

# Guillaume Pare

## List of Publications by Year in descending order

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270  
papers

36,363  
citations

10650

74  
h-index

4305

179  
g-index

299  
all docs

299  
docs citations

299  
times ranked

51155  
citing authors

#	ARTICLE	IF	CITATIONS
1	Dysbetalipoproteinemia: Differentiating Multifactorial Remnant Cholesterol Disease From Genetic ApoE Deficiency. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2022, 107, 538-548.	1.8	11
2	Validation of the classification for type 2 diabetes into five subgroups: a report from the ORIGIN trial. <i>Diabetologia</i> , 2022, 65, 206-215.	2.9	31
3	GWAS and ExWAS of blood mitochondrial DNA copy number identifies 71 loci and highlights a potential causal role in dementia. <i>ELife</i> , 2022, 11, .	2.8	42
4	Variations in risks from smoking between high-income, middle-income, and low-income countries: an analysis of data from 179â€™000 participants from 63 countries. <i>The Lancet Global Health</i> , 2022, 10, e216-e226.	2.9	16
5	Genetically proxied therapeutic inhibition of antihypertensive drug targets and risk of common cancers: A mendelian randomization analysis. <i>PLoS Medicine</i> , 2022, 19, e1003897.	3.9	30
6	Caffeine blocks SREBP2-induced hepatic PCSK9 expression to enhance LDLR-mediated cholesterol clearance. <i>Nature Communications</i> , 2022, 13, 770.	5.8	47
7	Lipoprotein(a) Cholesterol Masquerading as Low-Density Lipoprotein Cholesterol. <i>Journal of the American College of Cardiology</i> , 2022, 79, 1047-1049.	1.2	1
8	Cohort profile: genomic data for 26 622 individuals from the Canadian Longitudinal Study on Aging (CLSA). <i>BMJ Open</i> , 2022, 12, e059021.	0.8	17
9	Identification of genetic effects underlying type 2 diabetes in South Asian and European populations. <i>Communications Biology</i> , 2022, 5, 329.	2.0	21
10	Mitochondrial DNA Copy Number as a Marker and Mediator of Stroke Prognosis. <i>Neurology</i> , 2022, 98, .	1.5	10
11	Elevated Lipoprotein(a) and Risk of Atrial Fibrillation. <i>Journal of the American College of Cardiology</i> , 2022, 79, 1579-1590.	1.2	42
12	Biomarkers of Prevalent and Incident Cognitive Dysfunction in People with Dysglycemia- Data from the ORIGIN Trial. <i>Journal of Alzheimer's Disease</i> , 2022, , 1-8.	1.2	0
13	Genome-wide studies reveal factors associated with circulating uromodulin and its relationships to complex diseases. <i>JCI Insight</i> , 2022, 7, .	2.3	12
14	ACLY and CKD: A Mendelian Randomization Analysis. <i>Kidney International Reports</i> , 2022, 7, 1673-1681.	0.4	1
15	Multi-ancestry genetic study of type 2 diabetes highlights the power of diverse populations for discovery and translation. <i>Nature Genetics</i> , 2022, 54, 560-572.	9.4	250
16	Genome-wide association study identifies 48 common genetic variants associated with handedness. <i>Nature Human Behaviour</i> , 2021, 5, 59-70.	6.2	79
17	Renal Impairment and Risk of Acute Stroke: The INTERSTROKE Study. <i>Neuroepidemiology</i> , 2021, 55, 206-215.	1.1	2
18	The loss-of-function PCSK9Q152H variant increases ER chaperones GRP78 and GRP94 and protects against liver injury. <i>Journal of Clinical Investigation</i> , 2021, 131, .	3.9	29

