Cornelia M Van Duijn

List of Publications by Year
in descending order

854	144,871	57 177	127 336
papers	citations	h-index	g-index

942	942	942	102670
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	The probabilistic model of Alzheimer disease: the amyloid hypothesis revised. Nature Reviews Neuroscience, 2022, 23, 53-66.	4.9	203
2	Gene-mapping study of extremes of cerebral small vessel disease reveals TRIM47 as a strong candidate. Brain, 2022, 145, 1992-2007.	3.7	6
3	Fat metabolism is associated with telomere length in six population-based studies. Human Molecular Genetics, 2022, 31, 1159-1170.	1.4	7
4	A multi-omics study of circulating phospholipid markers of blood pressure. Scientific Reports, 2022, 12, 574.	1.6	10
5	Interpretation and actionability of genetic variants in cardiomyopathies: a position statement from the European Society of Cardiology Council on cardiovascular genomics. European Heart Journal, 2022, 43, 1901-1916.	1.0	32
6	Matrix metalloproteinase 10 is linked to the risk of progression to dementia of the Alzheimerâ€™s type. Brain, 2022, 145, 2507-2517.	3.7	16
7	Elucidating the relationship between migraine risk and brain structure using genetic data. Brain, 2022, 145, 3214-3224.	3.7	7
8	Genome-wide study of DNA methylation shows alterations in metabolic, inflammatory, and cholesterol pathways in ALS. Science Translational Medicine, 2022, 14, eabj0264.	5.8	38
9	Multiâ€phenotype analyses of hemostatic traits with cardiovascular events reveal novel genetic associations. Journal of Thrombosis and Haemostasis, 2022, 20, 1331-1349.	1.9	12
10	Polygenic prediction of educational attainment within and between families from genome-wide association analyses in 3 million individuals. Nature Genetics, 2022, 54, 437-449.	9.4	215
11	Understanding the complex genetic architecture connecting rheumatoid arthritis, osteoporosis and inflammation: discovering causal pathways. Human Molecular Genetics, 2022, , .	1.4	3
12	DNA methylation in peripheral tissues and left-handedness. Scientific Reports, 2022, 12, 5606.	1.6	12
13	New insights into the genetic etiology of Alzheimerâ€™s disease and related dementias. Nature Genetics, 2022, 54, 412-436.	9.4	700
14	Association of Diabetes Medication With Open-Angle Glaucoma, Age-Related Macular Degeneration, and Cataract in the Rotterdam Study. JAMA Ophthalmology, 2022, 140, 674.	1.4	15
15	Association of Rare APOE Missense Variants V236E and R251G With Risk of Alzheimer Disease. JAMA Neurology, 2022, 79, 652.	4.5	31
16	Microbiomics, Metabolomics, Predicted Metagenomics, and Hepatic Steatosis in a Populationâ€Based Study of 1,355 Adults. Hepatology, 2021, 73, 968-982.	3.6	43
17	Genome-wide association study identifies 48 common genetic variants associated with handedness. Nature Human Behaviour, 2021, 5, 59-70.	6.2	79
18	Association of Circulating Metabolites in Plasma or Serum and Risk of Stroke. Neurology, 2021, 96, .	1.5	24

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19	Multi-ancestry genome-wide association study accounting for gene-psychosocial factor interactions identifies novel loci for blood pressure traits. Human Genetics and Genomics Advances, 2021, 2, 100013.	1.0	2
20	Circulating metabolites are associated with brain atrophy and white matter hyperintensities. Alzheimer's and Dementia, 2021, 17, 205-214.	0.4	17
21	Sex-dimorphic genetic effects and novel loci for fasting glucose and insulin variability. Nature Communications, 2021, 12, 24.	5.8	87
22	DNA methylation signatures of aggression and closely related constructs: A meta-analysis of epigenome-wide studies across the lifespan. Molecular Psychiatry, 2021, 26, 2148-2162.	4.1	21
23	Large-scale association analyses identify host factors influencing human gut microbiome composition. Nature Genetics, 2021, 53, 156-165.	9.4	676
24	Genome-wide meta-analysis identifies 127 open-angle glaucoma loci with consistent effect across ancestries. Nature Communications, 2021, 12, 1258.	5.8	196
25	Genome-wide association study in almost 195,000 individuals identifies 50 previously unidentified genetic loci for eye color. Science Advances, 2021, 7, .	4.7	36
26	First participant diagnosed with Creutzfeldt-Jakob disease in the population-based Rotterdam Study was classified with mild cognitive impairment. BMJ Case Reports, 2021, 14, e235509.	0.2	0
27	Plasma Brain-Derived Neurotropic Factor Levels Are Associated with Aging and Smoking But Not with Future Dementia in the Rotterdam Study. Journal of Alzheimer's Disease, 2021, 80, 1139-1149.	1.2	8
28	Multiomics integrative analysis identifies APOE allele-specific blood biomarkers associated to Alzheimerâ€™s disease etiopathogenesis. Aging, 2021, 13, 9277-9329.	1.4	15
29	Multi-ancestry genome-wide geneâ€“sleep interactions identify novel loci for blood pressure. Molecular Psychiatry, 2021, 26, 6293-6304.	4.1	13
30	Plasma amyloid Î² levels are driven by genetic variants near APOE, BACE1, APP, PSEN2: A genomeâ€wide association study in over 12,000 nonâ€demented participants. Alzheimer's and Dementia, 2021, 17, 1663-1674.	0.4	20
31	Genetic variation affects morphological retinal phenotypes extracted from UK Biobank optical coherence tomography images. PLoS Genetics, 2021, 17, e1009497.	1.5	50
32	The trans-ancestral genomic architecture of glycemic traits. Nature Genetics, 2021, 53, 840-860.	9.4	341
33	Common variants in Alzheimerâ€™s disease and risk stratification by polygenic risk scores. Nature Communications, 2021, 12, 3417.	5.8	140
34	Evaluation of Shared Genetic Susceptibility to High and Low Myopia and Hyperopia. JAMA Ophthalmology, 2021, 139, 601.	1.4	22
35	Identification of 371 genetic variants for age at first sex and birth linked to externalising behaviour. Nature Human Behaviour, 2021, 5, 1717-1730.	6.2	62
36	Characteristics of p.Gln368Ter Myocilin Variant and Influence of Polygenic Risk on Glaucoma Penetrance in the UK Biobank. Ophthalmology, 2021, 128, 1300-1311.	2.5	27

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37	Genome-wide association study of frontotemporal dementia identifies a C9ORF72 haplotype with a median of 12-G4C2 repeats that predisposes to pathological repeat expansions. Translational Psychiatry, 2021, 11, 451.	2.4	6
38	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
39	Metabolic profileÂchanges in serum of migraine patients detected using 1H-NMR spectroscopy. Journal of Headache and Pain, 2021, 22, 142.	2.5	7
40	Association of low-frequency and rare coding variants with information processing speed. Translational Psychiatry, 2021, 11, 613.	2.4	2
41	The power of genetic diversity in genome-wide association studies of lipids. Nature, 2021, 600, 675-679.	13.7	353
42	Gut microbiomeâ€related metabolites in plasma are associated with general cognition. Alzheimer's and Dementia, 2021, 17, .	0.4	0
43	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
44	Novel Rare Genetic Variants Associated with Airflow Obstruction in the General Population. American Journal of Respiratory and Critical Care Medicine, 2020, 201, 485-488.	2.5	2
45	Metabolomics Profile in Depression: A Pooled Analysis of 230 Metabolic Markers in 5283 Cases With Depression and 10,145 Controls. Biological Psychiatry, 2020, 87, 409-418.	0.7	129
46	Heritability estimates for 361 blood metabolites across 40 genome-wide association studies. Nature Communications, 2020, 11, 39.	5.8	64
47	Lipidomic profiling identifies signatures of metabolic risk. EBioMedicine, 2020, 51, 102520.	2.7	56
48	Association of lysophosphatidic acids with cerebrospinal fluid biomarkers and progression to Alzheimerâ€™s disease. Alzheimer's Research and Therapy, 2020, 12, 124.	3.0	12
49	Genetic correlations and genome-wide associations of cortical structure in general population samples of 22,824 adults. Nature Communications, 2020, 11, 4796.	5.8	61
50	A cross-omics integrative study of metabolic signatures of chronic obstructive pulmonary disease. BMC Pulmonary Medicine, 2020, 20, 193.	0.8	15
51	Metabolic Age Based on the BBMRI-NL ¹ H-NMR Metabolomics Repository as Biomarker of Age-related Disease. Circulation Genomic and Precision Medicine, 2020, 13, 541-547.	1.6	50
52	Genome-wide identification of genes regulating DNA methylation using genetic anchors for causal inference. Genome Biology, 2020, 21, 220.	3.8	27
53	Identification of novel risk loci and causal insights for sporadic Creutzfeldt-Jakob disease: a genome-wide association study. Lancet Neurology, The, 2020, 19, 840-848.	4.9	42
54	Genetic Studies of Leptin Concentrations Implicate Leptin in the Regulation of Early Adiposity. Diabetes, 2020, 69, 2806-2818.	0.3	26

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55	Association of common genetic variants with brain microbleeds. Neurology, 2020, 95, e3331-e3343.	1.5	40
56	Genetic Determinants of Electrocardiographic P-Wave Duration and Relation to Atrial Fibrillation. Circulation Genomic and Precision Medicine, 2020, 13, 387-395.	1.6	16
57	Genetic Predisposition to Coronary Artery Disease in Type 2 Diabetes Mellitus. Circulation Genomic and Precision Medicine, 2020, 13, e002769.	1.6	5
58	Crossâ€omics studies of the role of apolipoprotein E in Alzheimerâ€™s disease and dementia: Searching common pathways in patients, populations and cellular models. Alzheimer's and Dementia, 2020, 16, e040282.	0.4	0
59	Exome sequencing identifies three novel ADâ€associated genes. Alzheimer's and Dementia, 2020, 16, e041592.	0.4	6
60	Genomeâ€wide metaâ€analysis of lateâ€onset Alzheimerâ€™s disease using rare variant imputation in 65,602 subjects identifies risk loci with roles in memory, neurodevelopment, and cardiometabolic traits: The international genomics of Alzheimerâ€™s project (IGAP). Alzheimer's and Dementia, 2020, 16, e044193.	0.4	1
61	SORL1 â€variant carriers in ADESâ€ADSP: A higher level of variant pathogenicity associates with earlier age at onset of Alzheimer's disease. Alzheimer's and Dementia, 2020, 16, e044492.	0.4	1
62	Clostridium shows a higher abundance in less neurovascular and neurodegenerative changes: A microbiomeâ€wide association study. Alzheimer's and Dementia, 2020, 16, e044743.	0.4	4
63	Analysis of individual families implicates noncoding DNA variation and multiple biological pathways in Alzheimerâ€™s disease risk. Alzheimer's and Dementia, 2020, 16, e046456.	0.4	0
64	Cerebral small vessel disease genomics and its implications across the lifespan. Nature Communications, 2020, 11, 6285.	5.8	89
65	Habitual sleep disturbances and migraine: a Mendelian randomization study. Annals of Clinical and Translational Neurology, 2020, 7, 2370-2380.	1.7	18
66	Gene-educational attainment interactions in a multi-ancestry genome-wide meta-analysis identify novel blood pressure loci. Molecular Psychiatry, 2020, 26, 2111-2125.	4.1	17
67	Refining Attention-Deficit/Hyperactivity Disorder and Autism Spectrum Disorder Genetic Loci by Integrating Summary Data From Genome-wide Association, Gene Expression, and DNA Methylation Studies. Biological Psychiatry, 2020, 88, 470-479.	0.7	14
68	CDH6 and HAGH protein levels in plasma associate with Alzheimerâ€™s disease in APOE Îµ4 carriers. Scientific Reports, 2020, 10, 8233.	1.6	17
69	Smoking-by-genotype interaction in type 2 diabetes risk and fasting glucose. PLoS ONE, 2020, 15, e0230815.	1.1	10
70	Multi-ancestry GWAS of the electrocardiographic PR interval identifies 202 loci underlying cardiac conduction. Nature Communications, 2020, 11, 2542.	5.8	59
71	Integrating Metabolomics, Genomics, and Disease Pathways in Age-Related Macular Degeneration. Ophthalmology, 2020, 127, 1693-1709.	2.5	43
72	Common Genetic Variation Indicates Separate Causes for Periventricular and Deep White Matter Hyperintensities. Stroke, 2020, 51, 2111-2121.	1.0	71

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73	Genome-wide association meta-analysis of corneal curvature identifies novel loci and shared genetic influences across axial length and refractive error. Communications Biology, 2020, 3, 133.	2.0	22
74	Prion protein codon 129 polymorphism in mild cognitive impairment and dementia: the Rotterdam Study. Brain Communications, 2020, 2, fcaa030.	1.5	3
75	A genome-wide cross-phenotype meta-analysis of the association of blood pressure with migraine. Nature Communications, 2020, 11, 3368.	5.8	49
76	Measurement and genetic architecture of lifetime depression in the Netherlands as assessed by LIDAS (Lifetime Depression Assessment Self-report). Psychological Medicine, 2020, , 1-10.	2.7	4
77	Genome-wide Association Analysis in Humans Links Nucleotide Metabolism to Leukocyte Telomere Length. American Journal of Human Genetics, 2020, 106, 389-404.	2.6	118
78	Integration of epidemiologic, pharmacologic, genetic and gut microbiome data in a drugâ€“metabolite atlas. Nature Medicine, 2020, 26, 110-117.	15.2	54
79	Multitrait analysis of glaucoma identifies new risk loci and enables polygenic prediction of disease susceptibility and progression. Nature Genetics, 2020, 52, 160-166.	9.4	192
80	Exome Sequencing Analysis Identifies Rare Variants in ATM and RPL8 That Are Associated With Shorter Telomere Length. Frontiers in Genetics, 2020, 11, 337.	1.1	4
81	Cross-trait analyses with migraine reveal widespread pleiotropy and suggest a vascular component to migraine headache. International Journal of Epidemiology, 2020, 49, 1022-1031.	0.9	34
82	Smoking-by-genotype interaction in type 2 diabetes risk and fasting glucose. , 2020, 15, e0230815.		0
83	Smoking-by-genotype interaction in type 2 diabetes risk and fasting glucose. , 2020, 15, e0230815.		0
84	Smoking-by-genotype interaction in type 2 diabetes risk and fasting glucose. , 2020, 15, e0230815.		0
85	Smoking-by-genotype interaction in type 2 diabetes risk and fasting glucose. , 2020, 15, e0230815.		0
86	Linkage analysis and whole exome sequencing identify a novel candidate gene in a Dutch multiple sclerosis family. Multiple Sclerosis Journal, 2019, 25, 909-917.	1.4	19
87	Quality control and integration of genotypes from two calling pipelines for whole genome sequence data in the Alzheimer's disease sequencing project. Genomics, 2019, 111, 808-818.	1.3	26
88	EIF2AK3 variants in Dutch patients with Alzheimer's disease. Neurobiology of Aging, 2019, 73, 229.e11-229.e18.	1.5	25
89	A metabolic profile of all-cause mortality risk identified in an observational study of 44,168 individuals. Nature Communications, 2019, 10, 3346.	5.8	188
90	A genome-wide association study identifies genetic loci associated with specific lobar brain volumes. Communications Biology, 2019, 2, 285.	2.0	27

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91	A meta-analysis of genome-wide association studies identifies multiple longevity genes. Nature Communications, 2019, 10, 3669.	5.8	214
92	Metabolomics reveals a link between homocysteine and lipid metabolism and leukocyte telomere length: the ENGAGE consortium. Scientific Reports, 2019, 9, 11623.	1.6	13
93	Determining Possible Shared Genetic Architecture Between Myopia and Primary Open-Angle Glaucoma. , 2019, 60, 3142.		10
94	Association of Altered Liver Enzymes With Alzheimer Disease Diagnosis, Cognition, Neuroimaging Measures, and Cerebrospinal Fluid Biomarkers. JAMA Network Open, 2019, 2, e197978.	2.8	142
95	Genome-wide Association Study of Change in Fasting Glucose over time in 13,807 non-diabetic European Ancestry Individuals. Scientific Reports, 2019, 9, 9439.	1.6	5
96	Disentangling the genetics of lean mass. American Journal of Clinical Nutrition, 2019, 109, 276-287.	2.2	38
97	Associations of autozygosity with a broad range of human phenotypes. Nature Communications, 2019, 10, 4957.	5.8	84
98	Association of Genetic Variants With Primary Open-Angle Glaucoma Among Individuals With African Ancestry. JAMA - Journal of the American Medical Association, 2019, 322, 1682.	3.8	50
99	Multi-ancestry sleep-by-SNP interaction analysis in 126,926 individuals reveals lipid loci stratified by sleep duration. Nature Communications, 2019, 10, 5121.	5.8	62
100	GWAS on longitudinal growth traits reveals different genetic factors influencing infant, child, and adult BMI. Science Advances, 2019, 5, eaaw3095.	4.7	86
101	Phenome-wide investigation of health outcomes associated with genetic predisposition to loneliness. Human Molecular Genetics, 2019, 28, 3853-3865.	1.4	62
102	Target genes, variants, tissues and transcriptional pathways influencing human serum urate levels. Nature Genetics, 2019, 51, 1459-1474.	9.4	251
103	Genome-wide association meta-analysis of 30,000 samples identifies seven novel loci for quantitative ECG traits. European Journal of Human Genetics, 2019, 27, 952-962.	1.4	29
104	Multiancestry Genome-Wide Association Study of Lipid Levels Incorporating Gene-Alcohol Interactions. American Journal of Epidemiology, 2019, 188, 1033-1054.	1.6	85
105	Multi-ancestry study of blood lipid levels identifies four loci interacting with physical activity. Nature Communications, 2019, 10, 376.	5.8	64
106	Occupational exposure to gases/fumes and mineral dust affect DNA methylation levels of genes regulating expression. Human Molecular Genetics, 2019, 28, 2477-2485.	1.4	9
107	Analysis of Whole-Exome Sequencing Data for Alzheimer Disease Stratified by APOE Genotype. JAMA Neurology, 2019, 76, 1099.	4.5	32
108	Exome-Derived Adiponectin-Associated Variants Implicate Obesity and Lipid Biology. American Journal of Human Genetics, 2019, 105, 15-28.	2.6	21

