William K Scott

List of Publications by Year
in descending order

220 papers	18,821 citations	28736 57 h-index	15698 129 g-index
236 all docs	236 docs citations	236 times ranked	23477 citing authors

#	Article	IF	Citations
1	Genetic variants in the SHISA6 gene are associated with delayed cognitive impairment in two family datasets. Alzheimer's and Dementia, 2023, 19, 611-620.	0.4	4
2	Neuropathological lesions and their contribution to dementia and cognitive impairment in a heterogeneous clinical population. Alzheimer's and Dementia, 2022, 18, 2403-2412.	0.4	4
3	The genetic architecture of Alzheimer disease risk in the Ohio and Indiana Amish. Human Genetics and Genomics Advances, 2022, 3, 100114.	1.0	1
4	Lower Levels of Education Are Associated with Cognitive Impairment in the Old Order Amish. Journal of Alzheimer's Disease, 2021, 79, 451-458.	1.2	8
5	Methylome-wide Analysis Reveals Epigenetic Marks Associated With Resistance to Tuberculosis in Human Immunodeficiency Virusâ€“Infected Individuals From East Africa. Journal of Infectious Diseases, 2021, 224, 695-704.	1.9	1
6	Increased APOE Îµ4 expression is associated with the difference in Alzheimer's disease risk from diverse ancestral backgrounds. Alzheimer's and Dementia, 2021, 17, 1179-1188.	0.4	33
7	Transdisciplinary Perspectives on Precision Medicine. Health Equity, 2021, 5, 288-298.	0.8	1
8	Genomewide Association Studies of <scp>LRRK2</scp> Modifiers of Parkinson's Disease. Annals of Neurology, 2021, 90, 76-88.	2.8	30
9	Association of Smoking, Alcohol Consumption, Blood Pressure, Body Mass Index, and Glycemic Risk Factors With Age-Related Macular Degeneration. JAMA Ophthalmology, 2021, 139, 1299.	1.4	29
10	Plasma Metabolomics of Intermediate and Neovascular Age-Related Macular Degeneration Patients. Cells, 2021, 10, 3141.	1.8	13
11	Assessment of ADâ€related plasma biomarkers in diverse ancestral populations. Alzheimer's and Dementia, 2021, 17, .	0.4	0
12	Neuropathologic lesions and comorbidity in Alzheimer disease and related dementias in a heterogeneous clinical population. Alzheimer's and Dementia, 2021, 17, e056249.	0.4	0
13	Association of a locus on chromosome 17 with earlier age at onset of cognitive impairment in a familial Amish dataset. Alzheimer's and Dementia, 2021, 17, e056288.	0.4	0
14	Genomeâ€wide association for protective variants in Alzheimerâ€™s disease in the Midwestern Amish. Alzheimer's and Dementia, 2021, 17, e056363.	0.4	0
15	Preferential preservation of constructional praxis delayed recall compared to word list delayed recall in the Amish. Alzheimer's and Dementia, 2021, 17, e056386.	0.4	0
16	Genetic risk score for Alzheimer's disease in the Amish highlights differences in the genetic architecture compared to other European ancestry populations.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e053304.	0.4	0
17	Suggestive linkage and association of preserved cognition to chromosome 18 in genetically at-risk Amish.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e056306.	0.4	0
18	Genome-wide association study of cognitive status and decline in the Amish.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e056525.	0.4	0