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19	Whole exome sequencing reveals a biallelic frameshift mutation in GRXCR2 in hearing impairment in Cameroon. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2021, 9, e1609.	0.6	6
20	Polygenic Risk Score for Low-Density Lipoprotein Cholesterol Is Associated With Risk of Ischemic Heart Disease and Enriches for Individuals With Familial Hypercholesterolemia. <i>Circulation Genomic and Precision Medicine</i> , 2021, 14, e003106.	1.6	21
21	Acute Phase Response and Non-Reproducible Elevated Concentrations with a High-Sensitivity Cardiac Troponin I Assay. <i>Journal of Clinical Medicine</i> , 2021, 10, 1014.	1.0	14
22	Testosterone and sex hormone-binding globulin in dysglycemic women at high cardiovascular risk: A report from the Outcome Reduction with an Initial Glargine Intervention trial. <i>Diabetes and Vascular Disease Research</i> , 2021, 18, 147916412110024.	0.9	6
23	Polygenic Risk Score for Alzheimer's Disease in Caribbean Hispanics. <i>Annals of Neurology</i> , 2021, 90, 366-376.	2.8	15
24	NT-proBNP versus routine clinical risk factors as a predictor of cardiovascular events or death in people with dysglycemia – A brief report from the ORIGIN trial. <i>Journal of Diabetes and Its Complications</i> , 2021, 35, 107928.	1.2	2
25	Causal Effect of MMP-1 (Matrix Metalloproteinase-1), MMP-8, and MMP-12 Levels on Ischemic Stroke. <i>Stroke</i> , 2021, 52, e316-e320.	1.0	18
26	Genome-Wide Association Study Identifies First Locus Associated with Susceptibility to Cerebral Venous Thrombosis. <i>Annals of Neurology</i> , 2021, 90, 777-788.	2.8	10
27	The genomics of heart failure: design and rationale of the HERMES consortium. <i>ESC Heart Failure</i> , 2021, 8, 5531-5541.	1.4	11
28	Factor V Leiden and the Risk of Bleeding in Patients With Acute Coronary Syndromes Treated With Antiplatelet Therapy: Pooled Analysis of 3 Randomized Clinical Trials. <i>Journal of the American Heart Association</i> , 2021, 10, e021115.	1.6	2
29	Epigenetic Age and the Risk of Incident Atrial Fibrillation. <i>Circulation</i> , 2021, 144, 1899-1911.	1.6	35
30	Calibrated rare variant genetic risk scores for complex disease prediction using large exome sequence repositories. <i>Nature Communications</i> , 2021, 12, 5852.	5.8	19
31	Postneoadjuvant Pure and Predominantly Pure Intralymphatic Breast Carcinoma. <i>American Journal of Surgical Pathology</i> , 2021, 45, 537-542.	2.1	3
32	Polygenic risk score for Alzheimer's disease in Caribbean Hispanics. <i>Alzheimer's and Dementia</i> , 2021, 17, e055031.	0.4	0
33	Implications of OPRM1 and CYP2B6 variants on treatment outcomes in methadone-maintained patients in Ontario: Exploring sex differences. <i>PLoS ONE</i> , 2021, 16, e0261201.	1.1	4
34	Genetic risk for dengue hemorrhagic fever and dengue fever in multiple ancestries. <i>EBioMedicine</i> , 2020, 51, 102584.	2.7	10
35	Triglycerides, hypertension, and smoking predict cardiovascular disease in dysbetalipoproteinemia. <i>Journal of Clinical Lipidology</i> , 2020, 14, 46-52.	0.6	7
36	Novel Biomarkers for Change in Renal Function in People With Dysglycemia. <i>Diabetes Care</i> , 2020, 43, 433-439.	4.3	8