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109	A catalog of genetic loci associated with kidney function from analyses of a million individuals. Nature Genetics, 2019, 51, 957-972.	9.4	549
110	Search for Early Pancreatic Cancer Blood Biomarkers in Five European Prospective Population Biobanks Using Metabolomics. Endocrinology, 2019, 160, 1731-1742.	1.4	19
111	An integrative cross-omics analysis of DNA methylation sites of glucose and insulin homeostasis. Nature Communications, 2019, 10, 2581.	5.8	62
112	Maternal and fetal genetic effects on birth weight and their relevance to cardio-metabolic risk factors. Nature Genetics, 2019, 51, 804-814.	9.4	402
113	Revisiting the Role of Insulin-Like Growth Factor-I Receptor Stimulating Activity and the Apolipoprotein E in Alzheimerâ€™s Disease. Frontiers in Aging Neuroscience, 2019, 11, 20.	1.7	24
114	Independent Multiple Factor Association Analysis for Multiblock Data in Imaging Genetics. Neuroinformatics, 2019, 17, 583-592.	1.5	2
115	The Epistasis Project: A Multi-Cohort Study of the Effects of BDNF, DBH, and SORT1 Epistasis on Alzheimerâ€™s Disease Risk. Journal of Alzheimer's Disease, 2019, 68, 1535-1547.	1.2	11
116	Association of variants inHTRA1andNOTCH3with MRI-defined extremes of cerebral small vessel disease in older subjects. Brain, 2019, 142, 1009-1023.	3.7	37
117	Limited overlap in significant hits between genome-wide association studies on two airflow obstruction definitions in the same population. BMC Pulmonary Medicine, 2019, 19, 58.	0.8	4
118	Loss of DPP6 in neurodegenerative dementia: a genetic player in the dysfunction of neuronal excitability. Acta Neuropathologica, 2019, 137, 901-918.	3.9	37
119	A multi-ancestry genome-wide study incorporating geneâ€“smoking interactions identifies multiple new loci for pulse pressure and mean arterial pressure. Human Molecular Genetics, 2019, 28, 2615-2633.	1.4	31
120	Multi-ancestry genome-wide geneâ€“smoking interaction study of 387,272 individuals identifies new loci associated with serum lipids. Nature Genetics, 2019, 51, 636-648.	9.4	112
121	Large-scale plasma metabolome analysis reveals alterations in HDL metabolism in migraine. Neurology, 2019, 92, e1899-e1911.	1.5	42
122	Association of metformin, sulfonylurea and insulin use with brain structure and function and risk of dementia and Alzheimerâ€™s disease: Pooled analysis from 5 cohorts. PLoS ONE, 2019, 14, e0212293.	1.1	65
123	Epigenome-wide Association Study of Attention-Deficit/Hyperactivity Disorder Symptoms in Adults. Biological Psychiatry, 2019, 86, 599-607.	0.7	47
124	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
125	Protein-coding variants implicate novel genes related to lipid homeostasis contributing to body-fat distribution. Nature Genetics, 2019, 51, 452-469.	9.4	89
126	Relationship between gut microbiota and circulating metabolites in population-based cohorts. Nature Communications, 2019, 10, 5813.	5.8	168

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127	DNA methylation is associated with lung function in never smokers. Respiratory Research, 2019, 20, 268.	1.4	14
128	Multi-trait genome-wide association study identifies new loci associated with optic disc parameters. Communications Biology, 2019, 2, 435.	2.0	22
129	Genetic architecture of subcortical brain structures in 38,851 individuals. Nature Genetics, 2019, 51, 1624-1636.	9.4	192
130	Genetic and lifestyle risk factors for MRI-defined brain infarcts in a population-based setting. Neurology, 2019, 92, .	1.5	30
131	Altered bile acid profile associates with cognitive impairment in Alzheimer's diseaseâ€"An emerging role for gut microbiome. Alzheimer's and Dementia, 2019, 15, 76-92.	0.4	396
132	Altered bile acid profile in mild cognitive impairment and Alzheimer's disease: Relationship to neuroimaging and CSF biomarkers. Alzheimer's and Dementia, 2019, 15, 232-244.	0.4	198
133	Increased High-Density Lipoprotein Levels Associated with Age-Related Macular Degeneration. Ophthalmology, 2019, 126, 393-406.	2.5	88
134	Candidate CSPG4 mutations and induced pluripotent stem cell modeling implicate oligodendrocyte progenitor cell dysfunction in familial schizophrenia. Molecular Psychiatry, 2019, 24, 757-771.	4.1	51
135	Genetic variation underlying cognition and its relation with neurological outcomes and brain imaging. Aging, 2019, 11, 1440-1456.	1.4	3
136	Association of branchedâ€chain amino acids and other circulating metabolites with risk of incident dementia and Alzheimer's disease: A prospective study in eight cohorts. Alzheimer's and Dementia, 2018, 14, 723-733.	0.4	182
137	Disentangling the biological pathways involved in early features of Alzheimer's disease in the Rotterdam Study. , 2018, 14, 848-857.		36
138	Genome-wide association meta-analysis of individuals of European ancestry identifies new loci explaining a substantial fraction of hair color variation and heritability. Nature Genetics, 2018, 50, 652-656.	9.4	86
139	Refining the accuracy of validated target identification through coding variant fine-mapping in type 2 diabetes. Nature Genetics, 2018, 50, 559-571.	9.4	356
140	COPD GWAS variant at 19q13.2 in relation with DNA methylation and gene expression. Human Molecular Genetics, 2018, 27, 396-405.	1.4	24
141	Understanding the role of the chromosome 15q25.1 in COPD through epigenetics and transcriptomics. European Journal of Human Genetics, 2018, 26, 709-722.	1.4	21
142	Circulating metabolites and general cognitive ability and dementia: Evidence from 11 cohort studies. Alzheimer's and Dementia, 2018, 14, 707-722.	0.4	143
143	Meta-analysis of epigenome-wide association studies of cognitive abilities. Molecular Psychiatry, 2018, 23, 2133-2144.	4.1	68
144	Life-Course Genome-wide Association Study Meta-analysis of Total Body BMD and Assessment of Age-Specific Effects. American Journal of Human Genetics, 2018, 102, 88-102.	2.6	252

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145	Meta-analysis of genome-wide association studies identifies 8 novel loci involved in shape variation of human head hair. Human Molecular Genetics, 2018, 27, 559-575.	1.4	51
146	Whole-Exome Sequencing in Age-Related Macular Degeneration Identifies Rare Variants in COL8A1, a Component of Bruchâ€™s Membrane. Ophthalmology, 2018, 125, 1433-1443.	2.5	35
147	DNA methylation signatures of educational attainment. Npj Science of Learning, 2018, 3, 7.	1.5	42
148	The effect of APOE and other common genetic variants on the onset of Alzheimer's disease and dementia: a community-based cohort study. Lancet Neurology, The, 2018, 17, 434-444.	4.9	177
149	Metabolic profiling of intra- and extracranial carotid artery atherosclerosis. Atherosclerosis, 2018, 272, 60-65.	0.4	24
150	Heritability and genome-wide associations studies of cerebral blood flow in the general population. Journal of Cerebral Blood Flow and Metabolism, 2018, 38, 1598-1608.	2.4	14
151	F3â€02â€02: CIRCULATING METABOLITESâ€™ ASSOCIATION WITH ALZHEIMER'S DISEASEâ€“ASSOCIATED GENETIC VARIANTS. Alzheimer's and Dementia, 2018, 14, P997.	0.4	0
152	P1â€156: GENEâ€BASED ANALYSES IN WHOLE GENOME SEQUENCING OF FAMILIAL LATEâ€ONSET ALZHEIMER'S DISEASE. Alzheimer's and Dementia, 2018, 14, P336.	0.4	0
153	O5â€04â€05: GENETIC VARIATION UNDERLYING COGNITION AND ITS RELATION WITH NEUROLOGICAL OUTCOMES. Alzheimer's and Dementia, 2018, 14, P1652.	0.4	0
154	P3â€134: CIRCULATING METABOLITES ARE ASSOCIATED WITH WHITE MATTER HYPERINTENSITIES. Alzheimer's and Dementia, 2018, 14, P1119.	0.4	0
155	P1â€297: METABOLIC BLOODâ€BASED BIOMARKERS RELATE TO BRAIN ATROPHY AND WHITE MATTER HYPERINTENSITIES IN ALZHEIMER'S DISEASE. Alzheimer's and Dementia, 2018, 14, P401.	0.4	0
156	P1â€298: CEREBROSPINAL FLUID AND PLASMA LEVELS OF LYSOPHOSPHATIDIC ACIDS (LPAS) ASSOCIATE WITH CEREBROSPINAL FLUID AÎ²â€42 AND Pâ€TAU. Alzheimer's and Dementia, 2018, 14, P403.	0.4	0
157	P2â€108: WHOLEâ€GENOME SEQUENCING IN NONâ€HISPANIC WHITE FAMILIES IMPLICATES RARE VARIATION IN LATEâ€ONSET ALZHEIMER'S DISEASE RISK. Alzheimer's and Dementia, 2018, 14, P710.	0.4	0
158	Blood Metabolomic Measures Associate With Present and Future Glycemic Control in Type 2 Diabetes. Journal of Clinical Endocrinology and Metabolism, 2018, 103, 4569-4579.	1.8	25
159	From blood to lung tissue: effect of cigarette smoke on DNA methylation and lung function. Respiratory Research, 2018, 19, 212.	1.4	47
160	Alzheimer's disease in Down syndrome: An overlooked population for prevention trials. Alzheimer's and Dementia: Translational Research and Clinical Interventions, 2018, 4, 703-713.	1.8	63
161	P4â€042: HIGHâ€DIMENSIONAL ANALYSIS OF RNA EXPRESSION WITH CORTICAL THICKNESS. Alzheimer's and Dementia, 2018, 14, P1449.	0.4	0
162	GWAS and colocalization analyses implicate carotid intima-media thickness and carotid plaque loci in cardiovascular outcomes. Nature Communications, 2018, 9, 5141.	5.8	119

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163	Long-term Air Pollution Exposure, Genome-wide DNA Methylation and Lung Function in the LifeLines Cohort Study. Environmental Health Perspectives, 2018, 126, 027004.	2.8	71
164	Genome-wide association study of 23,500 individuals identifies 7 loci associated with brain ventricular volume. Nature Communications, 2018, 9, 3945.	5.8	31
165	Whole-Genome Linkage Scan Combined With Exome Sequencing Identifies Novel Candidate Genes for Carotid Intima-Media Thickness. Frontiers in Genetics, 2018, 9, 420.	1.1	3
166	ExomeChip-Wide Analysis of 95 626 Individuals Identifies 10 Novel Loci Associated With QT and JT Intervals. Circulation Genomic and Precision Medicine, 2018, 11, e001758.	1.6	27
167	Large-scale whole-exome sequencing association studies identify rare functional variants influencing serum urate levels. Nature Communications, 2018, 9, 4228.	5.8	43
168	Genome-wide association study of primary open-angle glaucoma in continental and admixed African populations. Human Genetics, 2018, 137, 847-862.	1.8	40
169	Low-Frequency and Rare-Coding Variation Contributes to Multiple Sclerosis Risk. Cell, 2018, 175, 1679-1687.e7.	13.5	115
170	Occupational exposure to pesticides is associated with differential DNA methylation. Occupational and Environmental Medicine, 2018, 75, 427-435.	1.3	61
171	Autosomal genetic variation is associated with DNA methylation in regions variably escaping X-chromosome inactivation. Nature Communications, 2018, 9, 3738.	5.8	24
172	Genetic analysis of over 1 million people identifies 535 new loci associated with blood pressure traits. Nature Genetics, 2018, 50, 1412-1425.	9.4	924
173	Genome-wide association meta-analysis highlights light-induced signaling as a driver for refractive error. Nature Genetics, 2018, 50, 834-848.	9.4	239
174	Common and Rare Coding Genetic Variation Underlying the Electrocardiographic PR Interval. Circulation Genomic and Precision Medicine, 2018, 11, e002037.	1.6	19
175	Study of 300,486 individuals identifies 148 independent genetic loci influencing general cognitive function. Nature Communications, 2018, 9, 2098.	5.8	484
176	Cross-ancestry genome-wide association analysis of corneal thickness strengthens link between complex and Mendelian eye diseases. Nature Communications, 2018, 9, 1864.	5.8	63
177	Age at onset of genetic (E200K) and sporadic Creutzfeldt-Jakob diseases is modulated by the CYP4X1 gene. Journal of Neurology, Neurosurgery and Psychiatry, 2018, 89, 1243-1249.	0.9	14
178	PR interval genome-wide association meta-analysis identifies 50 loci associated with atrial and atrioventricular electrical activity. Nature Communications, 2018, 9, 2904.	5.8	71
179	A Genome-Wide Linkage Study for Chronic Obstructive Pulmonary Disease in a Dutch Genetic Isolate Identifies Novel Rare Candidate Variants. Frontiers in Genetics, 2018, 9, 133.	1.1	8
180	Exome Chip Analysis Identifies Low-Frequency and Rare Variants in MRPL38 for White Matter Hyperintensities on Brain Magnetic Resonance Imaging. Stroke, 2018, 49, 1812-1819.	1.0	17

#	Article	IF	Citations
181	Exome-chip meta-analysis identifies novel loci associated with cardiac conduction, including ADAMTS6. Genome Biology, 2018, 19, 87.	3.8	47
182	Reversal of Agingâ€Induced Increases in Aortic Stiffness by Targeting Cytoskeletal Proteinâ€Protein Interfaces. Journal of the American Heart Association, 2018, 7, .	1.6	17
183	A combined linkage, microarray and exome analysis suggests MAP3K11 as a candidate gene for left ventricular hypertrophy. BMC Medical Genomics, 2018, 11, 22.	0.7	4
184	Male-specific epistasis between WWC1 and TLN2 genes is associated with Alzheimer's disease. Neurobiology of Aging, 2018, 72, 188.e3-188.e12.	1.5	24
185	Three VCP Mutations in Patients with Frontotemporal Dementia. Journal of Alzheimer's Disease, 2018, 65, 1139-1146.	1.2	19
186	Genome-wide identification of directed gene networks using large-scale population genomics data. Nature Communications, 2018, 9, 3097.	5.8	18
187	Standard process-oriented workflow introduces pre-analytical error when used in large study sample batches. Clinical Chemistry and Laboratory Medicine, 2018, 56, e277-e279.	1.4	0
188	Novel genetic associations for blood pressure identified via gene-alcohol interaction in up to 570K individuals across multiple ancestries. PLoS ONE, 2018, 13, e0198166.	1.1	94
189	Multiancestry genome-wide association study of 520,000 subjects identifies 32 loci associated with stroke and stroke subtypes. Nature Genetics, 2018, 50, 524-537.	9.4	1,124
190	Novel genetic loci associated with hippocampal volume. Nature Communications, 2017, 8, 13624.	5.8	250
191	Rare and low-frequency coding variants alter human adult height. Nature, 2017, 542, 186-190.	13.7	544
192	Discovery and functional prioritization of Parkinsonâ€™s disease candidate genes from large-scale whole exome sequencing. Genome Biology, 2017, 18, 22.	3.8	96
193	Gray matter heritability in familyâ€based and populationâ€based studies using voxelâ€based morphometry. Human Brain Mapping, 2017, 38, 2408-2423.	1.9	9
194	New insights into the genetics of primary open-angle glaucoma based on meta-analyses of intraocular pressure and optic disc characteristics.. Human Molecular Genetics, 2017, 26, ddw399.	1.4	120
195	Genetic susceptibility to multiple sclerosis: Brain structure and cognitive function in the general population. Multiple Sclerosis Journal, 2017, 23, 1697-1706.	1.4	7
196	Discovery of novel heart rate-associated loci using the Exome Chip. Human Molecular Genetics, 2017, 26, 2346-2363.	1.4	29
197	Haplotype reference consortium panel: Practical implications of imputations with large reference panels. Human Mutation, 2017, 38, 1025-1032.	1.1	43
198	Genome-wide meta-analysis of 241,258 adults accounting for smoking behaviour identifies novel loci for obesity traits. Nature Communications, 2017, 8, 14977.	5.8	169

#	Article	IF	Citations
199	Disease variants alter transcription factor levels and methylation of their binding sites. Nature Genetics, 2017, 49, 131-138.	9.4	390
200	Identification of context-dependent expression quantitative trait loci in whole blood. Nature Genetics, 2017, 49, 139-145.	9.4	363
201	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
202	Variants in TTC25 affect autistic trait in patients with autism spectrum disorder and general population. European Journal of Human Genetics, 2017, 25, 982-987.	1.4	5
203	Genome-wide association meta-analysis of 78,308 individuals identifies new loci and genes influencing human intelligence. Nature Genetics, 2017, 49, 1107-1112.	9.4	425
204	Characterization of pathogenic SORL1 genetic variants for association with Alzheimerâ€™s disease: a clinical interpretation strategy. European Journal of Human Genetics, 2017, 25, 973-981.	1.4	102
205	An Expanded Genome-Wide Association Study of Type 2 Diabetes in Europeans. Diabetes, 2017, 66, 2888-2902.	0.3	615
206	Parental family history of dementia in relation to subclinical brain disease and dementia risk. Neurology, 2017, 88, 1642-1649.	1.5	44
207	Metabolic network failures in Alzheimer's disease: A biochemical roadÂmap. Alzheimer's and Dementia, 2017, 13, 965-984.	0.4	362
208	Genome-wide Trans-ethnic Meta-analysis Identifies Seven Genetic Loci Influencing Erythrocyte Traits and a Role for RBPMS in Erythropoiesis. American Journal of Human Genetics, 2017, 100, 51-63.	2.6	45
209	An Analysis of Two Genome-wide Association Meta-analyses Identifies a New Locus for Broad Depression Phenotype. Biological Psychiatry, 2017, 82, 322-329.	0.7	84
210	Evaluation of the Myocilin Mutation Gln368Stop Demonstrates Reduced Penetrance for Glaucoma in European Populations. Ophthalmology, 2017, 124, 547-553.	2.5	23
211	Epigenome-wide association study of body mass index, and the adverse outcomes of adiposity. Nature, 2017, 541, 81-86.	13.7	743
212	The Rotterdam Study: 2018 update on objectives, design and main results. European Journal of Epidemiology, 2017, 32, 807-850.	2.5	379
213	Plasma Amyloid-Î² Levels, Cerebral Small Vessel Disease, and Cognition: The Rotterdam Study. Journal of Alzheimer's Disease, 2017, 60, 977-987.	1.2	43
214	Genome-wide meta-analysis associates HLA-DQA1/DRB1 and LPA and lifestyle factors with human longevity. Nature Communications, 2017, 8, 910.	5.8	118
215	New Blood Pressureâ€“Associated Loci Identified in Meta-Analyses of 475â€‰000 Individuals. Circulation: Cardiovascular Genetics, 2017, 10, .	5.1	48
216	A Mendelian Randomization Study of Metabolite Profiles, Fasting Glucose, and Type 2 Diabetes. Diabetes, 2017, 66, 2915-2926.	0.3	40

