Christine Van Broeckhoven

List of Publications by Year in descending order

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799 papers	76,635 citations	492 129 h-index	911 241 g-index
872	872	872	51248
all docs	docs citations	times ranked	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.	21.4	3,741
2	Genome-wide association study identifies variants at CLU and PICALM associated with Alzheimer's disease. Nature Genetics, 2009, 41, 1088-1093.	21.4	2,697
3	Genome-wide association study identifies variants at CLU and CR1 associated with Alzheimer's disease. Nature Genetics, 2009, 41, 1094-1099.	21.4	2,155
4	Genetic meta-analysis of diagnosed Alzheimer's disease identifies new risk loci and implicates Aβ, tau, immunity and lipid processing. Nature Genetics, 2019, 51, 414-430.	21.4	1,962
5	NanoPack: visualizing and processing long-read sequencing data. Bioinformatics, 2018, 34, 2666-2669.	4.1	1,713
6	Common variants at ABCA7, MS4A6A/MS4A4E, EPHA1, CD33 and CD2AP are associated with Alzheimer's disease. Nature Genetics, 2011, 43, 429-435.	21.4	1,708
7	Null mutations in progranulin cause ubiquitin-positive frontotemporal dementia linked to chromosome 17q21. Nature, 2006, 442, 920-924.	27.8	1,386
8	Atherosclerosis, apolipoprotein E, and prevalence of dementia and Alzheimer's disease in the Rotterdam Study. Lancet, The, 1997, 349, 151-154.	13.7	1,304
9	De Novo Mutations in the Sodium-Channel Gene SCN1A Cause Severe Myoclonic Epilepsy of Infancy. American Journal of Human Genetics, 2001, 68, 1327-1332.	6.2	1,111
10	The <i>C9orf72</i> GGGGCC Repeat Is Translated into Aggregating Dipeptide-Repeat Proteins in FTLD/ALS. Science, 2013, 339, 1335-1338.	12.6	1,095
11	VEGF is a modifier of amyotrophic lateral sclerosis in mice and humans and protects motoneurons against ischemic death. Nature Genetics, 2003, 34, 383-394.	21.4	794
12	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
13	Mutation of POLG is associated with progressive external ophthalmoplegia characterized by mtDNA deletions. Nature Genetics, 2001, 28, 211-212.	21.4	748
14	Extra-pair paternity results from female preference for high-quality males in the blue tit. Nature, 1992, 357, 494-496.	27.8	720
15	Presenile dementia and cerebral haemorrhage linked to a mutation at codon 692 of the β–amyloid precursor protein gene. Nature Genetics, 1992, 1, 218-221.	21.4	715
16	New insights into the genetic etiology of Alzheimer's disease and related dementias. Nature Genetics, 2022, 54, 412-436.	21.4	700
17	The genetic landscape of Alzheimer disease: clinical implications and perspectives. Genetics in Medicine, 2016, 18, 421-430.	2.4	695
18	TREM2 mutations implicated in neurodegeneration impair cell surface transport and phagocytosis. Science Translational Medicine, 2014, 6, 243ra86.	12.4	600

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19	A C9orf72 promoter repeat expansion in a Flanders-Belgian cohort with disorders of the frontotemporal lobar degeneration-amyotrophic lateral sclerosis spectrum: a gene identification study. Lancet Neurology, The, 2012, 11, 54-65.	10.2	565
20	APOE and Alzheimer disease: a major gene with semi-dominant inheritance. Molecular Psychiatry, 2011, 16, 903-907.	7.9	529
21	TDP-43 transgenic mice develop spastic paralysis and neuronal inclusions characteristic of ALS and frontotemporal lobar degeneration. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 3858-3863.	7.1	491
22	Common variants at 7p21 are associated with frontotemporal lobar degeneration with TDP-43 inclusions. Nature Genetics, 2010, 42, 234-239.	21.4	479
23	Smoking and risk of dementia and Alzheimer's disease in a population-based cohort study: the Rotterdam Study. Lancet, The, 1998, 351, 1840-1843.	13.7	475
24	Collaborative Analysis of α-Synuclein Gene Promoter Variability and Parkinson Disease. JAMA - Journal of the American Medical Association, 2006, 296, 661.	7.4	467
