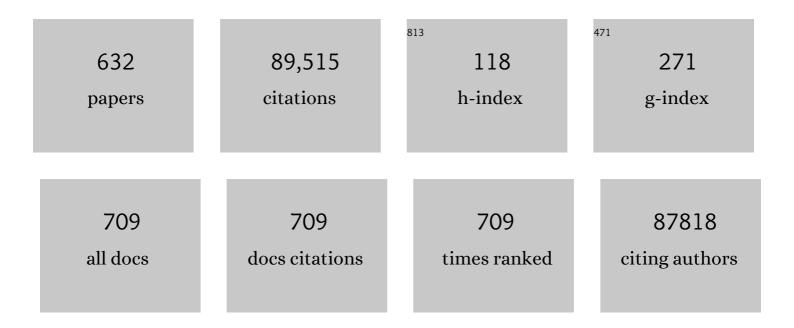
List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Worldwide trends in body-mass index, underweight, overweight, and obesity from 1975 to 2016: a pooled analysis of 2416 population-based measurement studies in 128Å·9 million children, adolescents, and adults. Lancet, The, 2017, 390, 2627-2642.	6.3	5,010
2	Genetic studies of body mass index yield new insights for obesity biology. Nature, 2015, 518, 197-206.	13.7	3,823
3	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
4	Worldwide trends in diabetes since 1980: a pooled analysis of 751 population-based studies with 4·4 million participants. Lancet, The, 2016, 387, 1513-1530.	6.3	2,842
5	<i>TREM2</i> Variants in Alzheimer's Disease. New England Journal of Medicine, 2013, 368, 117-127.	13.9	2,385
6	Genome-wide association study identifies variants at CLU and CR1 associated with Alzheimer's disease. Nature Genetics, 2009, 41, 1094-1099.	9.4	2,155
7	Myocardial infarction and coronary deaths in the World Health Organization MONICA Project. Registration procedures, event rates, and case-fatality rates in 38 populations from 21 countries in four continents Circulation, 1994, 90, 583-612.	1.6	2,056
8	Genetic meta-analysis of diagnosed Alzheimer's disease identifies new risk loci and implicates Aβ, tau, immunity and lipid processing. Nature Genetics, 2019, 51, 414-430.	9.4	1,962
9	Deletion polymorphism in the gene for angiotensin-converting enzyme is a potent risk factor for myocardial infarction. Nature, 1992, 359, 641-644.	13.7	1,880
10	α-synuclein locus duplication as a cause of familial Parkinson's disease. Lancet, The, 2004, 364, 1167-1169.	6.3	1,858
11	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
12	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
13	Worldwide trends in blood pressure from 1975 to 2015: a pooled analysis of 1479 population-based measurement studies with $19\hat{A}\cdot1$ million participants. Lancet, The, 2017, 389, 37-55.	6.3	1,667
14	Large-scale association analysis identifies new risk loci for coronary artery disease. Nature Genetics, 2013, 45, 25-33.	9.4	1,439
15	New genetic loci link adipose and insulin biology to body fat distribution. Nature, 2015, 518, 187-196.	13.7	1,328
16	Worldwide trends in hypertension prevalence and progress in treatment and control from 1990 to 2019: a pooled analysis of 1201 population-representative studies with 104 million participants. Lancet, The, 2021, 398, 957-980.	6.3	1,289
17	Contribution of trends in survival and coronar y-event rates to changes in coronary heart disease mortality: 10-year results from 37 WHO MONICA Project populations. Lancet, The, 1999, 353, 1547-1557.	6.3	1,280
18	Defeating Alzheimer's disease and other dementias: a priority for European science and society. Lancet Neurology, The, 2016, 15, 455-532.	4.9	1,242

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19	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
20	Genome-wide Analysis of Genetic Loci Associated With Alzheimer Disease. JAMA - Journal of the American Medical Association, 2010, 303, 1832.	3.8	1,064
21	Fruit and Vegetable Consumption and Risk of Coronary Heart Disease: A Meta-Analysis of Cohort Studies. Journal of Nutrition, 2006, 136, 2588-2593.	1.3	933
22	Plasma Fibrinogen Level and the Risk of Major Cardiovascular Diseases and Nonvascular Mortality. JAMA - Journal of the American Medical Association, 2005, 294, 1799-809.	3.8	925
23	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
24	C-Reactive Protein, Fibrinogen, and Cardiovascular Disease Prediction. New England Journal of Medicine, 2012, 367, 1310-1320.	13.9	909
25	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
26	EUROASPIRE IV: A European Society of Cardiology survey on the lifestyle, risk factor and therapeutic management of coronary patients from 24 European countries. European Journal of Preventive Cardiology, 2016, 23, 636-648.	0.8	772