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19	Novel Variants in LRRK2 and GBA Identified in Latino Parkinson Disease Cohort Enriched for Caribbean Origin. Frontiers in Neurology, 2020, 11, 573733.	1.1	6
20	Reduction of neurogranin immunostaining in the hippocampus of postâ€mortem brain of Alzheimerâ€™s disease. Alzheimer's and Dementia, 2020, 16, e040707.	0.4	0
21	Longitudinal assessment of cognitive decline in the Amish. Alzheimer's and Dementia, 2020, 16, e043440.	0.4	0
22	Increased APOEâ€e4 expression is associated with reactive A1 astrocytes and may confer the difference in Alzheimer disease risk from different ancestral backgrounds. Alzheimer's and Dementia, 2020, 16, e045415.	0.4	0
23	Joint linkage and association mapping of preserved cognition in the oldâ€order Amish. Alzheimer's and Dementia, 2020, 16, e046416.	0.4	0
24	Spectrum of Genetic Variants Associated with Anterior Segment Dysgenesis in South Florida. Genes, 2020, 11, 350.	1.0	14
25	AMISH EYE STUDY. Retina, 2019, 39, 1540-1550.	1.0	17
26	Motivations for Participation in Parkinson Disease Genetic Research Among Hispanics versus Non-Hispanics. Frontiers in Genetics, 2019, 10, 658.	1.1	10
27	Rare variants and loci for age-related macular degeneration in the Ohio and Indiana Amish. Human Genetics, 2019, 138, 1171-1182.	1.8	7
28	Association of a Primary Open-Angle Glaucoma Genetic Risk Score With Earlier Age at Diagnosis. JAMA Ophthalmology, 2019, 137, 1190.	1.4	32
29	Genome-wide brain DNA methylation analysis suggests epigenetic reprogramming in Parkinson disease. Neurology: Genetics, 2019, 5, e342.	0.9	50
30	Functional variants in the LRRK2 gene confer shared effects on risk for Crohnâ€™s disease and Parkinsonâ€™s disease. Science Translational Medicine, 2018, 10, .	5.8	273
31	Human genetic variation in GLS2 is associated with development of complicated Staphylococcus aureus bacteremia. PLoS Genetics, 2018, 14, e1007667.	1.5	16
32	The Carnitine Shuttle Pathway is Altered in Patients With Neovascular Age-Related Macular Degeneration. , 2018, 59, 4978.		37
33	Testosterone Pathway Genetic Polymorphisms in Relation to Primary Open-Angle Glaucoma: An Analysis in Two Large Datasets. , 2018, 59, 629.		14
34	Evaluating genetic susceptibility to Staphylococcus aureus bacteremia in African Americans using admixture mapping. Genes and Immunity, 2017, 18, 95-99.	2.2	27
35	Genetic correlations between intraocular pressure, blood pressure and primary open-angle glaucoma: a multi-cohort analysis. European Journal of Human Genetics, 2017, 25, 1261-1267.	1.4	18
36	A population-specific reference panel empowers genetic studies of Anabaptist populations. Scientific Reports, 2017, 7, 6079.	1.6	16

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37	Age at natural menopause genetic risk score in relation to age at natural menopause and primary open-angle glaucoma in a US-based sample. Menopause, 2017, 24, 150-156.	0.8	6
38	Genomics of Human Pulmonary Tuberculosis: from Genes to Pathways. Current Genetic Medicine Reports, 2017, 5, 149-166.	1.9	30
39	Generation of disease-specific autopsy-confirmed iPSCs lines from postmortem isolated Peripheral Blood Mononuclear Cells. Neuroscience Letters, 2017, 637, 201-206.	1.0	6
40	Joint Analysis of Nuclear and Mitochondrial Variants in Age-Related Macular Degeneration Identifies Novel Loci TRPM1 and ABHD2/RLBP1. , 2017, 58, 4027.		21
41	Candidate genes on murine chromosome 8 are associated with susceptibility to Staphylococcus aureus infection in mice and are involved with Staphylococcus aureus septicemia in humans. PLoS ONE, 2017, 12, e0179033.	1.1	5
42	A chromosome 5q31.1 locus associates with tuberculin skin test reactivity in HIV-positive individuals from tuberculosis hyper-endemic regions in east Africa. PLoS Genetics, 2017, 13, e1006710.	1.5	28
43	Progression Rate From Intermediate to Advanced Age-Related Macular Degeneration Is Correlated With the Number of Risk Alleles at the CFH Locus. , 2016, 57, 6107.		18
44	Genetic Association Analysis of Drusen Progression. , 2016, 57, 2225.		12
45	The Application of Genetic Risk Scores in Age-Related Macular Degeneration: A Review. Journal of Clinical Medicine, 2016, 5, 31.	1.0	31
46	A Common Variant in MIR182 Is Associated With Primary Open-Angle Glaucoma in the NEIGHBORHOOD Consortium. , 2016, 57, 4528.		42
47	Assessing the Association of Mitochondrial Genetic Variation With Primary Open-Angle Glaucoma Using Gene-Set Analyses. , 2016, 57, 5046.		44
48	Linkage of familial essential tremor to chromosome 5q35. Movement Disorders, 2016, 31, 1059-1062.	2.2	15
49	DNA variants in CACNA1C modify Parkinson disease risk only when vitamin D level is deficient. Neurology: Genetics, 2016, 2, e72.	0.9	11
50	Heritability of Choroidal Thickness in the Amish. Ophthalmology, 2016, 123, 2537-2544.	2.5	24
51	Overlap between Parkinson disease and Alzheimer disease in ABCA7 functional variants. Neurology: Genetics, 2016, 2, e44.	0.9	31
52	A Locus at 5q33.3 Confers Resistance to Tuberculosis in Highly Susceptible Individuals. American Journal of Human Genetics, 2016, 98, 514-524.	2.6	78
53	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
54	Genome-wide association analysis identifies TXNRD2, ATXN2 and FOXC1 as susceptibility loci for primary open-angle glaucoma. Nature Genetics, 2016, 48, 189-194.	9.4	211