#	Article	IF	Citations
217	Bloodâ€based metabolic signatures in Alzheimer's disease. Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring, 2017, 8, 196-207.	1.2	56
218	Novel Blood Pressure Locus and Gene Discovery Using Genome-Wide Association Study and Expression Data Sets From Blood and the Kidney. Hypertension, 2017, 70, .	1.3	123
219	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
220	Large meta-analysis of genome-wide association studies identifies five loci for lean body mass. Nature Communications, 2017, 8, 80.	5.8	147
221	Metabolomics based markers predict type 2 diabetes in a 14-year follow-up study. Metabolomics, 2017, 13, 104.	1.4	82
222	Common variants at 2q11.2, 8q21.3, and 11q13.2 are associated with major mood disorders. Translational Psychiatry, 2017, 7, 1273.	2.4	9
223	DNA Methylation Analysis Identifies Loci for Blood Pressure Regulation. American Journal of Human Genetics, 2017, 101, 888-902.	2.6	154
224	[P3â€“248]: PLASMA AMYLOID BETA LEVELS, CEREBRAL SMALLâ€VESSEL DISEASES AND COGNITION: THE ROTTERDAM STUDY. Alzheimer's and Dementia, 2017, 13, P1035.	0.4	0
225	Large-scale GWAS identifies multiple loci for hand grip strength providing biological insights into muscular fitness. Nature Communications, 2017, 8, 16015.	5.8	149
226	Lack of evidence for a role of genetic variation in TMEM230 in the risk for Parkinson's disease in the Caucasian population. Neurobiology of Aging, 2017, 50, 167.e11-167.e13.	1.5	24
227	Nonsynonymous Variation in NKPD1 Increases Depressive Symptoms in European Populations. Biological Psychiatry, 2017, 81, 702-707.	0.7	26
228	Genome-wide association study on the FEV 1 /FVC ratio in never-smokers identifies HHIP and FAM13A. Journal of Allergy and Clinical Immunology, 2017, 139, 533-540.	1.5	45
229	[P3â€“226]: PROFILING PERIPHERAL METABOLIC DYSREGULATION IN ALZHEIMER's DISEASE: THE ADDED VALUE OF MULTIPLE SIGNATURES. Alzheimer's and Dementia, 2017, 13, P1024.	0.4	0
230	[P1â€“139]: PATHWAYâ€SPECIFIC GENETIC RISK SCORE ASSOCIATED WITH ALZHEIMER's DISEASE AND WHITE MATTER LESIONS IN COGNITIVELY NORMAL SUBJECTS. Alzheimer's and Dementia, 2017, 13, P295.	0.4	0
231	[O2â€“08â€“03]: WHOLEâ€GENOME SEQUENCING IN FAMILIAL LATEâ€ONSET ALZHEIMER's DISEASE IDENTIFIES RARE VARIATION IN AD CANDIDATE GENES. Alzheimer's and Dementia, 2017, 13, P571.	0.4	1
232	A Genome-Wide Scan for MicroRNA-Related Genetic Variants Associated With Primary Open-Angle Glaucoma. , 2017, 58, 5368.		25
233	Exome-Wide Meta-Analysis Identifies Rare 3â€²-UTR Variant in ERCC1/CD3EAP Associated with Symptoms of Sleep Apnea. Frontiers in Genetics, 2017, 8, 151.	1.1	7
234	Genetic African Ancestry Is Associated With Central Corneal Thickness and Intraocular Pressure in Primary Open-Angle Glaucoma. , 2017, 58, 3172.		11

#	Article	IF	Citations
235	Genome-wide physical activity interactions in adiposity â€• A meta-analysis of 200,452 adults. PLoS Genetics, 2017, 13, e1006528.	1.5	158
236	A Combined Linkage and Exome Sequencing Analysis for Electrocardiogram Parameters in the Erasmus Rucphen Family Study. Frontiers in Genetics, 2016, 7, 190.	1.1	5
237	An Empirical Comparison of Joint and Stratified Frameworks for Studying G Ã— E Interactions: Systolic Blood Pressure and Smoking in the CHARGE Geneâ€Lifestyle Interactions Working Group. Genetic Epidemiology, 2016, 40, 404-415.	0.6	18
238	Exome Genotyping Identifies Pleiotropic Variants Associated with Red Blood Cell Traits. American Journal of Human Genetics, 2016, 99, 8-21.	2.6	60
239	Burden of genetic risk variants in multiple sclerosis families in the Netherlands. Multiple Sclerosis Journal - Experimental, Translational and Clinical, 2016, 2, 205521731664872.	0.5	6
240	The Generation R Study: design and cohort update 2017. European Journal of Epidemiology, 2016, 31, 1243-1264.	2.5	608
241	Meta-analysis of 49â€…549 individuals imputed with the 1000 Genomes Project reveals an exonic damaging variant inANGPTL4determining fasting TG levels. Journal of Medical Genetics, 2016, 53, 441-449.	1.5	34
242	P1â€•381: Evaluation of The Absolute Genetic Risk of Alzheimer's Disease in The Aging Population. Alzheimer's and Dementia, 2016, 12, P578.	0.4	1
243	Heritability of the shape of subcortical brain structures in the general population. Nature Communications, 2016, 7, 13738.	5.8	78
244	Associations with intraocular pressure across Europe: The European Eye Epidemiology (E3) Consortium. European Journal of Epidemiology, 2016, 31, 1101-1111.	2.5	26
245	F1-02-03: Metabolites Associated with Cognitive Function in the Rotterdam Study and Erasmus Rucphen Family Study. , 2016, 12, P165-P165.		4
246	O1-09-01: Genomewide Linkage Analysis Identifies Novel Candidate Genes for Alzheimerâ€™s Disease. , 2016, 12, P196-P196.		0
247	O1â€09â€03: Whole Genome Sequencing in Familial Lateâ€Onset Alzheimerâ€™s Disease Identifies Variations in TTC3 and FSIP2. Alzheimer's and Dementia, 2016, 12, P197.	0.4	0
248	O1â€09â€04: Identification of Whole Exome Sequencing Variants Associated with Lateâ€Onset Alzheimer's Disease in the Cohorts for Heart and Aging Research in Genomic Epidemiology (Charge) Consortium. Alzheimer's and Dementia, 2016, 12, P197.	0.4	0
249	O2â€06â€03: Tissueâ€Specific Genomeâ€Wide Predictions of Genetically Regulated Expression in Alzheimer's Disease. Alzheimer's and Dementia, 2016, 12, P239.	0.4	0
250	P1â€118: Association of Lowâ€Frequency and Rare Coding Variants with Information Processing Speed. Alzheimer's and Dementia, 2016, 12, P448.	0.4	0
251	Evaluation of a Genetic Risk Score to Improve Risk Prediction for Alzheimerâ€™s Disease. Journal of Alzheimer's Disease, 2016, 53, 921-932.	1.2	77
252	Diagnostic Yield and Clinical Utility of Sequencing Familial Hypercholesterolemia Genes in Patients With Severe Hypercholesterolemia. Journal of the American College of Cardiology, 2016, 67, 2578-2589.	1.2	723

#	Article	IF	Citations
253	Human age estimation from blood using mRNA, DNA methylation, DNA rearrangement, and telomere length. Forensic Science International: Genetics, 2016, 24, 33-43.	1.6	102
254	Identification of additional risk loci for stroke and small vessel disease: a meta-analysis of genome-wide association studies. Lancet Neurology, The, 2016, 15, 695-707.	4.9	130
255	Genetic variants associated with subjective well-being, depressive symptoms, and neuroticism identified through genome-wide analyses. Nature Genetics, 2016, 48, 624-633.	9.4	870
256	A Genome-Wide Association Study in isolated populations reveals new genes associated to common food likings. Reviews in Endocrine and Metabolic Disorders, 2016, 17, 209-219.	2.6	22
257	Genetic variants in RBFOX3 are associated with sleep latency. European Journal of Human Genetics, 2016, 24, 1488-1495.	1.4	27
258	Genome-wide association study identifies 74 loci associated with educational attainment. Nature, 2016, 533, 539-542.	13.7	1,204
259	Fine-mapping the effects of Alzheimer's disease risk loci on brain morphology. Neurobiology of Aging, 2016, 48, 204-211.	1.5	31
260	A novel method for serum lipoprotein profiling using high performance capillary isotachophoresis. Analytica Chimica Acta, 2016, 944, 57-69.	2.6	5
261	Discovery and refinement of genetic loci associated with cardiometabolic risk using dense imputation maps. Nature Genetics, 2016, 48, 1303-1312.	9.4	66
262	Age-related accrual of methylomic variability is linked to fundamental ageing mechanisms. Genome Biology, 2016, 17, 191.	3.8	120
263	Genome-wide associations for birth weight and correlations with adult disease. Nature, 2016, 538, 248-252.	13.7	406
264	Novel genetic loci underlying human intracranial volume identified through genome-wide association. Nature Neuroscience, 2016, 19, 1569-1582.	7.1	213
265	Blood lipids influence DNA methylation in circulating cells. Genome Biology, 2016, 17, 138.	3.8	154
266	Genomewide metaâ€analysis identifies loci associated with <scp>IGF</scp> â€I and <scp>IGFBP</scp> â€3 levels with impact on ageâ€related traits. Aging Cell, 2016, 15, 811-824.	3.0	83
267	Whole-Exome Sequencing Identifies Loci Associated with Blood Cell Traits and Reveals a Role for Alternative GFI1B Splice Variants in Human Hematopoiesis. American Journal of Human Genetics, 2016, 99, 481-488.	2.6	45
268	Shared genetic contribution to ischemic stroke and Alzheimer's disease. Annals of Neurology, 2016, 79, 739-747.	2.8	56
269	A reference panel of 64,976 haplotypes for genotype imputation. Nature Genetics, 2016, 48, 1279-1283.	9.4	2,421
270	KLB is associated with alcohol drinking, and its gene product Î²-Klotho is necessary for FGF21 regulation of alcohol preference. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 14372-14377.	3.3	208

#	Article	IF	Citations
271	Multiethnic Exome-Wide Association Study of Subclinical Atherosclerosis. Circulation: Cardiovascular Genetics, 2016, 9, 511-520.	5.1	54
272	O2-10-02: Genetic Determinants of MRI Subcortical Brain Structures: 24 Novel Loci Identified Through Gwas in 26,000 Persons. , 2016, 12, P251-P251.		0
273	A principal component meta-analysis on multiple anthropometric traits identifies novel loci for body shape. Nature Communications, 2016, 7, 13357.	5.8	74
274	52 Genetic Loci Influencing MyocardialÂMass. Journal of the American College of Cardiology, 2016, 68, 1435-1448.	1.2	113
275	Meta-analysis identifies common and rare variants influencing blood pressure and overlapping with metabolic trait loci. Nature Genetics, 2016, 48, 1162-1170.	9.4	223
276	The genetics of blood pressure regulation and its target organs from association studies in 342,415 individuals. Nature Genetics, 2016, 48, 1171-1184.	9.4	362
277	The Well-Known Gene HHIP and Novel Gene MECR Are Implicated in Small Airway Obstruction. American Journal of Respiratory and Critical Care Medicine, 2016, 194, 1299-1302.	2.5	11
278	Meta-analysis of geneâ€“environment-wide association scans accounting for education level identifies additional loci for refractive error. Nature Communications, 2016, 7, 11008.	5.8	104
279	Genome-wide study for circulating metabolites identifies 62 loci and reveals novel systemic effects of LPA. Nature Communications, 2016, 7, 11122.	5.8	576
280	A high-quality human reference panel reveals the complexity and distribution of genomic structural variants. Nature Communications, 2016, 7, 12989.	5.8	99
281	Non-additive genome-wide association scan reveals a new gene associated with habitual coffee consumption. Scientific Reports, 2016, 6, 31590.	1.6	25
282	Genome-wide association study for acute otitis media in children identifies FNDC1 as disease contributing gene. Nature Communications, 2016, 7, 12792.	5.8	50
283	Genome-wide analysis identifies 12 loci influencing human reproductive behavior. Nature Genetics, 2016, 48, 1462-1472.	9.4	284
284	Genetic variants linked to education predict longevity. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 13366-13371.	3.3	110
285	Twenty-eight genetic loci associated with ST-T-wave amplitudes of the electrocardiogram. Human Molecular Genetics, 2016, 25, 2093-2103.	1.4	24
286	Meta-analysis of 375,000 individuals identifies 38 susceptibility loci for migraine. Nature Genetics, 2016, 48, 856-866.	9.4	520
287	Genetic risk of Parkinsonâ€™s disease in the general population. Parkinsonism and Related Disorders, 2016, 29, 54-59.	1.1	11
288	Genetic loci for serum lipid fractions and intracerebral hemorrhage. Atherosclerosis, 2016, 246, 287-292.	0.4	11

#	Article	IF	Citations
289	Rare Exome Sequence Variants in CLCN6 Reduce Blood Pressure Levels and Hypertension Risk. Circulation: Cardiovascular Genetics, 2016, 9, 64-70.	5.1	44
290	Gene-based pleiotropy across migraine with aura and migraine without aura patient groups. Cephalalgia, 2016, 36, 648-657.	1.8	47
291	Quantifying prion disease penetrance using large population control cohorts. Science Translational Medicine, 2016, 8, 322ra9.	5.8	289
292	Heritability and Genome-Wide Association Analyses of Human Gait Suggest Contribution of Common Variants. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2016, 71, 740-746.	1.7	15
293	International Genome-Wide Association Study Consortium Identifies Novel Loci Associated With Blood Pressure in Children and Adolescents. Circulation: Cardiovascular Genetics, 2016, 9, 266-278.	5.1	48
294	Serum apolipoprotein E is associated with long-term risk of Alzheimerâ€™s disease: The Rotterdam Study. Neuroscience Letters, 2016, 617, 139-142.	1.0	25
295	New loci for body fat percentage reveal link between adiposity and cardiometabolic disease risk. Nature Communications, 2016, 7, 10495.	5.8	245
296	Bone Mineral Density in SjÃ¶gren Syndrome Patients with and Without Distal Renal Tubular Acidosis. Calcified Tissue International, 2016, 98, 573-579.	1.5	9
297	Genome-wide meta-analysis uncovers novel loci influencing circulating leptin levels. Nature Communications, 2016, 7, 10494.	5.8	153
298	Association of the IGF1 gene with fasting insulin levels. European Journal of Human Genetics, 2016, 24, 1337-1343.	1.4	5
299	Genetic associations at 53 loci highlight cell types and biological pathways relevant for kidney function. Nature Communications, 2016, 7, 10023.	5.8	412
300	Heritability and Genome-Wide Association Analyses of Intracranial Carotid Artery Calcification. Stroke, 2016, 47, 912-917.	1.0	15
301	Refined mapping of autoimmune disease associated genetic variants with gene expression suggests an important role for non-coding RNAs. Journal of Autoimmunity, 2016, 68, 62-74.	3.0	64
302	Gene co-expression analysis identifies brain regions and cell types involved in migraine pathophysiology: a GWAS-based study using the Allen Human Brain Atlas. Human Genetics, 2016, 135, 425-439.	1.8	47
303	Transmission of human mtDNA heteroplasmy in the Genome of the Netherlands families: support for a variable-size bottleneck. Genome Research, 2016, 26, 417-426.	2.4	84
304	Novel Genetic Loci Associated With Retinal Microvascular Diameter. Circulation: Cardiovascular Genetics, 2016, 9, 45-54.	5.1	28
305	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
306	Harmonising and linking biomedical and clinical data across disparate data archives to enable integrative cross-biobank research. European Journal of Human Genetics, 2016, 24, 521-528.	1.4	27

#	Article	IF	Citations
307	Meta-analysis of Genome-Wide Association Studies for Extraversion: Findings from the Genetics of Personality Consortium. Behavior Genetics, 2016, 46, 170-182.	1.4	178
308	Prediction of male-pattern baldness from genotypes. European Journal of Human Genetics, 2016, 24, 895-902.	1.4	44
309	The GenABEL Project for statistical genomics. F1000Research, 2016, 5, 914.	0.8	55
310	Rare Functional Variant in TM2D3 is Associated with Late-Onset Alzheimer's Disease. PLoS Genetics, 2016, 12, e1006327.	1.5	47
311	Investigating the Causal Relationship of C-Reactive Protein with 32 Complex Somatic and Psychiatric Outcomes: A Large-Scale Cross-Consortium Mendelian Randomization Study. PLoS Medicine, 2016, 13, e1001976.	3.9	150
312	Evaluation of Presumably Disease Causing SCN1A Variants in a Cohort of Common Epilepsy Syndromes. PLoS ONE, 2016, 11, e0150426.	1.1	22
313	O4-05-02: Genome-wide association study of lobar brain volumes. , 2015, 11, P278-P278.		0
314	Serum NGAL is Associated with Distinct Plasma Amyloid-Î² Peptides According to the Clinical Diagnosis of Dementia in Down Syndrome. Journal of Alzheimer's Disease, 2015, 45, 733-743.	1.2	17
315	Fine mapping the CETP region reveals a common intronic insertion associated to HDL-C. Npj Aging and Mechanisms of Disease, 2015, 1, 15011.	4.5	8
316	Heritabilities, proportions of heritabilities explained by GWAS findings, and implications of cross-phenotype effects on PR interval. Human Genetics, 2015, 134, 1211-1219.	1.8	20
317	Pathologically confirmed autoimmune encephalitis in suspected Creutzfeldt-Jakob disease. Neurology: Neuroimmunology and NeuroInflammation, 2015, 2, e178.	3.1	47
318	Rare variants in Î³â€aminobutyric acid type <scp>A</scp> receptor genes in rolandic epilepsy and related syndromes. Annals of Neurology, 2015, 77, 972-986.	2.8	51
319	Large-Scale Genomic Analyses Link Reproductive Aging to Hypothalamic Signaling, Breast Cancer Susceptibility, and BRCA1-Mediated DNA Repair. Obstetrical and Gynecological Survey, 2015, 70, 758-762.	0.2	0
320	A Genome Wide Association Study Links Glutamate Receptor Pathway to Sporadic Creutzfeldt-Jakob Disease Risk. PLoS ONE, 2015, 10, e0123654.	1.1	28
321	The Influence of Age and Sex on Genetic Associations with Adult Body Size and Shape: A Large-Scale Genome-Wide Interaction Study. PLoS Genetics, 2015, 11, e1005378.	1.5	331
322	Discovery and Fine-Mapping of Glycaemic and Obesity-Related Trait Loci Using High-Density Imputation. PLoS Genetics, 2015, 11, e1005230.	1.5	77
323	Parent-of-Origin Effects of the APOB Gene on Adiposity in Young Adults. PLoS Genetics, 2015, 11, e1005573.	1.5	16
324	Adiposity as a cause of cardiovascular disease: a Mendelian randomization study. International Journal of Epidemiology, 2015, 44, 578-586.	0.9	123