25	Apolipoprotein E4 allele in a population–based study of early–onset Alzheimer's disease. Nature Genetics, 1994, 7, 74-78.	21.4	460
26	Amyloid beta protein precursor gene and hereditary cerebral hemorrhage with amyloidosis (Dutch). Science, 1990, 248, 1120-1122.	12.6	445
27	Highly efficient gene delivery by mRNA electroporation in human hematopoietic cells: superiority to lipofection and passive pulsing of mRNA and to electroporation of plasmid cDNA for tumor antigen loading of dendritic cells. Blood, 2001, 98, 49-56.	1.4	438
28	Genetic etiology of Parkinson disease associated with mutations in the SNCA, PARK2, PINK1, PARK7, and LRRK2 genes: a mutation update. Human Mutation, 2010, 31, 763-780.	2.5	428
29	Bidirectional transcripts of the expanded C9orf72 hexanucleotide repeat are translated into aggregating dipeptide repeat proteins. Acta Neuropathologica, 2013, 126, 881-893.	7.7	427
30	Genetic Association of Apolipoprotein E with Age-Related Macular Degeneration. American Journal of Human Genetics, 1998, 63, 200-206.	6.2	425
31	Locusâ€specific mutation databases for neurodegenerative brain diseases. Human Mutation, 2012, 33, 1340-1344.	2.5	414
32	Molecular genetics of earlyâ€onset Alzheimer's disease revisited. Alzheimer's and Dementia, 2016, 12, 733-748.	0.8	409
33	Genetic linkage studies suggest that Alzheimer's disease is not a single homogeneous disorder. Nature, 1990, 347, 194-197.	27.8	407
34	The peripheral myelin protein gene PMP–22 is contained within the Charcot–Marie–Tooth disease type 1A duplication. Nature Genetics, 1992, 1, 171-175.	21.4	404
35	Genome Scan Meta-Analysis of Schizophrenia and Bipolar Disorder, Part III: Bipolar Disorder. American Journal of Human Genetics, 2003, 73, 49-62.	6.2	400
36	Estimation of the Genetic Contribution of Presenilin-1 and -2 Mutations in a Population-Based Study of Presenile Alzheimer Disease. Human Molecular Genetics, 1998, 7, 43-51.	2.9	396

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37	Hot-spot residue in small heat-shock protein 22 causes distal motor neuropathy. Nature Genetics, 2004, 36, 597-601.	21.4	395
38	<scp>sTREM</scp> 2 cerebrospinal fluid levels are a potential biomarker for microglia activity in earlyâ€stage Alzheimer's disease and associate with neuronal injury markers. EMBO Molecular Medicine, 2016, 8, 466-476.	6.9	392
39	Risk Estimates of Dementia by Apolipoprotein E Genotypes From a Population-Based Incidence Study: The Rotterdam Study. Archives of Neurology, 1998, 55, 964.	4.5	378
40	A Multicenter Study of Glucocerebrosidase Mutations in Dementia With Lewy Bodies. JAMA Neurology, 2013, 70, 727.	9.0	374
41	Clinical Phenotypes of Different MPZ (P0) Mutations May Include Charcot–Marie–Tooth Type 1B, Dejerine–Sottas, and Congenital Hypomyelination. Neuron, 1996, 17, 451-460.	8.1	372
42	Estimation of the Mutation Frequencies in Charcot-Marie-Tooth Disease Type 1 and Hereditary Neuropathy with Liability to Pressure Palsies: A European Collaborative Study. European Journal of Human Genetics, 1996, 4, 25-33.	2.8	371
43	APP duplication is sufficient to cause early onset Alzheimer's dementia with cerebral amyloid angiopathy. Brain, 2006, 129, 2977-2983.	7.6	337
44	Phenotype variability in progranulin mutation carriers: a clinical, neuropsychological, imaging and genetic study. Brain, 2008, 131, 732-746.	7.6	331
45	The role of tau (MAPT) in frontotemporal dementia and related tauopathies. Human Mutation, 2004, 24, 277-295.	2.5	323
46	Mapping of a gene predisposing to early–onset Alzheimer's disease to chromosome 14q24.3. Nature Genetics, 1992, 2, 335-339.	21.4	321
47	Increased expression of BIN1 mediates Alzheimer genetic risk by modulating tau pathology. Molecular Psychiatry, 2013, 18, 1225-1234.	7.9	321
48	Genetic insights in Alzheimer's disease. Lancet Neurology, The, 2013, 12, 92-104.	10.2	310