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55	The Relationship Between Reticular Pseudodrusen and Severity of AMD. Ophthalmology, 2016, 123, 921-923.	2.5	15
56	Polymorphisms in HLA Class II Genes Are Associated With Susceptibility toStaphylococcus aureusInfection in a White Population. Journal of Infectious Diseases, 2016, 213, 816-823.	1.9	44
57	Whole exome sequencing of extreme age-related macular degeneration phenotypes. Molecular Vision, 2016, 22, 1062-76.	1.1	12
58	Estimating cumulative pathway effects on risk for age-related macular degeneration using mixed linear models. BMC Bioinformatics, 2015, 16, 329.	1.2	9
59	PARK10 is a major locus for sporadic neuropathologically confirmed Parkinson disease. Neurology, 2015, 84, 972-980.	1.5	48
60	The relationship between obsessiveâ€compulsive symptoms and PARKIN genotype: The COREâ€PD study. Movement Disorders, 2015, 30, 278-283.	2.2	16
61	Vitamin D from different sources is inversely associated with Parkinson disease. Movement Disorders, 2015, 30, 560-566.	2.2	61
62	Examination of Candidate Exonic Variants for Association to Alzheimer Disease in the Amish. PLoS ONE, 2015, 10, e0118043.	1.1	13
63	Epiregulin (EREG) and human V-ATPase (TCIRG1): genetic variation, ethnicity and pulmonary tuberculosis susceptibility in Guinea-Bissau and The Gambia. Genes and Immunity, 2014, 15, 370-377.	2.2	11
64	Dusp3 and Psme3 Are Associated with Murine Susceptibility to Staphylococcus aureus Infection and Human Sepsis. PLoS Pathogens, 2014, 10, e1004149.	2.1	28
65	Glutamate Receptor Gene GRIN2A, Coffee, and Parkinson Disease. PLoS Genetics, 2014, 10, e1004774.	1.5	7
66	Cognitive and Motor Function in Long-DurationPARKIN-Associated Parkinson Disease. JAMA Neurology, 2014, 71, 62.	4.5	49
67	DNA Copy Number Variants of Known Glaucoma Genes in Relation to Primary Open-Angle Glaucoma. Investigative Ophthalmology and Visual Science, 2014, 55, 8251-8258.	3.3	27
68	Set-Based Joint Test of Interaction Between SNPs in the VEGF Pathway and Exogenous Estrogen Finds Association With Age-Related Macular Degeneration. , 2014, 55, 4873.		5
69	Rare VariantAPOC3R19X Is Associated With Cardio-Protective Profiles in a Diverse Population-Based Survey as Part of the Epidemiologic Architecture for Genes Linked to Environment Study. Circulation: Cardiovascular Genetics, 2014, 7, 848-853.	5.1	31
70	Autophagy is redundant for the host defense against systemic Candida albicans infections. European Journal of Clinical Microbiology and Infectious Diseases, 2014, 33, 711-722.	1.3	35
71	A genome-wide association study of variants associated with acquisition of Staphylococcus aureus bacteremia in a healthcare setting. BMC Infectious Diseases, 2014, 14, 83.	1.3	36
72	Role of autophagy genetic variants for the risk of Candida infections. Medical Mycology, 2014, 52, 333-341.	0.3	17

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73	Large-scale meta-analysis of genome-wide association data identifies six new risk loci for Parkinson's disease. Nature Genetics, 2014, 46, 989-993.	9.4	1,685
74	Hypothesis-independent pathway analysis implicates GABA and Acetyl-CoA metabolism in primary open-angle glaucoma and normal-pressure glaucoma. Human Genetics, 2014, 133, 1319-1330.	1.8	32
75	Rare Complement Factor H Variant Associated With Age-Related Macular Degeneration in the Amish. , 2014, 55, 4455.		47
76	Absence of C9ORF72 expanded or intermediate repeats in autopsyâ€confirmed Parkinson's disease. Movement Disorders, 2014, 29, 827-830.	2.2	24
77	Parkinson disease loci in the mid-western Amish. Human Genetics, 2013, 132, 1213-1221.	1.8	14
78	Linkage and association of successful aging to the 6q25 region in large Amish kindreds. Age, 2013, 35, 1467-1477.	3.0	25
79	C9ORF72 Intermediate Repeat Copies Are a Significant Risk Factor for Parkinson Disease. Annals of Human Genetics, 2013, 77, 351-363.	0.3	69
80	Functional genomics identifies type I interferon pathway as central for host defense against Candida albicans. Nature Communications, 2013, 4, 1342.	5.8	157
81	Variants at chromosome 10q26 locus and the expression of HTRA1 in the retina. Experimental Eye Research, 2013, 112, 102-105.	1.2	26
82	Genetic Factors in Nonsmokers with Ageâ€Related Macular Degeneration Revealed Through Genomeâ€Wide Geneâ€Environment Interaction Analysis. Annals of Human Genetics, 2013, 77, 215-231.	0.3	43
83	Seven new loci associated with age-related macular degeneration. Nature Genetics, 2013, 45, 433-439.	9.4	687
84	High-Resolution Survey in Familial Parkinson Disease Genes Reveals Multiple Independent Copy Number Variation Events in PARK2. Human Mutation, 2013, 34, 1071-1074.	1.1	13
85	Coding Variants in ARMS2 and the Risk of Age-Related Macular Degeneration. JAMA Ophthalmology, 2013, 131, 804.	1.4	8
86	Whole exome sequencing of rare variants in EIF4G1 and VPS35 in Parkinson disease. Neurology, 2013, 80, 982-989.	1.5	68
87	Evaluating Power and Type 1 Error in Large Pedigree Analyses of Binary Traits. PLoS ONE, 2013, 8, e62615.	1.1	7
88	Estrogen pathway polymorphisms in relation to primary open angle glaucoma: an analysis accounting for gender from the United States. Molecular Vision, 2013, 19, 1471-81.	1.1	40
89	Comprehensive Research Synopsis and Systematic Meta-Analyses in Parkinson's Disease Genetics: The PDGene Database. PLoS Genetics, 2012, 8, e1002548.	1.5	495
90	Haplotype Association Mapping Identifies a Candidate Gene Region in Mice Infected With Staphylococcus aureus. G3: Genes, Genomes, Genetics, 2012, 2, 693-700.	0.8	14

