L Adrienne Cupples

List of Publications by Year in descending order

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646 papers

103,140 citations

146 h-index 289 g-index

679 all docs

679 docs citations

times ranked

679

82084 citing authors

#	Article	IF	CITATIONS
1	Genetic studies of body mass index yield new insights for obesity biology. Nature, 2015, 518, 197-206.	13.7	3,823
2	Biological, clinical and population relevance of 95 loci for blood lipids. Nature, 2010, 466, 707-713.	13.7	3,249
3	Discovery and refinement of loci associated with lipid levels. Nature Genetics, 2013, 45, 1274-1283.	9.4	2,641
4	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
5	Abdominal Visceral and Subcutaneous Adipose Tissue Compartments. Circulation, 2007, 116, 39-48.	1.6	2,349
6	A comprehensive 1000 Genomes–based genome-wide association meta-analysis of coronary artery disease. Nature Genetics, 2015, 47, 1121-1130.	9.4	2,054
7	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
8	Plasma HDL cholesterol and risk of myocardial infarction: a mendelian randomisation study. Lancet, The, 2012, 380, 572-580.	6.3	1,937
9	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
10	Hundreds of variants clustered in genomic loci and biological pathways affect human height. Nature, 2010, 467, 832-838.	13.7	1,789
11	Large-scale association analysis identifies 13 new susceptibility loci for coronary artery disease. Nature Genetics, 2011, 43, 333-338.	9.4	1,685
12	Effects of age, sex, and ethnicity on the association between apolipoprotein E genotype and Alzheimer disease. A meta-analysis. APOE and Alzheimer Disease Meta Analysis Consortium. JAMA - Journal of the American Medical Association, 1997, 278, 1349-56.	3.8	1,658
13	New genetic loci link adipose and insulin biology to body fat distribution. Nature, 2015, 518, 187-196.	13.7	1,328
14	Common variants at 30 loci contribute to polygenic dyslipidemia. Nature Genetics, 2009, 41, 56-65.	9.4	1,234
15	Heart rate and cardiovascular mortality: The Framingham study. American Heart Journal, 1987, 113, 1489-1494.	1.2	1,170
16	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
17	Sequencing of 53,831 diverse genomes from the NHLBI TOPMed Program. Nature, 2021, 590, 290-299.	13.7	1,069
18	The neuronal sortilin-related receptor SORL1 is genetically associated with Alzheimer disease. Nature Genetics, 2007, 39, 168-177.	9.4	1,045

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19	Loss-of-Function Mutations in (i>APOC3, (i>Triglycerides, and Coronary Disease. New England Journal of Medicine, 2014, 371, 22-31.	13.9	936
20	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
21	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
22	Genetic Associations with Valvular Calcification and Aortic Stenosis. New England Journal of Medicine, 2013, 368, 503-512.	13.9	767
23	Common variants associated with plasma triglycerides and risk for coronary artery disease. Nature Genetics, 2013, 45, 1345-1352.	9.4	754
24	The Third Generation Cohort of the National Heart, Lung, and Blood Institute's Framingham Heart Study: Design, Recruitment, and Initial Examination. American Journal of Epidemiology, 2007, 165, 1328-1335.	1.6	752
25	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
26	Genotype Score in Addition to Common Risk Factors for Prediction of Type 2 Diabetes. New England Journal of Medicine, 2008, 359, 2208-2219.	13.9	696
27	Relation of pooled logistic regression to time dependent cox regression analysis: The framingham heart study. Statistics in Medicine, 1990, 9, 1501-1515.	0.8	681
28	Variants in MTNR1B influence fasting glucose levels. Nature Genetics, 2009, 41, 77-81.	9.4	662
29	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
30	Risk Factors for Longitudinal Bone Loss in Elderly Men and Women: The Framingham Osteoporosis Study. Journal of Bone and Mineral Research, 2010, 15, 710-720.	3.1	620
31	Potassium, magnesium, and fruit and vegetable intakes are associated with greater bone mineral density in elderly men and women. American Journal of Clinical Nutrition, 1999, 69, 727-736.	2.2	603
32	Exome sequencing identifies rare LDLR and APOA5 alleles conferring risk for myocardial infarction. Nature, 2015, 518, 102-106.	13.7	581
33	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
34	Abdominal Aortic Calcific Deposits Are an Important Predictor of Vascular Morbidity and Mortality. Circulation, 2001, 103, 1529-1534.	1.6	546
35	New indices to classify location, severity and progression of calcific lesions in the abdominal aorta: a 25-year follow-up study. Atherosclerosis, 1997, 132, 245-250.	0.4	541
36	Homocysteine as a Predictive Factor for Hip Fracture in Older Persons. New England Journal of Medicine, 2004, 350, 2042-2049.	13.9	539

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37	Evidence for a Gene Influencing Blood Pressure on Chromosome 17. Hypertension, 2000, 36, 477-483.	1.3	534
38	Association between alcohol and cardiovascular disease: Mendelian randomisation analysis based on individual participant data. BMJ, The, 2014, 349, g4164-g4164.	3.0	528
39	Parental transmission of type 2 diabetes: the Framingham Offspring Study. Diabetes, 2000, 49, 2201-2207.	0.3	513
40	Disclosure of <i>APOE </i> Genotype for Risk of Alzheimer's Disease. New England Journal of Medicine, 2009, 361, 245-254.	13.9	490
41	Depression as a Risk Factor for Alzheimer Disease. Archives of Neurology, 2003, 60, 753.	4.9	485
42	Wholeâ€genome sequencing identifies EN1 as a determinant of bone density and fracture. Nature, 2015, 526, 112-117.	13.7	483
43	Exome-wide association study of plasma lipids in >300,000 individuals. Nature Genetics, 2017, 49, 1758-1766.	9.4	470
44	Insulin resistance, oxidative stress, hypertension, and leukocyte telomere length in men from the Framingham Heart Study. Aging Cell, 2006, 5, 325-330.	3.0	465
45	Regional obesity and risk of cardiovascular disease; the Framingham study. Journal of Clinical Epidemiology, 1991, 44, 183-190.	2.4	459
46	Effect of Dietary Protein on Bone Loss in Elderly Men and Women: The Framingham Osteoporosis Study. Journal of Bone and Mineral Research, 2000, 15, 2504-2512.	3.1	446
47	Head injury and the risk of AD in the MIRAGE study. Neurology, 2000, 54, 1316-1323.	1.5	440
48	Bone loss and the progression of abdominal aortic calcification over a 25 year period: The Framingham heart study. Calcified Tissue International, 2001, 68, 271-276.	1.5	416
49	Association of Cholesteryl Ester Transfer Protein– <i>Taq</i> IB Polymorphism With Variations in Lipoprotein Subclasses and Coronary Heart Disease Risk. Arteriosclerosis, Thrombosis, and Vascular Biology, 2000, 20, 1323-1329.	1.1	385
50	FTO genotype is associated with phenotypic variability of body mass index. Nature, 2012, 490, 267-272.	13.7	383
51	Meta-analysis Confirms CR1, CLU, and PICALM as Alzheimer Disease Risk Loci and Reveals Interactions With APOE Genotypes. Archives of Neurology, 2010, 67, 1473.	4.9	376
52	Inherited causes of clonal haematopoiesis in 97,691 whole genomes. Nature, 2020, 586, 763-768.	13.7	376
53	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
54	Dietary vitamin K intakes are associated with hip fracture but not with bone mineral density in elderly men and women. American Journal of Clinical Nutrition, 2000, 71, 1201-1208.	2.2	353

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55	The power of genetic diversity in genome-wide association studies of lipids. Nature, 2021, 600, 675-679.	13.7	353
56	NIDDM and Blood Pressure as Risk Factors for Poor Cognitive Performance: The Framingham Study. Diabetes Care, 1997, 20, 1388-1395.	4.3	339
57	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
58	Risk of Dementia Among White and African American Relatives of Patients With Alzheimer Disease. JAMA - Journal of the American Medical Association, 2002, 287, 329.	3.8	330
59	Multiple loci influence erythrocyte phenotypes in the CHARGE Consortium. Nature Genetics, 2009, 41, 1191-1198.	9.4	324
60	Television viewing and change in body fat from preschool to early adolescence: The Framingham Children's Study. International Journal of Obesity, 2003, 27, 827-833.	1.6	319
61	The landscape of recombination in African Americans. Nature, 2011, 476, 170-175.	13.7	319
62	Cardiac failure and sudden death in the Framingham Study. American Heart Journal, 1988, 115, 869-875.	1.2	312
63	Genetic assessment of age-associated Alzheimer disease risk: Development and validation of a polygenic hazard score. PLoS Medicine, 2017, 14, e1002258.	3.9	311
64	Cardiorespiratory Events Recorded on Home Monitors SUBTITLE Comparison of Healthy Infants With Those at Increased Risk for SIDS / SUBTITLE > . JAMA - Journal of the American Medical Association, 2001, 285, 2199.	3.8	310
65	Common variants in 22 loci are associated with QRS duration and cardiac ventricular conduction. Nature Genetics, 2010, 42, 1068-1076.	9.4	308
66	A Randomized Trial of Treatment Options for Alcohol-Abusing Workers. New England Journal of Medicine, 1991, 325, 775-782.	13.9	304
67	Forty-Three Loci Associated with Plasma Lipoprotein Size, Concentration, and Cholesterol Content in Genome-Wide Analysis. PLoS Genetics, 2009, 5, e1000730.	1.5	300
68	Genome-wide association study in 79,366 European-ancestry individuals informs the genetic architecture of 25-hydroxyvitamin D levels. Nature Communications, 2018, 9, 260.	5.8	295
69	Parental history is an independent risk factor for coronary artery disease: The Framingham Study. American Heart Journal, 1990, 120, 963-969.	1.2	294
70	A genome-wide association study for blood lipid phenotypes in the Framingham Heart Study. BMC Medical Genetics, 2007, 8, S17.	2.1	289
71	Genetic variation near IRS1 associates with reduced adiposity and an impaired metabolic profile. Nature Genetics, 2011, 43, 753-760.	9.4	289
72	Elevated Plasma Lipoprotein(a) and Coronary Heart Disease in Men Aged 55 Years and Younger. JAMA - Journal of the American Medical Association, 1996, 276, 544.	3.8	287

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73	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
74	Bone Mass and the Risk of Breast Cancer among Postmenopausal Women. New England Journal of Medicine, 1997, 336, 611-617.	13.9	283
75	Association Between the UGT1A1*28 Allele, Bilirubin Levels, and Coronary Heart Disease in the Framingham Heart Study. Circulation, 2006, 114, 1476-1481.	1.6	283
76	Dietary Silicon Intake Is Positively Associated With Bone Mineral Density in Men and Premenopausal Women of the Framingham Offspring Cohort. Journal of Bone and Mineral Research, 2003, 19, 297-307.	3.1	281
77	Does early physical activity predict body fat change throughout childhood? Preventive Medicine, 2003, 37, 10-17.	1.6	281
78	High-Density Lipoprotein Subpopulation Profile and Coronary Heart Disease Prevalence in Male Participants of the Framingham Offspring Study. Arteriosclerosis, Thrombosis, and Vascular Biology, 2004, 24, 2181-2187.	1.1	275
79	Apolipoprotein E genotype and cardiovascular disease in the Framingham Heart Study. Atherosclerosis, 2001, 154, 529-537.	0.4	271
80	Genome-Wide Association Study for Coronary Artery Calcification With Follow-Up in Myocardial Infarction. Circulation, 2011, 124, 2855-2864.	1.6	269
81	Risk of dementia among relatives of Alzheimer's disease patients in the MIRAGE study. Neurology, 1996, 46, 641-650.	1.5	261
82	Blood pressure as a risk factor for cardiovascular disease. The Framingham Study-30 years of follow-up Hypertension, 1989, 13, 113-8.	1.3	259
83	Sex and Age Differences in Lipoprotein Subclasses Measured by Nuclear Magnetic Resonance Spectroscopy: The Framingham Study. Clinical Chemistry, 2004, 50, 1189-1200.	1.5	259
84	Value of High-Density Lipoprotein (HDL) Subpopulations in Predicting Recurrent Cardiovascular Events in the Veterans Affairs HDL Intervention Trial. Arteriosclerosis, Thrombosis, and Vascular Biology, 2005, 25, 2185-2191.	1.1	258
85	Remnant-like particle (RLP) cholesterol is an independent cardiovascular disease risk factor in women: results from the Framingham Heart Study. Atherosclerosis, 2001, 154, 229-236.	0.4	257
86	Weight loss in early stage of Huntington's disease. Neurology, 2002, 59, 1325-1330.	1.5	255
87	Collaborative Meta-analysis: Associations of 150 Candidate Genes With Osteoporosis and Osteoporotic Fracture. Annals of Internal Medicine, 2009, 151, 528.	2.0	250
88	Genome-Wide Association for Abdominal Subcutaneous and Visceral Adipose Reveals a Novel Locus for Visceral Fat in Women. PLoS Genetics, 2012, 8, e1002695.	1.5	245
89	New loci for body fat percentage reveal link between adiposity and cardiometabolic disease risk. Nature Communications, 2016, 7, 10495.	5.8	245
90	Chemokine receptor mutant CX3CR1-M280 has impaired adhesive function and correlates with protection from cardiovascular disease in humans. Journal of Clinical Investigation, 2003, 111, 1241-1250.	3.9	245