49	Mean age-of-onset of familial alzheimer disease caused by presenilin mutations correlates with both increased Al²42 and decreased Al²40. Human Mutation, 2006, 27, 686-695.	2.5	306
50	hnRNP A3 binds to GGGGCC repeats and is a constituent of p62-positive/TDP43-negative inclusions in the hippocampus of patients with C9orf72 mutations. Acta Neuropathologica, 2013, 125, 413-423.	7.7	302
51	Frontotemporal dementia and its subtypes: a genome-wide association study. Lancet Neurology, The, 2014, 13, 686-699.	10.2	302
52	Association of LRRK2 exonic variants with susceptibility to Parkinson's disease: a case–control study. Lancet Neurology, The, 2011, 10, 898-908.	10.2	294
53	A novel Alzheimer disease locus located near the gene encoding tau protein. Molecular Psychiatry, 2016, 21, 108-117.	7.9	260
54	novoSNP, a novel computational tool for sequence variation discovery. Genome Research, 2005, 15, 436-442.	5.5	254

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55	<i>POLG</i> mutations in neurodegenerative disorders with ataxia but no muscle involvement. Neurology, 2004, 63, 1251-1257.	1.1	252
56	Apolipoprotein E epsilon4 and the risk of dementia with stroke. A population-based investigation. JAMA - Journal of the American Medical Association, 1997, 277, 818-821.	7.4	252
57	A Panâ€ <scp>E</scp> uropean Study of the <i>C9orf72</i> Repeat Associated with <scp>FTLD</scp> : Geographic Prevalence, Genomic Instability, and Intermediate Repeats. Human Mutation, 2013, 34, 363-373.	2.5	247
58	Potent amyloidogenicity and pathogenicity of AÎ ² 43. Nature Neuroscience, 2011, 14, 1023-1032.	14.8	245
59	The genetics and neuropathology of frontotemporal lobar degeneration. Acta Neuropathologica, 2012, 124, 353-372.	7.7	242
60	Drosophila screen connects nuclear transport genes to DPR pathology in c9ALS/FTD. Scientific Reports, 2016, 6, 20877.	3.3	239
61	Mutations in DNAJC5, Encoding Cysteine-String Protein Alpha, Cause Autosomal-Dominant Adult-Onset Neuronal Ceroid Lipofuscinosis. American Journal of Human Genetics, 2011, 89, 241-252.	6.2	236
62	Mutations in the peripheral myelin genes and associated genes in inherited peripheral neuropathies. Human Mutation, 1999, 13, 11-28.	2.5	232
63	ALS Genes in the Genomic Era and their Implications for FTD. Trends in Genetics, 2018, 34, 404-423.	6.7	229
64	Mutations in SEPT9 cause hereditary neuralgic amyotrophy. Nature Genetics, 2005, 37, 1044-1046.	21.4	222
65	FUS pathology defines the majority of tau- and TDP-43-negative frontotemporal lobar degeneration. Acta Neuropathologica, 2010, 120, 33-41.	7.7	222
66	Recessive POLG mutations presenting with sensory and ataxic neuropathy in compound heterozygote patients with progressive external ophthalmoplegia. Neuromuscular Disorders, 2003, 13, 133-142.	0.6	216
67	The Thr124Met mutation in the peripheral myelin protein zero (MPZ) gene is associated with a clinically distinct Charcot–Marie–Tooth phenotype. Brain, 1999, 122, 281-290.	7.6	215
68	Pathogenesis of polyglutamine disorders: aggregation revisited. Human Molecular Genetics, 2003, 12, R173-R186.	2.9	212
69	A novel presenilin 1 mutation associated with Pick's disease but not βâ€amyloid plaques. Annals of Neurology, 2004, 55, 617-626.	5.3	210
70	Genetic variation in DPP6 is associated with susceptibility to amyotrophic lateral sclerosis. Nature Genetics, 2008, 40, 29-31.	21.4	205
71	Presenilins and Alzheimer disease. Nature Genetics, 1995, 11, 230-232.	21.4	204
72	Germline mosaicism and Duchenne muscular dystrophy mutations. Nature, 1987, 329, 554-556.	27.8	201

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73	The C9orf72 repeat size correlates with onset age of disease, DNA methylation and transcriptional downregulation of the promoter. Molecular Psychiatry, 2016, 21, 1112-1124.	7.9	201
74	α‣ynuclein promoter confers susceptibility to Parkinson's disease. Annals of Neurology, 2004, 56, 591-595.	5.3	200