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91	Toll-like Receptor 1 Polymorphisms Increase Susceptibility to Candidemia. Journal of Infectious Diseases, 2012, 205, 934-943.	1.9	116
92	Cytokine Gene Polymorphisms and the Outcome of Invasive Candidiasis: A Prospective Cohort Study. Clinical Infectious Diseases, 2012, 54, 502-510.	2.9	68
93	THE ARMS2 A69S VARIANT AND BILATERAL ADVANCED AGE-RELATED MACULAR DEGENERATION. Retina, 2012, 32, 1486-1491.	1.0	22
94	Genomeâ€Wide Association and Linkage Study in the Amish Detects a Novel Candidate Lateâ€Onset Alzheimer Disease Gene. Annals of Human Genetics, 2012, 76, 342-351.	0.3	40
95	A novel ARMS2 splice variant is identified in human retina. Experimental Eye Research, 2012, 94, 187-191.	1.2	11
96	Populationâ€Based Caseâ€Control Association Studies. Current Protocols in Human Genetics, 2012, 74, Unit1.17.	3.5	9
97	MCP1 SNPs and Pulmonary Tuberculosis in Cohorts from West Africa, the USA and Argentina: Lack of Association or Epistasis with IL12B Polymorphisms. PLoS ONE, 2012, 7, e32275.	1.1	16
98	Retinoblastoma treatment: impact of the glycolytic inhibitor 2-deoxy-d-glucose on molecular genomics expression in LHBETATAG retinal tumors. Clinical Ophthalmology, 2012, 6, 817.	0.9	4
99	Metaâ€analysis of Parkinson's Disease: Identification of a novel locus, RIT2. Annals of Neurology, 2012, 71, 370-384.	2.8	264
100	Human genetic susceptibility toCandidainfections. Medical Mycology, 2012, 50, 785-794.	0.3	37
101	The impact of caspase-12 on susceptibility to candidemia. European Journal of Clinical Microbiology and Infectious Diseases, 2012, 31, 277-280.	1.3	14
102	Mitochondrial Haplogroup X is associated with successful aging in the Amish. Human Genetics, 2012, 131, 201-208.	1.8	23
103	Identifying Consensus Disease Pathways in Parkinson's Disease Using an Integrative Systems Biology Approach. PLoS ONE, 2011, 6, e16917.	1.1	72
104	Genome-Wide Gene-Environment Study Identifies Glutamate Receptor Gene GRIN2A as a Parkinson's Disease Modifier Gene via Interaction with Coffee. PLoS Genetics, 2011, 7, e1002237.	1.5	206
105	The relation between depression and parkin genotype: The CORE-PD study. Parkinsonism and Related Disorders, 2011, 17, 740-744.	1.1	38
106	Interleukin 12B (IL12B) Genetic Variation and Pulmonary Tuberculosis: A Study of Cohorts from The Gambia, Guinea-Bissau, United States and Argentina. PLoS ONE, 2011, 6, e16656.	1.1	33
107	Vitamin D Receptor Gene as a Candidate Gene for Parkinson Disease. Annals of Human Genetics, 2011, 75, 201-210.	0.3	95
108	A Genomeâ€Wide Linkage Screen in the Amish with Parkinson Disease Points to Chromosome 6. Annals of Human Genetics, 2011, 75, 351-358.	0.3	9