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91	Bone mineral density and dietary patterns in older adults: the Framingham Osteoporosis Study,,. American Journal of Clinical Nutrition, 2002, 76, 245-252.	2.2	244
92	Association Between Estrogen Receptor \hat{l}_{\pm} Gene Variation and Cardiovascular Disease. JAMA - Journal of the American Medical Association, 2003, 290, 2263.	3.8	243
93	Total serum bilirubin and risk of cardiovascular disease in the Framingham offspring study. American Journal of Cardiology, 2001, 87, 1196-1200.	0.7	240
94	Genome-wide meta-analysis identifies six novel loci associated with habitual coffee consumption. Molecular Psychiatry, 2015, 20, 647-656.	4.1	235
95	A prospective investigation of elevated lipoprotein (a) detected by electrophoresis and cardiovascular disease in women. The Framingham Heart Study Circulation, 1994, 90, 1688-1695.	1.6	230
96	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
97	Comparison of baseline and repeated measure covariate techniques in the Framingham heart study. Statistics in Medicine, 1988, 7, 205-218.	0.8	228
98	Large-Scale Gene-Centric Meta-analysis across 32 Studies Identifies Multiple Lipid Loci. American Journal of Human Genetics, 2012, 91, 823-838.	2.6	227
99	Increased rate of suicide among patients with Huntington's disease Journal of Neurology, Neurosurgery and Psychiatry, 1984, 47, 1283-1287.	0.9	226
100	A Comprehensive Genetic Association Study of Alzheimer Disease in African Americans. Archives of Neurology, 2011, 68, 1569.	4.9	221
101	Postmenopausal hormone therapy and Alzheimer's disease risk: interaction with age. Journal of Neurology, Neurosurgery and Psychiatry, 2005, 76, 103-105.	0.9	219
102	Best Practices and Joint Calling of the HumanExome BeadChip: The CHARGE Consortium. PLoS ONE, 2013, 8, e68095.	1.1	219
103	Preschool Physical Activity Level and Change in Body Fatness in Young Children: The Framingham Children's Study. American Journal of Epidemiology, 1995, 142, 982-988.	1.6	217
104	Pharmacogenetic meta-analysis of genome-wide association studies of LDL cholesterol response to statins. Nature Communications, 2014, 5, 5068.	5.8	216
105	Genome-wide association meta-analysis for total serum bilirubin levels. Human Molecular Genetics, 2009, 18, 2700-2710.	1.4	214
106	Familial aggregation of stroke. The Framingham Study Stroke, 1993, 24, 1366-1371.	1.0	212
107	Decreased Neuronal and Increased Oligodendroglial Densities in Huntington's Disease Caudate Nucleus. Journal of Neuropathology and Experimental Neurology, 1991, 50, 729-742.	0.9	211
108	CCL2 Polymorphisms Are Associated With Serum Monocyte Chemoattractant Protein-1 Levels and Myocardial Infarction in the Framingham Heart Study. Circulation, 2005, 112, 1113-1120.	1.6	210

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109	Genome-wide meta-analysis of observational studies shows common genetic variants associated with macronutrient intake. American Journal of Clinical Nutrition, 2013, 97, 1395-1402.	2.2	210
110	Vitamin K intake and bone mineral density in women and men. American Journal of Clinical Nutrition, 2003, 77, 512-516.	2.2	209
111	CUBN Is a Gene Locus for Albuminuria. Journal of the American Society of Nephrology: JASN, 2011, 22, 555-570.	3.0	208
112	Clinical and neuropathologic assessment of severity in Huntington's disease. Neurology, 1988, 38, 341-341.	1.5	207
113	Functional Variant of CYP4A11 20-Hydroxyeicosatetraenoic Acid Synthase Is Associated With Essential Hypertension. Circulation, 2005, 111, 63-69.	1.6	206
114	Colas, but not other carbonated beverages, are associated with low bone mineral density in older women: The Framingham Osteoporosis Study. American Journal of Clinical Nutrition, 2006, 84, 936-942.	2.2	203
115	Health Behavior Changes After Genetic Risk Assessment for Alzheimer Disease: The REVEAL Study. Alzheimer Disease and Associated Disorders, 2008, 22, 94-97.	0.6	203
116	Use of >100,000 NHLBI Trans-Omics for Precision Medicine (TOPMed) Consortium whole genome sequences improves imputation quality and detection of rare variant associations in admixed African and Hispanic/Latino populations. PLoS Genetics, 2019, 15, e1008500.	1.5	203
117	Factors Associated With Slow Progression in Huntington's Disease. Archives of Neurology, 1991, 48, 800-804.	4.9	196
118	A Genetic Risk Score Is Associated With Incident Cardiovascular Disease and Coronary Artery Calcium. Circulation: Cardiovascular Genetics, 2012, 5, 113-121.	5.1	196
119	Validation and Comparison of Two Frailty Indexes: The MOBILIZE Boston Study. Journal of the American Geriatrics Society, 2009, 57, 1532-1539.	1.3	193
120	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
121	An Integration of Genome-Wide Association Study and Gene Expression Profiling to Prioritize the Discovery of Novel Susceptibility Loci for Osteoporosis-Related Traits. PLoS Genetics, 2010, 6, e1000977.	1.5	191
122	Meta-analysis of genome-wide association studies from the CHARGE consortium identifies common variants associated with carotid intima media thickness and plaque. Nature Genetics, 2011, 43, 940-947.	9.4	191
123	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
124	Serum Albumin and Risk of Myocardial Infarction and All-Cause Mortality in the Framingham Offspring Study. Circulation, 2002, 106, 2919-2924.	1.6	189
125	Association of JAG1 with Bone Mineral Density and Osteoporotic Fractures: A Genome-wide Association Study and Follow-up Replication Studies. American Journal of Human Genetics, 2010, 86, 229-239.	2.6	188
126	Dietary Fat Intake Determines the Effect of a Common Polymorphism in the Hepatic Lipase Gene Promoter on High-Density Lipoprotein Metabolism. Circulation, 2002, 106, 2315-2321.	1.6	186

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127	Heritability of Age at Natural Menopause in the Framingham Heart Study. Journal of Clinical Endocrinology and Metabolism, 2005, 90, 3427-3430.	1.8	186
128	Association of Low-Density Lipoprotein Cholesterol–Related Genetic Variants With Aortic Valve Calcium and Incident Aortic Stenosis. JAMA - Journal of the American Medical Association, 2014, 312, 1764.	3.8	184
129	Physical Activity and Stroke Risk: The Framingham Study. American Journal of Epidemiology, 1994, 140, 608-620.	1.6	182
130	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
131	Second Hip Fracture in Older Men and Women. Archives of Internal Medicine, 2007, 167, 1971.	4.3	175
132	A Genome-Wide Scan for Loci Linked to Plasma Levels of Glucose and HbA1c in a Community-Based Sample of Caucasian Pedigrees: The Framingham Offspring Study. Diabetes, 2002, 51, 833-840.	0.3	173
133	Low-frequency and rare exome chip variants associate with fasting glucose and type 2 diabetes susceptibility. Nature Communications, 2015, 6, 5897.	5.8	173
134	Directional dominance on stature and cognition inÂdiverse human populations. Nature, 2015, 523, 459-462.	13.7	173
135	Polyunsaturated fatty acids modulate the effects of the APOA1 G-A polymorphism on HDL-cholesterol concentrations in a sex-specific manner: the Framingham Study. American Journal of Clinical Nutrition, 2002, 75, 38-46.	2.2	172
136	The Framingham Heart Study 100K SNP genome-wide association study resource: overview of 17 phenotype working group reports. BMC Medical Genetics, 2007, 8 , S1.	2.1	169
137	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
138	Metabolic Risk Factors Worsen Continuously across the Spectrum of Nondiabetic Glucose Tolerance: The Framingham Offspring Study. Annals of Internal Medicine, 1998, 128, 524.	2.0	166
139	Genetic Risk Reclassification for Type 2 Diabetes by Age Below or Above 50 Years Using 40 Type 2 Diabetes Risk Single Nucleotide Polymorphisms. Diabetes Care, 2011, 34, 121-125.	4.3	165
140	Quantitative neuropathological changes in presymptomatic Huntington's disease. Annals of Neurology, 2001, 49, 29-34.	2.8	163
141	Loss-of-function variants in endothelial lipase are a cause of elevated HDL cholesterol in humans. Journal of Clinical Investigation, 2009, 119, 1042-50.	3.9	162
142	Genomewide Linkage Analysis to Serum Creatinine, GFR, and Creatinine Clearance in a Community-Based Population: The Framingham Heart Study. Journal of the American Society of Nephrology: JASN, 2004, 15, 2457-2461.	3.0	161
143	Design of the Coronary ARtery Disease Genome-Wide Replication And Meta-Analysis (CARDIoGRAM) Study. Circulation: Cardiovascular Genetics, 2010, 3, 475-483.	5.1	159
144	Meta-analysis of gene-environment interaction: joint estimation of SNP and SNP \tilde{A} — environment regression coefficients. Genetic Epidemiology, 2011, 35, 11-18.	0.6	158

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145	Genome-wide physical activity interactions in adiposity ― A meta-analysis of 200,452 adults. PLoS Genetics, 2017, 13, e1006528.	1.5	158
146	Coronary risk associated with age and sex of parental heart disease in the Framingham Study. American Journal of Cardiology, 1989, 64, 555-559.	0.7	156
147	Assessing the contribution of rare variants to complex trait heritability from whole-genome sequence data. Nature Genetics, 2022, 54, 263-273.	9.4	156
148	Influence of the APOA5 locus on plasma triglyceride, lipoprotein subclasses, and CVD risk in the Framingham Heart Study. Journal of Lipid Research, 2004, 45, 2096-2105.	2.0	155
149	Framingham Heart Study 100K project: genome-wide associations for cardiovascular disease outcomes. BMC Medical Genetics, 2007, 8, S5.	2.1	155
150	Genome-wide association to body mass index and waist circumference: the Framingham Heart Study 100K project. BMC Medical Genetics, 2007, 8, S18.	2.1	154
151	High-normal blood pressure progression to hypertension in the Framingham Heart Study Hypertension, 1991, 17, 22-27.	1.3	153
152	Estimating risk curves for first-degree relatives of patients with Alzheimer's disease: The REVEAL study. Genetics in Medicine, 2004, 6, 192-196.	1.1	153
153	APOA2, Dietary Fat, and Body Mass Index. Archives of Internal Medicine, 2009, 169, 1897.	4.3	150
154	A Genome Scan for Modifiers of Age at Onset in Huntington Disease: The HD MAPS Study. American Journal of Human Genetics, 2003, 73, 682-687.	2.6	148
155	Effects of beer, wine, and liquor intakes on bone mineral density in older men and women. American Journal of Clinical Nutrition, 2009, 89, 1188-1196.	2.2	148
156	Diabetes and Deficits in Cortical Bone Density, Microarchitecture, and Bone Size: Framingham HR-pQCT Study. Journal of Bone and Mineral Research, 2018, 33, 54-62.	3.1	148
157	The relative importance of selected risk factors for various manifestations of cardiovascular disease among men and women from 35 to 64 years old: 30 years of follow-up in the Framingham Study. Circulation, 1987, 75, V65-73.	1.6	148
158	Common Variants in the Adiponectin Gene (<i>ADIPOQ</i>) Associated With Plasma Adiponectin Levels, Type 2 Diabetes, and Diabetes-Related Quantitative Traits. Diabetes, 2008, 57, 3353-3359.	0.3	147
159	Interaction of normal and expanded CAG repeat sizes influences age at onset of Huntington disease. , 2003, 119A, 279-282.		146
160	Dynamic incorporation of multiple in silico functional annotations empowers rare variant association analysis of large whole-genome sequencing studies at scale. Nature Genetics, 2020, 52, 969-983.	9.4	146
161	Absence of duplication of chromosome 21 genes in familial and sporadic Alzheimer's disease. Science, 1987, 238, 664-666.	6.0	145
162	Assessment of genetic risk for alzheimer's disease among first-degree relatives. Annals of Neurology, 1989, 25, 485-493.	2.8	145

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163	Risk variable clustering in the insulin resistance syndrome. The Framingham Offspring Study. Diabetes, 1997, 46, 1594-1600.	0.3	145
164	Genomewide Linkage Analysis of Body Mass Index across 28 Years of the Framingham Heart Study. American Journal of Human Genetics, 2002, 71, 1044-1050.	2.6	144
165	Meta-Analysis of Genome-Wide Scans Provides Evidence for Sex- and Site-Specific Regulation of Bone Mass. Journal of Bone and Mineral Research, 2006, 22, 173-183.	3.1	144
166	FTO genetic variants, dietary intake and body mass index: insights from 177 330 individuals. Human Molecular Genetics, 2014, 23, 6961-6972.	1.4	143
167	Associations between Vitamin K Biochemical Measures and Bone Mineral Density in Men and Women. Journal of Clinical Endocrinology and Metabolism, 2004, 89, 4904-4909.	1.8	142
168	Genetic Risk Assessment for Adult Children of People With Alzheimer's Disease: The Risk Evaluation and Education for Alzheimer's Disease (REVEAL) Study. Journal of Geriatric Psychiatry and Neurology, 2005, 18, 250-255.	1.2	142
169	Small Dense LDL Cholesterol and Coronary Heart Disease: Results from the Framingham Offspring Study. Clinical Chemistry, 2010, 56, 967-976.	1.5	142
170	Intake of Fruits, Vegetables, and Dairy Products in Early Childhood and Subsequent Blood Pressure Change. Epidemiology, 2005, 16, 4-11.	1.2	140
171	Deep-coverage whole genome sequences and blood lipids among 16,324 individuals. Nature Communications, 2018, 9, 3391.	5.8	140
172	Leukocyte Telomere Length and Carotid Artery Intimal Medial Thickness. Arteriosclerosis, Thrombosis, and Vascular Biology, 2008, 28, 1165-1171.	1.1	139
173	Optimizing the Tracking of Falls in Studies of Older Participants: Comparison of Quarterly Telephone Recall With Monthly Falls Calendars in the MOBILIZE Boston Study. American Journal of Epidemiology, 2010, 171, 1031-1036.	1.6	139
174	Candidate Gene Association Resource (CARe). Circulation: Cardiovascular Genetics, 2010, 3, 267-275.	5.1	139
175	Glycemic Index, Glycemic Load, and Cereal Fiber Intake and Risk of Type 2 Diabetes in US Black Women. Archives of Internal Medicine, 2007, 167, 2304.	4.3	138
176	Parental eating attitudes and the development of obesity in children. The Framingham Children's Study. International Journal of Obesity, 2000, 24, 1319-1325.	1.6	136
177	APOE, vascular pathology, and the AD brain. Neurology, 2005, 65, 259-265.	1.5	135
178	Low Plasma Vitamin B12 Is Associated With Lower BMD: The Framingham Osteoporosis Study. Journal of Bone and Mineral Research, 2005, 20, 152-158.	3.1	134
179	Association Between the PPARA L162V Polymorphism and Plasma Lipid Levels. Arteriosclerosis, Thrombosis, and Vascular Biology, 2002, 22, 805-810.	1.1	132
180	Whole-genome sequence–based analysis of high-density lipoprotein cholesterol. Nature Genetics, 2013, 45, 899-901.	9.4	132