75	Genetic contribution of <i>FUS</i> to frontotemporal lobar degeneration. Neurology, 2010, 74, 366-371.	1.1	197
76	Molecular biological characterization of an azole-resistant Candida glabrata isolate. Antimicrobial Agents and Chemotherapy, 1997, 41, 2229-2237.	3.2	196
77	Molecular genetics of Alzheimer's disease: An update. Annals of Medicine, 2008, 40, 562-583.	3.8	196
78	ITPR2 as a susceptibility gene in sporadic amyotrophic lateral sclerosis: a genome-wide association study. Lancet Neurology, The, 2007, 6, 869-877.	10.2	195
79	Serum biomarker for progranulinâ€associated frontotemporal lobar degeneration. Annals of Neurology, 2009, 65, 603-609.	5.3	195
80	Progress in unraveling the genetic etiology of Parkinson disease in a genomic era. Trends in Genetics, 2015, 31, 140-149.	6.7	193
81	Further evidence that neurofilament light chain gene mutations can cause Charcot-Marie-Tooth disease type 2E. Annals of Neurology, 2001, 49, 245-249.	5.3	188
82	Germinal mosaicism increases the recurrence risk for 'new' Duchenne muscular dystrophy mutations Journal of Medical Genetics, 1989, 26, 553-559.	3.2	187
83	Alzheimer risk associated with a copy number variation in the complement receptor 1 increasing C3b/C4b binding sites. Molecular Psychiatry, 2012, 17, 223-233.	7.9	179
84	Pathogenic APP mutations near the gamma-secretase cleavage site differentially affect Abeta secretion and APP C-terminal fragment stability. Human Molecular Genetics, 2001, 10, 1665-1671.	2.9	178
85	De-novo mutation in hereditary motor and sensory neuropathy type I. Lancet, The, 1992, 339, 1081-1082.	13.7	177
86	Promoter Mutations That Increase Amyloid Precursor-Protein Expression Are Associated with Alzheimer Disease. American Journal of Human Genetics, 2006, 78, 936-946.	6.2	173
87	Genetic variation in the KIF1B locus influences susceptibility to multiple sclerosis. Nature Genetics, 2008, 40, 1402-1403.	21.4	173
88	Convergent genetic and expression data implicate immunity in Alzheimer's disease. Alzheimer's and Dementia, 2015, 11, 658-671.	0.8	173
89	Molecular genetic analysis of familial early-onset Alzheimer's disease linked to chromosome 14q24.3. Human Molecular Genetics, 1995, 4, 2363-2371.	2.9	171
90	De novoSCN1Amutations are a major cause of severe myoclonic epilepsy of infancy. Human Mutation, 2003, 21, 615-621.	2.5	170

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91	The presenilin genes: a new gene family involved in Alzheimer disease pathology. Human Molecular Genetics, 1996, 5, 1449-1455.	2.9	169
92	Presenilin mutations in Alzheimer's disease. Human Mutation, 1998, 11, 183-190.	2.5	169
93	Human Meiotic Recombination Products Revealed by Sequencing a Hotspot for Homologous Strand Exchange in Multiple HNPP Deletion Patients. American Journal of Human Genetics, 1998, 62, 1023-1033.	6.2	168
94	Dense-Core Plaques in Tg2576 and PSAPP Mouse Models of Alzheimer's Disease Are Centered on Vessel Walls. American Journal of Pathology, 2005, 167, 527-543.	3.8	168
95	Disruption of endocytic trafficking in frontotemporal dementia with CHMP2B mutations. Human Molecular Genetics, 2010, 19, 2228-2238.	2.9	163
96	Progranulin null mutations in both sporadic and familial frontotemporal dementia. Human Mutation, 2007, 28, 846-855.	2.5	162
97	Genetic variability in <i>progranulin</i> contributes to risk for clinically diagnosed Alzheimer disease. Neurology, 2008, 71, 656-664.	1.1	158
98	Investigating the role of rare heterozygous TREM2 variants in Alzheimer's disease and frontotemporal dementia. Neurobiology of Aging, 2014, 35, 726.e11-726.e19.	3.1	158
99	Flemish and Dutch Mutations in Amyloid β Precursor Protein Have Different Effects on Amyloid β Secretion. Neurobiology of Disease, 1998, 5, 281-286.	4.4	157
100	The molecular basis of the frontotemporal lobar degeneration–amyotrophic lateral sclerosis spectrum. Annals of Medicine, 2012, 44, 817-828.	3.8	157