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109	Successful Aging Shows Linkage to Chromosomes 6, 7, and 14 in the Amish. Annals of Human Genetics, 2011, 75, 516-528.	0.3	27
110	Genetic Variation in the Dectin-1/CARD9 Recognition Pathway and Susceptibility to Candidemia. Journal of Infectious Diseases, 2011, 204, 1138-1145.	1.9	80
111	Regional and Temporal Differences in Gene Expression of LH_{BETA}T_{AG}Retinoblastoma Tumors. , 2011, 52, 5359.		14
112	Neuropsychological Profile of Parkin Mutation Carriers with and without Parkinson Disease: The CORE-PD Study. Journal of the International Neuropsychological Society, 2011, 17, 91-100.	1.2	21
113	CD4 Intragenic SNPs Associate With HIV-2 Plasma Viral Load and CD4 Count in a Community-Based Study From Guinea-Bissau, West Africa. Journal of Acquired Immune Deficiency Syndromes (1999), 2011, 56, 1-8.	0.9	29
114	Dissection of Chromosome 16p12 Linkage Peak Suggests a Possible Role forCACNG3Variants in Age-Related Macular Degeneration Susceptibility. , 2011, 52, 1748.		10
115	Using Genetic Variation and Environmental Risk Factor Data to Identify Individuals at High Risk for Age-Related Macular Degeneration. PLoS ONE, 2011, 6, e17784.	1.1	40
116	Genotype at Polymorphism rs11200638 and HTRA1 Expression Level. JAMA Ophthalmology, 2010, 128, 1491.	2.6	14
117	Variants in toll-like receptors 2 and 9 influence susceptibility to pulmonary tuberculosis in Caucasians, African-Americans, and West Africans. Human Genetics, 2010, 127, 65-73.	1.8	143
118	Analysis of the indel at the ARMS2 3â€²UTR in age-related macular degeneration. Human Genetics, 2010, 127, 595-602.	1.8	59
119	A rare novel deletion of the tyrosine hydroxylase gene in Parkinson disease. Human Mutation, 2010, 31, E1767-E1771.	1.1	29
120	Genomeâ€Wide Association Study Confirms SNPs in SNCA and the MAPT Region as Common Risk Factors for Parkinson Disease. Annals of Human Genetics, 2010, 74, 97-109.	0.3	417
121	Analysis of Single Nucleotide Polymorphisms in the NOS2A Gene and Interaction with Smoking in Ageâ€Related Macular Degeneration. Annals of Human Genetics, 2010, 74, 195-201.	0.3	16
122	Common genetic variation in the HLA region is associated with late-onset sporadic Parkinson's disease. Nature Genetics, 2010, 42, 781-785.	9.4	692
123	Dementia Revealed: Novel Chromosome 6 Locus for Late-Onset Alzheimer Disease Provides Genetic Evidence for Folate-Pathway Abnormalities. PLoS Genetics, 2010, 6, e1001130.	1.5	130
124	Frequency of Known Mutations in Early-Onset Parkinson Disease. Archives of Neurology, 2010, 67, 1116-22.	4.9	121
125	A New Locus for Familial FSGS on Chromosome 2P. Journal of the American Society of Nephrology: JASN, 2010, 21, 1390-1397.	3.0	7
126	Inverse Association of Female Hormone Replacement Therapy with Age-Related Macular Degeneration and Interactions withARMS2Polymorphisms. , 2010, 51, 1873.		33