#	Article	IF	Citations
325	Meta-analysis of Genome-wide Association Studies for Neuroticism, and the Polygenic Association With Major Depressive Disorder. JAMA Psychiatry, 2015, 72, 642.	6.0	289
326	Shared genetic basis for migraine and ischemic stroke. Neurology, 2015, 84, 2132-2145.	1.5	91
327	Genome-wide association study identifies novel genetic variants contributing to variation in blood metabolite levels. Nature Communications, 2015, 6, 7208.	5.8	178
328	A priori collaboration in population imaging: The Uniform Neuroâ€Imaging of Virchowâ€Robin Spaces Enlargement consortium. Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring, 2015, 1, 513-520.	1.2	46
329	P2-024: Whole-exome sequencing in dutch families with Alzheimer's disease. , 2015, 11, P490-P490.		0
330	O3-13-01: Whole genome sequencing of late-onset Alzheimer's disease patients from genetic isolate. , 2015, 11, P250-P251.		0
331	O4-05-03: Whole exome sequence analysis of white matter hyperintensities on cranial MRI. , 2015, 11, P278-P279.		1
332	ARHGEF12 influences the risk of glaucoma by increasing intraocular pressure. Human Molecular Genetics, 2015, 24, 2689-2699.	1.4	79
333	New genetic loci link adipose and insulin biology to body fat distribution. Nature, 2015, 518, 187-196.	13.7	1,328
334	Genetic studies of body mass index yield new insights for obesity biology. Nature, 2015, 518, 197-206.	13.7	3,823
335	The dystrophin gene and cognitive function in the general population. European Journal of Human Genetics, 2015, 23, 837-843.	1.4	6
336	Low-frequency and rare exome chip variants associate with fasting glucose and type 2 diabetes susceptibility. Nature Communications, 2015, 6, 5897.	5.8	173
337	Metaâ€analysis of Genomeâ€Wide Association Studies Identifies Novel Loci Associated With Optic Disc Morphology. Genetic Epidemiology, 2015, 39, 207-216.	0.6	72
338	Common genetic variants influence human subcortical brain structures. Nature, 2015, 520, 224-229.	13.7	772
339	Association of Alzheimer's disease GWAS loci with MRI markers of brain aging. Neurobiology of Aging, 2015, 36, 1765.e7-1765.e16.	1.5	82
340	Age- and Sex-Specific Causal Effects of Adiposity on Cardiovascular Risk Factors. Diabetes, 2015, 64, 1841-1852.	0.3	63
341	Population-specific genotype imputations using minimac or IMPUTE2. Nature Protocols, 2015, 10, 1285-1296.	5.5	84
342	Directional dominance on stature and cognition inÂdiverse human populations. Nature, 2015, 523, 459-462.	13.7	173

#	Article	IF	Citations
343	Insight in Genome-Wide Association of Metabolite Quantitative Traits by Exome Sequence Analyses. PLoS Genetics, 2015, 11, e1004835.	1.5	70
344	Increasing Prevalence of Myopia in Europe and the Impact of Education. Ophthalmology, 2015, 122, 1489-1497.	2.5	329
345	Improving Phenotypic Prediction by Combining Genetic and Epigenetic Associations. American Journal of Human Genetics, 2015, 97, 75-85.	2.6	116
346	Genetic risk of neurodegenerative diseases is associated with mild cognitive impairment and conversion to dementia. Alzheimer's and Dementia, 2015, 11, 1277-1285.	0.4	76
347	Genome-wide association study for refractive astigmatism reveals genetic co-determination with spherical equivalent refractive error: the CREAM consortium. Human Genetics, 2015, 134, 131-146.	1.8	24
348	Prevalence of refractive error in Europe: the European Eye Epidemiology (E3) Consortium. European Journal of Epidemiology, 2015, 30, 305-315.	2.5	306
349	Characteristics of de novo structural changes in the human genome. Genome Research, 2015, 25, 792-801.	2.4	115
350	Multiethnic Genome-Wide Association Study of Cerebral White Matter Hyperintensities on MRI. Circulation: Cardiovascular Genetics, 2015, 8, 398-409.	5.1	162
351	Genome-wide association analysis on five isolated populations identifies variants of the HLA-DOA gene associated with white wine liking. European Journal of Human Genetics, 2015, 23, 1717-1722.	1.4	12
352	Genome-wide patterns and properties of de novo mutations in humans. Nature Genetics, 2015, 47, 822-826.	9.4	384
353	GWAS and Meta-Analysis in Aging/Longevity. Advances in Experimental Medicine and Biology, 2015, 847, 107-125.	0.8	22
354	GWAS of Longevity in CHARGE Consortium Confirms APOE and FOXO3 Candidacy. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2015, 70, 110-118.	1.7	250
355	The impact of low-frequency and rare variants on lipid levels. Nature Genetics, 2015, 47, 589-597.	9.4	310
356	PLD3 variants in population studies. Nature, 2015, 520, E2-E3.	13.7	49
357	Expression and Gene Variation Studies Deny Association of Human HSD3B1 Gene With Aldosterone Production or Blood Pressure. American Journal of Hypertension, 2015, 28, 113-120.	1.0	7
358	Polygenic Overlap Between C-Reactive Protein, Plasma Lipids, and Alzheimer Disease. Circulation, 2015, 131, 2061-2069.	1.6	145
359	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. Nature Genetics, 2015, 47, 1294-1303.	9.4	357
360	Common polygenic variation enhances risk prediction for Alzheimerâ€™s disease. Brain, 2015, 138, 3673-3684.	3.7	359

#	Article	IF	Citations
361	Effects of Metformin on Metabolite Profiles and LDL Cholesterol in Patients With Type 2 Diabetes. Diabetes Care, 2015, 38, 1858-1867.	4.3	97
362	White Matter Lesion Progression. Stroke, 2015, 46, 3048-3057.	1.0	27
363	Genome of the Netherlands population-specific imputations identify an ABCA6 variant associated with cholesterol levels. Nature Communications, 2015, 6, 6065.	5.8	45
364	Genetic Determinants of Unruptured Intracranial Aneurysms in the General Population. Stroke, 2015, 46, 2961-2964.	1.0	13
365	Wholeâ€genome sequencing identifies EN1 as a determinant of bone density and fracture. Nature, 2015, 526, 112-117.	13.7	483
366	WNT10A exonic variant increases the risk of keratoconus by decreasing corneal thickness. Human Molecular Genetics, 2015, 24, 5060-5068.	1.4	58
367	The Rotterdam Study: 2016 objectives and design update. European Journal of Epidemiology, 2015, 30, 661-708.	2.5	358
368	Genetic fine mapping and genomic annotation defines causal mechanisms at type 2 diabetes susceptibility loci. Nature Genetics, 2015, 47, 1415-1425.	9.4	365
369	Convergent genetic and expression data implicate immunity in Alzheimer's disease. Alzheimer's and Dementia, 2015, 11, 658-671.	0.4	173
370	A novel common variant in DCST2 is associated with length in early life and height in adulthood. Human Molecular Genetics, 2015, 24, 1155-1168.	1.4	109
371	Genome-wide Studies of Verbal Declarative Memory in Nondemented Older People: The Cohorts for Heart and Aging Research in Genomic Epidemiology Consortium. Biological Psychiatry, 2015, 77, 749-763.	0.7	67
372	Identifying genetic risk variants for coronary heart disease in familial hypercholesterolemia: an extreme genetics approach. European Journal of Human Genetics, 2015, 23, 381-387.	1.4	15
373	Genome-wide meta-analysis identifies six novel loci associated with habitual coffee consumption. Molecular Psychiatry, 2015, 20, 647-656.	4.1	235
374	Interferon gamma receptor 2 gene variants are associated with liver fibrosis in the general population: the Rotterdam Study. Gut, 2015, 64, 692-694.	6.1	3
375	Modulation of Genetic Associations with Serum Urate Levels by Body-Mass-Index in Humans. PLoS ONE, 2015, 10, e0119752.	1.1	64
376	Association Analysis of Bitter Receptor Genes in Five Isolated Populations Identifies a Significant Correlation between TAS2R43 Variants and Coffee Liking. PLoS ONE, 2014, 9, e92065.	1.1	41
377	Genome-Wide Meta-Analysis of Myopia and Hyperopia Provides Evidence for Replication of 11 Loci. PLoS ONE, 2014, 9, e107110.	1.1	40
378	The Generation R Study: Biobank update 2015. European Journal of Epidemiology, 2014, 29, 911-927.	2.5	189

#	Article	IF	Citations
379	Improved imputation quality of low-frequency and rare variants in European samples using the â€˜Genome of The Netherlandsâ€™. European Journal of Human Genetics, 2014, 22, 1321-1326.	1.4	92
380	O4-04-05: ASSOCIATION OF ALZHEIMER DISEASE GWAS LOCI WITH MRI-MARKERS OF BRAIN AGING. , 2014, 10, P258-P258.		0
381	Meta-analysis of genome-wide association studies identifies novel loci that influence cupping and the glaucomatous process. Nature Communications, 2014, 5, 4883.	5.8	89
382	Common genetic variants associated with cognitive performance identified using the proxy-phenotype method. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 13790-13794.	3.3	244
383	Leukocyte telomere length associates with prospective mortality independent of immune-related parameters and known genetic markers. International Journal of Epidemiology, 2014, 43, 878-886.	0.9	95
384	Gene-centric Meta-analysis in 87,736 Individuals of European Ancestry Identifies Multiple Blood-Pressure-Related Loci. American Journal of Human Genetics, 2014, 94, 349-360.	2.6	158
385	Fraction of exhaled nitric oxide values in childhood are associated with 17q11.2-q12 and 17q12-q21 variants. Journal of Allergy and Clinical Immunology, 2014, 134, 46-55.	1.5	33
386	A metabolomic profile is associated with the risk of incident coronary heart disease. American Heart Journal, 2014, 168, 45-52.e7.	1.2	74
387	Association of Low-Frequency and Rare Coding-Sequence Variants with Blood Lipids and Coronary Heart Disease in 56,000 Whites and Blacks. American Journal of Human Genetics, 2014, 94, 223-232.	2.6	287
388	Whole-Exome Sequencing Identifies Rare and Low-Frequency Coding Variants Associated with LDL Cholesterol. American Journal of Human Genetics, 2014, 94, 233-245.	2.6	193
389	Genetic Susceptibility, Dietary Antioxidants, and Long-Term Incidence of Age-Related Macular Degeneration in Two Populations. Ophthalmology, 2014, 121, 667-675.	2.5	59
390	Scientific reporting is suboptimal for aspects that characterize genetic risk prediction studies: a review of published articles based on the Genetic RIsk Prediction Studies statement. Journal of Clinical Epidemiology, 2014, 67, 487-499.	2.4	7
391	The Genome of the Netherlands: design, and project goals. European Journal of Human Genetics, 2014, 22, 221-227.	1.4	246
392	Insulin-Like Growth Factor-I Receptor Stimulating Activity is Associated with Dementia. Journal of Alzheimer's Disease, 2014, 42, 137-142.	1.2	25
393	Exome sequencing and functional analyses suggest that SIX6 is a gene involved in an altered proliferationâ€“differentiation balance early in life and optic nerve degeneration at old age. Human Molecular Genetics, 2014, 23, 1320-1332.	1.4	63
394	Predicting Stroke Through Genetic Risk Functions. Stroke, 2014, 45, 403-412.	1.0	62
395	Multilocus Genetic Risk Score Associates With Ischemic Stroke in Caseâ€“Control and Prospective Cohort Studies. Stroke, 2014, 45, 394-402.	1.0	56
396	Novel loci affecting iron homeostasis and their effects in individuals at risk for hemochromatosis. Nature Communications, 2014, 5, 4926.	5.8	192

#	Article	IF	Citations
397	Association of adiponectin and leptin with relative telomere length in seven independent cohorts including 11,448 participants. European Journal of Epidemiology, 2014, 29, 629-638.	2.5	23
398	Gene-Age Interactions in Blood Pressure Regulation: A Large-Scale Investigation with the CHARGE, Global BPgen, and ICBP Consortia. American Journal of Human Genetics, 2014, 95, 24-38.	2.6	109
399	Effects of Long-Term Averaging of Quantitative Blood Pressure Traits on the Detection of Genetic Associations. American Journal of Human Genetics, 2014, 95, 49-65.	2.6	73
400	Parent-of-origin-specific allelic associations among 106 genomic loci for age at menarche. Nature, 2014, 514, 92-97.	13.7	548
401	Genome-wide analysis of multi-ancestry cohorts identifies new loci influencing intraocular pressure and susceptibility to glaucoma. Nature Genetics, 2014, 46, 1126-1130.	9.4	212
402	Direct-to-Consumer Personal Genome Testing for Age-Related Macular Degeneration. , 2014, 55, 6167.		18
403	Genome-wide association meta-analysis of human longevity identifies a novel locus conferring survival beyond 90 years of age. Human Molecular Genetics, 2014, 23, 4420-4432.	1.4	227
404	Defining the role of common variation in the genomic and biological architecture of adult human height. Nature Genetics, 2014, 46, 1173-1186.	9.4	1,818
405	Harmonization of Neuroticism and Extraversion phenotypes across inventories and cohorts in the Genetics of Personality Consortium: an application of Item Response Theory. Behavior Genetics, 2014, 44, 295-313.	1.4	103
406	Telomere length in circulating leukocytes is associated with lung function and disease. European Respiratory Journal, 2014, 43, 983-992.	3.1	103
407	Genome-wide trans-ancestry meta-analysis provides insight into the genetic architecture of type 2 diabetes susceptibility. Nature Genetics, 2014, 46, 234-244.	9.4	959
408	Genetic association study of QT interval highlights role for calcium signaling pathways in myocardial repolarization. Nature Genetics, 2014, 46, 826-836.	9.4	281
409	Trans-ethnic meta-analysis of white blood cell phenotypes. Human Molecular Genetics, 2014, 23, 6944-6960.	1.4	60
410	TMEM106B Influences Volume of Left-Sided Temporal Lobe and Interhemispheric Structures in the General Population. Biological Psychiatry, 2014, 76, 503-508.	0.7	21
411	Genetics of the human metabolome, what is next?. Biochimica Et Biophysica Acta - Molecular Basis of Disease, 2014, 1842, 1923-1931.	1.8	28
412	The sex-specific associations of the aromatase gene with Alzheimerâ€™s disease and its interaction with IL10 in the Epistasis Project. European Journal of Human Genetics, 2014, 22, 216-220.	1.4	35
413	Serum MHPG Strongly Predicts Conversion to Alzheimer's Disease in Behaviorally Characterized Subjects with Down Syndrome. Journal of Alzheimer's Disease, 2014, 43, 871-891.	1.2	32
414	O4-08-03: GENETIC RISK OF NEURODEGENERATIVE DISEASES IS ASSOCIATED WITH MILD COGNITIVE IMPAIRMENT AND CONVERSION TO DEMENTIA: THE ROTTERDAM STUDY. , 2014, 10, P267-P267.		0

#	Article	IF	Citations
415	P1-167: SERUM LIPOCALIN-2: MONITORING THE NEUROPATHOLOGICAL PROGRESSION OF AD IN DOWN SYNDROME?. , 2014, 10, P361-P361.		0
416	P4-282: PLD3 ASSOCIATES TO PROLINE A PROPOSED BIOMARKER IN MAPSTONE ET AL. , 2014, 10, P887-P888.		2
417	Susceptibility to Chronic Mucus Hypersecretion, a Genome Wide Association Study. PLoS ONE, 2014, 9, e91621.	1.1	25
418	Gene-Wide Analysis Detects Two New Susceptibility Genes for Alzheimer's Disease. PLoS ONE, 2014, 9, e94661.	1.1	155
419	Associations of NINJ2 Sequence Variants with Incident Ischemic Stroke in the Cohorts for Heart and Aging in Genomic Epidemiology (CHARGE) Consortium. PLoS ONE, 2014, 9, e99798.	1.1	11
420	The Challenges of Genome-Wide Interaction Studies: Lessons to Learn from the Analysis of HDL Blood Levels. PLoS ONE, 2014, 9, e109290.	1.1	13
421	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. Neurobiology of Aging, 2013, 34, 1309.e1-1309.e7.	1.5	29
422	Heritability of dietary food intake patterns. Acta Diabetologica, 2013, 50, 721-726.	1.2	18
423	Discovery and refinement of loci associated with lipid levels. Nature Genetics, 2013, 45, 1274-1283.	9.4	2,641
424	Common variants associated with plasma triglycerides and risk for coronary artery disease. Nature Genetics, 2013, 45, 1345-1352.	9.4	754
425	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
426	Meta-analysis of genome-wide association studies in five cohorts reveals common variants in RBFOX1, a regulator of tissue-specific splicing, associated with refractive error. Human Molecular Genetics, 2013, 22, 2754-2764.	1.4	60
427	A genomeâ€wide association study of sleep habits and insomnia. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2013, 162, 439-451.	1.1	104
428	Loci influencing blood pressure identified using a cardiovascular gene-centric array. Human Molecular Genetics, 2013, 22, 1663-1678.	1.4	141
429	Automated workflow-based exploitation of pathway databases provides new insights into genetic associations of metabolite profiles. BMC Genomics, 2013, 14, 865.	1.2	14
430	The Rotterdam Study: 2014 objectives and design update. European Journal of Epidemiology, 2013, 28, 889-926.	2.5	282
431	The vast complexity of primary open angle glaucoma: Disease genes, risks, molecular mechanisms and pathobiology. Progress in Retinal and Eye Research, 2013, 37, 31-67.	7.3	149
432	Refining genome-wide linkage intervals using a meta-analysis of genome-wide association studies identifies loci influencing personality dimensions. European Journal of Human Genetics, 2013, 21, 876-882.	1.4	24