#	Article	IF	CITATIONS
181	RISK FACTORS FOR LONG-TERM CORONARY PROGNOSIS AFTER INITIAL MYOCARDIAL INFARCTION: THE FRAMINGHAM STUDY. American Journal of Epidemiology, 1989, 130, 469-480.	1.6	130
182	Genome-wide association study for subclinical atherosclerosis in major arterial territories in the NHLBI's Framingham Heart Study. BMC Medical Genetics, 2007, 8, S4.	2.1	130
183	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
184	The Bsml Vitamin D Receptor Restriction Fragment Length Polymorphism (bb) Influences the Effect of Calcium Intake on Bone Mineral Density. Journal of Bone and Mineral Research, 1997, 12, 1049-1057.	3.1	129
185	Alcohol drinking determines the effect of the APOE locus on LDL-cholesterol concentrations in men: the Framingham Offspring Study. American Journal of Clinical Nutrition, 2001, 73, 736-745.	2.2	127
186	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
187	Right coronary artery stenosis: An independent predictor of atrial fibrillation after coronary artery bypass surgery. Journal of the American College of Cardiology, 1995, 25, 198-202.	1.2	126
188	Meta-analysis of genome-wide association studies confirms a susceptibility locus for knee osteoarthritis on chromosome 7q22. Annals of the Rheumatic Diseases, 2011, 70, 349-355.	0.5	126
189	Polyunsaturated Fatty Acids Interact with the PPARA-L162V Polymorphism to Affect Plasma Triglyceride and Apolipoprotein C-III Concentrations in the Framingham Heart Study. Journal of Nutrition, 2005, 135, 397-403.	1.3	123
190	Protective effects of fish intake and interactive effects of long-chain polyunsaturated fatty acid intakes on hip bone mineral density in older adults: the Framingham Osteoporosis Study. American Journal of Clinical Nutrition, 2011, 93, 1142-1151.	2.2	123
191	Disentangling the Genetic Determinants of Human Aging: Biological Age as an Alternative to the Use of Survival Measures. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2005, 60, 574-587.	1.7	122
192	Frailty and the Degradation of Complex Balance Dynamics During a Dual-Task Protocol. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2009, 64A, 1304-1311.	1.7	120
193	Insights into the genetic architecture of osteoarthritis from stage 1 of the arcOGEN study. Annals of the Rheumatic Diseases, 2011, 70, 864-867.	0.5	119
194	Genome Screen for Quantitative Trait Loci Contributing to Normal Variation in Bone Mineral Density: The Framingham Study. Journal of Bone and Mineral Research, 2002, 17, 1718-1727.	3.1	118
195	Evidence for the GluR6 gene associated with younger onset age of Huntington's disease. Neurology, 1999, 53, 1330-1330.	1.5	118
196	Identification of a Sudden Cardiac Death Susceptibility Locus at 2q24.2 through Genome-Wide Association in European Ancestry Individuals. PLoS Genetics, 2011, 7, e1002158.	1.5	117
197	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
198	Multiethnic genome-wide meta-analysis of ectopic fat depots identifies loci associated with adipocyte development and differentiation. Nature Genetics, 2017, 49, 125-130.	9.4	116

#	Article	IF	CITATIONS
199	Inverse association of carotenoid intakes with 4-y change in bone mineral density in elderly men and women: the Framingham Osteoporosis Study. American Journal of Clinical Nutrition, 2009, 89, 416-424.	2.2	115
200	Transmission and ageâ€atâ€onset patterns in familial Alzheimer's disease. Neurology, 1990, 40, 395-395.	1.5	115
201	Elevated Remnant-Like Particle Cholesterol and Triglyceride Levels in Diabetic Men and Women in the Framingham Offspring Study. Diabetes Care, 2002, 25, 989-994.	4.3	112
202	Plasma B Vitamins, Homocysteine, and Their Relation with Bone Loss and Hip Fracture in Elderly Men and Women. Journal of Clinical Endocrinology and Metabolism, 2008, 93, 2206-2212.	1.8	112
203	Multi-ancestry genome-wide gene–smoking interaction study of 387,272 individuals identifies new loci associated with serum lipids. Nature Genetics, 2019, 51, 636-648.	9.4	112
204	Acute effects of a high-fat meal with and without red wine on endothelial function in healthy subjects. American Journal of Cardiology, 1999, 84, 660-664.	0.7	111
205	Association of the Sst-I polymorphism at the APOC3 gene locus with variations in lipid levels, lipoprotein subclass profiles and coronary heart disease risk: the Framingham offspring study. Atherosclerosis, 2001, 158, 173-181.	0.4	111
206	Association of Leukocyte Telomere Length With Circulating Biomarkers of the Renin-Angiotensin-Aldosterone System. Circulation, 2008, 117, 1138-1144.	1.6	111
207	Enhanced Statistical Tests for GWAS in Admixed Populations: Assessment using African Americans from CARe and a Breast Cancer Consortium. PLoS Genetics, 2011, 7, e1001371.	1.5	110
208	Protective Effect of Total Carotenoid and Lycopene Intake on the Risk of Hip Fracture: A 17-Year Follow-Up From the Framingham Osteoporosis Study. Journal of Bone and Mineral Research, 2009, 24, 1086-1094.	3.1	109
209	Genetic Association for Renal Traits among Participants of African Ancestry Reveals New Loci for Renal Function. PLoS Genetics, 2011, 7, e1002264.	1.5	109
210	Genome-Wide Association of Body Fat Distribution in African Ancestry Populations Suggests New Loci. PLoS Genetics, 2013, 9, e1003681.	1.5	109
211	Genetic Variation at the Scavenger Receptor Class B Type I Gene Locus Determines Plasma Lipoprotein Concentrations and Particle Size and Interacts with Type 2 Diabetes: The Framingham Study. Journal of Clinical Endocrinology and Metabolism, 2003, 88, 2869-2879.	1.8	108
212	Dietary Intake of n-6 Fatty Acids Modulates Effect of Apolipoprotein A5 Gene on Plasma Fasting Triglycerides, Remnant Lipoprotein Concentrations, and Lipoprotein Particle Size. Circulation, 2006, 113, 2062-2070.	1.6	107
213	Genetic variation at the low-density lipoprotein receptor-related protein 5 (LRP5) locus modulates Wnt signaling and the relationship of physical activity with bone mineral density in men. Bone, 2007, 40, 587-596.	1.4	107
214	Association of Variants in <i>RETN</i> With Plasma Resistin Levels and Diabetes-Related Traits in the Framingham Offspring Study. Diabetes, 2009, 58, 750-756.	0.3	107
215	Association of Obesity With Mortality Over 24 Years of Weight History. JAMA Network Open, 2018, 1, e184587.	2.8	107
216	Multiple Loci Are Associated with White Blood Cell Phenotypes. PLoS Genetics, 2011, 7, e1002113.	1.5	106

#	Article	IF	CITATIONS
217	Association of clonal hematopoiesis with chronic obstructive pulmonary disease. Blood, 2022, 139, 357-368.	0.6	106
218	Effects of Race and Hypertension on Flow-Mediated and Nitroglycerin-Mediated Dilation of the Brachial Artery. Hypertension, 2001, 38, 1349-1354.	1.3	105
219	A Novel Aging Phenotype of Slow Gait, Impaired Executive Function, and Depressive Symptoms: Relationship to Blood Pressure and Other Cardiovascular Risks. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2009, 64A, 994-1001.	1.7	105
220	Subsensory vibrations to the feet reduce gait variability in elderly fallers. Gait and Posture, 2009, 30, 383-387.	0.6	104
221	Epigenetic Patterns in Blood Associated With Lipid Traits Predict Incident Coronary Heart Disease Events and Are Enriched for Results From Genome-Wide Association Studies. Circulation: Cardiovascular Genetics, 2017, 10, .	5.1	104
222	Association of the Câ^'514T Polymorphism in the Hepatic Lipase Gene With Variations in Lipoprotein Subclass Profiles. Arteriosclerosis, Thrombosis, and Vascular Biology, 2000, 20, 815-822.	1.1	103
223	Efficient Variant Set Mixed Model Association Tests for Continuous and Binary Traits in Large-Scale Whole-Genome Sequencing Studies. American Journal of Human Genetics, 2019, 104, 260-274.	2.6	103
224	Metacarpal Cortical Area and Risk of Coronary Heart Disease: The Framingham Study. American Journal of Epidemiology, 2004, 159, 589-595.	1.6	102
225	Who seeks genetic susceptibility testing for Alzheimer's disease? Findings from a multisite, randomized clinical trial. Genetics in Medicine, 2004, 6, 197-203.	1.1	101
226	Genome-wide search for genes affecting serum uric acid levels: the Framingham Heart Study. Metabolism: Clinical and Experimental, 2005, 54, 1435-1441.	1.5	101
227	Parental Occurrence of Premature Cardiovascular Disease Predicts Increased Coronary Artery and Abdominal Aortic Calcification in the Framingham Offspring and Third Generation Cohorts. Circulation, 2007, 116, 1473-1481.	1.6	101
228	Genome-wide association study for serum urate concentrations and gout among African Americans identifies genomic risk loci and a novel URAT1 loss-of-function allele. Human Molecular Genetics, 2011, 20, 4056-4068.	1.4	101
229	Remnant lipoprotein cholesterol and triglyceride reference ranges from the Framingham Heart Study. Clinical Chemistry, 1998, 44, 1224-1232.	1.5	100
230	Evidence for a gene influencing the TG/HDL-C ratio on chromosome 7q32.3-qter: a genome-wide scan in the Framingham Study. Human Molecular Genetics, 2000, 9, 1315-1320.	1.4	100
231	Polymorphisms in the Insulin-Degrading Enzyme Gene Are Associated With Type 2 Diabetes in Men From the NHLBI Framingham Heart Study. Diabetes, 2003, 52, 1562-1567.	0.3	100
232	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
233	Association Between Apolipoprotein E Genotype and Alzheimer Disease in African American Subjects. Archives of Neurology, 2002, 59, 594.	4.9	98
234	APOA5 gene variation modulates the effects of dietary fat intake on body mass index and obesity risk in the Framingham Heart Study. Journal of Molecular Medicine, 2007, 85, 119-128.	1.7	98

#	Article	IF	Citations
235	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
236	Discovery and fine-mapping of adiposity loci using high density imputation of genome-wide association studies in individuals of African ancestry: African Ancestry Anthropometry Genetics Consortium. PLoS Genetics, 2017, 13, e1006719.	1.5	98
237	Antihypertensive Therapy Increases Cerebral Blood Flow and Carotid Distensibility in Hypertensive Elderly Subjects. Hypertension, 2005, 45, 216-221.	1.3	97
238	Aging, Brain Disease, and Reserve: Implications for Delirium. American Journal of Geriatric Psychiatry, 2010, 18, 117-127.	0.6	97
239	Comparison of Techniques for Estimating Nutrient Intake: The Framingham Study. Epidemiology, 1992, 3, 171-177.	1.2	95
240	Power and type I error rate of false discovery rate approaches in genome-wide association studies. BMC Genetics, 2005, 6, S134.	2.7	95
241	Calcium intake is not associated with increased coronary artery calcification: the Framingham Study. American Journal of Clinical Nutrition, 2012, 96, 1274-1280.	2.2	95
242	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
243	Chapter 2 Genetics of the Framingham Heart Study Population. Advances in Genetics, 2008, 62, 33-65.	0.8	93
244	Protective effect of high protein and calcium intake on the risk of hip fracture in the framingham offspring cohort. Journal of Bone and Mineral Research, 2010, 25, 2770-2776.	3.1	93
245	Analysis commons, a team approach to discovery in a big-data environment for genetic epidemiology. Nature Genetics, 2017, 49, 1560-1563.	9.4	93
246	Age, gender, and body mass effects on quantitative trait loci for bone mineral density: the framingham studya~†. Bone, 2003, 33, 308-316.	1.4	91
247	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
248	Association of the A-204C polymorphism in the cholesterol $7\hat{l}$ ±-hydroxylase gene with variations in plasma low density lipoprotein cholesterol levels in the Framingham Offspring Study. Journal of Lipid Research, 1999, 40, 1883-1889.	2.0	91
249	Genomewide Linkage Analysis for Internal Carotid Artery Intimal Medial Thickness: Evidence for Linkage to Chromosome 12. American Journal of Human Genetics, 2004, 74, 253-261.	2.6	90
250	Long- and short-term risk of sudden coronary death. Circulation, 1992, 85, I11-8.	1.6	90
251	Alcohol Consumption and Risk of Breast Cancer: The Framingham Study Revisited. American Journal of Epidemiology, 1999, 149, 93-101.	1.6	89
252	Obesity Modulates the Association among <i>APOE</i> Genotype, Insulin, and Glucose in Men. Obesity, 2003, 11, 1502-1508.	4.0	89