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127	Two Genes on A/J Chromosome 18 Are Associated with Susceptibility to Staphylococcus aureus Infection by Combined Microarray and QTL Analyses. PLoS Pathogens, 2010, 6, e1001088.	2.1	61
128	Overall Diet Quality and Age-Related Macular Degeneration. Ophthalmic Epidemiology, 2010, 17, 58-65.	0.8	36
129	Predictors of Parkin Mutations in Early-Onset Parkinson Disease. Archives of Neurology, 2010, 67, 731-8.	4.9	81
130	Self-report of cognitive impairment and mini-mental state examination performance in PRKN, LRRK2, and GBA carriers with early onset Parkinson's disease. Journal of Clinical and Experimental Neuropsychology, 2010, 32, 775-779.	0.8	50
131	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
132	Motor Phenotype of LRRK2 G2019S Carriers in Early-Onset Parkinson Disease. Archives of Neurology, 2009, 66, 1517-22.	4.9	63
133	Localization of Age-Related Macular Degeneration-Associated ARMS2 in Cytosol, Not Mitochondria. , 2009, 50, 3084.		85
134	NOS2A, TLR4, and IFNGR1 interactions influence pulmonary tuberculosis susceptibility in African-Americans. Human Genetics, 2009, 126, 643-653.	1.8	73
135	Genome-wide Linkage Screen in Familial Parkinson Disease Identifies Loci on Chromosomes 3 and 18. American Journal of Human Genetics, 2009, 84, 499-504.	2.6	11
136	Molecular and Contextual Markers of Hepatitis C Virus and Drug Abuse. Molecular Diagnosis and Therapy, 2009, 13, 153-179.	1.6	5
137	Molecular and contextual markers of hepatitis C virus and drug abuse. Molecular Diagnosis and Therapy, 2009, 13, 153-79.	1.6	3
138	Nitric oxide synthase genes and their interactions with environmental factors in Parkinsonâ€™s disease. Neurogenetics, 2008, 9, 249-262.	0.7	91
139	Variation in the miRNA-433 Binding Site of FGF20 Confers Risk for Parkinson Disease by Overexpression of Î±-Synuclein. American Journal of Human Genetics, 2008, 82, 283-289.	2.6	437
140	Geneâ€€Gene Interaction Between FGF20 and MAOB in Parkinson Disease. Annals of Human Genetics, 2008, 72, 157-162.	0.3	34
141	Pesticide exposure and risk of Parkinson's disease: A family-based case-control study. BMC Neurology, 2008, 8, 6.	0.8	221
142	Phenotype Analysis of Patients With the Risk Variant LOC387715 (A69S) in Age-related Macular Degeneration. American Journal of Ophthalmology, 2008, 145, 303-307.e1.	1.7	32
143	Peripheral Reticular Pigmentary Change Is Associated with Complement Factor H Polymorphism (Y402H) in Age-Related Macular Degeneration. Ophthalmology, 2008, 115, 520-524.	2.5	34
144	Deletion of CFHR3 and CFHR1 genes in age-related macular degeneration. Human Molecular Genetics, 2008, 17, 971-977.	1.4	85

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145	C3 R102G polymorphism increases risk of age-related macular degeneration. Human Molecular Genetics, 2008, 17, 1821-1824.	1.4	120
146	Haplotypes Spanning the Complement Factor H Gene Are Protective against Age-Related Macular Degeneration. , 2007, 48, 4277.		22
147	Smoking, Caffeine, and Nonsteroidal Anti-inflammatory Drugs in Families With Parkinson Disease. Archives of Neurology, 2007, 64, 576.	4.9	107
148	Protective effect of complement factor B and complement component 2 variants in age-related macular degeneration. Human Molecular Genetics, 2007, 16, 1986-1992.	1.4	175
149	Neovascular Age-Related Macular Degeneration and Its Association With LOC387715 and Complement Factor H Polymorphism. JAMA Ophthalmology, 2007, 125, 63.	2.6	58
150	Populationâ€Based Caseâ€Control Association Studies. Current Protocols in Human Genetics, 2007, 52, Unit 1.17.	3.5	6
151	Independent Effects of Complement Factor H Y402H Polymorphism and Cigarette Smoking on Risk of Age-Related Macular Degeneration. Ophthalmology, 2007, 114, 1151-1156.	2.5	80
152	Methods for interaction analyses using family-based case-control data: conditional logistic regression versus generalized estimating equations. Genetic Epidemiology, 2007, 31, 883-893.	0.6	36
153	Construction and validation of a Parkinson's disease mutation genotyping array for the Parkin gene. Movement Disorders, 2007, 22, 932-937.	2.2	16
154	Cigarette Smoking Strongly Modifies the Association of LOC387715 and Age-Related Macular Degeneration. American Journal of Human Genetics, 2006, 78, 852-864.	2.6	316
155	Revealing the role of glutathione S-transferase omega in age-at-onset of Alzheimer and Parkinson diseases. Neurobiology of Aging, 2006, 27, 1087-1093.	1.5	60
156	Complement Factor H Increases Risk for Atrophic Age-Related Macular Degeneration. Ophthalmology, 2006, 113, 1504-1507.	2.5	58
157	Functional Candidate Genes in Age-Related Macular Degeneration: Significant Association withVEGF,VLDLR, andLRP6. , 2006, 47, 329.		178
158	Combinatorial Mismatch Scan (CMS) for loci associated with dementia in the Amish. BMC Medical Genetics, 2006, 7, 19.	2.1	11
159	NOS2Aand the modulating effect of cigarette smoking in Parkinson's disease. Annals of Neurology, 2006, 60, 366-373.	2.8	38
160	Family-based caseâ€“control study of MAOA and MAOB polymorphisms in Parkinson disease. Movement Disorders, 2006, 21, 2175-2180.	2.2	33
161	A genome-wide linkage analysis of dementia in the Amish. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2006, 141B, 160-166.	1.1	42
162	Maternal lineages and Alzheimer disease risk in the Old Order Amish. Human Genetics, 2005, 118, 115-122.	1.8	29

