <b>2019</b> , 51, 414-430	36.3	917
144	A comprehensive screening of copy number variability in dementia with Lewy bodies. <i>Neurobiology of Aging</i> , <b>2019</b> , 75, 223.e1-223.e10	5.6	10
143	Complement receptor 1 gene (CR1) intragenic duplication and risk of Alzheimer's disease. <i>Human Genetics</i> , <b>2018</b> , 137, 305-314	6.3	15
142	Mendelian adult-onset leukodystrophy genes in Alzheimer's disease: critical influence of CSF1R and NOTCH3. <i>Neurobiology of Aging</i> , <b>2018</b> , 66, 179.e17-179.e29	5.6	23
141	Polygenic risk score in postmortem diagnosed sporadic early-onset Alzheimer's disease. <i>Neurobiology of Aging</i> , <b>2018</b> , 62, 244.e1-244.e8	5.6	25
140	Genotyping of the Alzheimer's Disease Genome-Wide Association Study Index Single Nucleotide Polymorphisms in the Brains for Dementia Research Cohort. <i>Journal of Alzheimer's Disease</i> , <b>2018</b> , 64, 355-362	4.3	6
139	Analysis of shared heritability in common disorders of the brain. <i>Science</i> , <b>2018</b> , 360,	33.3	666
138	Whole-exome sequencing of the BDR cohort: evidence to support the role of the PILRA gene in Alzheimer's disease. <i>Neuropathology and Applied Neurobiology</i> , <b>2018</b> , 44, 506-521	5.2	27
137	Investigating the genetic architecture of dementia with Lewy bodies: a two-stage genome-wide association study. <i>Lancet Neurology</i> , <b>2018</b> , 17, 64-74	24.1	121
136	P3-126: USING RNA-SEQ TO IDENTIFY RNA EDITING IN CONTROL AND ALZHEIMER'S DISEASE POSTMORTEM BRAIN TISSUE <b>2018</b> , 14, P1115-P1116		
135	Observations of extensive gene expression differences in the cerebellum and potential relevance to Alzheimer's disease. <i>BMC Research Notes</i> , <b>2018</b> , 11, 646	2.3	4
134	A candidate regulatory variant at the TREM gene cluster associates with decreased Alzheimer's disease risk and increased TREML1 and TREM2 brain gene expression. <i>Alzheimer's and Dementia</i> , <b>2017</b> , 13, 663-673	1.2	35

133	Methylation Profiling RIN3 and MEF2C Identifies Epigenetic Marks Associated with Sporadic Early Onset Alzheimer's Disease. <i>Journal of Alzheimer's Disease Reports</i> , <b>2017</b> , 1, 97-108	3.3	14
132	Rare coding variants in PLCG2, ABI3, and TREM2 implicate microglial-mediated innate immunity in Alzheimer's disease. <i>Nature Genetics</i> , <b>2017</b> , 49, 1373-1384	36.3	508
131	Analysis of C9orf72 repeat expansions in a large international cohort of dementia with Lewy bodies. <i>Neurobiology of Aging</i> , <b>2017</b> , 49, 214.e13-214.e15	5.6	10
130	Mutation analysis of sporadic early-onset Alzheimer's disease using the NeuroX array. <i>Neurobiology of Aging</i> , <b>2017</b> , 49, 215.e1-215.e8	5.6	15
129	A novel Alzheimer disease locus located near the gene encoding tau protein. <i>Molecular Psychiatry</i> , <b>2016</b> , 21, 108-17	15.1	175
128	ABCA7 p.G215S as potential protective factor for Alzheimer's disease. <i>Neurobiology of Aging</i> , <b>2016</b> , 46, 235.e1-9	5.6	33
127	Genome-wide analysis of genetic correlation in dementia with Lewy bodies, Parkinson's and Alzheimer's diseases. <i>Neurobiology of Aging</i> , <b>2016</b> , 38, 214.e7-214.e10	5.6	49
126	Screening exons 16 and 17 of the amyloid precursor protein gene in sporadic early-onset Alzheimer's disease. <i>Neurobiology of Aging</i> , <b>2016</b> , 39, 220.e1-7	5.6	9
125	Evaluating pathogenic dementia variants in posterior cortical atrophy. <i>Neurobiology of Aging</i> , <b>2016</b> , 37, 38-44	5.6	19
124	Influence of Coding Variability in APP-A $\beta$ Metabolism Genes in Sporadic Alzheimer's Disease. <i>PLoS ONE</i> , <b>2016</b> , 11, e0150079	3.7	26