#	Article	IF	Citations
433	Variant ofTREM2Associated with the Risk of Alzheimer's Disease. New England Journal of Medicine, 2013, 368, 107-116.	13.9	2,085
434	Genome-wide association analyses identify 18 new loci associated with serum urate concentrations. Nature Genetics, 2013, 45, 145-154.	9.4	675
435	Nine Loci for Ocular Axial Length Identified through Genome-wide Association Studies, Including Shared Loci with Refractive Error. American Journal of Human Genetics, 2013, 93, 264-277.	2.6	139
436	Common Variants in Mendelian Kidney Disease Genes and Their Association with Renal Function. Journal of the American Society of Nephrology: JASN, 2013, 24, 2105-2117.	3.0	33
437	Plasma phosphatidylcholine and sphingomyelin concentrations are associated with depression and anxiety symptoms in a Dutch family-based lipidomics study. Journal of Psychiatric Research, 2013, 47, 357-362.	1.5	115
438	Alzheimer's Disease Genes and Cognition in the Nondemented General Population. Biological Psychiatry, 2013, 73, 429-434.	0.7	79
439	Prediction of Age-related Macular Degeneration in the General Population. Ophthalmology, 2013, 120, 2644-2655.	2.5	84
440	Assessment of the 9p21.3 locus in severity of coronary artery disease in the presence and absence of type 2 diabetes. BMC Medical Genetics, 2013, 14, 11.	2.1	24
441	Genome-wide meta-analyses of multiancestry cohorts identify multiple new susceptibility loci for refractive error and myopia. Nature Genetics, 2013, 45, 314-318.	9.4	398
442	Identification of seven loci affecting mean telomere length and their association with disease. Nature Genetics, 2013, 45, 422-427.	9.4	808
443	Distinguishing true from false positives in genomic studies: p values. European Journal of Epidemiology, 2013, 28, 131-138.	2.5	36
444	Genome-wide meta-analysis identifies 11 new loci for anthropometric traits and provides insights into genetic architecture. Nature Genetics, 2013, 45, 501-512.	9.4	578
445	A Genome-Wide Association Study of Depressive Symptoms. Biological Psychiatry, 2013, 73, 667-678.	0.7	149
446	Seven new loci associated with age-related macular degeneration. Nature Genetics, 2013, 45, 433-439.	9.4	687
447	Identification of heart rateâ€“associated loci and their effects on cardiac conduction and rhythm disorders. Nature Genetics, 2013, 45, 621-631.	9.4	282
448	Meta-analysis of telomere length in 19â€‰713 subjects reveals high heritability, stronger maternal inheritance and a paternal age effect. European Journal of Human Genetics, 2013, 21, 1163-1168.	1.4	380
449	GWAS of 126,559 Individuals Identifies Genetic Variants Associated with Educational Attainment. Science, 2013, 340, 1467-1471.	6.0	750
450	Genome-wide meta-analysis identifies new susceptibility loci for migraine. Nature Genetics, 2013, 45, 912-917.	9.4	338

#	Article	IF	Citations
451	Genome-wide analysis of BMI in adolescents and young adults reveals additional insight into the effects of genetic loci over the life course. Human Molecular Genetics, 2013, 22, 3597-3607.	1.4	116
452	Identification of a rare coding variant in complement 3 associated with age-related macular degeneration. Nature Genetics, 2013, 45, 1375-1379.	9.4	158
453	Low-density lipoprotein receptor mutations generate synthetic genome-wide associations. European Journal of Human Genetics, 2013, 21, 563-566.	1.4	7
454	Loci influencing blood pressure identified using a cardiovascular gene-centric array. Human Molecular Genetics, 2013, 22, 3394-3395.	1.4	1
455	Sex-stratified Genome-wide Association Studies Including 270,000 Individuals Show Sexual Dimorphism in Genetic Loci for Anthropometric Traits. PLoS Genetics, 2013, 9, e1003500.	1.5	371
456	Fine-Mapping the Genetic Association of the Major Histocompatibility Complex in Multiple Sclerosis: HLA and Non-HLA Effects. PLoS Genetics, 2013, 9, e1003926.	1.5	250
457	The Role of Adiposity in Cardiometabolic Traits: A Mendelian Randomization Analysis. PLoS Medicine, 2013, 10, e1001474.	3.9	178
458	Risk Scores of Common Genetic Variants for Lipid Levels Influence Atherosclerosis and Incident Coronary Heart Disease. Arteriosclerosis, Thrombosis, and Vascular Biology, 2013, 33, 2233-2239.	1.1	44
459	Genome-wide association study of intraocular pressure identifies the GLCCI1/ICA1 region as a glaucoma susceptibility locus. Human Molecular Genetics, 2013, 22, 4653-4660.	1.4	29
460	Genetic variation associated with circulating monocyte count in the eMERGE Network. Human Molecular Genetics, 2013, 22, 2119-2127.	1.4	56
461	Genetic influences on plasma CFH and CFHR1 concentrations and their role in susceptibility to age-related macular degeneration. Human Molecular Genetics, 2013, 22, 4857-4869.	1.4	77
462	Genetic Loci for Coronary Calcification and Serum Lipids Relate to Aortic and Carotid Calcification. Circulation: Cardiovascular Genetics, 2013, 6, 47-53.	5.1	19
463	Genome-wide association study meta-analysis of chronic widespread pain: evidence for involvement of the 5p15.2 region. Annals of the Rheumatic Diseases, 2013, 72, 427-436.	0.5	112
464	Association of Adiposity Genetic Variants With Menarche Timing in 92,105 Women of European Descent. American Journal of Epidemiology, 2013, 178, 451-460.	1.6	51
465	Genome-wide association analyses identify multiple loci associated with central corneal thickness and keratoconus. Nature Genetics, 2013, 45, 155-163.	9.4	269
466	Migraine is not associated with enhanced atherosclerosis. Cephalalgia, 2013, 33, 228-235.	1.8	57
467	New loci associated with birth weight identify genetic links between intrauterine growth and adult height and metabolism. Nature Genetics, 2013, 45, 76-82.	9.4	293
468	Identification of a Candidate Gene for Astigmatism. , 2013, 54, 1260.		31

#	Article	IF	Citations
469	Insights into the Genetic Architecture of Early Stage Age-Related Macular Degeneration: A Genome-Wide Association Study Meta-Analysis. PLoS ONE, 2013, 8, e53830.	1.1	108
470	Genome-Wide Association Study of Retinopathy in Individuals without Diabetes. PLoS ONE, 2013, 8, e54232.	1.1	22
471	The Molecular Genetic Architecture of Self-Employment. PLoS ONE, 2013, 8, e60542.	1.1	41
472	Genetic Loci for Retinal Arteriolar Microcirculation. PLoS ONE, 2013, 8, e65804.	1.1	27
473	Development and Application of Genomic Control Methods for Genome-Wide Association Studies Using Non-Additive Models. PLoS ONE, 2013, 8, e81431.	1.1	14
474	Region-Based Association Analysis of Human Quantitative Traits in Related Individuals. PLoS ONE, 2013, 8, e65395.	1.1	30
475	Best Practices and Joint Calling of the HumanExome BeadChip: The CHARGE Consortium. PLoS ONE, 2013, 8, e68095.	1.1	219
476	Common Genetic Determinants of Intraocular Pressure and Primary Open-Angle Glaucoma. PLoS Genetics, 2012, 8, e1002611.	1.5	164
477	Genome-Wide Profiling of Blood Pressure in Adults and Children. Hypertension, 2012, 59, 241-247.	1.3	31
478	Evidence of Inbreeding Depression on Human Height. PLoS Genetics, 2012, 8, e1002655.	1.5	79
479	Genome-Wide Association Study Identifies Novel Loci Associated with Circulating Phospho- and Sphingolipid Concentrations. PLoS Genetics, 2012, 8, e1002490.	1.5	181
480	Genome-Wide Association and Functional Follow-Up Reveals New Loci for Kidney Function. PLoS Genetics, 2012, 8, e1002584.	1.5	166
481	Novel Loci for Adiponectin Levels and Their Influence on Type 2 Diabetes and Metabolic Traits: A Multi-Ethnic Meta-Analysis of 45,891 Individuals. PLoS Genetics, 2012, 8, e1002607.	1.5	419
482	A Genetic Epidemiologic Study of Candidate Genes Involved in the Optic Nerve Head Morphology. , 2012, 53, 1485.		9
483	Integration of genome-wide association studies with biological knowledge identifies six novel genes related to kidney function. Human Molecular Genetics, 2012, 21, 5329-5343.	1.4	64
484	Genome-Wide Association Study of Vascular Dementia. Stroke, 2012, 43, 315-319.	1.0	51
485	Eight genetic loci associated with variation in lipoprotein-associated phospholipase A2 mass and activity and coronary heart disease: meta-analysis of genome-wide association studies from five community-based studies. European Heart Journal, 2012, 33, 238-251.	1.0	89
486	Large-scale association analyses identify new loci influencing glycemic traits and provide insight into the underlying biological pathways. Nature Genetics, 2012, 44, 991-1005.	9.4	746

#	Article	IF	Citations
487	Common variants at 6q22 and 17q21 are associated with intracranial volume. Nature Genetics, 2012, 44, 539-544.	9.4	126
488	Common variants at 12q14 and 12q24 are associated with hippocampal volume. Nature Genetics, 2012, 44, 545-551.	9.4	212
489	Common variants at 12q15 and 12q24 are associated with infant head circumference. Nature Genetics, 2012, 44, 532-538.	9.4	130
490	Common Genetic Variation in the 3â€²- BCL11B Gene Desert Is Associated With Carotid-Femoral Pulse Wave Velocity and Excess Cardiovascular Disease Risk. Circulation: Cardiovascular Genetics, 2012, 5, 81-90.	5.1	90
491	Cerebrospinal fluid biomarker supported diagnosis of Creutzfeldtâ€“Jakob disease and rapid dementias: a longitudinal multicentre study over 10 years. Brain, 2012, 135, 3051-3061.	3.7	135
492	CYP1A2 and coffee intake and the modifying effect of sex, age, and smoking. American Journal of Clinical Nutrition, 2012, 96, 182-187.	2.2	38
493	Large scale international replication and meta-analysis study confirms association of the 15q14 locus with myopia. The CREAM consortium. Human Genetics, 2012, 131, 1467-1480.	1.8	67
494	Linkage analysis for plasma amyloid beta levels in persons with hypertension implicates AÎ²-40 levels to presenilin 2. Human Genetics, 2012, 131, 1869-1876.	1.8	7
495	A Large-Scale Population-Based Analysis of Common Genetic Variation in the Thyroid Hormone Receptor Alpha Locus and Bone. Thyroid, 2012, 22, 223-224.	2.4	7
496	Genome-wide meta-analysis identifies 56 bone mineral density loci and reveals 14 loci associated with risk of fracture. Nature Genetics, 2012, 44, 491-501.	9.4	1,100
497	Large-scale association analysis provides insights into the genetic architecture and pathophysiology of type 2 diabetes. Nature Genetics, 2012, 44, 981-990.	9.4	1,748
498	The Generation R Study: design and cohort update 2012. European Journal of Epidemiology, 2012, 27, 739-756.	2.5	486
499	A genome-wide association meta-analysis identifies new childhood obesity loci. Nature Genetics, 2012, 44, 526-531.	9.4	352
500	Transferrin and HFE genes interact in Alzheimer's disease risk: the Epistasis Project. Neurobiology of Aging, 2012, 33, 202.e1-202.e13.	1.5	51
501	Genome-wide study links MTMR7 gene to variant Creutzfeldt-Jakob risk. Neurobiology of Aging, 2012, 33, 1487.e21-1487.e28.	1.5	40
502	Rapid variance componentsâ€“based method for whole-genome association analysis. Nature Genetics, 2012, 44, 1166-1170.	9.4	193
503	Association Between Chromosome 9p21 Variants and the Ankle-Brachial Index Identified by a Meta-Analysis of 21 Genome-Wide Association Studies. Circulation: Cardiovascular Genetics, 2012, 5, 100-112.	5.1	98
504	Accuracy of self-reported family history is strongly influenced by the accuracy of self-reported personal health status of relatives. Journal of Clinical Epidemiology, 2012, 65, 82-89.	2.4	24

#	Article	IF	Citations
505	FTO genotype is associated with phenotypic variability of body mass index. Nature, 2012, 490, 267-272.	13.7	383
506	Nucleotide Excision DNA Repair Is Associated With Age-Related Vascular Dysfunction. Circulation, 2012, 126, 468-478.	1.6	153
507	Meta-analyses identify 13 loci associated with age at menopause and highlight DNA repair and immune pathways. Nature Genetics, 2012, 44, 260-268.	9.4	303
508	Discovery and Fine Mapping of Serum Protein Loci through Transethnic Meta-analysis. American Journal of Human Genetics, 2012, 91, 744-753.	2.6	69
509	Heritability and Genome-Wide Association Study to Assess Genetic Differences between Advanced Age-related Macular Degeneration Subtypes. Ophthalmology, 2012, 119, 1874-1885.	2.5	73
510	Meta-analysis of genome-wide association studies identifies three new risk loci for atopic dermatitis. Nature Genetics, 2012, 44, 187-192.	9.4	311
511	A Genome-Wide Association Search for Type 2 Diabetes Genes in African Americans. PLoS ONE, 2012, 7, e29202.	1.1	197
512	A genome-wide approach accounting for body mass index identifies genetic variants influencing fasting glycemic traits and insulin resistance. Nature Genetics, 2012, 44, 659-669.	9.4	762
513	Genome-wide association analysis identifies susceptibility loci for migraine without aura. Nature Genetics, 2012, 44, 777-782.	9.4	294
514	Interaction of insulin and PPAR-Î± genes in Alzheimerâ€™s disease: the Epistasis Project. Journal of Neural Transmission, 2012, 119, 473-479.	1.4	20
515	Human Prion Diseases in The Netherlands (1998â€“2009): Clinical, Genetic and Molecular Aspects. PLoS ONE, 2012, 7, e36333.	1.1	44
516	Candidate Gene-Based Association Study of Antipsychotic-Induced Movement Disorders in Long-Stay Psychiatric Patients: A Prospective Study. PLoS ONE, 2012, 7, e36561.	1.1	22
517	Antipsychotic-Induced Movement Disorders in Long-Stay Psychiatric Patients and 45 Tag SNPs in 7 Candidate Genes: A Prospective Study. PLoS ONE, 2012, 7, e50970.	1.1	27
518	Interactions between PPAR-Î± and inflammation-related cytokine genes on the development of Alzheimer's disease, observed by the Epistasis Project. International Journal of Molecular Epidemiology and Genetics, 2012, 3, 39-47.	0.4	13
519	Multicenter cohort association study of SLC2A1 single nucleotide polymorphisms and age-related macular degeneration. Molecular Vision, 2012, 18, 657-74.	1.1	5
520	Meta-Analysis of Genome-Wide Association Studies in >80 000 Subjects Identifies Multiple Loci for C-Reactive Protein Levels. Circulation, 2011, 123, 731-738.	1.6	461
521	O3-01-01: Genome-wide association studies of hippocampal volume: The CHARGE consortium. , 2011, 7, S495-S496.		0
522	Genetic variation near IRS1 associates with reduced adiposity and an impaired metabolic profile. Nature Genetics, 2011, 43, 753-760.	9.4	289

#	Article	IF	Citations
523	Genome-wide association study identifies loci influencing concentrations of liver enzymes in plasma. Nature Genetics, 2011, 43, 1131-1138.	9.4	501
524	Strengthening the reporting of Genetic RIsk Prediction Studies (GRIPS): explanation and elaboration. Journal of Clinical Epidemiology, 2011, 64, e1-e22.	2.4	9
525	The Neuronal Transporter Gene SLC6A15 Confers Risk to Major Depression. Neuron, 2011, 70, 252-265.	3.8	189
526	Clinical Implications of Old and New Genes for Open-Angle Glaucoma. Ophthalmology, 2011, 118, 2389-2397.	2.5	34
527	Association of HSP70 and its Co-Chaperones with Alzheimer's Disease. Journal of Alzheimer's Disease, 2011, 25, 93-102.	1.2	21
528	Variation in the von Willebrand factor gene is associated with von Willebrand factor levels and with the risk for cardiovascular disease. Blood, 2011, 117, 1393-1399.	0.6	55
529	Strengthening the reporting of genetic risk prediction studies (GRIPS): explanation and elaboration. European Journal of Clinical Investigation, 2011, 41, 1010-1035.	1.7	30
530	Genomeâ€wide association study identifies a single major locus contributing to survival into old age; the APOE locus revisited. Aging Cell, 2011, 10, 686-698.	3.0	249
531	Common variants at ABCA7, MS4A6A/MS4A4E, EPHA1, CD33 and CD2AP are associated with Alzheimer's disease. Nature Genetics, 2011, 43, 429-435.	9.4	1,708
532	Genetic architecture of circulating lipid levels. European Journal of Human Genetics, 2011, 19, 813-819.	1.4	23
533	Strengthening the reporting of genetic risk prediction studies (GRIPS): explanation and elaboration. European Journal of Human Genetics, 2011, 19, 615-615.	1.4	12
534	Meta-analysis of genome-wide association for migraine in six population-based European cohorts. European Journal of Human Genetics, 2011, 19, 901-907.	1.4	87
535	Genetics of cortisol secretion and depressive symptoms: A candidate gene and genome wide association approach. Psychoneuroendocrinology, 2011, 36, 1053-1061.	1.3	85
536	Genome-Wide Association Study for Coronary Artery Calcification With Follow-Up in Myocardial Infarction. Circulation, 2011, 124, 2855-2864.	1.6	269
537	Strengthening the reporting of genetic risk prediction studies (GRIPS): explanation and elaboration. European Journal of Epidemiology, 2011, 26, 313-337.	2.5	14
538	Strengthening the reporting of genetic risk prediction studies: the GRIPS statement. European Journal of Epidemiology, 2011, 26, 255-259.	2.5	13
539	PredictABEL: an R package for the assessment of risk prediction models. European Journal of Epidemiology, 2011, 26, 261-264.	2.5	231
540	The Rotterdam Study: 2012 objectives and design update. European Journal of Epidemiology, 2011, 26, 657-686.	2.5	273