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163	Searching for epistatic interactions in nuclear families using conditional linkage analysis. BMC Genetics, 2005, 6, S148.	2.7	5
164	Potential for expanded power in linkage studies using the ALLEGRO and MERLIN software programs. Journal of Medical Genetics, 2005, 42, e68-e68.	1.5	5
165	Linkage Disequilibrium Inflates Type I Error Rates in Multipoint Linkage Analysis when Parental Genotypes Are Missing. Human Heredity, 2005, 59, 220-227.	0.4	74
166	An autosomal genomic screen for dementia in an extended Amish family. Neuroscience Letters, 2005, 379, 199-204.	1.0	31
167	Comparing Age-related Macular Degeneration Phenotype in Probands From Singleton and Multiplex Families. American Journal of Ophthalmology, 2005, 139, 820-825.	1.7	27
168	Complement Factor H Variant Increases the Risk of Age-Related Macular Degeneration. Science, 2005, 308, 419-421.	6.0	2,232
169	Joint effects of smoking history and APOE genotypes in age-related macular degeneration. Molecular Vision, 2005, 11, 941-9.	1.1	40
170	Glutathione S-transferase omega-1 modifies age-at-onset of Alzheimer disease and Parkinson disease. Human Molecular Genetics, 2004, 13, 573-573.	1.4	5
171	Linkage disequilibrium and haplotype tagging polymorphisms in the Tau H1 haplotype. Neurogenetics, 2004, 5, 147-155.	0.7	30
172	Ordered subset linkage analysis supports a susceptibility locus for age-related macular degeneration on chromosome 16p12. BMC Genetics, 2004, 5, 18.	2.7	48
173	Analysis of European mitochondrial haplogroups with Alzheimer disease risk. Neuroscience Letters, 2004, 365, 28-32.	1.0	264
174	Fibroblast Growth Factor 20 Polymorphisms and Haplotypes Strongly Influence Risk of Parkinson Disease. American Journal of Human Genetics, 2004, 74, 1121-1127.	2.6	136
175	S2-01-01 Dementia in mid-Western U.S. amish families. Neurobiology of Aging, 2004, 25, S24-S25.	1.5	3
176	Parkin mutations and susceptibility alleles in late-onset Parkinson's disease. Annals of Neurology, 2003, 53, 624-629.	2.8	201
177	The Q7R Saitohin gene polymorphism is not associated with Alzheimer disease. Neuroscience Letters, 2003, 347, 143-146.	1.0	12
178	Design of the Genetics of Early Onset Cardiovascular Disease (GENECARD) study. American Heart Journal, 2003, 145, 602-613.	1.2	55
179	Mitochondrial Polymorphisms Significantly Reduce the Risk of Parkinson Disease. American Journal of Human Genetics, 2003, 72, 804-811.	2.6	507
180	Ordered-Subsets Linkage Analysis Detects Novel Alzheimer Disease Loci on Chromosomes 2q34 and 15q22. American Journal of Human Genetics, 2003, 73, 1041-1051.	2.6	99

#	Article	IF	Citations
181	Glutathione S-transferase omega-1 modifiesage-at-onset of Alzheimer disease and Parkinson disease. Human Molecular Genetics, 2003, 12, 3259-3267.	1.4	208
182	Association Study of Parkin Gene Polymorphisms With Idiopathic Parkinson Disease. Archives of Neurology, 2003, 60, 975.	4.9	51
183	Detailed Analysis of Allelic Variation in theABCA4Gene in Age-Related Maculopathy. , 2003, 44, 2868.		33
184	Linkage of a Gene Causing Familial Membranoproliferative Glomerulonephritis Type III to Chromosome 1. Journal of the American Society of Nephrology: JASN, 2002, 13, 2052-2057.	3.0	32
185	Age at Onset in Two Common Neurodegenerative Diseases Is Genetically Controlled. American Journal of Human Genetics, 2002, 70, 985-993.	2.6	291
186	Linkage of Parkinsonism and Alzheimer's disease with Lewy body pathology to chromosome 12. Annals of Neurology, 2002, 52, 524-524.	2.8	1
187	Life After the Screen: Making Sense of Many Pâ€Values. Genetic Epidemiology, 2001, 21, S546-51.	0.6	5
188	Fine mapping and genetic heterogeneity in the pure form of autosomal dominant familial spastic paraplegia. Neurogenetics, 2001, 3, 91-97.	0.7	13
189	Association of Single-Nucleotide Polymorphisms of the Tau Gene With Late-Onset Parkinson Disease. JAMA - Journal of the American Medical Association, 2001, 286, 2245.	3.8	171
190	Dissecting A Complex Disease Using Modern Techniques of Molecular Biology. Laboratory Medicine, 2001, 32, 594-598.	0.8	0
191	Complete Genomic Screen in Parkinson Disease. JAMA - Journal of the American Medical Association, 2001, 286, 2239.	3.8	257
192	Identification of Novel Genes in Late-Onset Alzheimer's Disease. Experimental Gerontology, 2000, 35, 1343-1352.	1.2	183
193	Linkage of the CCR5Î”32 Mutation with a Functional Polymorphism of CD45RA. Journal of Immunology, 2000, 165, 148-157.	0.4	10
194	Fine Mapping of the Chromosome 12 Late-Onset Alzheimer Disease Locus: Potential Genetic and Phenotypic Heterogeneity. American Journal of Human Genetics, 2000, 66, 922-932.	2.6	113
195	An Î±-2-macroglobulin insertion-deletion polymorphism in Alzheimer disease. Nature Genetics, 1999, 22, 19-21.	9.4	115
196	No association between the HLA-A2 allele and Alzheimer disease. Neurogenetics, 1999, 2, 177-182.	0.7	19
197	The Î±-synuclein gene is not a major risk factor in familial Parkinson disease. Neurogenetics, 1999, 2, 191-192.	0.7	31
198	Analysis of association between Alzheimer disease and the K variant of butyrylcholinesterase (BCHE-K). Neuroscience Letters, 1999, 269, 115-119.	1.0	37