#	Article	IF	CITATIONS
253	Congestive heart failure in patients with coronary artery disease: The gender paradox. American Heart Journal, 1997, 134, 207-212.	1.2	88
254	Head circumference, atrophy, and cognition. Neurology, 2010, 75, 137-142.	1.5	88
255	Small Dense Lowâ€Density Lipoprotein Cholesterol Is the Most Atherogenic Lipoprotein Parameter in the Prospective Framingham Offspring Study. Journal of the American Heart Association, 2021, 10, e019140.	1.6	88
256	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
257	Polymorphisms in the PON gene cluster are associated with Alzheimer disease. Human Molecular Genetics, 2006, 15, 77-85.	1.4	87
258	A 100K Genome-Wide Association Scan for Diabetes and Related Traits in the Framingham Heart Study: Replication and Integration With Other Genome-Wide Datasets. Diabetes, 2007, 56, 3063-3074.	0.3	87
259	Common genetic variation in multiple metabolic pathways influences susceptibility to low HDL-cholesterol and coronary heart disease. Journal of Lipid Research, 2010, 51, 3524-3532.	2.0	87
260	Genome scan for quantity of hand osteoarthritis: The Framingham study. Arthritis and Rheumatism, 2002, 46, 946-952.	6.7	86
261	Estrogen Receptor α Gene Variation Is Associated With Risk of Myocardial Infarction in More Than Seven Thousand Men From Five Cohorts. Circulation Research, 2006, 98, 590-592.	2.0	86
262	Long-term effects of serum cholesterol on bone mineral density in women and men: the Framingham Osteoporosis Study. Bone, 2004, 34, 557-561.	1.4	85
263	Multiancestry Genome-Wide Association Study of Lipid Levels Incorporating Gene-Alcohol Interactions. American Journal of Epidemiology, 2019, 188, 1033-1054.	1.6	85
264	Preexisting cardiovascular conditions and long-term prognosis after initial myocardial infarction: The Framingham Study. American Heart Journal, 1993, 125, 863-872.	1.2	84
265	Gene $\tilde{A}-$ dietary pattern interactions in obesity: analysis of up to 68 317 adults of European ancestry. Human Molecular Genetics, 2015, 24, 4728-4738.	1.4	84
266	Variation in estrogen-related genes and cross-sectional and longitudinal blood pressure in the Framingham Heart Study. Journal of Hypertension, 2005, 23, 2193-2200.	0.3	83
267	Reasons for Seeking Genetic Susceptibility Testing Among First-Degree Relatives of People With Alzheimer Disease. Alzheimer Disease and Associated Disorders, 2003, 17, 86-93.	0.6	82
268	Low Plasma Vitamin B12 Is Associated With Lower BMD: The Framingham Osteoporosis Study. Journal of Bone and Mineral Research, 2005, 20, 152-158.	3.1	82
269	Tracking of glycated hemoglobin in the original cohort of the framingham heart study. Journal of Clinical Epidemiology, 1996, 49, 411-417.	2.4	80
270	Genome-wide association with diabetes-related traits in the Framingham Heart Study. BMC Medical Genetics, 2007, 8, S16.	2.1	80

#	Article	IF	Citations
271	Genetics Analysis Workshop 16 Problem 2: the Framingham Heart Study data. BMC Proceedings, 2009, 3, S3.	1.8	80
272	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
273	Cerebrovascular hemodynamics, gait, and falls in an elderly population. Neurology, 2010, 74, 1627-1633.	1.5	80
274	Adiponectin: An independent risk factor for coronary heart disease in men in the Framingham offspring Study. Atherosclerosis, 2011, 217, 543-548.	0.4	80
275	Genetic Loci Influencing Lung Function. American Journal of Respiratory and Critical Care Medicine, 2002, 165, 795-799.	2.5	79
276	Evidence for Heritability of Abdominal Aortic Calcific Deposits in the Framingham Heart Study. Circulation, 2002, 106, 337-341.	1.6	79
277	A common polymorphism decreases low-density lipoprotein receptor exon 12 splicing efficiency and associates with increased cholesterol. Human Molecular Genetics, 2007, 16, 1765-1772.	1.4	79
278	Genome-wide Comparison of African-Ancestry Populations from CARe and Other Cohorts Reveals Signals of Natural Selection. American Journal of Human Genetics, 2011, 89, 368-381.	2.6	79
279	Deep coverage whole genome sequences and plasma lipoprotein(a) in individuals of European and African ancestries. Nature Communications, 2018, 9, 2606.	5.8	79
280	Association of the A-204C polymorphism in the cholesterol 7alpha-hydroxylase gene with variations in plasma low density lipoprotein cholesterol levels in the Framingham Offspring Study. Journal of Lipid Research, 1999, 40, 1883-9.	2.0	78
281	Association of APOE genotype with carotid atherosclerosis in men and women. Journal of Lipid Research, 2004, 45, 1868-1875.	2.0	77
282	Genetic Susceptibility Loci for Subtypes of Breast Cancer in an African American Population. Cancer Epidemiology Biomarkers and Prevention, 2013, 22, 127-134.	1.1	77
283	Dietary Patterns of Men and Women Suggest Targets for Health Promotion: The Framingham Nutrition Studies. American Journal of Health Promotion, 1996, 11, 42-52.	0.9	76
284	Parental Age at Child's Birth and Son's Risk of Prostate Cancer: The Framingham Study. American Journal of Epidemiology, 1999, 150, 1208-1212.	1.6	76
285	Performance of random forest when SNPs are in linkage disequilibrium. BMC Bioinformatics, 2009, 10, 78.	1.2	76
286	Dietary Intakes of Arachidonic Acid and α-Linolenic Acid Are Associated with Reduced Risk of Hip Fracture in Older Adults. Journal of Nutrition, 2011, 141, 1146-1153.	1.3	76
287	Association of a Common Polymorphism in the Methylenetetrahydrofolate Reductase (MTHFR) Gene With Bone Phenotypes Depends on Plasma Folate Status. Journal of Bone and Mineral Research, 2003, 19, 410-418.	3.1	75
288	Meta-Analysis Investigating Associations Between Healthy Diet and Fasting Glucose and Insulin Levels and Modification by Loci Associated With Glucose Homeostasis in Data From 15 Cohorts. American Journal of Epidemiology, 2013, 177, 103-115.	1.6	74

#	Article	IF	Citations
289	A principal component meta-analysis on multiple anthropometric traits identifies novel loci for body shape. Nature Communications, 2016, 7, 13357.	5.8	74
290	Lipoprotein(a) levels, apo(a) isoform size, and coronary heart disease risk in the Framingham Offspring Study. Journal of Lipid Research, 2011, 52, 1181-1187.	2.0	73
291	Genetics of coronary artery calcification among African Americans, a meta-analysis. BMC Medical Genetics, 2013, 14, 75.	2.1	73
292	Loss of Heterozygosity or Allele Imbalance in Histologically Normal Breast Epithelium Is Distinct from Loss of Heterozygosity or Allele Imbalance in Co-Existing Carcinomas. American Journal of Pathology, 2002, 161, 283-290.	1.9	72
293	A genome scan for joint-specific hand osteoarthritis susceptibility: The Framingham Study. Arthritis and Rheumatism, 2004, 50, 2489-2496.	6.7	72
294	Genome-wide significance for a modifier of age at neurological onset in Huntington's Disease at 6q23-24: the HD MAPS study. BMC Medical Genetics, 2006, 7, 71.	2.1	72
295	Vascular Calcification in Middle Age and Long-Term Risk of Hip Fracture: The Framingham Study. Journal of Bone and Mineral Research, 2007, 22, 1449-1454.	3.1	72
296	Validation of two-dimensional models for estimation of portion size in nutrition research. Journal of the American Dietetic Association, 1992, 92, 738-41.	1.3	72
297	Association Between Concurrent and Remote Blood Pressure and Disability in Older Adults. Hypertension, 2007, 50, 1026-1032.	1.3	71
298	Does dietary protein reduce hip fracture risk in elders? The Framingham osteoporosis study. Osteoporosis International, 2011, 22, 345-349.	1.3	71
299	Interactions of Interleukin-6 Promoter Polymorphisms With Dietary and Lifestyle Factors and Their Association With Bone Mass in Men and Women From the Framingham Osteoporosis Study. Journal of Bone and Mineral Research, 2004, 19, 552-559.	3.1	70
300	The Significant Increase in Cardiovascular Disease Risk inAPOEÉ, 4 Carriers is Evident Only in Men Who Smoke: Potential Relationship Between Reduced Antioxidant Status and ApoE4. Annals of Human Genetics, 2005, 69, 613-622.	0.3	70
301	Correlation between Waardenburg syndrome phenotype and genotype in a population of individuals with identified PAX3 mutations. Human Genetics, 1998, 102, 499-506.	1.8	69
302	Nonsteroidal anti-inflammatory drug use and Alzheimer's disease risk: the MIRAGE Study. BMC Geriatrics, 2005, 5, 2.	1.1	69
303	Leptin Receptor Polymorphisms Interact with Polyunsaturated Fatty Acids to Augment Risk of Insulin Resistance and Metabolic Syndrome in Adults. Journal of Nutrition, 2010, 140, 238-244.	1.3	69
304	"l know what you told me, but this is what I think:―Perceived risk of Alzheimer disease among individuals who accurately recall their genetics-based risk estimate. Genetics in Medicine, 2010, 12, 219-227.	1.1	69
305	A genetic risk score based on direct associations with coronary heart disease improves coronary heart disease risk prediction in the Atherosclerosis Risk in Communities (ARIC), but not in the Rotterdam and Framingham Offspring, Studies. Atherosclerosis, 2012, 223, 421-426.	0.4	69
306	Discovery and Fine Mapping of Serum Protein Loci through Transethnic Meta-analysis. American Journal of Human Genetics, 2012, 91, 744-753.	2.6	69

#	Article	IF	Citations
307	Consumption of meat is associated with higher fasting glucose and insulin concentrations regardless of glucose and insulin genetic risk scores: a meta-analysis of 50,345 Caucasians. American Journal of Clinical Nutrition, 2015, 102, 1266-1278.	2.2	69
308	Genome-Wide Association Study for Incident Myocardial Infarction and Coronary Heart Disease in Prospective Cohort Studies: The CHARGE Consortium. PLoS ONE, 2016, 11, e0144997.	1.1	69
309	A high-resolution HLA reference panel capturing global population diversity enables multi-ancestry fine-mapping in HIV host response. Nature Genetics, 2021, 53, 1504-1516.	9.4	69
310	Issues in Conducting Epidemiologic Research Among Elders: Lessons From The MOBILIZE Boston Study. American Journal of Epidemiology, 2008, 168, 1444-1451.	1.6	68
311	Risk Factors for Hospitalization Among Community-Dwelling Primary Care Older Patients. Medical Care, 2008, 46, 726-731.	1.1	68
312	Estrogen Receptor \hat{l}^2 Polymorphisms Are Associated With Bone Mass in Women and Men: The Framingham Study. Journal of Bone and Mineral Research, 2003, 19, 773-781.	3.1	67
313	Evidence for a modifier of onset age in Huntington disease linked to the HD gene in 4p16. Neurogenetics, 2004, 5, 109-114.	0.7	67
314	The Relationship Between Aldosterone, Oxidative Stress, and Inflammation in Chronic, Stable Human Heart Failure. Journal of Cardiac Failure, 2006, 12, 122-127.	0.7	67
315	Matrix Gla Protein Is Associated With Risk Factors for Atherosclerosis but not With Coronary Artery Calcification. Arteriosclerosis, Thrombosis, and Vascular Biology, 2006, 26, 2769-2774.	1.1	67
316	Gain-of-Function Lipoprotein Lipase Variant rs13702 Modulates Lipid Traits through Disruption of a MicroRNA-410 Seed Site. American Journal of Human Genetics, 2013, 92, 5-14.	2.6	67
317	Genetically abnormal clones in histologically normal breast tissue. American Journal of Pathology, 1998, 152, 1591-8.	1.9	67
318	Circulating CD34+ progenitor cell frequency is associated with clinical and genetic factors. Blood, 2013, 121, e50-e56.	0.6	65
319	Integromic Analysis of Genetic Variation and Gene Expression Identifies Networks for Cardiovascular Disease Phenotypes. Circulation, 2015, 131, 536-549.	1.6	65
320	Segregation analysis reveals evidence of a major gene for Alzheimer disease. American Journal of Human Genetics, 1991, 48, 1026-33.	2.6	65
321	Genome-wide association and functional studies identify a role for matrix Gla protein in osteoarthritis of the hand. Annals of the Rheumatic Diseases, 2017, 76, 2046-2053.	0.5	64
322	Multi-ancestry study of blood lipid levels identifies four loci interacting with physical activity. Nature Communications, 2019, 10, 376.	5.8	64
323	Variants at the APOA5 locus, association with carotid atherosclerosis, and modification by obesity: the Framingham Study. Journal of Lipid Research, 2006, 47, 990-996.	2.0	63
324	Hip structural geometry in old and old-old age: Similarities and differences between men and women. Bone, 2007, 41, 722-732.	1.4	63

#	Article	IF	CITATIONS
325	Overall and Central Obesity and Risk of Type 2 Diabetes in U.S. Black Women*. Obesity, 2007, 15, 1860-1866.	1.5	63
326	Association of Leukocyte Telomere Length With Echocardiographic Left Ventricular Mass. Circulation, 2009, 120, 1195-1202.	1.6	63
327	<i>PCSK9</i> Loss-of-Function Variants, Low-Density Lipoprotein Cholesterol, and Risk of Coronary Heart Disease and Stroke. Circulation: Cardiovascular Genetics, 2017, 10, e001632.	5.1	63
328	Abdominal aortic calcific deposits are associated with increased risk for congestive heart failure: The Framingham Heart Study. American Heart Journal, 2002, 144, 733-739.	1.2	62
329	Bias due to missing exposure data using complete-case analysis in the proportional hazards regression model. Statistics in Medicine, 2003, 22, 545-557.	0.8	62
330	Prediction of Intermittent Claudication, Ischemic Stroke, and Other Cardiovascular Disease by Detection of Abdominal Aortic Calcific Deposits by Plain Lumbar Radiographs. American Journal of Cardiology, 2008, 101, 326-331.	0.7	62
331	Plasma Levels of HDL Subpopulations and Remnant Lipoproteins Predict the Extent of Angiographically-Defined Coronary Artery Disease in Postmenopausal Women. Arteriosclerosis, Thrombosis, and Vascular Biology, 2008, 28, 575-579.	1.1	62
332	Effect of Alzheimer disease genetic risk disclosure on dietary supplement use. American Journal of Clinical Nutrition, 2010, 91, 1402-1407.	2.2	62
333	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
334	Evidence of cortical metabolic dysfunction in early Huntington's disease by single-photon-emission computed tomography. Movement Disorders, 1996, 11, 671-677.	2.2	61
335	Mapping of Quantitative Ultrasound of the Calcaneus Bone to Chromosome 1 by Genome-Wide Linkage Analysis. Osteoporosis International, 2002, 13, 796-802.	1.3	60
336	Sex-Specific Association between Estrogen Receptor-α Gene Variation and Measures of Adiposity: The Framingham Heart Study. Journal of Clinical Endocrinology and Metabolism, 2005, 90, 6257-6262.	1.8	60
337	Association of Estrogen Receptor \hat{l}^2 Gene Polymorphisms With Left Ventricular Mass and Wall Thickness in Women. American Journal of Hypertension, 2005, 18, 1388-1395.	1.0	60
338	Maximizing Clinical Research Participation in Vulnerable Older Persons: Identification of Barriers and Motivators. Journal of the American Geriatrics Society, 2008, 56, 1522-1527.	1.3	60
339	Association of Distinct Variants in SORL1 With Cerebrovascular and Neurodegenerative Changes Related to Alzheimer Disease. Archives of Neurology, 2008, 65, 1640.	4.9	60
340	Trans-ethnic meta-analysis of white blood cell phenotypes. Human Molecular Genetics, 2014, 23, 6944-6960.	1.4	60
341	A fully adjusted twoâ€stage procedure for rankâ€normalization in genetic association studies. Genetic Epidemiology, 2019, 43, 263-275.	0.6	60
342	Complement component 3 polymorphisms interact with polyunsaturated fatty acids to modulate risk of metabolic syndrome. American Journal of Clinical Nutrition, 2009, 90, 1665-1673.	2.2	59