101	High-density SNP haplotyping suggests altered regulation of tau gene expression in progressive supranuclear palsy. Human Molecular Genetics, 2005, 14, 3281-3292.	2.9	156
102	Sequencing an Ashkenazi reference panel supports population-targeted personal genomics and illuminates Jewish and European origins. Nature Communications, 2014, 5, 4835.	12.8	156
103	Common pathobiochemical hallmarks of progranulin-associated frontotemporal lobar degeneration and neuronal ceroid lipofuscinosis. Acta Neuropathologica, 2014, 127, 845-60.	7.7	156
104	The dopamine D4 receptor gene 48-base-pair-repeat polymorphism and mood disorders: A meta-analysis. Biological Psychiatry, 2005, 57, 999-1003.	1.3	155
105	Gene-Wide Analysis Detects Two New Susceptibility Genes for Alzheimer's Disease. PLoS ONE, 2014, 9, e94661.	2.5	155
106	CAG repeat expansion in the TATA box-binding protein gene causes autosomal dominant cerebellar ataxia. Brain, 2001, 124, 1939-1947.	7.6	154
107	Altered brain white matter integrity in healthy carriers of the <i>APOE</i> Îμ4 allele. Neurology, 2006, 66, 1029-1033.	1.1	153
108	Meta-analysis of the Association Between Variants in SORL1 and Alzheimer Disease. Archives of Neurology, 2011, 68, 99.	4.5	153

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109	Two divergent types of nerve pathology in patients with different P sub 0 mutations in Charcot-Marie-Tooth disease. Neurology, 1996, 47, 761-765.	1.1	152
110	Mutations in <i>GDAP1</i> . Neurology, 2002, 59, 1865-1872.	1.1	152
111	The gene for autosomal dominant cerebellar ataxia with pigmentary macular dystrophy maps to chromosome 3p12–p21.1. Nature Genetics, 1995, 10, 84-88.	21.4	151
112	Current insights into the C9orf72 repeat expansion diseases of the FTLD/ALS spectrum. Trends in Neurosciences, 2013, 36, 450-459.	8.6	151
113	Loss of <i>TBK1</i> is a frequent cause of frontotemporal dementia in a Belgian cohort. Neurology, 2015, 85, 2116-2125.	1.1	151
114	Novel missense mutation in the early growth response 2 gene associated with Dejerine–Sottas syndrome phenotype. Neurology, 1999, 52, 1827-1827.	1.1	150
115	Variability of 5-HT2C receptor cys23ser polymorphism among European populations and vulnerability to affective disorder. Molecular Psychiatry, 2001, 6, 579-585.	7.9	150
116	A deletion in <i>SCN1B</i> is associated with febrile seizures and early-onset absence epilepsy. Neurology, 2003, 61, 854-856.	1.1	150
117	Detection of expanded CAG repeats in Bipolar Affective Disorder using the repeat expansion detection (RED) method. Neurobiology of Disease, 1995, 2, 55-62.	4.4	148
118	<i>Progranulin</i> genetic variability contributes to amyotrophic lateral sclerosis. Neurology, 2008, 71, 253-259.	1.1	148
119	Novel POLG mutations in progressive external ophthalmoplegia mimicking mitochondrial neurogastrointestinal encephalomyopathy. European Journal of Human Genetics, 2003, 11, 547-549.	2.8	145
120	A Genomewide Screen for Late-Onset Alzheimer Disease in a Genetically Isolated Dutch Population. American Journal of Human Genetics, 2007, 81, 17-31.	6.2	145
121	Alzheimer and Parkinson Diagnoses in Progranulin Null Mutation Carriers in an Extended Founder Family. Archives of Neurology, 2007, 64, 1436.	4.5	143
122	Structural variants identified by Oxford Nanopore PromethION sequencing of the human genome. Genome Research, 2019, 29, 1178-1187.	5.5	143
123	A novel GABRG2 mutation associated with febrile seizures. Neurology, 2006, 67, 687-690.	1.1	142
124	Earlyâ€onset Alzheimer's disease in 2 large Belgian families. Neurology, 1991, 41, 62-62.	1.1	141
125	Common variants in Alzheimer's disease and risk stratification by polygenic risk scores. Nature Communications, 2021, 12, 3417.	12.8	140
126	Glucocorticoid Receptor Gene-Based SNP Analysis in Patients with Recurrent Major Depression. Neuropsychopharmacology, 2006, 31, 620-627.	5.4	139