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37	Are large simple trials for dementia prevention possible?. Age and Ageing, 2020, 49, 154-160.	0.7	17
38	Plasma ACE2 and risk of death or cardiometabolic diseases: a case-cohort analysis. Lancet, The, 2020, 396, 968-976.	6.3	119
39	Association of Factor V Leiden With Subsequent Atherothrombotic Events. Circulation, 2020, 142, 546-555.	1.6	11
40	Discovery of rare variants associated with blood pressure regulation through meta-analysis of 1.3 million individuals. Nature Genetics, 2020, 52, 1314-1332.	9.4	91
41	Genome-Wide Association Study Meta-Analysis of Stroke in 22 000 Individuals of African Descent Identifies Novel Associations With Stroke. Stroke, 2020, 51, 2454-2463.	1.0	26
42	Novel Outcome Biomarkers Identified With Targeted Proteomic Analyses of Plasma From Critically Ill Coronavirus Disease 2019 Patients. , 2020, 2, e0189.		44
43	Beyond the Brain. Stroke, 2020, 51, 3007-3017.	1.0	20
44	Tracing risk of multiple cardiovascular diseases to smoking-related genes. European Heart Journal, 2020, 41, 3311-3313.	1.0	2
45	A simplified diagnosis algorithm for dysbetalipoproteinemia. Journal of Clinical Lipidology, 2020, 14, 431-437.	0.6	37
46	Fine-tuning of Genome-Wide Polygenic Risk Scores and Prediction of Gestational Diabetes in South Asian Women. Scientific Reports, 2020, 10, 8941.	1.6	25
47	<i>PPARA</i> Polymorphism Influences the Cardiovascular Benefit of Fenofibrate in Type 2 Diabetes: Findings From ACCORD-Lipid. Diabetes, 2020, 69, 771-783.	0.3	28
48	Global Assessment of Mendelian Stroke Genetic Prevalence in 101 635 Individuals From 7 Ethnic Groups. Stroke, 2020, 51, 1290-1293.	1.0	20
49	High-Sensitivity Cardiac Troponin T for Risk Stratification in Patients With Embolic Stroke of Undetermined Source. Stroke, 2020, 51, 2386-2394.	1.0	18
50	Hypolipidaemia among patients with PMM2-CDG is associated with low circulating PCSK9 levels: a case report followed by observational and experimental studies. Journal of Medical Genetics, 2020, 57, 11-17.	1.5	8
51	Influence of Genetic Ancestry on Human Serum Proteome. American Journal of Human Genetics, 2020, 106, 303-314.	2.6	19
52	Identification of Circulating Proteins Associated With Blood Pressure Using Mendelian Randomization. Circulation Genomic and Precision Medicine, 2020, 13, e002605.	1.6	8
53	ACE and Type 2 Diabetes Risk: A Mendelian Randomization Study. Diabetes Care, 2020, 43, 835-842.	4.3	28
54	Effects of lifelong testosterone exposure on health and disease using Mendelian randomization. ELife, 2020, 9, .	2.8	32