27	New insights into the genetic etiology of Alzheimer's disease and related dementias. Nature Genetics, 2022, 54, 412-436.	9.4	700
28	Association of Cardiometabolic Multimorbidity With Mortality. JAMA - Journal of the American Medical Association, 2015, 314, 52.	3.8	624
29	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
30	World Health Organization cardiovascular disease risk charts: revised models to estimate risk in 21 global regions. The Lancet Global Health, 2019, 7, e1332-e1345.	2.9	554
31	Rare and low-frequency coding variants alter human adult height. Nature, 2017, 542, 186-190.	13.7	544
32	APOE and Alzheimer disease: a major gene with semi-dominant inheritance. Molecular Psychiatry, 2011, 16, 903-907.	4.1	529
33	Genome-wide association analyses identify new risk variants and the genetic architecture of amyotrophic lateral sclerosis. Nature Genetics, 2016, 48, 1043-1048.	9.4	494
34	SCORE2 risk prediction algorithms: new models to estimate 10-year risk of cardiovascular disease in Europe. European Heart Journal, 2021, 42, 2439-2454.	1.0	491
35	Study of 300,486 individuals identifies 148 independent genetic loci influencing general cognitive function. Nature Communications, 2018, 9, 2098.	5.8	484
36	Exome-wide association study of plasma lipids in >300,000 individuals. Nature Genetics, 2017, 49, 1758-1766.	9.4	470

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37	Rare variant in scavenger receptor BI raises HDL cholesterol and increases risk of coronary heart disease. Science, 2016, 351, 1166-1171.	6.0	438
38	Unhealthy Effects of Atmospheric Temperature and Pressure on the Occurrence of Myocardial Infarction and Coronary Deaths. Circulation, 1999, 100, E1-7.	1.6	412
39	Apolipoprotein E, ɛ4 allele as a major risk factor for sporadic early and late-onset forms of Alzheimer's disease: analysis of the 19q13.2 chromosomal region. Human Molecular Genetics, 1994, 3, 569-574.	1.4	400
40	Risk Factors for Coronary Heart Disease in Patients Treated for Human Immunodeficiency Virus Infection Compared with the General Population. Clinical Infectious Diseases, 2003, 37, 292-298.	2.9	364
41	Common polygenic variation enhances risk prediction for Alzheimer's disease. Brain, 2015, 138, 3673-3684.	3.7	359
42	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
43	Cigarette Smoking Is Associated with Unhealthy Patterns of Nutrient Intake: a Meta-analysis. Journal of Nutrition, 1998, 128, 1450-1457.	1.3	352
44	C-Reactive Protein, Interleukin-6, and Fibrinogen as Predictors of Coronary Heart Disease. Arteriosclerosis, Thrombosis, and Vascular Biology, 2003, 23, 1255-1261.	1.1	348
45	Genetic contributions to variation in general cognitive function: a meta-analysis of genome-wide association studies in the CHARGE consortium (N=53 949). Molecular Psychiatry, 2015, 20, 183-192.	4.1	344
46	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
47	Trans-ethnic association study of blood pressure determinants in over 750,000 individuals. Nature Genetics, 2019, 51, 51-62.	9.4	328
48	Increased expression of BIN1 mediates Alzheimer genetic risk by modulating tau pathology. Molecular Psychiatry, 2013, 18, 1225-1234.	4.1	321
49	Interleukin-18 and the Risk of Coronary Heart Disease in European Men. Circulation, 2003, 108, 2453-2459.	1.6	317
50	A Polymorphism in CALHM1 Influences Ca2+ Homeostasis, Aβ Levels, and Alzheimer's Disease Risk. Cell, 2008, 133, 1149-1161.	13.5	310
51	Fruit and vegetable consumption and risk of stroke: A meta-analysis of cohort studies. Neurology, 2005, 65, 1193-1197.	1.5	302
52	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
53	Protein-altering variants associated with body mass index implicate pathways that control energy intake and expenditure in obesity. Nature Genetics, 2018, 50, 26-41.	9.4	286
54	The γ e4 allele of the apolipoprotein E gene as a potential protective factor for exudative age-related macular degeneration. American Journal of Ophthalmology, 1998, 125, 353-359.	1.7	265