#	Article	IF	Citations
199	Texture segregation shows only a very small lower-hemifield advantage. Vision Research, 1999, 39, 1171-1175.	0.7	6
200	Further Evidence Linking Late-Onset Alzheimer Disease With Chromosome 12. JAMA - Journal of the American Medical Association, 1999, 281, 513-514.	3.8	37
201	Association between bleomycin hydrolase and Alzheimer's disease in caucasians. Annals of Neurology, 1998, 44, 808-811.	2.8	48
202	No genetic association between the LRP receptor and sporadic or late-onset familial Alzheimer disease. Neurogenetics, 1998, 1, 179-183.	0.7	39
203	Lack of association between apolipoprotein E genotype and sporadic amyotrophic lateral sclerosis. Neurogenetics, 1998, 1, 213-216.	0.7	46
204	Complete Genomic Screen in Late-Onset Familial Alzheimerâ€™s Disease. Neurobiology of Aging, 1998, 19, S39-S42.	1.5	54
205	No association of Î±1-antichymotrypsin flanking region polymorphism and Alzheimer disease risk in early- and late-onset Alzheimer disease patients. Neuroscience Letters, 1998, 250, 79-82.	1.0	2
206	Evidence for anticipation in autosomal dominant limb-girdle muscular dystrophy.. Journal of Medical Genetics, 1998, 35, 305-308.	1.5	13
207	Absence of Mutation in the Â- and Â-Synuclein Genes in Familial Autosomal Dominant Parkinson's Disease. DNA Research, 1998, 5, 401-402.	1.5	24
208	Heredity and Alzheimer's disease. Current Opinion in Psychiatry, 1998, 11, 417-423.	3.1	2
209	Risk of Institutionalization Among Community Long-Term Care Clients With Dementia. Gerontologist, The, 1997, 37, 46-51.	2.3	77
210	Preparedness of Educational Speech-Language Pathologists to Provide Services to Students with Traumatic Brain Injury. Communication Disorders Quarterly, 1997, 18, 49-63.	0.7	12
211	Genetic Complexity and Parkinson's Disease. Science, 1997, 277, 387-390.	6.0	70
212	LETTERS TO THE EDITOR. Genomics, 1997, 40, 382-384.	1.3	10
213	Major depression and all-cause mortality among white adults in the United States. Annals of Epidemiology, 1997, 7, 213-218.	0.9	157
214	Functional health status as a predictor of mortality in men and women over 65. Journal of Clinical Epidemiology, 1997, 50, 291-296.	2.4	175
215	Locus heterogeneity, anticipation and reduction of the chromosome 2p minimal candidate region in autosomal dominant familial spastic paraplegia. Neurogenetics, 1997, 1, 95-102.	0.7	21
216	No association or linkage between an intronic polymorphism of presenilin-1 and sporadic or late-onset familial Alzheimer disease. Genetic Epidemiology, 1997, 14, 307-315.	0.6	31

#	Article	IF	Citations
217	False positive rates in a genomic screen for complex quantitative traits. Genetic Epidemiology, 1997, 14, 891-896.	0.6	1
218	Naming in dementia secondary to Parkinson's, Huntington's, and Alzheimer's diseases. Journal of Communication Disorders, 1996, 29, 183-197.	0.8	59
219	A Genomic Search for Alzheimer's Disease Genes. , 0, , 31-43.		2
220	Complex Genetic Interactions. , 0, , 397-421.		0