#	Article	IF	CITATIONS
343	Visceral Adipose Tissue Is Associated With Bone Microarchitecture in the Framingham Osteoporosis Study. Journal of Bone and Mineral Research, 2017, 32, 143-150.	3.1	59
344	Genome screen for a combined bone phenotype using principal component analysis: the Framingham study. Bone, 2004, 34, 547-556.	1.4	58
345	Resequencing and Clinical Associations of the 9p21.3 Region. Circulation, 2013, 127, 799-810.	1.6	58
346	Sequence Kernel Association Test for Survival Traits. Genetic Epidemiology, 2014, 38, 191-197.	0.6	58
347	Age-adjusted survival curves with application in the Framingham study. Statistics in Medicine, 1995, 14, 1731-1744.	0.8	57
348	<i>PPARG</i> by Dietary Fat Interaction Influences Bone Mass in Mice and Humans. Journal of Bone and Mineral Research, 2008, 23, 1398-1408.	3.1	56
349	Identification of homogeneous genetic architecture of multiple genetically correlated traits by block clustering of genome-wide associations. Journal of Bone and Mineral Research, 2011, 26, 1261-1271.	3.1	56
350	Genetic variation associated with circulating monocyte count in the eMERGE Network. Human Molecular Genetics, 2013, 22, 2119-2127.	1.4	56
351	A Model for Predicting the Risk of Cancer Consequent to Retroviral Gene Therapy. Human Gene Therapy, 1992, 3, 479-486.	1.4	55
352	Alcohol Consumption and Risk of Intermittent Claudication in the Framingham Heart Study. Circulation, 2000, 102, 3092-3097.	1.6	55
353	Multiancestry Study of Gene–Lifestyle Interactions for Cardiovascular Traits in 610 475 Individuals From 124 Cohorts. Circulation: Cardiovascular Genetics, 2017, 10, .	5.1	55
354	Meta-analysis of loci associated with age at natural menopause in African-American women. Human Molecular Genetics, 2014, 23, 3327-3342.	1.4	54
355	Multiethnic Exome-Wide Association Study of Subclinical Atherosclerosis. Circulation: Cardiovascular Genetics, 2016, 9, 511-520.	5.1	54
356	Comparison of Metered-Dose Inhaler Attached to an Aerochamber with an Updraft Nebulizer for the Administration of Metaproterenol in Hospitalized Patients. Journal of Asthma, 1985, 22, 87-92.	0.9	53
357	Absence of association between genetic variation in the promoter of the microsomal triglyceride transfer protein gene and plasma lipoproteins in the Framingham Offspring Study. Atherosclerosis, 2000, 148, 337-343.	0.4	53
358	Quantitative analysis of allele imbalance supports atypical ductal hyperplasia lesions as direct breast cancer precursors. Journal of Pathology, 2006, 209, 307-316.	2.1	53
359	Gene-nutrient interactions with dietary fat modulate the association between genetic variation of the ACSL1 gene and metabolic syndrome. Journal of Lipid Research, 2010, 51, 1793-1800.	2.0	53
360	Assessment of cortical and striatal involvement in 523 Huntington disease brains. Neurology, 2012, 79, 1708-1715.	1.5	52

#	Article	IF	CITATIONS
361	Genome-wide association study of age at menarche in African-American women. Human Molecular Genetics, 2013, 22, 3329-3346.	1.4	52
362	Usefulness of fibrinogenolytic and procoagulant markers during thrombolytic therapy in predicting clinical outcomes in acute myocardial infarction. American Journal of Cardiology, 1996, 78, 503-510.	0.7	51
363	Genetic Variability of Adult Body Mass Index: A Longitudinal Assessment in Framingham Families. Obesity, 2002, 10, 675-681.	4.0	51
364	Effect of serum albumin and bilirubin on the risk of myocardial infarction (the Framingham Offspring) Tj ETQq0 0	0 rgBT /O	verlock 10 Tf
365	The PLIN4 Variant rs8887 Modulates Obesity Related Phenotypes in Humans through Creation of a Novel miR-522 Seed Site. PLoS ONE, 2011, 6, e17944.	1.1	51
366	Epidemiology of sudden coronary death: population at risk. Canadian Journal of Cardiology, 1990, 6, 439-44.	0.8	51
367	Evidence for a Gene Influencing Serum Bilirubin on Chromosome 2q Telomere: A Genomewide Scan in the Framingham Study. American Journal of Human Genetics, 2003, 72, 1029-1034.	2.6	50
368	Genome-wide pleiotropy of osteoporosis-related phenotypes: The framingham study. Journal of Bone and Mineral Research, 2010, 25, 1555-1563.	3.1	50
369	Polymorphisms in the <i>TOX3/LOC643714</i> Locus and Risk of Breast Cancer in African-American Women. Cancer Epidemiology Biomarkers and Prevention, 2010, 19, 1320-1327.	1.1	50
370	QCT measures of bone strength at the thoracic and lumbar spine: The Framingham study. Journal of Bone and Mineral Research, 2012, 27, 654-663.	3.1	50
371	A comparison of time dependent Cox regression, pooled logistic regression and cross sectional pooling with simulations and an application to the Framingham Heart Study. BMC Medical Research Methodology, 2016, 16, 148.	1.4	50
372	Inverse relationship between age at onset of Huntington disease and paternal age suggests involvement of genetic imprinting. American Journal of Human Genetics, 1992, 50, 528-35.	2.6	50
373	Secular Trends in Diet and Risk Factors for Cardiovascular Disease. Journal of the American Dietetic Association, 1995, 95, 171-179.	1.3	49
374	Determinants of Progression from Microalbuminuria to Proteinuria in Patients Who Have Type 1 Diabetes and Are Treated with Angiotensin-Converting Enzyme Inhibitors. Clinical Journal of the American Society of Nephrology: CJASN, 2007, 2, 461-469.	2.2	49
375	Proximal hip geometry is linked to several chromosomal regions: Genome-wide linkage results from the Framingham Osteoporosis Study. Bone, 2007, 40, 743-750.	1.4	49
376	Analysis of loss-of-function variants and 20 risk factor phenotypes in 8,554 individuals identifies loci influencing chronic disease. Nature Genetics, 2015, 47, 640-642.	9.4	49
377	Rare and Coding Region Genetic Variants Associated With Risk of Ischemic Stroke. JAMA Neurology, 2015, 72, 781.	4.5	49
378	Prediction of risk for hemodynamic compromise during percutaneous transluminal coronary angioplasty. American Journal of Cardiology, 1992, 70, 1540-1545.	0.7	48

#	Article	IF	Citations
379	Statin use and the risk of Alzheimer's disease: The MIRAGE Study. , 2006, 2, 96-103.		48
380	Estrogen receptor- $\hat{l}\pm$ variants are associated with lipoprotein size distribution and particle levels in women: The Framingham Heart Study. Atherosclerosis, 2006, 185, 210-218.	0.4	48
381	Estimating the probability of de novo HD cases from transmissions of expanded penetrant CAG alleles in the Huntington disease gene from male carriers of high normal alleles (27–35 CAG). American Journal of Medical Genetics, Part A, 2009, 149A, 1375-1381.	0.7	48
382	Additive Effect of Polymorphisms in the IL-6, LTA, and TNF-α Genes and Plasma Fatty Acid Level Modulate Risk for the Metabolic Syndrome and Its Components. Journal of Clinical Endocrinology and Metabolism, 2010, 95, 1386-1394.	1.8	48
383	Genome-Wide Association of Pericardial Fat Identifies a Unique Locus for Ectopic Fat. PLoS Genetics, 2012, 8, e1002705.	1.5	48
384	The normal Huntington disease (HD) allele, or a closely linked gene, influences age at onset of HD. American Journal of Human Genetics, 1993, 53, 125-30.	2.6	48
385	Comparison of Alzheimer's disease risk factors in white and African American families. Neurology, 2003, 60, 1372-1374.	1.5	47
386	Assessment of gene-by-sex interaction effect on bone mineral density. Journal of Bone and Mineral Research, 2012, 27, 2051-2064.	3.1	47
387	Higher Magnesium Intake Is Associated with Lower Fasting Glucose and Insulin, with No Evidence of Interaction with Select Genetic Loci, in a Meta-Analysis of 15 CHARGE Consortium Studies. Journal of Nutrition, 2013, 143, 345-353.	1.3	47
388	A genome scan for loci linked to quantitative insulin traits in persons without diabetes: the Framingham Offspring Study. Diabetologia, 2003, 46, 579-587.	2.9	46
389	Genetic Contribution to Biological Aging: The Framingham Study. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2004, 59, B218-B226.	1.7	46
390	Identification of Novel Candidate Genes for Alzheimer's Disease by Autozygosity Mapping using Genome Wide SNP Data. Journal of Alzheimer's Disease, 2011, 23, 349-359.	1.2	46
391	Multi-Ethnic Analysis of Lipid-Associated Loci: The NHLBI CARe Project. PLoS ONE, 2012, 7, e36473.	1.1	46
392	Development of a unidimensional composite measure of neuropsychological functioning in older cardiac surgery patients with good measurement precision. Journal of Clinical and Experimental Neuropsychology, 2010, 32, 1041-1049.	0.8	45
393	Genome of the Netherlands population-specific imputations identify an ABCA6 variant associated with cholesterol levels. Nature Communications, 2015, 6, 6065.	5.8	45
394	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
395	Genome-wide linkage analysis to urinary microalbuminuria in a community-based sample: The Framingham Heart Study. Kidney International, 2005, 67, 70-74.	2.6	44
396	The Insulin Gene Variable Number Tandem Repeat and Risk of Type 2 Diabetes in a Population-Based Sample of Families and Unrelated Men and Women. Journal of Clinical Endocrinology and Metabolism, 2005, 90, 1137-1143.	1.8	44

#	Article	IF	CITATIONS
397	The RAGE Gly82Ser polymorphism is not associated with cardiovascular disease in the Framingham offspring study. Atherosclerosis, 2005, 182, 301-305.	0.4	44
398	Genome-wide meta-analysis of macronutrient intake of 91,114 European ancestry participants from the cohorts for heart and aging research in genomic epidemiology consortium. Molecular Psychiatry, 2019, 24, 1920-1932.	4.1	44
399	Impact of Rare and Common Genetic Variants on Diabetes Diagnosis by Hemoglobin A1c in Multi-Ancestry Cohorts: The Trans-Omics for Precision Medicine Program. American Journal of Human Genetics, 2019, 105, 706-718.	2.6	44
400	Association of the A/T54 polymorphism in the intestinal fatty acid binding protein with variations in plasma lipids in The Framingham Offspring Study. Atherosclerosis, 2001, 159, 417-424.	0.4	43
401	A Genome-Wide Scan of Pulmonary Function Measures in the National Heart, Lung, and Blood Institute Family Heart Study. American Journal of Respiratory and Critical Care Medicine, 2003, 167, 1528-1533.	2.5	43
402	Heritability of prevalent vertebral fracture and volumetric bone mineral density and geometry at the lumbar spine in three generations of the framingham study. Journal of Bone and Mineral Research, 2012, 27, 954-958.	3.1	43
403	Haplotype Structure of the <i>ENPP1</i> Gene and Nominal Association of the K121Q Missense Single Nucleotide Polymorphism With Glycemic Traits in the Framingham Heart Study. Diabetes, 2008, 57, 1971-1977.	0.3	42
404	Education Attenuates the Effect of Medial Temporal Lobe Atrophy on Cognitive Function in Alzheimer's Disease: The MIRAGE Study. Journal of Alzheimer's Disease, 2009, 17, 855-862.	1.2	42
405	QCT Volumetric Bone Mineral Density and Vascular and Valvular Calcification: The Framingham Study. Journal of Bone and Mineral Research, 2015, 30, 1767-1774.	3.1	42
406	Novel Genetic Variants Associated With Increased Vertebral Volumetric BMD, Reduced Vertebral Fracture Risk, and Increased Expression of $\langle i \rangle SLC1A3 \langle j \rangle$ and $\langle i \rangle EPHB2 \langle j \rangle$. Journal of Bone and Mineral Research, 2016, 31, 2085-2097.	3.1	42
407	Association of the Ala54-thr Polymorphism in the Intestinal Fatty Acid-Binding Protein With 2-h Postchallenge Insulin Levels in the Framingham Offspring Study. Diabetes Care, 2001, 24, 1161-1166.	4.3	41
408	LpA-I, LpA-I:A-II HDL and CHD-risk: The Framingham Offspring Study and the Veterans Affairs HDL Intervention Trial. Atherosclerosis, 2006, 188, 59-67.	0.4	41
409	Vertebral Size, Bone Density, and Strength in Men and Women Matched for Age and Areal Spine BMD. Journal of Bone and Mineral Research, 2014, 29, 562-569.	3.1	41
410	Multi-ethnic fine-mapping of 14 central adiposity loci. Human Molecular Genetics, 2014, 23, 4738-4744.	1.4	41
411	Interrater agreement for diagnosis of Alzheimer's disease. Neurology, 1994, 44, 652-652.	1.5	40
412	Gene-Centric Meta-Analysis of Lipid Traits in African, East Asian and Hispanic Populations. PLoS ONE, 2012, 7, e50198.	1.1	40
413	Allele Imbalance, or Loss of Heterozygosity, in Normal Breast Epithelium of Sporadic Breast Cancer Cases and BRCA1 Gene Mutation Carriers Is Increased Compared With Reduction Mammoplasty Tissues. Journal of Clinical Oncology, 2005, 23, 8613-8619.	0.8	39
414	Comparing testâ€specific distress of susceptibility versus deterministic genetic testing for Alzheimer's disease. Alzheimer's and Dementia, 2008, 4, 406-413.	0.4	39