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55	Analytical strategies to include the X chromosome in variance heterogeneity analyses: Evidence for trait-specific polygenic variance structure. <i>Genetic Epidemiology</i> , 2019, 43, 815-830.	0.6	14
56	Identification of Novel Causal Blood Biomarkers Linking Metabolically Favorable Adiposity With Type 2 Diabetes Risk. <i>Diabetes Care</i> , 2019, 42, 1800-1808.	4.3	12
57	The Burden of Atherosclerotic Cardiovascular Disease in South Asians Residing in Canada: A Reflection From the South Asian Heart Alliance. <i>CJC Open</i> , 2019, 1, 271-281.	0.7	14
58	Novel Drug Targets for Ischemic Stroke Identified Through Mendelian Randomization Analysis of the Blood Proteome. <i>Circulation</i> , 2019, 140, 819-830.	1.6	84
59	Benefits and limitations of genome-wide association studies. <i>Nature Reviews Genetics</i> , 2019, 20, 467-484.	7.7	1,226
60	Lipoprotein(a): An underrecognized genetic risk factor for malignant coronary artery disease in young Indians. <i>Indian Heart Journal</i> , 2019, 71, 184-198.	0.2	40
61	Subsequent Event Risk in Individuals With Established Coronary Heart Disease. <i>Circulation Genomic and Precision Medicine</i> , 2019, 12, e002470.	1.6	17
62	Association of Chromosome 9p21 With Subsequent Coronary Heart Disease Events. <i>Circulation Genomic and Precision Medicine</i> , 2019, 12, e002471.	1.6	22
63	Metformin-induced increases in GDF15 are important for suppressing appetite and promoting weight loss. <i>Nature Metabolism</i> , 2019, 1, 1202-1208.	5.1	181
64	Genetics of early-onset coronary artery disease. <i>Current Opinion in Cardiology</i> , 2019, 34, 706-713.	0.8	6
65	Linking Spontaneous Coronary Artery Dissection, Cervical Artery Dissection, and Fibromuscular Dysplasia. <i>Journal of the American College of Cardiology</i> , 2019, 73, 67-69.	1.2	6
66	Moyamoya Disease Susceptibility Variant <i>RNF213</i> p.R4810K Increases the Risk of Ischemic Stroke Attributable to Large-Artery Atherosclerosis. <i>Circulation</i> , 2019, 139, 295-298.	1.6	64
67	A Mendelian Randomization-Based Approach to Identify Early and Sensitive Diagnostic Biomarkers of Disease. <i>Clinical Chemistry</i> , 2019, 65, 427-436.	1.5	16
68	Blood HER2 and Uromodulin as Causal Mediators of CKD. <i>Journal of the American Society of Nephrology: JASN</i> , 2018, 29, 1326-1335.	3.0	21
69	A novel mutation in <i>GPIHBP1</i> causes familial chylomicronemia syndrome. <i>Journal of Clinical Lipidology</i> , 2018, 12, 506-510.	0.6	10
70	The Genetic Link Between Diabetes and Atherosclerosis. <i>Canadian Journal of Cardiology</i> , 2018, 34, 565-574.	0.8	15
71	Polygenic Contribution in Individuals With Early-Onset Coronary Artery Disease. <i>Circulation Genomic and Precision Medicine</i> , 2018, 11, e001849.	1.6	41
72	Genomic DNA Methylation Signatures Enable Concurrent Diagnosis and Clinical Genetic Variant Classification in Neurodevelopmental Syndromes. <i>American Journal of Human Genetics</i> , 2018, 102, 156-174.	2.6	135