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55	Trans-ancestry meta-analyses identify rare and common variants associated with blood pressure and hypertension. Nature Genetics, 2016, 48, 1151-1161.	9.4	261
56	A novel Alzheimer disease locus located near the gene encoding tau protein. Molecular Psychiatry, 2016, 21, 108-117.	4.1	260
57	High frequency of potentially pathogenic SORL1 mutations in autosomal dominant early-onset Alzheimer disease. Molecular Psychiatry, 2012, 17, 875-879.	4.1	253
58	Deletion polymorphism in angiotensin-converting enzyme gene associated with parental history of myocardial infarction. Lancet, The, 1993, 341, 991-992.	6.3	251
59	Novel genetic loci associated with hippocampal volume. Nature Communications, 2017, 8, 13624.	5.8	250
60	Predictors of Restenosis After Coronary Stent Implantation. Journal of the American College of Cardiology, 1998, 31, 1291-1298.	1.2	239
61	A case-control study of lipoprotein particles in two populations at contrasting risk for coronary heart disease. The ECTIM Study Arteriosclerosis and Thrombosis: A Journal of Vascular Biology, 1992, 12, 701-707.	3.8	231
62	Fruits, vegetables and coronary heart disease. Nature Reviews Cardiology, 2009, 6, 599-608.	6.1	229
63	APOE genotype, cholesterol level, lipid-lowering treatment, and dementia: The Three-City Study. Neurology, 2005, 64, 1531-1538.	1.5	223
64	Common and rare variant association analyses in amyotrophic lateral sclerosis identify 15 risk loci with distinct genetic architectures and neuron-specific biology. Nature Genetics, 2021, 53, 1636-1648.	9.4	223
65	Meta-analysis of 65,734 Individuals Identifies TSPAN15 and SLC44A2 as Two Susceptibility Loci for Venous Thromboembolism. American Journal of Human Genetics, 2015, 96, 532-542.	2.6	222
66	Contributions of Depressive Mood and Circulating Inflammatory Markers to Coronary Heart Disease in Healthy European Men. Circulation, 2005, 111, 2299-2305.	1.6	220
67	Height and body-mass index trajectories of school-aged children and adolescents from 1985 to 2019 in 200 countries and territories: a pooled analysis of 2181 population-based studies with 65 million participants. Lancet, The, 2020, 396, 1511-1524.	6.3	219
68	Loci associated with ischaemic stroke and its subtypes (SiGN): a genome-wide association study. Lancet Neurology, The, 2016, 15, 174-184.	4.9	217
69	Are the Framingham and PROCAM coronary heart disease risk functions applicable to different European populations? The PRIME Study. European Heart Journal, 2003, 24, 1903-1911.	1.0	216
70	Novel genetic loci underlying human intracranial volume identified through genome-wide association. Nature Neuroscience, 2016, 19, 1569-1582.	7.1	213
71	Common variants at 12q14 and 12q24 are associated with hippocampal volume. Nature Genetics, 2012, 44, 545-551.	9.4	212
72	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

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73	Polymorphisms in the promoter regions of MMP-2, MMP-3, MMP-9 and MMP-12 genes as determinants of aneurysmal coronary artery disease. Journal of the American College of Cardiology, 2002, 40, 43-48.	1.2	208
74	Genome-Wide Scan Identifies TNIP1, PSORS1C1, and RHOB as Novel Risk Loci for Systemic Sclerosis. PLoS Genetics, 2011, 7, e1002091.	1.5	205
75	Endothelial Cell Markers and the Risk of Coronary Heart Disease. Circulation, 2004, 109, 1343-1348.	1.6	203
76	Genomeâ€wide association studies of cerebral white matter lesion burden. Annals of Neurology, 2011, 69, 928-939.	2.8	201
77	Lipoprotein (a) as a predictor of coronary heart disease: the PRIME Study. Atherosclerosis, 2002, 163, 377-384.	0.4	196
78	Impact of apolipoprotein E polymorphism on lipoproteins and risk of myocardial infarction. The ECTIM Study Arteriosclerosis and Thrombosis: A Journal of Vascular Biology, 1994, 14, 1412-1419.	3.8	195