123	Rationalising the role of Keratin 9 as a biomarker for Alzheimer's disease. <i>Scientific Reports</i> , <b>2016</b> , 6, 22942	4.2	19
122	Shared genetic contribution to Ischaemic Stroke and Alzheimer's Disease. <i>Annals of Neurology</i> , <b>2016</b> , 79, 739-747	9.4	42
121	Acute and chronic performance evaluation of a novel epicardial pacing lead placed by percutaneous subxiphoid approach in a canine model. <i>Circulation: Arrhythmia and Electrophysiology</i> , <b>2015</b> , 8, 659-66	6.4	10
120	Common polygenic variation enhances risk prediction for Alzheimer's disease. <i>Brain</i> , <b>2015</b> , 138, 3673-84	11.2	227
119	Convergent genetic and expression data implicate immunity in Alzheimer's disease. <i>Alzheimer's and Dementia</i> , <b>2015</b> , 11, 658-71	1.2	146
118	Blood type gene locus has no influence on ACE association with Alzheimer's disease. <i>Neurobiology of Aging</i> , <b>2015</b> , 36, 1767.e1-1767.e2	5.6	2
117	A Fluorescence Resonance Energy Transfer Assay For Monitoring $\beta$ Synuclein Aggregation in a Caenorhabditis Elegans Model For Parkinson's Disease. <i>CNS and Neurological Disorders - Drug Targets</i> , <b>2015</b> , 14, 1054-68	2.6	8
116	Reverse engineering of Alzheimer's disease based on biomarker pathways analysis. <i>Neurobiology of Aging</i> , <b>2014</b> , 35, 2029-38	5.6	5

115	Review: The genetics of Alzheimer's disease; putting flesh on the bones. <i>Neuropathology and Applied Neurobiology</i> , <b>2014</b> , 40, 97-105	5.2	48
114	Exome sequencing identifies 2 novel presenilin 1 mutations (p.L166V and p.S230R) in British early-onset Alzheimer's disease. <i>Neurobiology of Aging</i> , <b>2014</b> , 35, 2422.e13-6	5.6	26
113	Rare coding variants in the phospholipase D3 gene confer risk for Alzheimer's disease. <i>Nature</i> , <b>2014</b> , 505, 550-554	50.4	345
112	Missense variant in TREML2 protects against Alzheimer's disease. <i>Neurobiology of Aging</i> , <b>2014</b> , 35, 1510.e19-2684	5.6	26
111	The sex-specific associations of the aromatase gene with Alzheimer's disease and its interaction with IL10 in the Epistasis Project. <i>European Journal of Human Genetics</i> , <b>2014</b> , 22, 216-20	5.3	32
110	ApoE variant p.V236E is associated with markedly reduced risk of Alzheimer's disease. <i>Molecular Neurodegeneration</i> , <b>2014</b> , 9, 11	19	33
109	Genetic analysis implicates APOE, SNCA and suggests lysosomal dysfunction in the etiology of dementia with Lewy bodies. <i>Human Molecular Genetics</i> , <b>2014</b> , 23, 6139-46	5.6	152
108	Allele-specific RNA interference rescues the long-QT syndrome phenotype in human-induced pluripotency stem cell cardiomyocytes. <i>European Heart Journal</i> , <b>2014</b> , 35, 1078-87	9.5	85
107	Investigating the role of rare coding variability in Mendelian dementia genes (APP, PSEN1, PSEN2, GRN, MAPT, and PRNP) in late-onset Alzheimer's disease. <i>Neurobiology of Aging</i> , <b>2014</b> , 35, 2881.e1-2881.e6	5.6	45
106	Gene-wide analysis detects two new susceptibility genes for Alzheimer's disease. <i>PLoS ONE</i> , <b>2014</b> , 9, e94661	3.7	90
105	Practical detection of a definitive biomarker panel for Alzheimer's disease; comparisons between matched plasma and cerebrospinal fluid. <i>International Journal of Molecular Epidemiology and Genetics</i> , <b>2014</b> , 5, 53-70	0.9	10
104	Discovery by the Epistasis Project of an epistatic interaction between the GSTM3 gene and the HHEX/IDE/KIF11 locus in the risk of Alzheimer's disease. <i>Neurobiology of Aging</i> , <b>2013</b> , 34, 1309.e1-7	5.6	24