#	Article	IF	Citations
541	STROBE-ME too!. European Journal of Epidemiology, 2011, 26, 761-762.	2.5	1
542	Association of heat shock proteins with Parkinsonâ€™s disease. European Journal of Epidemiology, 2011, 26, 933-935.	2.5	8
543	The relationship between fertility and lifespan in humans. Age, 2011, 33, 615-622.	3.0	34
544	A Methodological Perspective on Genetic Risk Prediction Studies in Type 2 Diabetes: Recommendations for Future Research. Current Diabetes Reports, 2011, 11, 511-518.	1.7	28
545	A polymorphism in the regulatory region of PRNPis associated with increased risk of sporadic Creutzfeldt-Jakob disease. BMC Medical Genetics, 2011, 12, 73.	2.1	18
546	Genomeâ€wide association studies of cerebral white matter lesion burden. Annals of Neurology, 2011, 69, 928-939.	2.8	201
547	Linkage and association analyses of glaucoma related traits in a large pedigree from a Dutch genetically isolated population. Journal of Medical Genetics, 2011, 48, 802-809.	1.5	38
548	Association of Hypertension Drug Target Genes With Blood Pressure and Hypertension in 86 588 Individuals. Hypertension, 2011, 57, 903-910.	1.3	181
549	Reducing the Genetic Risk of Age-Related Macular Degeneration With Dietary Antioxidants, Zinc, and ï‰-3 Fatty Acids. JAMA Ophthalmology, 2011, 129, 758.	2.6	177
550	Common genetic variants associated with open-angle glaucoma. Human Molecular Genetics, 2011, 20, 2464-2471.	1.4	152
551	Genetic variation in the reninâ€”angiotensin system, use of reninâ€”angiotensin system inhibitors and the risk of myocardial infarction. JRAAS - Journal of the Renin-Angiotensin-Aldosterone System, 2011, 12, 208-214.	1.0	8
552	Genetic architecture of open angle glaucoma and related determinants. Journal of Medical Genetics, 2011, 48, 190-196.	1.5	21
553	Genome-wide association studies in Asians confirm the involvement of ATOH7 and TGFBR3, and further identify CARD10 as a novel locus influencing optic disc area. Human Molecular Genetics, 2011, 20, 1864-1872.	1.4	91
554	Large common deletions associate with mortality at old age. Human Molecular Genetics, 2011, 20, 4290-4296.	1.4	35
555	Genome-Based Prediction of Breast Cancer Risk in the General Population: A Modeling Study Based on Meta-Analyses of Genetic Associations. Cancer Epidemiology Biomarkers and Prevention, 2011, 20, 9-22.	1.1	29
556	Genome-wide association and genetic functional studies identify autism susceptibility candidate 2 gene (AUTS2) in the regulation of alcohol consumption. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 7119-7124.	3.3	258
557	Genome-wide association study identifies six new loci influencing pulse pressure and mean arterial pressure. Nature Genetics, 2011, 43, 1005-1011.	9.4	403
558	Total Zinc Intake May Modify the Glucose-Raising Effect of a Zinc Transporter (SLC30A8) Variant: A 14-Cohort Meta-analysis. Diabetes, 2011, 60, 2407-2416.	0.3	91

#	Article	IF	Citations
559	Common variants near FRK/COL10A1 and VEGFA are associated with advanced age-related macular degeneration. Human Molecular Genetics, 2011, 20, 3699-3709.	1.4	232
560	Multiple Loci Are Associated with White Blood Cell Phenotypes. PLoS Genetics, 2011, 7, e1002113.	1.5	106
561	A Genome-Wide Screen for Interactions Reveals a New Locus on 4p15 Modifying the Effect of Waist-to-Hip Ratio on Total Cholesterol. PLoS Genetics, 2011, 7, e1002333.	1.5	29
562	Physical Activity Attenuates the Influence of FTO Variants on Obesity Risk: A Meta-Analysis of 218,166 Adults and 19,268 Children. PLoS Medicine, 2011, 8, e1001116.	3.9	446
563	Runs of Homozygosity Do Not Influence Survival to Old Age. PLoS ONE, 2011, 6, e22580.	1.1	8
564	Perspectives on the Use of Multiple Sclerosis Risk Genes for Prediction. PLoS ONE, 2011, 6, e26493.	1.1	17
565	Prospective study of insulin-like growth factor-I, insulin-like growth factor-binding protein 3, genetic variants in the IGF1 and IGFBP3 genes and risk of coronary artery disease. International Journal of Molecular Epidemiology and Genetics, 2011, 2, 261-85.	0.4	8
566	Genome-Wide Association Studies of MRI-Defined Brain Infarcts. Stroke, 2010, 41, 210-217.	1.0	82
567	Translational Research in Genomics of Alzheimer's Disease: A Review of Current Practice and Future Perspectives. Journal of Alzheimer's Disease, 2010, 20, 967-980.	1.2	16
568	The Role of Body Mass Index, Insulin, and Adiponectin in the Relation Between Fat Distribution and Bone Mineral Density. Calcified Tissue International, 2010, 86, 116-125.	1.5	68
569	HFE gene mutations increase the risk of coronary heart disease in women. European Journal of Epidemiology, 2010, 25, 643-649.	2.5	11
570	ProbABEL package for genome-wide association analysis of imputed data. BMC Bioinformatics, 2010, 11, 134.	1.2	354
571	Shared genetic factors in the co-occurrence of symptoms of depression and cardiovascular risk factors. Journal of Affective Disorders, 2010, 122, 247-252.	2.0	19
572	Interleukinâ€1 gene cluster variants with innate cytokine production profiles and osteoarthritis in subjects from the Genetics, Osteoarthritis and Progression Study. Arthritis and Rheumatism, 2010, 62, 1119-1126.	6.7	10
573	Ped_Outlier software for automatic identification of within-family outliers. Computational Biology and Chemistry, 2010, 34, 242-243.	1.1	0
574	The dopamine Î²-hydroxylase -1021C/T polymorphism is associated with the risk of Alzheimer's disease in the Epistasis Project. BMC Medical Genetics, 2010, 11, 162.	2.1	50
575	A meta-analysis of genome-wide data from five European isolates reveals an association of COL22A1, SYT1, and GABRR2with serum creatinine level. BMC Medical Genetics, 2010, 11, 41.	2.1	48
576	An epidemiological perspective on the future of direct-to-consumer personal genome testing. Investigative Genetics, 2010, 1, 10.	3.3	38

#	Article	IF	Citations
577	Comparison of participant information and informed consent forms of five European studies in genetic isolated populations. European Journal of Human Genetics, 2010, 18, 296-302.	1.4	31
578	Biological, clinical and population relevance of 95 loci for blood lipids. Nature, 2010, 466, 707-713.	13.7	3,249
579	Hundreds of variants clustered in genomic loci and biological pathways affect human height. Nature, 2010, 467, 832-838.	13.7	1,789
580	Meta-analyses of genome-wide association studies identify multiple loci associated with pulmonary function. Nature Genetics, 2010, 42, 45-52.	9.4	549
581	Genome-wide association study of PR interval. Nature Genetics, 2010, 42, 153-159.	9.4	400
582	Genome-wide association study of intracranial aneurysm identifies three new risk loci. Nature Genetics, 2010, 42, 420-425.	9.4	262
583	Variants in ADCY5 and near CCNL1 are associated with fetal growth and birth weight. Nature Genetics, 2010, 42, 430-435.	9.4	223
584	New loci associated with kidney function and chronic kidney disease. Nature Genetics, 2010, 42, 376-384.	9.4	710
585	Sequence variants at CHRNB3â€“CHRNA6 and CYP2A6 affect smoking behavior. Nature Genetics, 2010, 42, 448-453.	9.4	649
586	Twelve type 2 diabetes susceptibility loci identified through large-scale association analysis. Nature Genetics, 2010, 42, 579-589.	9.4	1,631
587	Genome-wide association study of migraine implicates a common susceptibility variant on 8q22.1. Nature Genetics, 2010, 42, 869-873.	9.4	332
588	A genome-wide association study identifies a susceptibility locus for refractive errors and myopia at 15q14. Nature Genetics, 2010, 42, 897-901.	9.4	200
589	A genome-wide association study for myopia and refractive error identifies a susceptibility locus at 15q25. Nature Genetics, 2010, 42, 902-905.	9.4	204
590	Meta-analysis identifies 13 new loci associated with waist-hip ratio and reveals sexual dimorphism in the genetic basis of fat distribution. Nature Genetics, 2010, 42, 949-960.	9.4	836
591	Association analyses of 249,796 individuals reveal 18 new loci associated with body mass index. Nature Genetics, 2010, 42, 937-948.	9.4	2,634
592	Thirty new loci for age at menarche identified by a meta-analysis of genome-wide association studies. Nature Genetics, 2010, 42, 1077-1085.	9.4	445
593	Common variants in 22 loci are associated with QRS duration and cardiac ventricular conduction. Nature Genetics, 2010, 42, 1068-1076.	9.4	308
594	Reply to â€œLack of support for association between the KIF1B rs10492972[C] variant and multiple sclerosisâ€• Nature Genetics, 2010, 42, 470-471.	9.4	9

#	Article	IF	Citations
595	Association between Type 2 Diabetes Loci and Measures of Fatness. PLoS ONE, 2010, 5, e8541.	1.1	17
596	The ERCC6 Gene and Age-Related Macular Degeneration. PLoS ONE, 2010, 5, e13786.	1.1	26
597	Early Age at Menopause is Associated with Increased risk of Dementia and Mortality in Women with Down Syndrome. Journal of Alzheimer's Disease, 2010, 19, 545-550.	1.2	56
598	Genetic Variation at the Phospholipid Transfer Protein Locus Affects Its Activity and High-Density Lipoprotein Size and Is a Novel Marker of Cardiovascular Disease Susceptibility. Circulation, 2010, 122, 470-477.	1.6	86
599	European lactase persistence genotype shows evidence of association with increase in body mass index. Human Molecular Genetics, 2010, 19, 1129-1136.	1.4	58
600	Novel Associations of Multiple Genetic Loci With Plasma Levels of Factor VII, Factor VIII, and von Willebrand Factor. Circulation, 2010, 121, 1382-1392.	1.6	311
601	Genome-wide Analysis of Genetic Loci Associated With Alzheimer Disease. JAMA - Journal of the American Medical Association, 2010, 303, 1832.	3.8	1,064
602	Association of Genome-Wide Variation With the Risk of Incident Heart Failure in Adults of European and African Ancestry. Circulation: Cardiovascular Genetics, 2010, 3, 256-266.	5.1	176
603	Separating the Mechanism-Based and Off-Target Actions of Cholesteryl Ester Transfer Protein Inhibitors With CETP Gene Polymorphisms. Circulation, 2010, 121, 52-62.	1.6	96
604	Improvement of Risk Prediction by Genomic Profiling: Reclassification Measures Versus the Area Under the Receiver Operating Characteristic Curve. American Journal of Epidemiology, 2010, 172, 353-361.	1.6	61
605	The PCLO gene and depressive disorders: replication in a population-based study. Human Molecular Genetics, 2010, 19, 731-734.	1.4	43
606	IGF-I Bioactivity in an Elderly Population: Relation to Insulin Sensitivity, Insulin Levels, and the Metabolic Syndrome. Diabetes, 2010, 59, 505-508.	0.3	35
607	Genetic Architecture of Plasma Adiponectin Overlaps With the Genetics of Metabolic Syndromeâ€"Related Traits. Diabetes Care, 2010, 33, 908-913.	4.3	68
608	Interactions of Dietary Whole-Grain Intake With Fasting Glucose- and Insulin-Related Genetic Loci in Individuals of European Descent: A meta-analysis of 14 cohort studies. Diabetes Care, 2010, 33, 2684-2691.	4.3	127
609	Interactions between dietary vitamin E intake and SIRT1 genetic variation influence body mass index. American Journal of Clinical Nutrition, 2010, 91, 1387-1393.	2.2	24
610	Genetic Variants Influencing Circulating Lipid Levels and Risk of Coronary Artery Disease. Arteriosclerosis, Thrombosis, and Vascular Biology, 2010, 30, 2264-2276.	1.1	369
611	Insulin VNTR and IGF-1 Promoter Region Polymorphisms Are Not Associated with Body Composition in Early Childhood: The Generation R Study. Hormone Research in Paediatrics, 2010, 73, 120-127.	0.8	10
612	Genomic Variation Associated With Mortality Among Adults of European and African Ancestry With Heart Failure. Circulation: Cardiovascular Genetics, 2010, 3, 248-255.	5.1	80

#	Article	IF	Citations
613	Common Genetic Variants Associate with Serum Phosphorus Concentration. Journal of the American Society of Nephrology: JASN, 2010, 21, 1223-1232.	3.0	123
614	Modeling of Environmental Effects in Genome-Wide Association Studies Identifies SLC2A2 and HP as Novel Loci Influencing Serum Cholesterol Levels. PLoS Genetics, 2010, 6, e1000798.	1.5	51
615	Genome-Wide Association Studies of Serum Magnesium, Potassium, and Sodium Concentrations Identify Six Loci Influencing Serum Magnesium Levels. PLoS Genetics, 2010, 6, e1001045.	1.5	185
616	Four Novel Loci (19q13, 6q24, 12q24, and 5q14) Influence the Microcirculation In Vivo. PLoS Genetics, 2010, 6, e1001184.	1.5	134
617	A Genome-Wide Association Study of Optic Disc Parameters. PLoS Genetics, 2010, 6, e1000978.	1.5	187
618	Use of Genome-Wide Expression Data to Mine the â€œGray Zoneâ€•of GWA Studies Leads to Novel Candidate Obesity Genes. PLoS Genetics, 2010, 6, e1000976.	1.5	62
619	Multiple Genetic Loci Influence Serum Urate Levels and Their Relationship With Gout and Cardiovascular Disease Risk Factors. Circulation: Cardiovascular Genetics, 2010, 3, 523-530.	5.1	285
620	A Genome-Wide Screen for Depression in Two Independent Dutch Populations. Biological Psychiatry, 2010, 68, 187-196.	0.7	27
621	A Meta-analysis of Four Genome-Wide Association Studies of Survival to Age 90 Years or Older: The Cohorts for Heart and Aging Research in Genomic Epidemiology Consortium. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2010, 65A, 478-487.	1.7	117
622	Genome-wide association analysis identifies multiple loci related to resting heart rate. Human Molecular Genetics, 2010, 19, 3885-3894.	1.4	133
623	No evidence for prion protein gene locus multiplication in Creutzfeldt-Jakob disease. Neuroscience Letters, 2010, 472, 16-18.	1.0	5
624	The apolipoprotein E gene and its age-specific effects on cognitive function. Neurobiology of Aging, 2010, 31, 1831-1833.	1.5	60
625	Clear detection of ADIPOQ locus as the major gene for plasma adiponectin: Results of genome-wide association analyses including 4659 European individuals. Atherosclerosis, 2010, 208, 412-420.	0.4	146
626	Linkage and Genomeâ€wide Association Analysis of Obesityâ€related Phenotypes: Association of Weight With the MGAT1 Gene. Obesity, 2010, 18, 803-808.	1.5	54
627	Infectious mononucleosis-linked HLA class I single nucleotide polymorphism is associated with multiple sclerosis. Multiple Sclerosis Journal, 2010, 16, 1303-1307.	1.4	4
628	Genome-Wide Association Scan Meta-Analysis Identifies Three Loci Influencing Adiposity and Fat Distribution. PLoS Genetics, 2009, 5, e1000508.	1.5	453
629	Genetic evidence for a role of adiponutrin in the metabolism of apolipoprotein B-containing lipoproteins. Human Molecular Genetics, 2009, 18, 4669-4676.	1.4	49
630	Breast-Feeding Modifies the Association of PPARÎ³2 Polymorphism Pro12Ala With Growth in Early Life. Diabetes, 2009, 58, 992-998.	0.3	19

#	Article	IF	Citations
631	Genomewide Association Studies of Stroke. New England Journal of Medicine, 2009, 360, 1718-1728.	13.9	420
632	Genetic Variants Associated With Cardiac Structure and Function. JAMA - Journal of the American Medical Association, 2009, 302, 168.	3.8	202
633	Evaluation of risk prediction updates from commercial genome-wide scans. Genetics in Medicine, 2009, 11, 588-594.	1.1	69
634	An empirical comparison of meta-analyses of published gene-disease associations versus consortium analyses. Genetics in Medicine, 2009, 11, 153-162.	1.1	11
635	NRXN3 Is a Novel Locus for Waist Circumference: A Genome-Wide Association Study from the CHARGE Consortium. PLoS Genetics, 2009, 5, e1000539.	1.5	230
636	A Genome-Wide Association Study Reveals Variants in ARL15 that Influence Adiponectin Levels. PLoS Genetics, 2009, 5, e1000768.	1.5	148
637	Genetic Determinants of Circulating Sphingolipid Concentrations in European Populations. PLoS Genetics, 2009, 5, e1000672.	1.5	184
638	Association of Cognitive Functioning with Retinal Nerve Fiber Layer Thickness. , 2009, 50, 4576.		44
639	SIRT1 Genetic Variation Is Related to BMI and Risk of Obesity. Diabetes, 2009, 58, 2828-2834.	0.3	118
640	Association of Novel Genetic Loci With Circulating Fibrinogen Levels. Circulation: Cardiovascular Genetics, 2009, 2, 125-133.	5.1	86
641	Common variants in the JAZF1 gene associated with height identified by linkage and genome-wide association analysis. Human Molecular Genetics, 2009, 18, 373-380.	1.4	88
642	Genome-wide association meta-analysis for total serum bilirubin levels. Human Molecular Genetics, 2009, 18, 2700-2710.	1.4	214
643	A Genome-Wide Association Scan of RR and QT Interval Duration in 3 European Genetically Isolated Populations. Circulation: Cardiovascular Genetics, 2009, 2, 322-328.	5.1	67
644	SIRT1 genetic variation and mortality in type 2 diabetes: interaction with smoking and dietary niacin. Free Radical Biology and Medicine, 2009, 46, 836-841.	1.3	44
645	Cerebrospinal fluid biomarkers in human genetic transmissible spongiform encephalopathies. Journal of Neurology, 2009, 256, 1620-1628.	1.8	77
646	The Rotterdam Study: 2010 objectives and design update. European Journal of Epidemiology, 2009, 24, 553-572.	2.5	235
647	Genetic Scoring Analysis: a way forward in Genome Wide Association Studies?. European Journal of Epidemiology, 2009, 24, 585-587.	2.5	22
648	Type 2 diabetes gene TCF7L2polymorphism is not associated with fetal and postnatal growth in two birth cohort studies. BMC Medical Genetics, 2009, 10, 67.	2.1	15