#	Article	IF	Citations
415	Conditional linkage and genome-wide association studies identify UGT1A1 as a major gene for anti-atherogenic serum bilirubin levelsâ€"The Framingham Heart Study. Atherosclerosis, 2009, 206, 228-233.	0.4	39
416	Loss-of-function genomic variants highlight potential therapeutic targets for cardiovascular disease. Nature Communications, 2020, 11, 6417.	5.8	39
417	Predictors of medical care utilization by independently living adults with spinal cord injuries. Archives of Physical Medicine and Rehabilitation, 1989, 70, 471-476.	0.5	37
418	Association of Decreased Paternal Age and Late-Onset Alzheimer's Disease. Archives of Neurology, 1991, 48, 599.	4.9	37
419	Evidence for linkage of red blood cell size and count: Genome-wide scans in the Framingham Heart Study. American Journal of Hematology, 2007, 82, 605-610.	2.0	37
420	Generalized estimating equations for genomeâ€wide association studies using longitudinal phenotype data. Statistics in Medicine, 2015, 34, 118-130.	0.8	37
421	Abdominal Aortic Calcification and Exostoses at the Hand and Lumbar Spine: The Framingham Study. Calcified Tissue International, 2006, 78, 1-8.	1.5	36
422	Incorporating ethnicity into genetic risk assessment for Alzheimer disease: the REVEAL study experience. Genetics in Medicine, 2008, 10, 207-214.	1.1	36
423	Dietary Acid Load Is Not Associated with Lower Bone Mineral Density Except in Older Men,. Journal of Nutrition, 2011, 141, 588-594.	1.3	36
424	Relations of Long-Term and Contemporary Lipid Levels and Lipid Genetic Risk Scores With Coronary Artery Calcium in the Framingham Heart Study. Journal of the American College of Cardiology, 2012, 60, 2364-2371.	1.2	36
425	A genome-wide copy number association study of osteoporotic fractures points to the 6p25.1 locus. Journal of Medical Genetics, 2014, 51, 122-131.	1.5	36
426	A randomized controlled trial of disclosing genetic risk information for Alzheimer disease via telephone. Genetics in Medicine, 2018, 20, 132-141.	1.1	36
427	Mendelian randomization supports bidirectional causality between telomere length and clonal hematopoiesis of indeterminate potential. Science Advances, 2022, 8, eabl6579.	4.7	36
428	Genetic transmission of Alzheimer's disease among families in a Dutch population based study Journal of Medical Genetics, 1993, 30, 640-646.	1.5	35
429	Population Nutrient Intake Approaches Dietary Recommendations. Journal of the American Dietetic Association, 1997, 97, 742-749.	1.3	35
430	Bias due to two-stage residual-outcome regression analysis in genetic association studies. Genetic Epidemiology, 2011, 35, 592-596.	0.6	35
431	Whole Exome Sequencing in Atrial Fibrillation. PLoS Genetics, 2016, 12, e1006284.	1.5	35
432	Apolipoprotein E isoform polymorphisms are not associated with insulin resistance: the Framingham Offspring Study. Diabetes Care, 2000, 23, 669-674.	4.3	34

#	Article	IF	CITATIONS
433	Linkage and association with pulmonary function measures on chromosome 6q27 in the Framingham Heart Study. Human Molecular Genetics, 2003, 12, 2745-2751.	1.4	34
434	Genome-wide linkage analysis to age at natural menopause in a community-based sample: the Framingham Heart Study. Fertility and Sterility, 2005, 84, 1674-1679.	0.5	34
435	The Type 2 Deiodinase (DIO2) A/G Polymorphism Is Not Associated with Glycemic Traits: The Framingham Heart Study. Thyroid, 2007, 17, 199-202.	2.4	34
436	Variation in Estrogen-Related Genes Associated with Cardiovascular Phenotypes and Circulating Estradiol, Testosterone, and Dehydroepiandrosterone Sulfate Levels. Journal of Clinical Endocrinology and Metabolism, 2008, 93, 2779-2785.	1.8	34
437	Associations of APOE gene polymorphisms with bone mineral density and fracture risk: a meta-analysis. Osteoporosis International, 2011, 22, 1199-1209.	1.3	34
438	Plasma phosphatidylcholine concentrations of polyunsaturated fatty acids are differentially associated with hip bone mineral density and hip fracture in older adults: The framingham osteoporosis study. Journal of Bone and Mineral Research, 2012, 27, 1222-1230.	3.1	34
439	Meta-analysis of 49â€549 individuals imputed with the 1000 Genomes Project reveals an exonic damaging variant in⟨i⟩ANGPTL4⟨ i⟩determining fasting TG levels. Journal of Medical Genetics, 2016, 53, 441-449.	1.5	34
440	Disclosing Pleiotropic Effects During Genetic Risk Assessment for Alzheimer Disease. Annals of Internal Medicine, 2016, 164, 155.	2.0	34
441	Whole genome sequence analyses of brain imaging measures in the Framingham Study. Neurology, 2018, 90, e188-e196.	1.5	34
442	Discovery of Genetic Variation on Chromosome 5q22 Associated with Mortality in Heart Failure. PLoS Genetics, 2016, 12, e1006034.	1.5	34
443	Influence of Alcohol Dehydrogenase 1C Polymorphism on the Alcohol—Cardiovascular Disease Association (from the Framingham Offspring Study). American Journal of Cardiology, 2005, 96, 227-232.	0.7	33
444	The Effect of Specific Medical Conditions on Functional Decline. Journal of the American Geriatrics Society, 1997, 45, 1459-1463.	1.3	32
445	Tobacco smoking, estrogen receptor $\hat{l}\pm$ gene variation and small low density lipoprotein level. Human Molecular Genetics, 2005, 14, 2405-2413.	1.4	32
446	Bivariate Genome-Wide Linkage Analysis of Femoral Bone Traits and Leg Lean Mass: Framingham Study. Journal of Bone and Mineral Research, 2009, 24, 710-718.	3.1	32
447	Mining the LIPG Allelic Spectrum Reveals the Contribution of Rare and Common Regulatory Variants to HDL Cholesterol. PLoS Genetics, 2011, 7, e1002393.	1.5	32
448	Serum paraoxonase activity is associated with variants in the PON gene cluster and risk of Alzheimer disease. Neurobiology of Aging, 2012, 33, 1015.e7-1015.e23.	1.5	32
449	Genome-wide association study for radiographic vertebral fractures: A potential role for the 16q24 BMD locus. Bone, 2014, 59, 20-27.	1.4	32
450	Sugar-sweetened beverage intake associations with fasting glucose and insulin concentrations are not modified by selected genetic variants in a ChREBP-FGF21 pathway: a meta-analysis. Diabetologia, 2018, 61, 317-330.	2.9	32

#	Article	IF	Citations
451	Whole genome sequence analysis of pulmonary function and COPD in 19,996 multi-ethnic participants. Nature Communications, 2020, 11, 5182.	5.8	32
452	Elderly Cohort Study Subjects Unable to Return for Follow-up Have Lower Bone Mass than Those Who Can Return. American Journal of Epidemiology, 2000, 151, 689-692.	1.6	31
453	Can Metacarpal Cortical Area Predict the Occurrence of Hip Fracture in Women and Men Over 3 Decades of Follow-Up? Results From the Framingham Osteoporosis Study. Journal of Bone and Mineral Research, 2001, 16, 2260-2266.	3.1	31
454	A multi-ancestry genome-wide study incorporating gene–smoking interactions identifies multiple new loci for pulse pressure and mean arterial pressure. Human Molecular Genetics, 2019, 28, 2615-2633.	1.4	31
455	Reliability of Information Collected by Proxy in Family Studies of Alzheimer's Disease. Neuroepidemiology, 2001, 20, 105-111.	1.1	30
456	BMI Modifies Associations of ILâ€6 Genotypes with Insulin Resistance: The Framingham Study. Obesity, 2006, 14, 1454-1461.	1.5	30
457	Direct assessment of plasma low density lipoprotein and high density lipoprotein cholesterol levels and coronary heart disease: Results from the Framingham Offspring Study. Atherosclerosis, 2010, 213, 251-255.	0.4	30
458	Risk Factors for Nonadherence with Pap Testing in HIV-Infected Women. Journal of Women's Health, 2011, 20, 1635-1643.	1.5	30
459	Heritability and Genetic Correlations for Bone Microarchitecture: The Framingham Study Families. Journal of Bone and Mineral Research, 2017, 32, 106-114.	3.1	30
460	A Longitudinal Study of Trunk Muscle Properties and Severity of Thoracic Kyphosis in Women and Men: The Framingham Study. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2019, 74, 420-427.	1.7	30
461	Bone Mass and the Risk of Colon Cancer among Postmenopausal Women. American Journal of Epidemiology, 2001, 153, 31-37.	1.6	29
462	A genome-wide search for genes affecting circulating fibrinogen levels in the Framingham Heart Study. Thrombosis Research, 2003, 110, 57-64.	0.8	29
463	Variants in the <i>CNR1</i> and the <i>FAAH</i> Genes and Adiposity Traits in the Community. Obesity, 2009, 17, 755-760.	1.5	29
464	MHC region and risk of systemic lupus erythematosus in African American women. Human Genetics, 2011, 130, 807-815.	1.8	29
465	A System for Phenotype Harmonization in the National Heart, Lung, and Blood Institute Trans-Omics for Precision Medicine (TOPMed) Program. American Journal of Epidemiology, 2021, 190, 1977-1992.	1.6	29
466	Genome sequencing unveils a regulatory landscape of platelet reactivity. Nature Communications, 2021, 12, 3626.	5.8	29
467	Detection of monoclonal microsatellite alterations in atypical breast hyperplasia Journal of Clinical Investigation, 1996, 98, 1095-1100.	3.9	29
468	Genetic determinants of telomere length from 109,122 ancestrally diverse whole-genome sequences in TOPMed. Cell Genomics, 2022, 2, 100084.	3.0	29

#	Article	IF	CITATIONS
469	Diet and plasma lipids in women. II. Macronutrients and plasma triglycerides, high-density lipoprotein, and the ratio of total to high-density lipoprotein cholesterol in women: The Framingham Nutrition Studies. Journal of Clinical Epidemiology, 1996, 49, 665-672.	2.4	28
470	Genetic Predisposition to Stroke in Relatives of Hypertensives. Stroke, 2000, 31, 487-492.	1.0	28
471	Genetic association between endothelial nitric oxide synthase and Alzheimer disease. Clinical Genetics, 2006, 70, 49-56.	1.0	28
472	PAI-1 Gene 4G/5G Polymorphism and Risk of Type 2 Diabetes in a Population-based Sample*. Obesity, 2006, 14, 753-758.	1.5	28
473	Heritability of Magnetic Resonance Imaging (MRI) Traits in Alzheimer Disease Cases and Their Siblings in the MIRAGE Study. Alzheimer Disease and Associated Disorders, 2007, 21, 85-91.	0.6	28
474	Bivariate Linkage Study of Proximal Hip Geometry and Body Size Indices: The Framingham Study. Calcified Tissue International, 2007, 81, 162-173.	1.5	28
475	A New Scale Measuring Psychologic Impact of Genetic Susceptibility Testing for Alzheimer Disease. Alzheimer Disease and Associated Disorders, 2009, 23, 50-56.	0.6	28
476	Height loss predicts subsequent hip fracture in men and women of the Framingham Study. Journal of Bone and Mineral Research, 2012, 27, 146-152.	3.1	28
477	Association of total protein intake with bone mineral density and bone loss in men and women from the Framingham Offspring Study. Public Health Nutrition, 2014, 17, 2570-2576.	1.1	28
478	A randomized noninferiority trial of condensed protocols for genetic risk disclosure of Alzheimer's disease. Alzheimer's and Dementia, 2015, 11, 1222-1230.	0.4	28
479	Meta-analysis of genome-wide association studies of HDL cholesterol response to statins. Journal of Medical Genetics, 2016, 53, 835-845.	1.5	28
480	Estimation of morbid risk and age at onset with missing information. American Journal of Human Genetics, 1991, 49, 76-87.	2.6	28
481	Multiple etiologies for Alzheimer disease are revealed by segregation analysis. American Journal of Human Genetics, 1994, 55, 991-1000.	2.6	28
482	Diet, menopause, and serum cholesterol levels in women: The Framingham Study. American Heart Journal, 1993, 125, 483-489.	1.2	27
483	Diet and plasma lipids in women. I. Macronutrients and plasma total and low-density lipoprotein cholesterol in women: The Framingham Nutrition Studies. Journal of Clinical Epidemiology, 1996, 49, 657-663.	2.4	27
484	Two-stage approach for identifying single-nucleotide polymorphisms associated with rheumatoid arthritis using random forests and Bayesian networks. BMC Proceedings, 2007, 1, S56.	1.8	27
485	ACC2 gene polymorphisms, metabolic syndrome, and gene-nutrient interactions with dietary fat. Journal of Lipid Research, 2010, 51, 3500-3507.	2.0	27
486	Approaches to detect genetic effects that differ between two strata in genome-wide meta-analyses: Recommendations based on a systematic evaluation. PLoS ONE, 2017, 12, e0181038.	1.1	27