103	Meta-analysis of 74,046 individuals identifies 11 new susceptibility loci for Alzheimer's disease. <i>Nature Genetics</i> , <b>2013</b> , 45, 1452-8	36.3	2714
102	TREM2 variants in Alzheimer's disease. <i>New England Journal of Medicine</i> , <b>2013</b> , 368, 117-27	59.2	1805
101	Linking protective GAB2 variants, increased cortical GAB2 expression and decreased Alzheimer's disease pathology. <i>PLoS ONE</i> , <b>2013</b> , 8, e64802	3.7	12
100	Sialic Acid Binding Immunoglobulin-Like Lectin-3 (CD33) <b>2013</b> , 181-190		
99	Erythropoietin-Producing Human Hepatocellular Carcinoma (EphA1) <b>2013</b> , 191-199		
98	Bridging Integrator 1 (BIN1) <b>2013</b> , 103-133		

97	PICALM <b>2013</b> , 53-76		
96	The Future Role of Biomarkers in Alzheimer's Disease Diagnostics <b>2013</b> , 231-248		
95	CD2-Associated Protein (CD2AP) <b>2013</b> , 201-208		
94	Complement Component (3b/4b) Receptor 1 (CR1) <b>2013</b> , 77-101		1
93	The Genetics of Alzheimer's Disease: Introduction and Perspective for the Future <b>2013</b> , 1-5		
92	Clusterin <b>2013</b> , 25-51		1
91	ATP-Binding Cassette, Subfamily A (ABC1), Member 7 (ABCA7) <b>2013</b> , 135-158		3
90	Membrane-Spanning 4-Domains Subfamily A, MS4A Cluster <b>2013</b> , 159-179		2
89	Transferrin and HFE genes interact in Alzheimer's disease risk: the Epistasis Project. <i>Neurobiology of Aging</i> , <b>2012</b> , 33, 202.e1-13	5.6	43
88	Cognitive phenotypes in Alzheimer's disease and genetic variants in ACE and IDE. <i>Neurobiology of Aging</i> , <b>2012</b> , 33, 1486.e1-2	5.6	8
87	Genetic variants influencing human aging from late-onset Alzheimer's disease (LOAD) genome-wide association studies (GWAS). <i>Neurobiology of Aging</i> , <b>2012</b> , 33, 1849.e5-18	5.6	38
86	Identification of SPARC-like 1 protein as part of a biomarker panel for Alzheimer's disease in cerebrospinal fluid. <i>Journal of Alzheimer's Disease</i> , <b>2012</b> , 28, 625-36	4.3	49
85	Interaction of insulin and PPAR- $\gamma$ genes in Alzheimer's disease: the Epistasis Project. <i>Journal of Neural Transmission</i> , <b>2012</b> , 119, 473-9	4.3	19
84	Commentary on 'Functional analysis of APOE locus genetic variation implicates regional enhancers in the regulation of both TOMM40 and APOE'. <i>Journal of Human Genetics</i> , <b>2012</b> , 57, 3-4	4.3	5
83	The role of variation at ABP, PSEN1, PSEN2, and MAPT in late onset Alzheimer's disease. <i>Journal of Alzheimer's Disease</i> , <b>2012</b> , 28, 377-87	4.3	47
82	Interactions between PPAR- $\gamma$ and inflammation-related cytokine genes on the development of Alzheimer's disease, observed by the Epistasis Project. <i>International Journal of Molecular Epidemiology and Genetics</i> , <b>2012</b> , 3, 39-47	0.9	13
81	Next generation sequencing of CLU, PICALM and CR1: pitfalls and potential solutions. <i>International Journal of Molecular Epidemiology and Genetics</i> , <b>2012</b> , 3, 262-75	0.9	2
80	Investigating statistical epistasis in complex disorders. <i>Journal of Alzheimer's Disease</i> , <b>2011</b> , 25, 635-44	4.3	5

79	A multi-center study of ACE and the risk of late-onset Alzheimer's disease. <i>Journal of Alzheimer's Disease</i> , <b>2011</b> , 24, 587-97	4.3	29
78	Multiple insulin degrading enzyme variants alter in vitro reporter gene expression. <i>PLoS ONE</i> , <b>2011</b> , 6, e21429	3.7	2
77	The three new pathways leading to Alzheimer's disease. <i>Neuropathology and Applied Neurobiology</i> , <b>2011</b> , 37, 353-7	5.2	87