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127	Current status on Alzheimer disease molecular genetics: from past, to present, to future. Human Molecular Genetics, 2010, 19, R4-R11.	2.9	138
128	Mutational analysis of the MPZ, PMP22 and Cx32 genes in patients of Spanish ancestry with Charcot-Marie-Tooth disease and hereditary neuropathy with liability to pressure palsies. Human Genetics, 1997, 99, 746-754.	3.8	137
129	Variant Alzheimer's disease with spastic paraparesis and cotton wool plaques is caused by PS-1 mutations that lead to exceptionally high amyloid-? concentrations. Annals of Neurology, 2000, 48, 806-808.	5.3	135
130	Nonfibrillar diffuse amyloid deposition due to a gamma42-secretase site mutation points to an essential role for N-truncated Abeta42 in Alzheimer's disease. Human Molecular Genetics, 2000, 9, 2589-2598.	2.9	135
131	Association between COMT (Val158Met) functional polymorphism and early onset in patients with major depressive disorder in a European multicenter genetic association study. Molecular Psychiatry, 2005, 10, 598-605.	7.9	134
132	Reduced functional brain activity response in cognitively intact apolipoprotein E ε4 carriers. Brain, 2006, 129, 1240-1248.	7.6	133
133	Lrrk2 R1441C parkinsonism is clinically similar to sporadic Parkinson disease. Neurology, 2008, 70, 1456-1460.	1.1	132
134	The role of ABCA7 in Alzheimer's disease: evidence from genomics, transcriptomics and methylomics. Acta Neuropathologica, 2019, 138, 201-220.	7.7	132
135	Molecular genetics of Alzheimer's disease. Annals of Medicine, 1998, 30, 560-565.	3.8	131
136	CHMP2B C-truncating mutations in frontotemporal lobar degeneration are associated with an aberrant endosomal phenotype in vitro. Human Molecular Genetics, 2008, 17, 313-322.	2.9	131
137	Origin of the de novo duplication in Charcot — Marie — Tooth disease type 1A: unequal nonsister chromatid exchange during spermatogenesis. Human Molecular Genetics, 1993, 2, 2031-2035.	2.9	130
138	APOE genotype does not modulate age of onset in families with chromosome 14 encoded Alzheimer's disease. Neuroscience Letters, 1994, 169, 179-180.	2.1	130
139	The apolipoprotein E ε2 allele is associated with an increased risk of earlyâ€onset alzheimer's disease and a reduced survival. Annals of Neurology, 1995, 37, 605-610.	5.3	129
140	Granulin mutations associated with frontotemporal lobar degeneration and related disorders: An update. Human Mutation, 2008, 29, 1373-1386.	2.5	126
141	Loss of ALS-associated TDP-43 in zebrafish causes muscle degeneration, vascular dysfunction, and reduced motor neuron axon outgrowth. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 4986-4991.	7.1	126
142	Spinocerebellar ataxia type 7 associated with pigmentary retinal dystrophy. European Journal of Human Genetics, 2004, 12, 2-15.	2.8	124
143	Mutations in ABCA7 in a Belgian cohort of Alzheimer's disease patients: a targeted resequencing study. Lancet Neurology, The, 2015, 14, 814-822.	10.2	124
144	Estimation of the size of the chromosome 17p11.2 duplication in Charcot-Marie-Tooth neuropathy type 1a (CMT1a). HMSN Collaborative Research Group Journal of Medical Genetics, 1992, 29, 5-11.	3.2	123

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145	TMEM106B is a genetic modifier of frontotemporal lobar degeneration with C9orf72 hexanucleotide repeat expansions. Acta Neuropathologica, 2014, 127, 407-418.	7.7	123
146	The pursuit of susceptibility genes for Alzheimer's disease: progress and prospects. Trends in Genetics, 2010, 26, 84-93.	6.7	122
147	Pathological mechanisms underlying TDP-43 driven neurodegeneration in FTLD-ALS spectrum disorders. Human Molecular Genetics, 2013, 22, R77-R87.	2.9	122
148	Autosornal recessive hereditary motor and sensory neuropathy with focally folded myelin sheaths. Neurology, 1996, 46, 1318-1318.	1.1	121