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73	Peripheral blood epi-signature of Claes-Jensen syndrome enables sensitive and specific identification of patients and healthy carriers with pathogenic mutations in KDM5C. <i>Clinical Epigenetics</i> , 2018, 10, 21.	1.8	58
74	Postoperative Remote Automated Monitoring: Need for and State of the Science. <i>Canadian Journal of Cardiology</i> , 2018, 34, 850-862.	0.8	43
75	Characterization of Patients with Embolic Strokes of Undetermined Source in the NAVIGATE ESUS Randomized Trial. <i>Journal of Stroke and Cerebrovascular Diseases</i> , 2018, 27, 1673-1682.	0.7	46
76	Cannabis use and risk of schizophrenia: a Mendelian randomization study. <i>Molecular Psychiatry</i> , 2018, 23, 1287-1292.	4.1	167
77	HDL Cholesterol, LDL Cholesterol, and Triglycerides as Risk Factors for CKD: A Mendelian Randomization Study. <i>American Journal of Kidney Diseases</i> , 2018, 71, 166-172.	2.1	90
78	BAFopathies™ DNA methylation epi-signatures demonstrate diagnostic utility and functional continuum of Coffinâ€™Siris and Nicolaidèsâ€™Baraitser syndromes. <i>Nature Communications</i> , 2018, 9, 4885.	5.8	83
79	Canadian Cardiovascular Society Position Statement on Familial Hypercholesterolemia: Update 2018. <i>Canadian Journal of Cardiology</i> , 2018, 34, 1553-1563.	0.8	105
80	Genetic Tools for Coronary Risk Assessment in Type 2 Diabetes: A Cohort Study From the ACCORD Clinical Trial. <i>Diabetes Care</i> , 2018, 41, 2404-2413.	4.3	32
81	Pharmacogenetics of Stroke. <i>Stroke</i> , 2018, 49, 2541-2548.	1.0	7
82	Genetic analysis of over 1 million people identifies 535 new loci associated with blood pressure traits. <i>Nature Genetics</i> , 2018, 50, 1412-1425.	9.4	924
83	Body mass index is negatively associated with telomere length: a collaborative cross-sectional meta-analysis of 87 observational studies. <i>American Journal of Clinical Nutrition</i> , 2018, 108, 453-475.	2.2	137
84	A robust method to estimate regional polygenic correlation under misspecified linkage disequilibrium structure. <i>Genetic Epidemiology</i> , 2018, 42, 636-647.	0.6	3
85	Rivaroxaban for Stroke Prevention after Embolic Stroke of Undetermined Source. <i>New England Journal of Medicine</i> , 2018, 378, 2191-2201.	13.9	730
86	Blood CSF1 and CXCL12 as Causal Mediators of Coronary Artery Disease. <i>Journal of the American College of Cardiology</i> , 2018, 72, 300-310.	1.2	69
87	Multi-ethnic genome-wide association study for atrial fibrillation. <i>Nature Genetics</i> , 2018, 50, 1225-1233.	9.4	552
88	Multiancestry genome-wide association study of 520,000 subjects identifies 32 loci associated with stroke and stroke subtypes. <i>Nature Genetics</i> , 2018, 50, 524-537.	9.4	1,124
89	Genome-wide association analysis identifies novel blood pressure loci and offers biological insights into cardiovascular risk. <i>Nature Genetics</i> , 2017, 49, 403-415.	9.4	492
90	Economic Considerations of Early Rule-In/Rule-Out Algorithms for The Diagnosis of Myocardial Infarction in The Emergency Department Using Cardiac Troponin and Glycemic Biomarkers. <i>Clinical Chemistry</i> , 2017, 63, 593-602.	1.5	11

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91	Alcohol and Cardiovascular Disease: How Much is Too Much?. <i>Current Atherosclerosis Reports</i> , 2017, 19, 13.	2.0	45
92	Identification of Cadherin 2 ( <i>CDH2</i> ) Mutations in Arrhythmogenic Right Ventricular Cardiomyopathy. <i>Circulation: Cardiovascular Genetics</i> , 2017, 10, .	5.1	123
93	Once versus twice daily aspirin after coronary bypass surgery: a randomized trial. <i>Journal of Thrombosis and Haemostasis</i> , 2017, 15, 889-896.	1.9	16
94	Molecular phenotype and bleeding risks of an inherited platelet disorder in a family with a <i>RUNX1</i> frameshift mutation. <i>Haemophilia</i> , 2017, 23, e204-e213.	1.0	14
95	Common coding variant in <i>SERPINA1</i> increases the risk for large artery stroke. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 3613-3618.	3.3	46
96	The 9p21.3 locus and cardiovascular risk in familial hypercholesterolemia. <i>Journal of Clinical Lipidology</i> , 2017, 11, 406-412.	0.6	17
97	Polygenic risk score predicts prevalence of cardiovascular disease in patients with familial hypercholesterolemia. <i>Journal of Clinical Lipidology</i> , 2017, 11, 725-732.e5.	0.6	90
98	Large-scale analyses of common and rare variants identify 12 new loci associated with atrial fibrillation. <i>Nature Genetics</i> , 2017, 49, 946-952.	9.4	279
99	PCSK9 genetic variants and risk of type 2 diabetes: a mendelian randomisation study. <i>Lancet Diabetes and Endocrinology</i> , 2017, 5, 97-105.	5.5	298
100	Endoplasmic Reticulum Stress and Ca <sup>2+</sup> Depletion Differentially Modulate the Sterol Regulatory Protein PCSK9 to Control Lipid Metabolism. <i>Journal of Biological Chemistry</i> , 2017, 292, 1510-1523.	1.6	31
101	Genetic Association Studies and Next Generation Sequencing in Stroke: Methods. , 2017, , 21-52.		0
102	S100A9 potentiates the activation of neutrophils by the etiological agent of gout, monosodium urate crystals. <i>Journal of Leukocyte Biology</i> , 2017, 102, 805-813.	1.5	15
103	Association between cannabis use and methadone maintenance treatment outcomes: an investigation into sex differences. <i>Biology of Sex Differences</i> , 2017, 8, 8.	1.8	30
104	Rule-In and Rule-Out of Myocardial Infarction Using Cardiac Troponin and Glycemic Biomarkers in Patients with Symptoms Suggestive of Acute Coronary Syndrome. <i>Clinical Chemistry</i> , 2017, 63, 403-414.	1.5	36
105	Relationships of Measured and Genetically Determined Height With the Cardiac Conduction System in Healthy Adults. <i>Circulation: Arrhythmia and Electrophysiology</i> , 2017, 10, .	2.1	19
106	Growth Differentiation Factor 15 as a Novel Biomarker for Metformin. <i>Diabetes Care</i> , 2017, 40, 280-283.	4.3	112
107	A machine-learning heuristic to improve gene score prediction of polygenic traits. <i>Scientific Reports</i> , 2017, 7, 12665.	1.6	69
108	Impact of Selection Bias on Estimation of Subsequent Event Risk. <i>Circulation: Cardiovascular Genetics</i> , 2017, 10, .	5.1	28