76	Common variants at ABCA7, MS4A6A/MS4A4E, EPHA1, CD33 and CD2AP are associated with Alzheimer's disease. <i>Nature Genetics</i> , <b>2011</b> , 43, 429-35	36.3	1421
75	Investigation of 15 of the top candidate genes for late-onset Alzheimer's disease. <i>Human Genetics</i> , <b>2011</b> , 129, 273-82	6.3	53
74	The role of IREB2 and transforming growth factor beta-1 genetic variants in COPD: a replication case-control study. <i>BMC Medical Genetics</i> , <b>2011</b> , 12, 24	2.1	35
73	Replication of EPHA1 and CD33 associations with late-onset Alzheimer's disease: a multi-centre case-control study. <i>Molecular Neurodegeneration</i> , <b>2011</b> , 6, 54	19	55
72	No evidence that extended tracts of homozygosity are associated with Alzheimer's disease. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2011</b> , 156B, 764-71	3.5	15
71	Replication of BIN1 association with Alzheimer's disease and evaluation of genetic interactions. <i>Journal of Alzheimer's Disease</i> , <b>2011</b> , 24, 751-8	4.3	55
70	Using Fisher's method with PLINK 'LD clumped' output to compare SNP effects across Genome-wide Association Study (GWAS) datasets. <i>International Journal of Molecular Epidemiology and Genetics</i> , <b>2011</b> , 2, 30-5	0.9	3
69	Genetic evidence implicates the immune system and cholesterol metabolism in the aetiology of Alzheimer's disease. <i>PLoS ONE</i> , <b>2010</b> , 5, e13950	3.7	276
68	Concordant association of insulin degrading enzyme gene (IDE) variants with IDE mRNA, Aβ, and Alzheimer's disease. <i>PLoS ONE</i> , <b>2010</b> , 5, e8764	3.7	40
67	Replication of CLU, CR1, and PICALM associations with Alzheimer disease. <i>Archives of Neurology</i> , <b>2010</b> , 67, 961-4		167
66	The dopamine β-hydroxylase -1021C/T polymorphism is associated with the risk of Alzheimer's disease in the Epistasis Project. <i>BMC Medical Genetics</i> , <b>2010</b> , 11, 162	2.1	43
65	Association of MMP-2 polymorphisms with severe and very severe COPD: a case control study of MMPs-1, 9 and 12 in a European population. <i>BMC Medical Genetics</i> , <b>2010</b> , 11, 7	2.1	53
64	Analysis of Genome-Wide Association Study (GWAS) data looking for replicating signals in Alzheimer's disease (AD). <i>International Journal of Molecular Epidemiology and Genetics</i> , <b>2010</b> , 1, 53-66	0.9	13
63	Using In silico LD clumping and meta-analysis of genome-wide datasets as a complementary tool to investigate and validate new candidate biomarkers in Alzheimer's disease. <i>International Journal of Molecular Epidemiology and Genetics</i> , <b>2010</b> , 1, 134-44	0.9	1
62	Prokineticin 1 modulates IL-8 expression via the calcineurin/NFAT signaling pathway. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , <b>2009</b> , 1793, 1315-24	4.9	29



61	Genome-wide association study identifies variants at CLU and PICALM associated with Alzheimer's disease. <i>Nature Genetics</i> , <b>2009</b> , 41, 1088-93	36.3	2018
60	The alpha-1-antitrypsin gene promoter in human A549 lung derived cells, and a novel transcription initiation site. <i>International Journal of Biochemistry and Cell Biology</i> , <b>2009</b> , 41, 1157-64	5.6	4
59	PSEN1 polymorphisms alter the rate of cognitive decline in sporadic Alzheimer's disease patients. <i>Neurobiology of Aging</i> , <b>2009</b> , 30, 1992-9	5.6	13
58	Replication by the Epistasis Project of the interaction between the genes for IL-6 and IL-10 in the risk of Alzheimer's disease. <i>Journal of Neuroinflammation</i> , <b>2009</b> , 6, 22	10.1	41
57	A SNP in the ACT gene associated with astrocytosis and rapid cognitive decline in AD. <i>Neurobiology of Aging</i> , <b>2008</b> , 29, 1167-76	5.6	11