#	Article	IF	Citations
649	Variation in the IGFâ€1 gene is associated with lymphocyte subset counts in neonates: The Generation R Study. Clinical Endocrinology, 2009, 70, 53-59.	1.2	3
650	The expression of type III hyperlipoproteinemia: involvement of lipolysis genes. European Journal of Human Genetics, 2009, 17, 620-628.	1.4	53
651	Suggestive linkage of ADHD to chromosome 18q22 in a young genetically isolated Dutch population. European Journal of Human Genetics, 2009, 17, 958-966.	1.4	17
652	Predicting human height by Victorian and genomic methods. European Journal of Human Genetics, 2009, 17, 1070-1075.	1.4	108
653	Loci influencing lipid levels and coronary heart disease risk in 16 European population cohorts. Nature Genetics, 2009, 41, 47-55.	9.4	776
654	Six new loci associated with body mass index highlight a neuronal influence on body weight regulation. Nature Genetics, 2009, 41, 25-34.	9.4	1,572
655	Variants in MTNR1B influence fasting glucose levels. Nature Genetics, 2009, 41, 77-81.	9.4	662
656	Multiple loci associated with indices of renal function and chronic kidney disease. Nature Genetics, 2009, 41, 712-717.	9.4	553
657	Genome-wide association study of blood pressure and hypertension. Nature Genetics, 2009, 41, 677-687.	9.4	1,224
658	Variants in ZFHX3 are associated with atrial fibrillation in individuals of European ancestry. Nature Genetics, 2009, 41, 879-881.	9.4	363
659	Twenty bone-mineral-density loci identified by large-scale meta-analysis of genome-wide association studies. Nature Genetics, 2009, 41, 1199-1206.	9.4	660
660	Multiple loci influence erythrocyte phenotypes in the CHARGE Consortium. Nature Genetics, 2009, 41, 1191-1198.	9.4	324
661	Complement Component C3 and Risk of Age-Related Macular Degeneration. Ophthalmology, 2009, 116, 474-480.e2.	2.5	89
662	A caseâ€“control study on the effect of p53 and p73 gene polymorphisms on gastric cancer risk and progression. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2009, 675, 60-65.	0.9	41
663	A genome-wide association study of northwestern Europeans involves the C-type natriuretic peptide signaling pathway in the etiology of human height variation. Human Molecular Genetics, 2009, 18, 3516-3524.	1.4	76
664	The GAB2 Gene and the Risk of Alzheimer's Disease: Replication and Meta-Analysis. Biological Psychiatry, 2009, 65, 995-999.	0.7	39
665	Replication by the Epistasis Project of the interaction between the genes for IL-6 and IL-10 in the risk of Alzheimer's disease. Journal of Neuroinflammation, 2009, 6, 22.	3.1	46
666	Genome-wide linkage analysis of serum creatinine in three isolated European populations. Kidney International, 2009, 76, 297-306.	2.6	71

#	Article	IF	Citations
667	Collaborative Meta-analysis: Associations of 150 Candidate Genes With Osteoporosis and Osteoporotic Fracture. Annals of Internal Medicine, 2009, 151, 528.	2.0	250
668	Genome-based prediction of common diseases: methodological considerations for future research. Genome Medicine, 2009, 1, 20.	3.6	133
669	Genetic Variation in the Renin-Angiotensin System and Arterial Stiffness. The Rotterdam Study. Clinical and Experimental Hypertension, 2009, 31, 389-399.	0.5	13
670	Replication of CD58 and CLEC16A as genome-wide significant risk genes for multiple sclerosis. Journal of Human Genetics, 2009, 54, 676-680.	1.1	65
671	Interaction between the Gly460Trp Î±-adducin gene variant and diuretics on the risk of myocardial infarction. Journal of Hypertension, 2009, 27, 61-68.	0.3	29
672	The effect of catechol-O-methyltransferase Met/Val functional polymorphism on smoking cessation: retrospective and prospective analyses in a cohort study. Pharmacogenetics and Genomics, 2009, 19, 45-51.	0.7	27
673	A Study of the SORL1 Gene in Alzheimer's Disease and Cognitive Function. Journal of Alzheimer's Disease, 2009, 18, 51-64.	1.2	36
674	Role of shared genetic and environmental factors in symptoms of depression and body composition. Psychiatric Genetics, 2009, 19, 32-38.	0.6	16
675	Genetic Factors Influence the Clustering of Depression among Individuals with Lower Socioeconomic Status. PLoS ONE, 2009, 4, e5069.	1.1	11
676	Apolipoprotein E gene is related to mortality only in normal weight individuals: The Rotterdam study. European Journal of Epidemiology, 2008, 23, 135-142.	2.5	10
677	The Generation R Study: design and cohort update until the age of 4Âyears. European Journal of Epidemiology, 2008, 23, 801-811.	2.5	154
678	Reninâ€angiotensin system inhibitors, angiotensin Iâ€converting enzyme gene insertion/deletion polymorphism, and cancer. Cancer, 2008, 112, 748-757.	2.0	117
679	Three Genome-wide Association Studies and a Linkage Analysis Identify HERC2 as a Human Iris Color Gene. American Journal of Human Genetics, 2008, 82, 411-423.	2.6	220
680	A Critical Appraisal of the Scientific Basis of Commercial Genomic Profiles Used to Assess Health Risks and Personalize Health Interventions. American Journal of Human Genetics, 2008, 82, 593-599.	2.6	258
681	Reply to Stephan etÂal.. American Journal of Human Genetics, 2008, 83, 131.	2.6	2
682	Familial aggregation of preeclampsia and intrauterine growth restriction in a genetically isolated population in The Netherlands. European Journal of Human Genetics, 2008, 16, 1437-1442.	1.4	8
683	An approach for cutting large and complex pedigrees for linkage analysis. European Journal of Human Genetics, 2008, 16, 854-860.	1.4	55
684	Susceptibility loci for intracranial aneurysm in European and Japanese populations. Nature Genetics, 2008, 40, 1472-1477.	9.4	247

#	Article	IF	Citations
685	Genetic variation in the KIF1B locus influences susceptibility to multiple sclerosis. Nature Genetics, 2008, 40, 1402-1403.	9.4	173
686	Survival in Elderly Persons with Down Syndrome. Journal of the American Geriatrics Society, 2008, 56, 2311-2316.	1.3	103
687	Variation in the IGF1 gene and growth in foetal life and infancy. The Generation R Study. Clinical Endocrinology, 2008, 68, 382-389.	1.2	17
688	Lack of association of two common polymorphisms on 9p21 with risk of coronary heart disease and myocardial infarction; results from a prospective cohort study. BMC Medicine, 2008, 6, 30.	2.3	26
689	Genome-based prediction of common diseases: advances and prospects. Human Molecular Genetics, 2008, 17, R166-R173.	1.4	287
690	Polymorphisms in the Vascular Endothelial Growth Factor Gene and Risk of Age-related Macular Degeneration. Ophthalmology, 2008, 115, 1899-1903.	2.5	51
691	Association of three genetic loci with uric acid concentration and risk of gout: a genome-wide association study. Lancet, The, 2008, 372, 1953-1961.	6.3	610
692	Predicting Type 2 Diabetes Based on Polymorphisms From Genome-Wide Association Studies. Diabetes, 2008, 57, 3122-3128.	0.3	265
693	A Systematic Review of Meta-Analyses on Gene Polymorphisms and Gastric Cancer Risk. Current Genomics, 2008, 9, 361-374.	0.7	50
694	Shared Constitutional Risks for Maternal Vascular-Related Pregnancy Complications and Future Cardiovascular Disease. Hypertension, 2008, 51, 1034-1041.	1.3	203
695	Identification of a common variant at the NOS1AP locus strongly associated to QT-interval duration. Human Molecular Genetics, 2008, 18, 347-357.	1.4	54
696	A variant of the IGF-I gene is associated with blood pressure but not with left heart dimensions at the age of 2 years: the Generation R Study. European Journal of Endocrinology, 2008, 159, 209-216.	1.9	6
697	Maternal Transmission of Multiple Sclerosis in a Dutch Population. Archives of Neurology, 2008, 65, 345-8.	4.9	58
698	Cataract Surgery and the Risk of Aging Macula Disorder: The Rotterdam Study. , 2008, 49, 4795.		60
699	Angiotensinogen M235T polymorphism and symptoms of depression in a population-based study and a family-based study. Psychiatric Genetics, 2008, 18, 162-166.	0.6	13
700	The interleukin-6 –174 G/C promoter polymorphism and arterial stiffness; the Rotterdam Study. Vascular Health and Risk Management, 2008, Volume 4, 863-869.	1.0	22
701	LPIN2 Is Associated With Type 2 Diabetes, Glucose Metabolism, and Body Composition. Diabetes, 2007, 56, 3020-3026.	0.3	52
702	Genetic Contributions to Glaucoma: Heritability of Intraocular Pressure, Retinal Nerve Fiber Layer Thickness, and Optic Disc Morphology. , 2007, 48, 3669.		110

#	Article	IF	Citations
703	Effects of the Renin-Angiotensin System Genes and Salt Sensitivity Genes on Blood Pressure and Atherosclerosis in the Total Population and Patients With Type 2 Diabetes. Diabetes, 2007, 56, 1905-1912.	0.3	13
704	Meta- and Pooled Analyses of the Methylenetetrahydrofolate Reductase C677T and A1298C Polymorphisms and Gastric Cancer Risk: A Huge-GSEC Review. American Journal of Epidemiology, 2007, 167, 505-516.	1.6	103
705	Insulin gene variable number of tandem repeats is not associated with weight from fetal life until infancy: the Generation R Study. European Journal of Endocrinology, 2007, 157, 741-748.	1.9	12
706	The Association Between H63D Mutations in HFE and Amyotrophic Lateral Sclerosis in a Dutch Population. Archives of Neurology, 2007, 64, 63.	4.9	69
707	The risk of myocardial infarction in patients with reduced activity of cytochrome P450 2C9. Pharmacogenetics and Genomics, 2007, 17, 473-479.	0.7	21
708	Heritability of blood pressure traits and the genetic contribution to blood pressure variance explained by four blood-pressure-related genes. Journal of Hypertension, 2007, 25, 565-570.	0.3	82
709	Plasma Î² amyloid and impaired CO2-induced cerebral vasomotor reactivity. Neurobiology of Aging, 2007, 28, 707-712.	1.5	12
710	A Genomewide Screen for Late-Onset Alzheimer Disease in a Genetically Isolated Dutch Population. American Journal of Human Genetics, 2007, 81, 17-31.	2.6	145
711	The impact of genotype frequencies on the clinical validity of genomic profiling for predicting common chronic diseases. Genetics in Medicine, 2007, 9, 528-535.	1.1	128
712	GenABEL: an R library for genome-wide association analysis. Bioinformatics, 2007, 23, 1294-1296.	1.8	1,711
713	Methylenetetrahydrofolate reductase C677T and A1298C polymorphisms and susceptibility to gastric adenocarcinoma in an Italian population. Biomarkers, 2007, 12, 635-644.	0.9	36
714	Estrogen Receptor Î± Gene Polymorphisms Associated with Incident Aging Macula Disorder. , 2007, 48, 1012.		27
715	A critical appraisal of epidemiological studies comes from basic knowledge: a reader's guide to assess potential for biases. World Journal of Emergency Surgery, 2007, 2, 7.	2.1	8
716	Angiotensinogen M235T polymorphism and the risk of myocardial infarction and stroke among hypertensive patients on ACE-inhibitors or Î²-blockers. European Journal of Human Genetics, 2007, 15, 478-484.	1.4	50
717	Mutation analysis of candidate genes within the 2q33.3 linkage area for familial early-onset generalised osteoarthritis. European Journal of Human Genetics, 2007, 15, 791-799.	1.4	22
718	Polymorphisms in metabolic genes, their combination and interaction with tobacco smoke and alcohol consumption and risk of gastric cancer: a case-control study in an Italian population. BMC Cancer, 2007, 7, 206.	1.1	85
719	Heritabilities, apolipoprotein E, and effects of inbreeding on plasma lipids in a genetically isolated population: The Erasmus Rucphen Family Study. European Journal of Epidemiology, 2007, 22, 99-105.	2.5	25
720	The Rotterdam Study: objectives and design update. European Journal of Epidemiology, 2007, 22, 819-829.	2.5	644

#	Article	IF	Citations
721	CYP1A1, CYP2E1, GSTM1, GSTT1, EPHX1 exons 3 and 4, and NAT2 polymorphisms, smoking, consumption of alcohol and fruit and vegetables and risk of head and neck cancer. Journal of Cancer Research and Clinical Oncology, 2007, 134, 93-100.	1.2	75
722	The first case of variant Creutzfeldt-Jakob disease in the Netherlands. Journal of Neurology, 2007, 254, 958-960.	1.8	7
723	A Genomic Background Based Method for Association Analysis in Related Individuals. PLoS ONE, 2007, 2, e1274.	1.1	233
724	Predictive testing for complex diseases using multiple genes: Fact or fiction?. Genetics in Medicine, 2006, 8, 395-400.	1.1	202
725	Promoter Mutations That Increase Amyloid Precursor-Protein Expression Are Associated with Alzheimer Disease. American Journal of Human Genetics, 2006, 78, 936-946.	2.6	173
726	A Common Polymorphism in the Complement Factor H Gene Is Associated With Increased Risk of Myocardial Infarction. Journal of the American College of Cardiology, 2006, 47, 1568-1575.	1.2	83
727	The UBQLN1 polymorphism, UBQ-8i, at 9q22 is not associated with Alzheimer's disease with onset before 70 years. Neuroscience Letters, 2006, 392, 72-74.	1.0	30
728	Î±-Adducin Polymorphism, Atherosclerosis, and Cardiovascular and Cerebrovascular Risk. Stroke, 2006, 37, 2930-2934.	1.0	45
729	No association between the angiotensin-converting enzyme gene and major depression: a caseâ€“control study and meta-analysis. Psychiatric Genetics, 2006, 16, 225-226.	0.6	13
730	The influence of the alpha-adducin G460W polymorphism and angiotensinogen M235T polymorphism on antihypertensive medication and blood pressure. European Journal of Human Genetics, 2006, 14, 860-866.	1.4	30
731	Strong linkage on 2q33.3 to familial early-onset generalized osteoarthritis and a consideration of two positional candidate genes. European Journal of Human Genetics, 2006, 14, 1280-1287.	1.4	26
732	Estrogen Receptor Î² (ESR2) Polymorphisms in Interaction With Estrogen Receptor Î± (ESR1) and Insulin-Like Growth Factor I (IGF1) Variants Influence the Risk of Fracture in Postmenopausal Women. Journal of Bone and Mineral Research, 2006, 21, 1443-1456.	3.1	73
733	Evidence for novel loci for late-onset Parkinsonâ€™s disease in a genetic isolate from the Netherlands. Human Genetics, 2006, 119, 51-60.	1.8	4
734	An Insulin-Like Growth Factor-I Promoter Polymorphism Is Associated With Increased Mortality in Subjects With Myocardial Infarction in an Elderly Caucasian Population. American Journal of Cardiology, 2006, 97, 1274-1276.	0.7	18
735	Epidemiology of neurological diseases in elderly people: what did we learn from the Rotterdam Study?. Lancet Neurology, The, 2006, 5, 545-550.	4.9	24
736	TGF-Î²1 Polymorphisms and Risk of Myocardial Infarction and Stroke. Stroke, 2006, 37, 2667-2671.	1.0	60
737	Complement Factor H Polymorphism, Complement Activators, and Risk of Age-Related Macular Degeneration. JAMA - Journal of the American Medical Association, 2006, 296, 301.	3.8	306
738	Drug-Gene Interaction Between the Insertion/Deletion Polymorphism of the Angiotensin-Converting Enzyme Gene and Antihypertensive Therapy. Annals of Pharmacotherapy, 2006, 40, 212-218.	0.9	12

#	Article	IF	Citations
739	Retinal Vessel Diameters and Risk of Impaired Fasting Glucose or Diabetes: The Rotterdam Study. Diabetes, 2006, 55, 506-510.	0.3	114
740	An IGF-I Gene Polymorphism Modifies the Risk of Diabetic Retinopathy. Diabetes, 2006, 55, 2387-2391.	0.3	24
741	Interleukin 6 â^'174 G/C Promoter Polymorphism and Risk of Coronary Heart Disease. Arteriosclerosis, Thrombosis, and Vascular Biology, 2006, 26, 212-217.	1.1	94
742	IGF-I gene promoter polymorphism is a predictor of survival after myocardial infarction in patients with type 2 diabetes. European Journal of Endocrinology, 2006, 155, 751-756.	1.9	13
743	Mortality in patients with hypertension on angiotensin-I converting enzyme (ACE)-inhibitor treatment is influenced by the ACE insertion/deletion polymorphism. Pharmacogenetics and Genomics, 2005, 15, 75-81.	0.7	31
744	Patients with an ApoE ϵ4 allele require lower doses of coumarin anticoagulants. Pharmacogenetics and Genomics, 2005, 15, 69-74.	0.7	52
745	Allelic variants of cytochrome P450 2C9 modify the interaction between nonsteroidal anti-inflammatory drugs and coumarin anticoagulants. Clinical Pharmacology and Therapeutics, 2005, 77, 479-485.	2.3	46
746	In search of genes involved in neurodegenerative disorders. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2005, 592, 89-101.	0.4	25
747	Novel parkin mutations detected in patients with early-onset Parkinson's disease. Movement Disorders, 2005, 20, 424-431.	2.2	60
748	Phenotypic Subtypes in Attention Deficit Hyperactivity Disorder in an Isolated Population. European Journal of Epidemiology, 2005, 20, 789-794.	2.5	4
749	The Effect of Genetic Drift in a Young Genetically Isolated Population. Annals of Human Genetics, 2005, 69, 288-295.	0.3	126
750	Large Meta-Analysis Establishes the ACE Insertion-Deletion Polymorphism as a Marker of Alzheimer's Disease. American Journal of Epidemiology, 2005, 162, 305-317.	1.6	190
751	The association between angiotensin-converting enzyme gene polymorphism and coronary calcification. Atherosclerosis, 2005, 182, 169-173.	0.4	22
752	ACE gene is associated with Alzheimer's disease and atrophy of hippocampus and amygdala. Neurobiology of Aging, 2005, 26, 1153-1159.	1.5	61
753	Heterogeneity of the genetic risk in age-related macular disease. Ophthalmology, 2005, 112, 482-487.	2.5	26
754	Linkage and Association Studies Identify a Novel Locus for Alzheimer Disease at 7q36 in a Dutch Population-Based Sample. American Journal of Human Genetics, 2005, 77, 643-652.	2.6	48
755	The dopamine D4 receptor gene 48-base-pair-repeat polymorphism and mood disorders: A meta-analysis. Biological Psychiatry, 2005, 57, 999-1003.	0.7	155
756	The effect of genetic drift in a young genetically isolated population. Annals of Human Genetics, 2005, 69, 288-95.	0.3	92