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109	<i>COL4A2</i> is associated with lacunar ischemic stroke and deep ICH. <i>Neurology</i> , 2017, 89, 1829-1839.	1.5	58
110	Genetic Interactions with Age, Sex, Body Mass Index, and Hypertension in Relation to Atrial Fibrillation: The AFGen Consortium. <i>Scientific Reports</i> , 2017, 7, 11303.	1.6	15
111	The defining DNA methylation signature of Kabuki syndrome enables functional assessment of genetic variants of unknown clinical significance. <i>Epigenetics</i> , 2017, 12, 923-933.	1.3	79
112	DNA methylation predicts stroke outcome better. <i>Neurology</i> , 2017, 89, 758-759.	1.5	4
113	Novel Blood Pressure Locus and Gene Discovery Using Genome-Wide Association Study and Expression Data Sets From Blood and the Kidney. <i>Hypertension</i> , 2017, 70, .	1.3	123
114	Genetic contribution to lipid levels in early life based on 158 loci validated in adults: the FAMILY study. <i>Scientific Reports</i> , 2017, 7, 68.	1.6	4
115	Clinical Validation of a Genome-Wide DNA Methylation Assay for Molecular Diagnosis of Imprinting Disorders. <i>Journal of Molecular Diagnostics</i> , 2017, 19, 848-856.	1.2	39
116	Mendelian Genes and Risk of Intracerebral Hemorrhage and Small-Vessel Ischemic Stroke in Sporadic Cases. <i>Stroke</i> , 2017, 48, 2263-2265.	1.0	12
117	Identification of epigenetic signature associated with alpha thalassemia/mental retardation X-linked syndrome. <i>Epigenetics and Chromatin</i> , 2017, 10, 10.	1.8	60
118	The association between age of onset of opioid use and comorbidity among opioid dependent patients receiving methadone maintenance therapy. <i>Addiction Science &amp; Clinical Practice</i> , 2017, 12, 9.	1.2	33
119	Gene Expression Profiles for the Identification of Prevalent Atrial Fibrillation. <i>Journal of the American Heart Association</i> , 2017, 6, .	1.6	6
120	Clinical evaluation of a hemochromatosis next-generation sequencing gene panel. <i>European Journal of Haematology</i> , 2017, 98, 228-234.	1.1	20
121	Factors independently associated with cardiac troponin I levels in young and healthy adults from the general population. <i>Clinical Research in Cardiology</i> , 2017, 106, 96-104.	1.5	21
122	Validation of the ORIGIN Cardiovascular Biomarker Panel and the Value of Adding Troponin I in Dysglycemic People. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2017, 102, 2251-2257.	1.8	13
123	Identification of new susceptibility loci for type 2 diabetes and shared etiological pathways with coronary heart disease. <i>Nature Genetics</i> , 2017, 49, 1450-1457.	9.4	218
124	Ranking and characterization of established BMI and lipid associated loci as candidates for gene-environment interactions. <i>PLoS Genetics</i> , 2017, 13, e1006812.	1.5	24
125	Determinants of Left Atrial Volume in Patients with Atrial Fibrillation. <i>PLoS ONE</i> , 2016, 11, e0164145.	1.1	5
126	Gender and <i>BCR-ABL</i> transcript type are correlated with molecular response to imatinib treatment in patients with chronic myeloid leukemia. <i>European Journal of Haematology</i> , 2016, 96, 360-366.	1.1	35