56	Prokineticin 1 signaling and gene regulation in early human pregnancy. <i>Endocrinology</i> , <b>2008</b> , 149, 2877-87	8.8	81
55	Genetic variants of microsomal epoxide hydrolase and glutamate-cysteine ligase in COPD. <i>European Respiratory Journal</i> , <b>2008</b> , 32, 931-7	13.6	30
54	A polymorphism of the alpha1-antitrypsin gene represents a risk factor for liver disease. <i>Hepatology</i> , <b>2008</b> , 47, 127-32	11.2	45
53	SERPINA3 (aka alpha-1-antichymotrypsin). <i>Frontiers in Bioscience - Landmark</i> , <b>2007</b> , 12, 2821-35	2.8	92
52	Sarcoidosis and MIF gene polymorphism: a case-control study in an Irish population. <i>European Respiratory Journal</i> , <b>2007</b> , 29, 325-9	13.6	5
51	Regulatory region single nucleotide polymorphisms of the apolipoprotein E gene and the rate of cognitive decline in Alzheimer's disease. <i>Human Molecular Genetics</i> , <b>2007</b> , 16, 2199-208	5.6	40
50	Variation in the tumour necrosis factor gene is not associated with susceptibility to COPD. <i>European Respiratory Journal</i> , <b>2007</b> , 30, 810-2	13.6	10
49	Effects of Alzheimer's peptide and alpha1-antichymotrypsin on astrocyte gene expression. <i>Neurobiology of Aging</i> , <b>2007</b> , 28, 51-61	5.6	10
48	The hydroxy-methyl-glutaryl CoA reductase promoter polymorphism is associated with Alzheimer's risk and cognitive deterioration. <i>Neuroscience Letters</i> , <b>2007</b> , 416, 66-70	3.3	23
47	Cryptic haplotypes of SERPINA1 confer susceptibility to chronic obstructive pulmonary disease. <i>Human Mutation</i> , <b>2006</b> , 27, 103-9	4.7	54
46	The SERPINE2 gene and chronic obstructive pulmonary disease. <i>American Journal of Human Genetics</i> , <b>2006</b> , 79, 184-6; author reply 186-7	11	27
45	Association of improved pulmonary phenotype in Irish cystic fibrosis patients with a 3' enhancer polymorphism in alpha-1-antitrypsin. <i>Pediatric Pulmonology</i> , <b>2006</b> , 41, 584-91	3.5	7
44	Cystic fibrosis, disease severity, and a macrophage migration inhibitory factor polymorphism. <i>American Journal of Respiratory and Critical Care Medicine</i> , <b>2005</b> , 172, 1412-5	10.2	73



43	A new promoter polymorphism in the alpha-1-antichymotrypsin gene is a disease modifier of Alzheimer's disease. <i>Neurobiology of Aging</i> , <b>2005</b> , 26, 449-53	5.6	33
42	Transforming growth factor beta1 regulates angiotensin II type I receptor gene expression in the extravillous trophoblast cell line SGHPL-4. <i>Molecular Human Reproduction</i> , <b>2005</b> , 11, 847-52	4.4	5
41	Haplotypes of the alpha-1 antitrypsin gene in healthy controls and Z deficiency patients. <i>Human Mutation</i> , <b>2004</b> , 24, 535-6	4.7	8
40	Allele-specific overexpression in astrocytes of an Alzheimer's disease associated alpha-1-antichymotrypsin promoter polymorphism. <i>Molecular Brain Research</i> , <b>2004</b> , 131, 88-92		8
39	Alpha-1-Antitrypsin Deficiency <b>2004</b> , 42-46		
38	Differential regulation of secretory leukocyte protease inhibitor and elafin by progesterone. <i>Biochemical and Biophysical Research Communications</i> , <b>2003</b> , 310, 594-9	3.4	56
37	Amyloid precursor protein (APP) and the biology of proteolytic processing: relevance to Alzheimer's disease. <i>International Journal of Biochemistry and Cell Biology</i> , <b>2003</b> , 35, 1505-35	5.6	152
36	Serpins: structure, function and molecular evolution. <i>International Journal of Biochemistry and Cell Biology</i> , <b>2003</b> , 35, 1536-47	5.6	150