#	Article	IF	CITATIONS
487	A longitudinal study of disc height narrowing and facet joint osteoarthritis at the thoracic and lumbar spine, evaluated by computed tomography: the Framingham Study. Spine Journal, 2018, 18, 2065-2073.	0.6	26
488	Common Coding Variants in <i>SCN10A</i> Are Associated With the Nav1.8 Late Current and Cardiac Conduction. Circulation Genomic and Precision Medicine, 2018, 11, e001663.	1.6	26
489	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
490	Association of mitochondrial DNA copy number with cardiometabolic diseases. Cell Genomics, $2021, 1, 100006$.	3.0	26
491	Impact of within-person variability on identifying children with hypercholesterolemia: Framingham children's study. Journal of Pediatrics, 1992, 121, 342-347.	0.9	25
492	A Genome-Wide Scan for Loci Affecting Normal Adult Height in the Framingham Heart Study. Human Heredity, 2003, 55, 191-201.	0.4	25
493	Polymorphisms in the gene encoding lipoprotein lipase in men with low HDL-C and coronary heart disease. Journal of Lipid Research, 2004, 45, 1885-1891.	2.0	25
494	Genetic variants on chromosome 5p12 are associated with risk of breast cancer in African American women: the Black Women's Health Study. Breast Cancer Research and Treatment, 2010, 123, 525-530.	1.1	25
495	Genetic determinants of the ankle-brachial index: A meta-analysis of a cardiovascular candidate gene 50K SNP panel in the candidate gene association resource (CARe) consortium. Atherosclerosis, 2012, 222, 138-147.	0.4	25
496	Genetic loci associated with ideal cardiovascular health: A meta-analysis of genome-wide association studies. American Heart Journal, 2016, 175, 112-120.	1.2	25
497	Association of Triglyceride-Related Genetic Variants With MitralÂAnnularÂCalcification. Journal of the American College of Cardiology, 2017, 69, 2941-2948.	1.2	25
498	Diet and Heart Disease Risk Factors in Adult American Men and Women: The Framingham Offspring-Spouse Nutrition Studies. International Journal of Epidemiology, 1993, 22, 1014-1025.	0.9	24
499	Ranking and characterization of established BMI and lipid associated loci as candidates for gene-environment interactions. PLoS Genetics, 2017, 13, e1006812.	1.5	24
500	Rare coding variants in 35 genes associate with circulating lipid levelsâ€"A multi-ancestry analysis of 170,000 exomes. American Journal of Human Genetics, 2022, 109, 81-96.	2.6	24
501	Genome-wide linkage analyses and candidate gene fine mapping for HDL3 cholesterol: the Framingham Study. Journal of Lipid Research, 2005, 46, 1416-1425.	2.0	23
502	Perceptions of Familial Risk in those Seeking a Genetic Risk Assessment for Alzheimer's Disease. Journal of Genetic Counseling, 2009, 18, 130-136.	0.9	23
503	Consent for genetic research in the Framingham Heart Study. American Journal of Medical Genetics, Part A, 2010, 152A, 1250-1256.	0.7	23
504	Genome-wide association of an integrated osteoporosis-related phenotype: Is there evidence for pleiotropic genes?. Journal of Bone and Mineral Research, 2012, 27, 319-330.	3.1	23

#	Article	IF	CITATIONS
505	Considerations in using linkage analysis as a presymptomatic test for Huntington's disease Journal of Medical Genetics, 1988, 25, 577-588.	1.5	22
506	Description of the Framingham Heart Study data for Genetic Analysis Workshop 13. BMC Genetics, 2003, 4, S2.	2.7	22
507	Heritability of Thoracic Spine Curvature and Genetic Correlations With Other Spine Traits: The Framingham Study. Journal of Bone and Mineral Research, 2016, 31, 2077-2084.	3.1	22
508	Thoracic Kyphosis and Physical Function: The Framingham Study. Journal of the American Geriatrics Society, 2017, 65, 2257-2264.	1.3	22
509	Lower Lean Mass Measured by Dual-Energy X-ray Absorptiometry (DXA) is Not Associated with Increased Risk of Hip Fracture in Women: The Framingham Osteoporosis Study. Calcified Tissue International, 2018, 103, 16-23.	1.5	22
510	A high throughput, functional screen of human Body Mass Index GWAS loci using tissue-specific RNAi Drosophila melanogaster crosses. PLoS Genetics, 2018, 14, e1007222.	1.5	22
511	Bone mass and the risk of prostate cancer: The Framingham study. American Journal of Medicine, 2002, 113, 734-739.	0.6	21
512	Association between well-characterized lipoprotein-related genetic variants and carotid intimal medial thickness and stenosis: The Framingham Heart Study. Atherosclerosis, 2006, 189, 222-228.	0.4	21
513	Sex and age specific effects of chromosomal regions linked to body mass index in the Framingham Study. BMC Genetics, 2006, 7, 7.	2.7	21
514	Non–Contrast-Enhanced Computerized Tomography and Analgesic-Related Kidney Disease: Report of the National Analgesic Nephropathy Study. Journal of the American Society of Nephrology: JASN, 2006, 17, 1472-1480.	3.0	21
515	Magnetic Resonance Imaging Traits in Siblings Discordant for Alzheimer Disease. Journal of Neuroimaging, 2008, 18, 268-275.	1.0	21
516	The Relationship of Estrogen Receptor- \hat{l}_{\pm} and - \hat{l}_{\pm}^2 Genes with Osteoarthritis of the Hand. Journal of Rheumatology, 2009, 36, 2772-2779.	1.0	21
517	Hip geometry variation is associated with bone mineralization pathway gene variants: The framingham study. Journal of Bone and Mineral Research, 2010, 25, 1564-1571.	3.1	21
518	Relation Between Serum Albumin and Carotid Atherosclerosis. Stroke, 2003, 34, 53-57.	1.0	20
519	The –174 IL-6 GG genotype is associated with a reduced risk of type 2 diabetes mellitus in a family sample from the National Heart, Lung and Blood Institute's Framingham Heart Study. Diabetologia, 2005, 48, 1492-1495.	2.9	20
520	Age-Related Changes in Echocardiographic Measurements. Hypertension, 2007, 49, 1000-1006.	1.3	20
521	Evidence for a gene influencing haematocrit on chromosome 6q23-24: genomewide scan in the Framingham Heart Study. Journal of Medical Genetics, 2005, 42, 75-79.	1.5	19
522	Estimation of fertility and fitness in Huntington disease in New England. American Journal of Medical Genetics Part A, 1989, 33, 248-254.	2.4	18

#	Article	IF	CITATIONS
523	Adult Height and Risk of Breast Cancer among White Women in a Case-Control Study. American Journal of Epidemiology, 1996, 143, 1123-1128.	1.6	18
524	Meta-analysis identifies loci affecting levels of the potential osteoarthritis biomarkers sCOMP and uCTX-II with genome wide significance. Journal of Medical Genetics, 2014, 51, 596-604.	1.5	18
525	Sequencing of 2 Subclinical Atherosclerosis Candidate Regions in 3669 Individuals. Circulation: Cardiovascular Genetics, 2014, 7, 359-364.	5.1	18
526	An Empirical Comparison of Joint and Stratified Frameworks for Studying G × E Interactions: Systolic Blood Pressure and Smoking in the CHARGE Gene‣ifestyle Interactions Working Group. Genetic Epidemiology, 2016, 40, 404-415.	0.6	18
527	Genome-wide association meta-analysis of fish and EPA+DHA consumption in 17 US and European cohorts. PLoS ONE, 2017, 12, e0186456.	1.1	18
528	Direct Versus Calculated LDL Cholesterol and C-Reactive Protein in Cardiovascular Disease Risk Assessment in the Framingham Offspring Study. Clinical Chemistry, 2019, 65, 1102-1114.	1.5	18
529	Statin-induced LDL cholesterol response and type 2 diabetes: a bidirectional two-sample Mendelian randomization study. Pharmacogenomics Journal, 2020, 20, 462-470.	0.9	18
530	Discovery and fine-mapping of height loci via high-density imputation of GWASs in individuals of African ancestry. American Journal of Human Genetics, 2021, 108, 564-582.	2.6	18
531	In utero exposure to diethylstilbestrol (DES) does not increase genomic instability in normal or neoplastic breast epithelium. Cancer, 2006, 107, 2122-2126.	2.0	17
532	Polymorphisms in the endothelial nitric oxide synthase gene and bone density/ultrasound and geometry in humans. Bone, 2008, 42, 53-60.	1.4	17
533	Association of TTR polymorphisms with hippocampal atrophy in Alzheimer disease families. Neurobiology of Aging, 2011, 32, 249-256.	1.5	17
534	Genetic and Clinical Correlates of Early-Outgrowth Colony-Forming Units. Circulation: Cardiovascular Genetics, 2011, 4, 296-304.	5.1	17
535	Ultraconserved Elements in the Human Genome: Association and Transmission Analyses of Highly Constrained Single-Nucleotide Polymorphisms. Genetics, 2012, 192, 253-266.	1.2	17
536	Chromosome Xq23 is associated with lower atherogenic lipid concentrations and favorable cardiometabolic indices. Nature Communications, 2021, 12, 2182.	5.8	17
537	Genome-wide association study for radiographic vertebral fractures: a potential role for the 16q24 BMD locus. Bone, 2014, 59, 20-7.	1.4	17
538	Absence of effect of seven functional mutations in the cyp2d6 gene in Parkinson's disease. Movement Disorders, 1999, 14, 590-595.	2.2	16
539	Refined QTLs of osteoporosis-related traits by linkage analysis with genome-wide SNPs: Framingham SHARe. Bone, 2010, 46, 1114-1121.	1.4	16
540	Genetic variation in TRPS1 may regulate hip geometry as well as bone mineral density. Bone, 2012, 50, 1188-1195.	1,4	16

#	Article	IF	Citations
541	Disclosing genetic risk for Alzheimer's dementia to individuals with mild cognitive impairment. Alzheimer's and Dementia: Translational Research and Clinical Interventions, 2020, 6, e12002.	1.8	16
542	Whole-genome association analyses of sleep-disordered breathing phenotypes in the NHLBI TOPMed program. Genome Medicine, 2021, 13, 136.	3.6	16
543	Using survival methods to estimate age-at-onset distributions for genetic diseases with an application to Huntington disease. Genetic Epidemiology, 1989, 6, 361-371.	0.6	15
544	Are Varicose Veins a Marker for Susceptibility to Coronary Heart Disease in Men? Results from the Normative Aging Study. Annals of Vascular Surgery, 2004, 18, 459-464.	0.4	15
545	Expectation Maximization Algorithm Based Haplotype Relative Risk (EM-HRR): Test of Linkage Disequilibrium Using Incomplete Case-Parents Trios. Human Heredity, 2005, 59, 125-135.	0.4	15
546	Association of dietary and biochemical measures of vitamin K with quantitative ultrasound of the heel in men and women. Osteoporosis International, 2006, 17, 600-607.	1.3	15
547	Trends in the association of parental history of obesity over 60 years. Obesity, 2014, 22, 919-924.	1.5	15
548	Drug-Gene Interactions of Antihypertensive Medications and Risk of Incident Cardiovascular Disease: A Pharmacogenomics Study from the CHARGE Consortium. PLoS ONE, 2015, 10, e0140496.	1.1	15
549	A genome wide linkage scan of metacarpal size and geometry in the Framingham Study. American Journal of Human Biology, 2008, 20, 663-670.	0.8	14
550	Estimating Interaction Between Genetic and Environmental Risk Factors. Epidemiology, 2008, 19, 83-93.	1.2	14
551	Proteomic Signatures of Lifestyle Risk Factors for Cardiovascular Disease: A Crossâ€Sectional Analysis of the Plasma Proteome in the Framingham Heart Study. Journal of the American Heart Association, 2021, 10, e018020.	1.6	14
552	Cancer Surveillance of Veterans in Massachusetts, USA, 1982–1988. International Journal of Epidemiology, 1991, 20, 7-12.	0.9	13
553	Informative-Transmission Disequilibrium Test (i-TDT): combined linkage and association mapping that includes unaffected offspring as well as affected offspring. Genetic Epidemiology, 2007, 31, 115-133.	0.6	13
554	Magnetic resonance imagingâ€measured atrophy and its relationship to cognitive functioning in vascular dementia and Alzheimer's disease patients. Alzheimer's and Dementia, 2011, 7, 493-500.	0.4	13
555	Genetic modifiers of response to glucose–insulin–potassium (GIK) infusion in acute coronary syndromes and associations with clinical outcomes in the IMMEDIATE trial. Pharmacogenomics Journal, 2015, 15, 488-495.	0.9	13
556	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
557	Population Genomic Analysis of 962 Whole Genome Sequences of Humans Reveals Natural Selection in Non-Coding Regions. PLoS ONE, 2015, 10, e0121644.	1.1	13
558	Segregation analysis for high density lipoprotein in the berkeley data. Genetic Epidemiology, 1993, 10, 629-634.	0.6	12