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127	Rivaroxaban for secondary stroke prevention in patients with embolic strokes of undetermined source: Design of the NAVIGATE ESUS randomized trial. <i>European Stroke Journal</i> , 2016, 1, 146-154.	2.7	83
128	Towards the genetic basis of cerebral venous thrombosis—the BEAST Consortium: a study protocol: Table A1. <i>BMJ Open</i> , 2016, 6, e012351.	0.8	23
129	A method to estimate the contribution of regional genetic associations to complex traits from summary association statistics. <i>Scientific Reports</i> , 2016, 6, 27644.	1.6	5
130	The defining DNA methylation signature of Floating-Harbor Syndrome. <i>Scientific Reports</i> , 2016, 6, 38803.	1.6	55
131	Telomere Length and Risk of Myocardial Infarction in a Multi-Ethnic Population. <i>Journal of the American College of Cardiology</i> , 2016, 67, 1863-1865.	1.2	19
132	Putting the Genome in Context: Gene-Environment Interactions in Type 2 Diabetes. <i>Current Diabetes Reports</i> , 2016, 16, 57.	1.7	28
133	Clinical Validation of Fragile X Syndrome Screening by DNA Methylation Array. <i>Journal of Molecular Diagnostics</i> , 2016, 18, 834-841.	1.2	37
134	Identification of a methylation profile for DNMT1-associated autosomal dominant cerebellar ataxia, deafness, and narcolepsy. <i>Clinical Epigenetics</i> , 2016, 8, 91.	1.8	66
135	Genetic determinants of warfarin maintenance dose and time in therapeutic treatment range: a RE-LY genomics substudy. <i>Pharmacogenomics</i> , 2016, 17, 1425-1439.	0.6	21
136	Frameshift mutation in the APOA5 gene causing hypertriglyceridemia in a Pakistani family: Management and considerations for cardiovascular risk. <i>Journal of Clinical Lipidology</i> , 2016, 10, 1272-1277.	0.6	6
137	Genetic Predictors of Cardiovascular Mortality During Intensive Glycemic Control in Type 2 Diabetes: Findings From the ACCORD Clinical Trial. <i>Diabetes Care</i> , 2016, 39, 1915-1924.	4.3	47
138	Impact of a Genetic Risk Score on Myocardial Infarction Risk Across Different Ethnic Populations. <i>Canadian Journal of Cardiology</i> , 2016, 32, 1440-1446.	0.8	18
139	A Digital Health Intervention to Lower Cardiovascular Risk. <i>JAMA Cardiology</i> , 2016, 1, 601.	3.0	45
140	Global and regional effects of potentially modifiable risk factors associated with acute stroke in 32 countries (INTERSTROKE): a case-control study. <i>Lancet</i> , 2016, 388, 761-775.	6.3	1,414
141	A Prospective Study to Investigate Predictors of Relapse among Patients with Opioid Use Disorder Treated with Methadone. <i>Substance Abuse: Research and Treatment</i> , 2016, 10, SART.S37030.	0.5	34
142	DNA methylation analysis in constitutional disorders: Clinical implications of the epigenome. <i>Critical Reviews in Clinical Laboratory Sciences</i> , 2016, 53, 147-165.	2.7	28
143	Loci associated with ischaemic stroke and its subtypes (SiGN): a genome-wide association study. <i>Lancet Neurology</i> , 2016, 15, 174-184.	4.9	217
144	Dabigatran etexilate and reduction in serum apolipoprotein B. <i>Heart</i> , 2016, 102, 57-62.	1.2	34