79	Common variation in PHACTR1 is associated with susceptibility to cervical artery dissection. Nature Genetics, 2015, 47, 78-83.	9.4	195
80	A genetic polymorphism of the peroxisome proliferator-activated receptor gamma gene influences plasma leptin levels in obese humans. Human Molecular Genetics, 1998, 7, 435-440.	1.4	193
81	Genetic architecture of subcortical brain structures in 38,851 individuals. Nature Genetics, 2019, 51, 1624-1636.	9.4	192
82	Whole exome sequencing study identifies novel rare and common Alzheimer's-Associated variants involved in immune response and transcriptional regulation. Molecular Psychiatry, 2020, 25, 1859-1875.	4.1	191
83	Associations of Fibrinogen, Factor VII and PAI-1 with Baseline Findings among 10,500 Male Participants in a Prospective Study of Myocardial Infarction. Thrombosis and Haemostasis, 1998, 80, 749-756.	1.8	184
84	Alzheimer risk associated with a copy number variation in the complement receptor 1 increasing C3b/C4b binding sites. Molecular Psychiatry, 2012, 17, 223-233.	4.1	179
85	Overall alcohol intake, beer, wine, and systemic markers of inflammation in western Europe: results from three MONICA samples (Augsburg, Glasgow, Lille). European Heart Journal, 2004, 25, 2092-2100.	1.0	178
86	The Role of Adiposity in Cardiometabolic Traits: A Mendelian Randomization Analysis. PLoS Medicine, 2013, 10, e1001474.	3.9	178
87	Application of non-HDL cholesterol for population-based cardiovascular risk stratification: results from the Multinational Cardiovascular Risk Consortium. Lancet, The, 2019, 394, 2173-2183.	6.3	177
88	CYP2D6 polymorphism, pesticide exposure, and Parkinson's disease. Annals of Neurology, 2004, 55, 430-434.	2.8	175
89	Convergent genetic and expression data implicate immunity in Alzheimer's disease. Alzheimer's and Dementia, 2015, 11, 658-671.	0.4	173
90	A new polymorphism in the APOE promoter associated with risk of developing Alzheimer's disease. Human Molecular Genetics, 1998, 7, 533-540.	1.4	170

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91	Value of HDL Cholesterol, Apolipoprotein A-I, Lipoprotein A-I, and Lipoprotein A-I/A-II in Prediction of Coronary Heart Disease. Arteriosclerosis, Thrombosis, and Vascular Biology, 2002, 22, 1155-1161.	1.1	169
92	Transcriptomic and genetic studies identify IL-33 as a candidate gene for Alzheimer's disease. Molecular Psychiatry, 2009, 14, 1004-1016.	4.1	167
93	Mosaic Loss of Chromosome Y in Blood Is Associated with Alzheimer Disease. American Journal of Human Genetics, 2016, 98, 1208-1219.	2.6	164
94	Genome-wide association study of intracranial aneurysms identifies 17 risk loci and genetic overlap with clinical risk factors. Nature Genetics, 2020, 52, 1303-1313.	9.4	163
95	Multiethnic Genome-Wide Association Study of Cerebral White Matter Hyperintensities on MRI. Circulation: Cardiovascular Genetics, 2015, 8, 398-409.	5.1	162
96	Tau deletion promotes brain insulin resistance. Journal of Experimental Medicine, 2017, 214, 2257-2269.	4.2	158
97	Circulating soluble adhesion molecules ICAM-1 and VCAM-1 and incident coronary heart disease: The PRIME Study. Atherosclerosis, 2003, 170, 169-176.	0.4	156
98	Household Income Is Associated With the Risk of Metabolic Syndrome in a Sex-Specific Manner. Diabetes Care, 2005, 28, 409-415.	4.3	156
99	Impact of the Peroxisome Proliferator Activated Receptor γ2 Pro12Ala polymorphism on adiposity, lipids and non-insulin-dependent diabetes mellitus. International Journal of Obesity, 2000, 24, 195-199.	1.6	155
100	Genome-wide association study confirms BST1 and suggests a locus on 12q24 as the risk loci for Parkinson's disease in the European population. Human Molecular Genetics, 2011, 20, 615-627.	1.4	155
101	Gene-Wide Analysis Detects Two New Susceptibility Genes for Alzheimer's Disease. PLoS ONE, 2014, 9, e94661.	1.1	155
102	The Emerging Risk Factors Collaboration: analysis of individual data on lipid, inflammatory and other markers in over 1.1 million participants in 104 prospective studies of cardiovascular diseases. European Journal of Epidemiology, 2007, 22, 839-869.	2.5	153