35	Oncostatin M induced alpha1-antitrypsin (AAT) gene expression in Hep G2 cells is mediated by a 3' enhancer. <i>Biochemical Journal</i> , <b>2002</b> , 365, 555-60	3.8	24
34	Gene regulation of the serine proteinase inhibitors $\alpha_1$ -antitrypsin and $\alpha_1$ -antichymotrypsin. <i>Biochemical Society Transactions</i> , <b>2002</b> , 30, 93-98	5.1	89
33	Polymorphism in the alpha(1)-antichymotrypsin (ACT) gene promoter: effect on expression in transfected glial and liver cell lines and plasma ACT concentrations. <i>Human Genetics</i> , <b>2001</b> , 109, 303-10	6.3	26
32	An alpha1-antitrypsin enhancer polymorphism is a genetic modifier of pulmonary outcome in cystic fibrosis. <i>European Journal of Human Genetics</i> , <b>2001</b> , 9, 273-8	5.3	33
31	HLA class I, II & III genes in confirmed late-onset Alzheimer's disease. <i>Neurobiology of Aging</i> , <b>2001</b> , 22, 71-7	5.6	25
30	A polymorphism in the tumor necrosis factor-alpha gene promoter region may predispose to a poor prognosis in COPD. <i>Chest</i> , <b>2000</b> , 118, 971-5	5.3	88
29	Evaluation of polymorphisms in the presenilin-1 gene and the butyrylcholinesterase gene as risk factors in sporadic Alzheimer's disease. <i>European Journal of Human Genetics</i> , <b>1999</b> , 7, 659-63	5.3	52
28	Elastase Inhibitors in the Lung: Expression and Functional Relationships <b>1999</b> , 69-94		1
27	Mutation in an alpha1-antitrypsin enhancer results in an interleukin-6 deficient acute-phase response due to loss of cooperativity between transcription factors. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , <b>1997</b> , 1362, 67-76	6.9	28
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24	The sensitivity of competitive hybridisation for the detection of mutant p53 alleles in a background of wild type. <i>Biochemical Society Transactions</i> , <b>1995</b> , 23, 129S	5.1	
23	Simple and efficient cDNA capture utilising a short gene-specific probe attached to magnetic beads. <i>Biochemical Society Transactions</i> , <b>1994</b> , 22, 453S	5.1	
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21	Transcription termination and DNase I hypersensitive sites in the 3' flanking sequence of the alpha-1-antitrypsin gene. <i>Biochemical Society Transactions</i> , <b>1992</b> , 20, 348S	5.1	
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19	Health, social functioning, and marital status: Stability and change among elderly recently widowed women. <i>International Journal of Geriatric Psychiatry</i> , <b>1992</b> , 7, 813-817	3.9	25
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15	Benzodiazepine withdrawal and rebound insomnia. <i>Acta Psychiatrica Scandinavica</i> , <b>1989</b> , 80, 297-8	6.5	4
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8	Complement receptor 1 gene (CR1) intragenic duplication and risk of Alzheimer's disease		1

7	A meta-analysis of epigenome-wide association studies in Alzheimer's disease highlights novel differentially methylated loci across cortex	8
6	Exome sequencing identifies rare damaging variants in the ATP8B4 and ABCA1 genes as novel risk factors for Alzheimer's Disease	1
5	New insights on the genetic etiology of Alzheimer's and related dementia	25
4	A comprehensive assessment of benign genetic variability for neurodegenerative disorders	7
3	Gene-Based Analysis in HRC Imputed Genome Wide Association Data Identifies Three Novel Genes For Alzheimer's Disease	1
2	Genetic variability in response to A $\beta$ deposition influences Alzheimer's risk	6
1	Genetics of Alzheimer's Disease 1-16	1