#	Article	IF	Citations
757	Alcohol intake in relation to brain magnetic resonance imaging findings in older persons without dementia. American Journal of Clinical Nutrition, 2004, 80, 992-997.	2.2	86
758	Interaction Between Hypertension, apoE, and Cerebral White Matter Lesions. Stroke, 2004, 35, 1057-1060.	1.0	167
759	Genetic Factors and Insulin Secretion: Gene Variants in the IGF Genes. Diabetes, 2004, 53, S26-S30.	0.3	68
760	The â^'514 Câ†'T Hepatic Lipase Promoter Region Polymorphism and Plasma Lipids: A Meta-Analysis. Journal of Clinical Endocrinology and Metabolism, 2004, 89, 3858-3863.	1.8	89
761	Genetic Testing Has No Place as a Routine Diagnostic Test in Sporadic and Familial Cases of Alzheimer's Disease. Journal of the American Geriatrics Society, 2004, 52, 2110-2113.	1.3	16
762	Polymorphisms in the prion protein gene and in the doppel gene increase susceptibility for Creutzfeldtâ€"Jakob disease. European Journal of Human Genetics, 2004, 12, 389-394.	1.4	40
763	Linkage disequilibrium in young genetically isolated Dutch population. European Journal of Human Genetics, 2004, 12, 527-534.	1.4	107
764	Insulin-like growth factor-I gene polymorphism and risk of heart failure (the Rotterdam Study). American Journal of Cardiology, 2004, 94, 384-386.	0.7	37
765	The Influence of an Insulin-Like Growth Factor I Gene Promoter Polymorphism on Hip Bone Geometry and the Risk of Nonvertebral Fracture in the Elderly: The Rotterdam Study. Journal of Bone and Mineral Research, 2004, 19, 1280-1290.	3.1	64
766	ApoE Gene Polymorphisms, BMD, and Fracture Risk in Elderly Men and Women: The Rotterdam Study. Journal of Bone and Mineral Research, 2004, 19, 1490-1496.	3.1	31
767	Attention-deficit/hyperactivity disorder (ADHD): parentsâ€™ judgment about school, teachersâ€™ judgment about home. European Child and Adolescent Psychiatry, 2004, 13, 315-320.	2.8	48
768	Association of the interleukin-1 gene cluster with radiographic signs of osteoarthritis of the hip. Arthritis and Rheumatism, 2004, 50, 1179-1186.	6.7	98
769	Plasma amyloid ?, apolipoprotein E, lacunar infarcts, and white matter lesions. Annals of Neurology, 2004, 55, 570-575.	2.8	112
770	Drug-Gene Interactions between Genetic Polymorphisms and Antihypertensive Therapy. Drugs, 2004, 64, 1801-1816.	4.9	36
771	A deletion in DJ-1 and the risk of dementiaâ€"a population-based survey. Neuroscience Letters, 2004, 372, 196-199.	1.0	7
772	Apolipoprotein E Ïµ4 allele is associated with left ventricular systolic dysfunction. American Heart Journal, 2004, 147, 685-689.	1.2	9
773	Cholesterol and age-related macular degeneration: is there a link?. American Journal of Ophthalmology, 2004, 137, 750-752.	1.7	102
774	The risk of bleeding complications in patients with cytochrome P450 CYP2C9*2 or CYP2C9*3 alleles on acenocoumarol or phenprocoumon. Thrombosis and Haemostasis, 2004, 92, 61-66.	1.8	89

#	Article	IF	Citations
775	Revisiting the Clinical Validity of Multiplex Genetic Testing in Complex Diseases. American Journal of Human Genetics, 2004, 74, 585-588.	2.6	72
776	Angiotensin-converting enzyme gene polymorphism and common carotid stiffness. Atherosclerosis, 2004, 174, 121-126.	0.4	45
777	Effectiveness of HMG-CoA reductase inhibitors is modified by the ACE insertion deletion polymorphism. Atherosclerosis, 2004, 175, 377-379.	0.4	21
778	The risk of overanticoagulation in patients with cytochrome P450 CYP2C9*2 or CYP2C9*3 alleles on acenocoumarol or phenprocoumon. Pharmacogenetics and Genomics, 2004, 14, 27-33.	5.7	91
779	Smoking-dependent effects of the angiotensin-converting enzyme gene insertion/deletion polymorphism on blood pressure. Journal of Hypertension, 2004, 22, 313-319.	0.3	31
780	Is genetic screening for hemochromatosis worthwhile?. European Journal of Epidemiology, 2003, 19, 101-108.	2.5	21
781	Cdx-2 Polymorphism in the Promoter Region of the Human Vitamin D Receptor Gene Determines Susceptibility to Fracture in the Elderly. Journal of Bone and Mineral Research, 2003, 18, 1632-1641.	3.1	120
782	Suggestive linkage to chromosome 19 in a large Cuban family with late-onset Parkinson's disease. Movement Disorders, 2003, 18, 1240-1249.	2.2	26
783	PRNP Val129 homozygosity increases risk for early-onset Alzheimer's disease. Annals of Neurology, 2003, 53, 409-412.	2.8	103
784	Early cognitive decline is associated with prion protein codon 129 polymorphism. Annals of Neurology, 2003, 54, 275-276.	2.8	43
785	Estrogen receptor ? gene haplotype is associated with radiographic osteoarthritis of the knee in elderly men and women. Arthritis and Rheumatism, 2003, 48, 1913-1922.	6.7	125
786	A population-based study of the effect of the HFE C282Y and H63D mutations on iron metabolism. European Journal of Human Genetics, 2003, 11, 225-231.	1.4	60
787	Mutations in the DJ-1 Gene Associated with Autosomal Recessive Early-Onset Parkinsonism. Science, 2003, 299, 256-259.	6.0	2,467
788	Association of 5' estrogen receptor alpha gene polymorphisms with bone mineral density, vertebral bone area and fracture risk. Human Molecular Genetics, 2003, 12, 1745-1754.	1.4	170
789	A Genome-Wide Search for Genes Involved in Type 2 Diabetes in a Recently Genetically Isolated Population From the Netherlands. Diabetes, 2003, 52, 3001-3004.	0.3	26
790	A polymorphism in the IGF-I gene influences the age-related decline in circulating total IGF-I levels. European Journal of Endocrinology, 2003, 148, 171-175.	1.9	83
791	Frontotemporal dementia in The Netherlands: patient characteristics and prevalence estimates from a population-based study. Brain, 2003, 126, 2016-2022.	3.7	423
792	Angiotensin-Converting Enzyme Gene Polymorphism and Carotid Artery Wall Thickness. Stroke, 2003, 34, 1634-1639.	1.0	85

#	Article	IF	Citations
793	Association between an Insulin-Like Growth Factor I Gene Promoter Polymorphism and Bone Mineral Density in the Elderly: The Rotterdam Study. Journal of Clinical Endocrinology and Metabolism, 2003, 88, 3878-3884.	1.8	57
794	A Genome-Wide Search for Linkage-Disequilibrium With Type 1 Diabetes in a Recent Genetically Isolated Population From the Netherlands. Diabetes, 2002, 51, 856-859.	0.3	37
795	The Role of Hemochromatosis C282Y and H63D Gene Mutations in Type 2 Diabetes: Findings from the Rotterdam Study and meta-analysis. Diabetes Care, 2002, 25, 2112-2113.	4.3	13
796	The effectiveness of hydroxy-methylglutaryl coenzyme A reductase inhibitors (statins) in the elderly is not influenced by apolipoprotein E genotype. Pharmacogenetics and Genomics, 2002, 12, 647-653.	5.7	21
797	Dense-Core Senile Plaques in the Flemish Variant of Alzheimer's Disease Are Vasocentric. American Journal of Pathology, 2002, 161, 507-520.	1.9	108
798	The Hemochromatosis N144H Mutation of SLC11A3 Gene in Patients with Type 2 Diabetes. Molecular Genetics and Metabolism, 2002, 75, 290-291.	0.5	10
799	Dominant Hemochromatosis Due to N144H Mutation of SLC11A3: Clinical and Biological Characteristics. Blood Cells, Molecules, and Diseases, 2002, 29, 439-443.	0.6	39
800	A pooled case-control study of the apolipoprotein E (APOE) gene in age-related maculopathy. Ophthalmic Genetics, 2002, 23, 209-223.	0.5	136
801	Alcohol consumption and risk of dementia: the Rotterdam Study. Lancet, The, 2002, 359, 281-286.	6.3	499
802	Association between genetic variation in the gene for insulin-like growth factor-I and low birthweight. Lancet, The, 2002, 359, 1036-1037.	6.3	191
803	Insulin-like growth factor-I genotype and birthweight. Lancet, The, 2002, 360, 945-946.	6.3	12
804	The Gene Encoding Nicastrin, a Major Î³-Secretase Component, Modifies Risk for Familial Early-Onset Alzheimer Disease in a Dutch Population-Based Sample. American Journal of Human Genetics, 2002, 70, 1568-1574.	2.6	45
805	Expression of Type III Hyperlipoproteinemia in Apolipoprotein E2 (Arg158â†'Cys) Homozygotes Is Associated With Hyperinsulinemia. Arteriosclerosis, Thrombosis, and Vascular Biology, 2002, 22, 294-299.	1.1	55
806	Mutations in the Hemochromatosis Gene (HFE) and Stroke. Stroke, 2002, 33, 2363-2366.	1.0	33
807	Localization of autosomal recessive early-onset parkinsonism to chromosome 1p36 (PARK7) in an independent dataset. Annals of Neurology, 2002, 51, 253-256.	2.8	74
808	Myocilin mutations in a population-based sample of cases with open-angle glaucoma: the Rotterdam Study. Graefe's Archive for Clinical and Experimental Ophthalmology, 2002, 240, 468-474.	1.0	27
809	Commentary: Genetic screening for common mutations: Lessons from hereditary hemochromatosis. European Journal of Epidemiology, 2002, 18, 3-4.	2.5	1
810	Editorial: Good prospects for genetic and molecular epidemiologic studies in the European Journal of Epidemiology. European Journal of Epidemiology, 2002, 18, 285-286.	2.5	8

#	Article	IF	Citations
811	REVIEW: Prospects of genetic epidemiology in the 21st century. European Journal of Epidemiology, 2002, 18, 607-616.	2.5	12
812	TheTNFRSF6 gene is not implicated in familial early-onset Alzheimer's disease. Human Genetics, 2001, 108, 552-553.	1.8	6
813	Amyloid Î² secretase gene (BACE) is neither mutated in nor associated with early-onset Alzheimer's disease. Neuroscience Letters, 2001, 313, 105-107.	1.0	47
814	Influence of the prion protein and the apolipoprotein E genotype on the Creutzfeldtâ€“Jakob Disease phenotype. Neuroscience Letters, 2001, 313, 69-72.	1.0	44
815	Nonsteroidal Antiinflammatory Drugs and the Risk of Alzheimer's Disease. New England Journal of Medicine, 2001, 345, 1515-1521.	13.9	1,148
816	The first two patients with dura mater associated Creutzfeldt-Jakob disease in the Netherlands. Journal of Neurology, 2001, 248, 877-880.	1.8	9
817	Interaction Between the Vitamin D Receptor Gene and Collagen Type IÎ±1 Gene in Susceptibility for Fracture. Journal of Bone and Mineral Research, 2001, 16, 379-385.	3.1	111
818	A mutation in SLC11A3 is associated with autosomal dominant hemochromatosis. Nature Genetics, 2001, 28, 213-214.	9.4	458
819	Age-related maculopathy: Its genetic basis. Eye, 2001, 15, 396-400.	1.1	19
820	Opportunities for Population-Based Studies of Complex Genetic Disorders after the Human Genome Project. Epidemiology, 2001, 12, 360-364.	1.2	2
821	Point Mutation in the Stalk of Angiotensin-Converting Enzyme Causes a Dramatic Increase in Serum Angiotensin-Converting Enzyme But No Cardiovascular Disease. Circulation, 2001, 104, 1236-1240.	1.6	51
822	Adjacent genes, for COL2A1 and the vitamin D receptor, are associated with separate features of radiographic osteoarthritis of the knee. Arthritis and Rheumatism, 2000, 43, 1456-1464.	6.7	80
823	Power of selective genotyping in genetic association analyses of quantitative traits. Behavior Genetics, 2000, 30, 141-146.	1.4	85
824	Presentation of amyloidosis in carriers of the codon 692 mutation in the amyloid precursor protein gene (APP692). Brain, 2000, 123, 2130-2140.	3.7	51
825	Genetic Testing Should Not Be Advocated as a Diagnostic Tool in Familial Forms of Dementia. American Journal of Human Genetics, 2000, 67, 1033-1035.	2.6	15
826	Genetic association of the presenilin-1 regulatory region with early-onset Alzheimer's disease in a population-based sample. European Journal of Human Genetics, 1999, 7, 801-806.	1.4	49
827	Apolipoprotein E genotype and progression of Alzheimer's disease: the Rotterdam Study. Journal of Neurology, 1999, 246, 304-308.	1.8	50
828	Heritabilities of radiologic osteoarthritis in peripheral joints and of disc degeneration of the spine. Arthritis and Rheumatism, 1999, 42, 1729-1735.	6.7	127

#	Article	IF	Citations
829	Antiepileptic drug regimens and major congenital abnormalities in the offspring. Annals of Neurology, 1999, 46, 739-746.	2.8	276
830	The Glu318Gly Substitution in Presenilin 1 Is Not Causally Related to Alzheimer Disease. American Journal of Human Genetics, 1999, 64, 290-292.	2.6	47
831	High Prevalence of Mutations in the Microtubule-Associated Protein Tau in a Population Study of Frontotemporal Dementia in the Netherlands. American Journal of Human Genetics, 1999, 64, 414-421.	2.6	410
832	Mutation screening of the tau gene in patients with early-onset Alzheimer's disease. Neuroscience Letters, 1999, 277, 137-139.	1.0	43
833	Antiepileptic drug regimens and major congenital abnormalities in the offspring. Annals of Neurology, 1999, 46, 739-746.	2.8	144
834	A study of familial aggregation of depression, dementia and Parkinson's disease. European Journal of Epidemiology, 1998, 14, 233-238.	2.5	28
835	Serum apolipoprotein E level is not increased in Alzheimer's disease: the Rotterdam study. Neuroscience Letters, 1998, 248, 21-24.	1.0	58
836	The â^'491 A/T polymorphism in the regulatory region of the Apolipoprotein E gene and early-onset Alzheimer's disease. Neuroscience Letters, 1998, 258, 65-68.	1.0	38
837	Genetic Association of Apolipoprotein E with Age-Related Macular Degeneration. American Journal of Human Genetics, 1998, 63, 200-206.	2.6	425
838	Risk Estimates of Dementia by Apolipoprotein E Genotypes From a Population-Based Incidence Study: The Rotterdam Study. Archives of Neurology, 1998, 55, 964.	4.9	378
839	Role of APOE in Dementia:A Critical Reappraisal. Pathophysiology of Haemostasis and Thrombosis: International Journal on Haemostasis and Thrombosis Research, 1998, 28, 195-201.	0.5	11
840	APOE genotyping in differential diagnosis of Alzheimer's disease. Lancet, The, 1996, 348, 334.	6.3	26
841	Clinical Features and Mortality in Patients with Early-Onset Alzheimerâ€™s Disease. European Neurology, 1996, 36, 103-106.	0.6	53
842	Apolipoprotein E genotype and concomitant clinical features in early-onset Alzheimer's disease. Journal of Neurology, 1996, 243, 465-468.	1.8	8
843	A large-scale population-based study of the association of vitamin D receptor gene polymorphisms with bone mineral density. Journal of Bone and Mineral Research, 1996, 11, 1241-1248.	3.1	200
844	The apolipoprotein E ?2 allele is associated with an increased risk of early-onset alzheimer's disease and a reduced survival. Annals of Neurology, 1995, 37, 605-610.	2.8	129
845	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
846	Apolipoprotein E genotype and association between smoking and early onset Alzheimer's disease. BMJ: British Medical Journal, 1995, 310, 627-631.	2.4	74

#	Article	IF	Citations
847	The apolipoprotein E ?4 allele does not influence the clinical expression of the amyloid precursor protein gene codon 693 or 692 mutations. Annals of Neurology, 1994, 36, 434-437.	2.8	58
848	Apolipoprotein E4 allele in a populationâ€“based study of earlyâ€“onset Alzheimer's disease. Nature Genetics, 1994, 7, 74-78.	9.4	460
849	Head Trauma and the Risk of Alzheimer's Disease. American Journal of Epidemiology, 1992, 135, 775-782.	1.6	102
850	Epidemiology of Alzheimer's Disease. Epidemiologic Reviews, 1992, 14, 59-82.	1.3	261
851	Presenile dementia and cerebral haemorrhage linked to a mutation at codon 692 of the Î²â€“amyloid precursor protein gene. Nature Genetics, 1992, 1, 218-221.	9.4	715
852	Evidence for Allelic Heterogeneity in Familial Early-Onset Alzheimer's Disease. British Journal of Psychiatry, 1991, 158, 471-474.	1.7	28
853	Is Parental Age Related to the Risk of Alzheimer's Disease?. British Journal of Psychiatry, 1990, 157, 273-275.	1.7	18
854	Serum levels of interleukin-6 are not elevated in patients with Alzheimer's disease. Neuroscience Letters, 1990, 108, 350-354.	1.0	77