103	Single nucleotide polymorphisms in the FADS gene cluster are associated with delta-5 and delta-6 desaturase activities estimated by serum fatty acid ratios. Journal of Lipid Research, 2010, 51, 2325-2333.	2.0	153
104	Excessive Daytime Sleepiness Is an Independent Risk Indicator for Cardiovascular Mortality in Community-Dwelling Elderly. Stroke, 2009, 40, 1219-1224.	1.0	152
105	Implication of the Immune System in Alzheimer's Disease: Evidence from Genome-Wide Pathway Analysis. Journal of Alzheimer's Disease, 2010, 20, 1107-1118.	1.2	152
106	Association Between Depressive Symptoms and Incident Cardiovascular Diseases. JAMA - Journal of the American Medical Association, 2020, 324, 2396.	3.8	152
107	Ischemic stroke is associated with the <i>ABO</i> locus: The EuroCLOT study. Annals of Neurology, 2013, 73, 16-31.	2.8	144
108	Association of Vascular Risk Factors With Cervical Artery Dissection and Ischemic Stroke in Young Adults. Circulation, 2011, 123, 1537-1544.	1.6	141

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109	Low-frequency and common genetic variation in ischemic stroke. Neurology, 2016, 86, 1217-1226.	1.5	141
110	β2-adrenoceptor gene polymorphism, body weight, and physical activity. Lancet, The, 1999, 353, 896.	6.3	140
111	Common variants in Alzheimer's disease and risk stratification by polygenic risk scores. Nature Communications, 2021, 12, 3417.	5.8	140
112	The apolipoprotein E alleles as major susceptibility factors for Creutzfeldt-Jakob disease. Lancet, The, 1994, 344, 1315-1318.	6.3	139
113	Effects of diabetes definition on global surveillance of diabetes prevalence and diagnosis: a pooled analysis of 96 population-based studies with 331â€~288 participants. Lancet Diabetes and Endocrinology,the, 2015, 3, 624-637.	5.5	139
114	Impact of genetic variation of PPARÎ <sup>3</sup> in humans. Molecular Genetics and Metabolism, 2004, 83, 93-102.	0.5	138
115	A Genome-Wide Association Study Identifies <i>LIPA</i> as a Susceptibility Gene for Coronary Artery Disease. Circulation: Cardiovascular Genetics, 2011, 4, 403-412.	5.1	130
116	Independent association of an APOE gene promoter polymorphism with increased risk of myocardial infarction and decreased APOE plasma concentrationsthe ECTIM Study. Human Molecular Genetics, 2000, 9, 57-61.	1.4	129
117	Pronounced impact of Th1/E47cs mutation compared with -491 AT mutation on neural APOE gene expression and risk of developing Alzheimer's disease. Human Molecular Genetics, 1998, 7, 1511-1516.	1.4	127
118	Patterns of alcohol consumption and ischaemic heart disease in culturally divergent countries: the Prospective Epidemiological Study of Myocardial Infarction (PRIME). BMJ: British Medical Journal, 2010, 341, c6077-c6077.	2.4	127
119	D Allele of the Angiotensin I–Converting Enzyme Is a Major Risk Factor for Restenosis After Coronary Stenting. Circulation, 1997, 96, 56-60.	1.6	127
120	Genetics of Venous Thrombosis: Insights from a New Genome Wide Association Study. PLoS ONE, 2011, 6, e25581.	1.1	127
121	Association of plasma amyloid $\hat{l}^2$ with risk of dementia. Neurology, 2009, 73, 847-853.	1.5	126
122	Adiposity as a cause of cardiovascular disease: a Mendelian randomization study. International Journal of Epidemiology, 2015, 44, 578-586.	0.9	123
123	The pursuit of susceptibility genes for Alzheimer's disease: progress and prospects. Trends in Genetics, 2010, 26, 84-93.	2.9	122
124	Genetic determinants of risk in pulmonary arterial hypertension: international genome-wide association studies and meta-analysis. Lancet Respiratory Medicine,the, 2019, 7, 227-238.	5.2	122
125	GWAS and colocalization analyses implicate carotid intima-media thickness and carotid plaque loci in cardiovascular outcomes. Nature Communications, 2018, 9, 5141.	5.8	119