#	Article	IF	Citations
559	Combined haplotype relative risk (CHRR): a general and simple genetic association test that combines trios and unrelated caseâ€controls. Genetic Epidemiology, 2009, 33, 54-62.	0.6	12
560	Sequencing of <i>SCN5A</i> Identifies Rare and Common Variants Associated With Cardiac Conduction: Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) Consortium. Circulation: Cardiovascular Genetics, 2014, 7, 365-373.	5.1	12
561	Cardiovascular disease prevalence and insulin resistance in the Kyushu–Okinawa Population Study and the Framingham Offspring Study. Journal of Clinical Lipidology, 2017, 11, 348-356.	0.6	12
562	Large-scale pharmacogenomic study of sulfonylureas and the QT, JT and QRS intervals: CHARGE Pharmacogenomics Working Group. Pharmacogenomics Journal, 2018, 18, 127-135.	0.9	12
563	Family study designs in the age of genome-wide association studies: experience from the Framingham Heart Study. Current Opinion in Lipidology, 2008, 19, 144-150.	1.2	11
564	Renin Angiotensin System Gene Polymorphisms and Cerebral Blood Flow Regulation. Stroke, 2010, 41, 635-640.	1.0	11
565	Revisiting heritability accounting for shared environmental effects and maternal inheritance. Human Genetics, 2015, 134, 169-179.	1.8	11
566	Rooted in risk: genetic predisposition for low-density lipoprotein cholesterol level associates with diminished low-density lipoprotein cholesterol response to statin treatment. Pharmacogenomics, 2016, 17, 1621-1628.	0.6	11
567	Whole exome sequencing in the Framingham Heart Study identifies rare variation in HYAL2 that influences platelet aggregation. Thrombosis and Haemostasis, 2017, 117, 1083-1092.	1.8	11
568	Role of Rare and Low-Frequency Variants in Gene-Alcohol Interactions on Plasma Lipid Levels. Circulation Genomic and Precision Medicine, 2020, 13, e002772.	1.6	11
569	Gene-lifestyle interactions in the genomics of human complex traits. European Journal of Human Genetics, 2022, 30, 730-739.	1.4	11
570	Genotype-by-Sex Interaction in the Regulation of High-Density Lipoprotein: The Framingham Heart Study. Human Biology, 2005, 77, 773-793.	0.4	10
571	Fucosyltransferase 3 polymorphism and atherothrombotic disease in the Framingham Offspring Study. American Heart Journal, 2007, 153, 636-639.	1.2	10
572	Factors Affecting Recall of Different Types of Personal Genetic Information about Alzheimer's Disease Risk: The REVEAL Study. Public Health Genomics, 2015, 18, 78-86.	0.6	10
573	Genetic analyses of longitudinal phenotype data: a comparison of univariate methods and a multivariate approach. BMC Genetics, 2003, 4, S29.	2.7	9
574	Association of apo A-IV 360 (Gln → His) polymorphism with plasma lipids and lipoproteins: the Framingham Offspring Study. Atherosclerosis, 2005, 179, 169-175.	0.4	9
575	Lipoprotein receptor-related protein 1 variants and dietary fatty acids: meta-analysis of European origin and African American studies. International Journal of Obesity, 2013, 37, 1211-1220.	1.6	9
576	A genetic risk score is associated with statin-induced low-density lipoprotein cholesterol lowering. Pharmacogenomics, 2016, 17, 583-591.	0.6	9

#	Article	IF	CITATIONS
577	Ethnic Differences in Glucose Homeostasis Markers between the Kyushu-Okinawa Population Study and the Framingham Offspring Study. Scientific Reports, 2016, 6, 36725.	1.6	9
578	A genome-wide interaction analysis of tricyclic/tetracyclic antidepressants and RR and QT intervals: a pharmacogenomics study from the Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) consortium. Journal of Medical Genetics, 2017, 54, 313-323.	1.5	9
579	Pharmacogenomics study of thiazide diuretics and QT interval in multi-ethnic populations: the cohorts for heart and aging research in genomic epidemiology. Pharmacogenomics Journal, 2018, 18, 215-226.	0.9	9
580	Genomeâ€Wide Interactions with Dairy Intake for Body Mass Index in Adults of European Descent. Molecular Nutrition and Food Research, 2018, 62, 1700347.	1.5	9
581	Do changes in DNA methylation mediate or interact with SNP variation? A pharmacoepigenetic analysis. BMC Genetics, 2018, 19, 70.	2.7	9
582	Estimating the probability for major gene Alzheimer disease. American Journal of Human Genetics, 1994, 54, 374-83.	2.6	9
583	Genetic Analysis Workshop 15: gene expression analysis and approaches to detecting multiple functional loci. BMC Proceedings, 2007, 1, S1.	1.8	8
584	Genetic Analysis Workshop 16: Strategies for genome-wide association study analyses. BMC Proceedings, 2009, 3, S1.	1.8	8
585	Cross-Calibration and Comparison of Variability in 2 Bone Densitometers in a Research Setting: The Framingham Experience. Journal of Clinical Densitometry, 2010, 13, 210-218.	0.5	8
586	CYP4A11 variant is associated with high-density lipoprotein cholesterol in women. Pharmacogenomics Journal, 2013, 13, 44-51.	0.9	8
587	Sequence Variation in <i>TMEM18</i> in Association With Body Mass Index. Circulation: Cardiovascular Genetics, 2014, 7, 344-349.	5.1	8
588	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
589	Robust analysis of secondary phenotypes in caseâ€control genetic association studies. Statistics in Medicine, 2016, 35, 4226-4237.	0.8	8
590	Evaluation of a Twoâ€Stage Approach in Transâ€Ethnic Metaâ€Analysis in Genomeâ€Wide Association Studies. Genetic Epidemiology, 2016, 40, 284-292.	0.6	8
591	Genetic variants modify the associations of concentrations of methylmalonic acid, vitamin B-12, vitamin B-6, and folate with bone mineral density. American Journal of Clinical Nutrition, 2021, 114, 578-587.	2.2	8
592	Healthy People 2000. The rationale and potential efficacy of preventive nutrition in heart disease: the Framingham Offspring-Spouse Study. Archives of Internal Medicine, 1993, 153, 1549-1556.	4.3	8
593	Integrating genetic, transcriptional, and biological information provides insights into obesity. International Journal of Obesity, 2019, 43, 457-467.	1.6	8
594	Data mining. Genetic Epidemiology, 2005, 29, S103-S109.	0.6	7

#	Article	IF	CITATIONS
595	Impact of non-ignorable missingness on genetic tests of linkage and/or association using case-parent trios. BMC Genetics, 2005, 6, S90.	2.7	7
596	Targeted sequencing of genome wide significant loci associated with bone mineral density (BMD) reveals significant novel and rare variants: the Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) targeted sequencing study. Human Molecular Genetics, 2016, 25, ddw289.	1.4	7
597	Evaluation of power of the Illumina HumanOmni5M-4v1 BeadChip to detect risk variants for human complex diseases. European Journal of Human Genetics, 2016, 24, 1029-1034.	1.4	7
598	Polygenic transcriptome risk scores for COPD and lung function improve cross-ethnic portability of prediction in the NHLBI TOPMed program. American Journal of Human Genetics, 2022, 109, 857-870.	2.6	7
599	Genetic Analysis Workshop 13: Introduction to workshop summaries. Genetic Epidemiology, 2003, 25, S1-S4.	0.6	6
600	Genetic variation at glucose and insulin trait loci and response to glucose–insulin–potassium (GIK) therapy: the IMMEDIATE trial. Pharmacogenomics Journal, 2015, 15, 55-62.	0.9	6
601	Common variants associated with changes in levels of circulating free fatty acids after administration of glucose–insulin–potassium (GIK) therapy in the IMMEDIATE trial. Pharmacogenomics Journal, 2017, 17, 76-83.	0.9	6
602	Robust, flexible, and scalable tests for Hardy–Weinberg equilibrium across diverse ancestries. Genetics, 2021, 218, .	1.2	6
603	Presence and transmission of mitochondrial heteroplasmic mutations in human populations of European and African ancestry. Mitochondrion, 2021, 60, 33-42.	1.6	6
604	Meta-analysis of genome-wide association studies identifies ancestry-specific associations underlying circulating total tau levels. Communications Biology, 2022, 5, 336.	2.0	6
605	The analysis of survival data with a non-susceptible fraction and dual censoring mechanisms. Statistics in Medicine, 2003, 22, 3249-3262.	0.8	5
606	Mapping quantitative traits in unselected families: algorithms and examples. Genetic Epidemiology, 2009, 33, 617-627.	0.6	5
607	A Polymorphism in a Gene Encoding Perilipin 4 Is Associated with Height but not with Bone Measures in Individuals from the Framingham Osteoporosis Study. Calcified Tissue International, 2012, 90, 96-107.	1.5	5
608	Imputing rare variants in families using a two-stage approach. BMC Proceedings, 2016, 10, 209-214.	1.8	5
609	Rare variant associations with waist-to-hip ratio in European-American and African-American women from the NHLBI-Exome Sequencing Project. European Journal of Human Genetics, 2016, 24, 1181-1187.	1.4	5
610	Generating survival times with time-varying covariates using the Lambert W Function. Communications in Statistics Part B: Simulation and Computation, 2022, 51, 135-153.	0.6	5
611	Searching for parent-of-origin effects on cardiometabolic traits in imprinted genomic regions. European Journal of Human Genetics, 2020, 28, 646-655.	1.4	5
612	Sociodemographic Patterns of Exposure to Civil Aircraft Noise in the United States. Environmental Health Perspectives, 2022, 130, 27009.	2.8	5

#	Article	IF	CITATIONS
613	Lymphocyte activation gene-3-associated protein networks are associated with HDL-cholesterol and mortality in the Trans-omics for Precision Medicine program. Communications Biology, 2022, 5, 362.	2.0	5
614	Estimation of familial risk in Alzheimer's disease. Annals of Neurology, 1990, 27, 338-340.	2.8	4
615	Predictors of Improved Left Ventricular Systolic Function in an Urban Cardiomyopathy Program. American Journal of Cardiology, 2006, 98, 1622-1626.	0.7	4
616	Ordered Stratification to Reduce Heterogeneity in Linkage to Diabetesâ€related Quantitative Traits. Obesity, 2008, 16, 2314-2322.	1.5	4
617	Genome-Wide Linkage and Association Scans for Quantitative Trait Loci of Serum Lactate Dehydrogenase—The Framingham Heart Study. Human Genomics and Proteomics, 2010, 2, 905237.	1.5	4
618	A fine-mapping study of central obesity loci incorporating functional annotation and imputation. European Journal of Human Genetics, 2018, 26, 1369-1377.	1.4	4
619	Revisit Population-based and Family-based Genotype Imputation. Scientific Reports, 2019, 9, 1800.	1.6	4
620	Genome-wide meta-analysis of variant-by-diuretic interactions as modulators of lipid traits in persons of European and African ancestry. Pharmacogenomics Journal, 2020, 20, 482-493.	0.9	4
621	Identification of novel and rare variants associated with handgrip strength using whole genome sequence data from the NHLBI Trans-Omics in Precision Medicine (TOPMed) Program. PLoS ONE, 2021, 16, e0253611.	1.1	4
622	Joint modeling of linkage and association using affected sib-pair data. BMC Proceedings, 2007, 1, S38.	1.8	3
623	Using linkage and association to identify and model genetic effects: summary of GAW15 Group 4. Genetic Epidemiology, 2007, 31, S34-S42.	0.6	3
624	Incorporating biological knowledge in the search for gene $\tilde{A}-$ gene interaction in genome-wide association studies. BMC Proceedings, 2009, 3, S81.	1.8	3
625	Heritability of serum γâ€glutamyltransferase level: genetic analysis from the Framingham Offspring Study. Liver International, 2009, 29, 776-777.	1.9	3
626	Consistency of linkage results across exams and methods in the Framingham Heart Study. BMC Genetics, 2003, 4, S30.	2.7	2
627	Comparing baseline and longitudinal measures in association studies. BMC Proceedings, 2014, 8, S84.	1.8	2
628	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
629	Exome sequence association study of levels and longitudinal change of cardiovascular risk factor phenotypes in European Americans and African Americans from the Atherosclerosis Risk in Communities Study. Genetic Epidemiology, 2021, 45, 651-663.	0.6	2
630	BinomiRare: A robust test for association of a rare genetic variant with a binary outcome for mixed models and any case-control proportion. Human Genetics and Genomics Advances, 2021, 2, 100040.	1.0	2

#	Article	IF	CITATIONS
631	Rare coding variants in RCN3 are associated with blood pressure. BMC Genomics, 2022, 23, 148.	1.2	2
632	Detecting linkage for a complex disease using simulated extended pedigrees. Genetic Epidemiology, 1997, 14, 981-986.	0.6	1
633	Power of concordant versus discordant sib pairs at different penetrance levels. Genetic Epidemiology, 1999, 17, S679-84.	0.6	1
634	Authors? reply. Diabetologia, 2003, 46, 1588-1588.	2.9	1
635	Comparisons of case-selection approaches based on allele sharing and/or disease severity index: application to the GAW14 simulated data. BMC Genetics, 2005, 6, S103.	2.7	1
636	Response to Letter Regarding Article, "Association of Leukocyte Telomere Length With Circulating Biomarkers of the Renin-Angiotensin-Aldosterone System: The Framingham Heart Study― Circulation, 2008, 118, .	1.6	1
637	Selection of the most informative individuals from families with multiple siblings for association studies. Genetic Epidemiology, 2009, 33, 299-307.	0.6	1
638	Genome-wide gene–environment interactions on quantitative traits using family data. European Journal of Human Genetics, 2016, 24, 1022-1028.	1.4	1
639	Revisiting methods for modeling longitudinal and survival data: Framingham Heart Study. BMC Medical Research Methodology, 2021, 21, 29.	1.4	1
640	Variant-specific inflation factors for assessing population stratification at the phenotypic variance level. Nature Communications, 2021, 12, 3506.	5 . 8	1
641	Response to Blood Pressure and Disability: First Steps in Future Studies. Hypertension, 2008, 51, .	1.3	0
642	Performance of statistical methods on CHARGE targeted sequencing data. BMC Genetics, 2014, 15, 104.	2.7	0
643	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
644	ANNORE: genetic fine-mapping with functional annotation. Human Molecular Genetics, 2021, 31, 32-40.	1.4	0
645	Lifestyle Risk Score: handling missingness of individual lifestyle components in meta-analysis of gene-by-lifestyle interactions. European Journal of Human Genetics, 2021, 29, 839-850.	1.4	0
646	Positive association of total protein intake and bone mineral density (BMD) in women from the Framingham Offspring Study. FASEB Journal, 2010, 24, lb285.	0.2	0