149	Rescue of Progranulin Deficiency Associated with Frontotemporal Lobar Degeneration by Alkalizing Reagents and Inhibition of Vacuolar ATPase. Journal of Neuroscience, 2011, 31, 1885-1894.	3.6	121
150	Large-scale replication and heterogeneity in Parkinson disease genetic loci. Neurology, 2012, 79, 659-667.	1.1	119
151	A blinded international study on the reliability of genetic testing for GGGGCC-repeat expansions in <i>C9orf72</i> reveals marked differences in results among 14 laboratories. Journal of Medical Genetics, 2014, 51, 419-424.	3.2	118
152	A Genome Wide Search for Susceptibility Loci in Three European Malignant Hyperthermia Pedigrees. Human Molecular Genetics, 1997, 6, 953-961.	2.9	117
153	Câ€ŧerminal neurogranin is increased in cerebrospinal fluid but unchanged in plasma in Alzheimer's disease. Alzheimer's and Dementia, 2015, 11, 1461-1469.	0.8	117
154	Mutations other than null mutations producing a pathogenic loss of progranulin in frontotemporal dementia. Human Mutation, 2007, 28, 416-416.	2.5	116
155	Transmission and ageâ€atâ€onset patterns in familial Alzheimer's disease. Neurology, 1990, 40, 395-395.	1.1	115
156	Estrogen use and early onset Alzheimer's disease: a population-based study. Journal of Neurology, Neurosurgery and Psychiatry, 1999, 67, 779-781.	1.9	114
157	Relationship between C9orf72 repeat size and clinical phenotype. Current Opinion in Genetics and Development, 2017, 44, 117-124.	3.3	114
158	WT1 MUTATION IN MALIGNANT MESOTHELIOMA AND WT1 IMMUNOREACTIVITY IN RELATION TOp53 AND GROWTH FACTOR RECEPTOR EXPRESSION, CELL-TYPE TRANSITION, AND PROGNOSIS. , 1997, 181, 67-74.		112
159	The influence of apoe status on episodic and semantic memory: Data from a population-based study Neuropsychology, 2006, 20, 645-657.	1.3	112
160	Reduced hippocampal volume in non-demented carriers of the apolipoprotein E ɛ4: Relation to chronological age and recognition memory. Neuroscience Letters, 2006, 396, 23-27.	2.1	112
161	Amyloid precursor protein gene mutation in early-onset Alzheimer's disease. Lancet, The, 1991, 337, 978.	13.7	110
162	Rapid screening of myelin genes in CMT1 patients by SSCP analysis: identification of new mutations and polymorphisms in the PO gene. Human Genetics, 1994, 94, 653-657.	3.8	110

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163	Prospective Belgian study of neurodegenerative and vascular dementia: APOE genotype effects. Journal of Neurology, Neurosurgery and Psychiatry, 2003, 74, 1148-1151.	1.9	110
164	No association of CSF biomarkers with APOEÂ4, plaque and tangle burden in definite Alzheimer's disease. Brain, 2007, 130, 2320-2326.	7.6	110
165	Loss of progranulin function in frontotemporal lobar degeneration. Trends in Genetics, 2008, 24, 186-194.	6.7	110
166	TMEM106B is associated with frontotemporal lobar degeneration in a clinically diagnosed patient cohort. Brain, 2011, 134, 808-815.	7.6	110
167	Dense-Core Senile Plaques in the Flemish Variant of Alzheimer's Disease Are Vasocentric. American Journal of Pathology, 2002, 161, 507-520.	3.8	108
168	Genetic and Clinical Features of Progranulin-Associated Frontotemporal Lobar Degeneration. Archives of Neurology, 2011, 68, 488.	4.5	108
169	Association of short-term cognitive decline and MCI-to-AD dementia conversion with CSF, MRI, amyloid- and 18F-FDG-PET imaging. NeuroImage: Clinical, 2019, 22, 101771.	2.7	108
170	Twinkle and POLG defects enhance age-dependent accumulation of mutations in the control region of mtDNA. Nucleic Acids Research, 2004, 32, 3053-3064.	14.5	107
171	Genetic variability in the mitochondrial serine protease <i>HTRA2</i> contributes to risk for Parkinson disease. Human Mutation, 2008, 29, 832-840.	2.5	107
172	Nonviral transfection of distinct types of human dendritic cells: high-efficiency gene transfer by electroporation into hematopoietic progenitor- but not monocyte-derived dendritic cells. Gene Therapy, 1998, 5, 700-707.	4.5	105
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