126	Fish Consumption Is Associated With Lower Heart Rates. Circulation, 2003, 108, 820-825.	1.6	118

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127	Contribution to Alzheimer's disease risk of rare variants in TREM2, SORL1, and ABCA7 in 1779 cases and 1273 controls. Neurobiology of Aging, 2017, 59, 220.e1-220.e9.	1.5	116
128	Leisure-time physical activity and regular walking or cycling to work are associated with adiposity and 5â€y weight gain in middle-aged men: the PRIME Study. International Journal of Obesity, 2001, 25, 940-948.	1.6	115
129	Association between beta-1 and beta-2 adrenergic receptor gene polymorphisms and the response to beta-blockade in patients with stable congestive heart failure. Pharmacogenetics and Genomics, 2005, 15, 137-142.	0.7	113
130	ApoE immunoreactivity and microglial cells in Alzheimer's disease brain. Neuroscience Letters, 1995, 195, 5-8.	1.0	112
131	Missense variant in TREML2 protects against Alzheimer's disease. Neurobiology of Aging, 2014, 35, 1510.e19-1510.e26.	1.5	110
132	Influence of Apolipoprotein E Genotype on the Risk of Cognitive Deterioration in Moderate Drinkers and Smokers. Epidemiology, 2000, 11, 280-284.	1.2	110
133	Alcohol intake and diet in France, the prominent role of lifestyle. European Heart Journal, 2004, 25, 1153-1162.	1.0	109
134	Cystatin C and Cardiovascular Disease. Journal of the American College of Cardiology, 2016, 68, 934-945.	1.2	109
135	Association between Parkinson's disease and polymorphisms in the nNOS and iNOS genes in a community-based case-control study. Human Molecular Genetics, 2003, 12, 79-86.	1.4	108
136	The Angiotensin II Type 1 Receptor Gene Polymorphism Is Associated With Coronary Artery Vasoconstriction. Journal of the American College of Cardiology, 1997, 29, 486-490.	1.2	107
137	Genetics of Alzheimer's disease: new evidences for an old hypothesis?. Current Opinion in Genetics and Development, 2011, 21, 295-301.	1.5	105
138	Characterization of a Unique Genetic Variant in the β1-adrenoceptor Gene and Evaluation of its Role in Idiopathic Dilated Cardiomyopathy. Journal of Molecular and Cellular Cardiology, 1999, 31, 1025-1032.	0.9	103
139	Genome-wide haplotype association study identifies the FRMD4A gene as a risk locus for Alzheimer's disease. Molecular Psychiatry, 2013, 18, 461-470.	4.1	103
140	Contribution of <i>APOE</i> promoter polymorphisms to Alzheimer's disease risk. Neurology, 2002, 59, 59-66.	1.5	102
141	Plasma cystatin-C and development of coronary heart disease: The PRIME Study. Atherosclerosis, 2006, 185, 375-380.	0.4	102
142	Patients with coronary artery disease and diabetes need improved management: a report from the EUROASPIRE IV survey: a registry from the EuroObservational Research Programme of the European Society of Cardiology. Cardiovascular Diabetology, 2015, 14, 133.	2.7	101
143	Multiple coronary heart disease risk factors are associated with menopause and influenced by substitutive hormonal therapy in a cohort of French women. Atherosclerosis, 1995, 118, 123-133.	0.4	100
144	Association between nutrition knowledge and nutritional intake in middle-aged men from Northern France. Public Health Nutrition, 2001, 4, 27-33.	1.1	100

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145	A genomic approach to therapeutic target validation identifies a glucose-lowering <i>GLP1R</i> variant protective for coronary heart disease. Science Translational Medicine, 2016, 8, 341ra76.	5.8	100
146	Confirmation of the ϵ4 allele of the apolipoprotein E gene as a risk factor for lateâ€onset Alzheimer's disease. Neurology, 1994, 44, 342-342.	1.5	100
147	Apolipoprotein E and Alzheimer disease: genotype-specific risks by age and sex. American Journal of Human Genetics, 1997, 60, 439-46.	2.6	100
148	A Genomeâ€Wide Association Study for Venous Thromboembolism: The Extended Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) Consortium. Genetic Epidemiology, 2013, 37, 512-521.	0.6	99
149	Functional screening of Alzheimer risk loci identifies PTK2B as an in vivo modulator and early marker of Tau pathology. Molecular Psychiatry, 2017, 22, 874-883.	4.1	98
150	Equalization of four cardiovascular risk algorithms after systematic recalibration: individual-participant meta-analysis of 86 prospective studies. European Heart Journal, 2019, 40, 621-631.	1.0	97
151	Frequency of fruit and vegetable consumption and coronary heart disease in France and Northern Ireland: the PRIME study. British Journal of Nutrition, 2004, 92, 963-972.	1.2	96
152	C-Reactive Protein, Interleukin 6, Fibrinogen and Risk of Sudden Death in European Middle-Aged Men: The PRIME Study. Arteriosclerosis, Thrombosis, and Vascular Biology, 2010, 30, 2047-2052.	1.1	96
153	SORL1 rare variants: a major risk factor for familial early-onset Alzheimer's disease. Molecular Psychiatry, 2016, 21, 831-836.	4.1	96
154	Association at LRP gene locus with sporadic late-onset Alzheimer's disease. Lancet, The, 1998, 351, 1787-1788.	6.3	95
155	High consumptions of grain, fish, dairy products and combinations of these are associated with a low prevalence of metabolic syndrome. Journal of Epidemiology and Community Health, 2007, 61, 810-817.	2.0	94
156	Evidence of a Role for Lactadherin in Alzheimer's Disease. American Journal of Pathology, 2007, 170, 921-929.	1.9	94
157	Effect of ACE inhibitors on angiographic restenosis after coronary stenting (PARIS): a randomised, double-blind, placebo-controlled trial. Lancet, The, 2001, 357, 1321-1324.	6.3	93
158	Genome-wide association analysis identifies a susceptibility locus for pulmonary arterial hypertension. Nature Genetics, 2013, 45, 518-521.	9.4	93
159	Association between the low density lipoprotein receptor-related protein (LRP) and Alzheimer's disease. Neuroscience Letters, 1997, 227, 68-70.	1.0	92
160	α-2 macroglobulin gene and Alzheimer disease. Nature Genetics, 1999, 22, 17-19.	9.4	91
161	A Functional Polymorphism in a STAT5B Site of the Human PPARÎ <sup>3</sup> 3 Gene Promoter Affects Height and Lipid Metabolism in a French Population. Arteriosclerosis, Thrombosis, and Vascular Biology, 2003, 23, 289-294.	1.1	91
162	The proteome and secretome of human arterial smooth muscle cells. Proteomics, 2005, 5, 585-596.	1.3	91

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163	Discovery of rare variants associated with blood pressure regulation through meta-analysis of 1.3 million individuals. Nature Genetics, 2020, 52, 1314-1332.	9.4	91
164	Association of occurrence of aneurysmal bleeding with meteorologic variations in the north of France Stroke, 1994, 25, 338-341.	1.0	89
165	Bilirubin and coronary heart disease risk in the Prospective Epidemiological Study of Myocardial Infarction (PRIME). European Journal of Cardiovascular Prevention and Rehabilitation, 2007, 14, 79-84.	3.1	89
166	Protein-coding variants implicate novel genes related to lipid homeostasis contributing to body-fat distribution. Nature Genetics, 2019, 51, 452-469.	9.4	89
167	Cerebral small vessel disease genomics and its implications across the lifespan. Nature Communications, 2020, 11, 6285.	5.8	89
168	Education, socioeconomic and lifestyle factors, and risk of coronary heart disease: the PRIME Study. International Journal of Epidemiology, 2005, 34, 268-275.	0.9	87
169	Distortion of Allelic Expression of Apolipoprotein E in Alzheimer's Disease. Human Molecular Genetics, 1997, 6, 2151-2154.	1.4	86
170	Gender differences in the implementation of cardiovascular prevention measures after an acute coronary event. Heart, 2010, 96, 1744-1749.	1.2	86
171	Impact of polymorphisms of the human β2-adrenoceptor gene on obesity in a French population. International Journal of Obesity, 2000, 24, 382-387.	1.6	84
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PHILIPPE AMOUYEL

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PHILIPPE AMOUYEL

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