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145	A Risk Assessment Tool Incorporating New Biomarkers for Cardiovascular Events in Acute Coronary Syndromes: The Organization to Assess Strategies in Ischemic Syndromes (OASIS) Risk Score. Canadian Journal of Cardiology, 2016, 32, 1332-1339.	0.8	7
146	Peripheral Blood <i>MCEMP1</i> Gene Expression as a Biomarker for Stroke Prognosis. Stroke, 2016, 47, 652-658.	1.0	48
147	Genome-wide meta-analysis of cerebral white matter hyperintensities in patients with stroke. Neurology, 2016, 86, 146-153.	1.5	91
148	PON1 Q192R genetic variant and response to clopidogrel and prasugrel: pharmacokinetics, pharmacodynamics, and a meta-analysis of clinical outcomes. Journal of Thrombosis and Thrombolysis, 2016, 41, 374-383.	1.0	32
149	Whole Blood Gene Expression Differentiates between Atrial Fibrillation and Sinus Rhythm after Cardioversion. PLoS ONE, 2016, 11, e0157550.	1.1	11
150	The Genetics of Opioid Addiction Risk Evaluation Tool (GREAT) for Treatment Response in Methadone Patients. International Journal of High Risk Behaviors & Addiction, 2016, 6, .	0.1	0
151	Abstract 521: c-MYC as a differentiating marker between angiosarcoma and atypical vascular lesion. , 2016, , .		0
152	Bleeding Risks Associated with Confirmed Platelet Dense Granule Deficiency and/or Impaired Aggregation Responses. Blood, 2016, 128, 3728-3728.	0.6	1
153	Impact of Chronic Pain on Treatment Prognosis for Patients with Opioid Use Disorder: A Systematic Review and Meta-analysis. Substance Abuse: Research and Treatment, 2015, 9, SART.S30120.	0.5	32
154	Opioid substitution and antagonist therapy trials exclude the common addiction patient: a systematic review and analysis of eligibility criteria. Trials, 2015, 16, 475.	0.7	33
155	Sex differences in substance use, health, and social functioning among opioid users receiving methadone treatment: a multicenter cohort study. Biology of Sex Differences, 2015, 6, 21.	1.8	62
156	Contribution of BDNF and DRD2 genetic polymorphisms to continued opioid use in patients receiving methadone treatment for opioid use disorder: an observational study. Addiction Science & Clinical Practice, 2015, 10, 19.	1.2	19
157	Sex differences in outcomes of methadone maintenance treatment for opioid use disorder: a systematic review and meta-analysis. CMAJ Open, 2015, 3, E344-E351.	1.1	49
158	Overlap Chronic Placental Inflammation Is Associated with a Unique Gene Expression Pattern. PLoS ONE, 2015, 10, e0133738.	1.1	19
159	Examining the Clinical Use of Hemochromatosis Genetic Testing. Canadian Journal of Gastroenterology and Hepatology, 2015, 29, 41-45.	0.8	8
160	Use of genetic data to guide therapy in arterial disease. Journal of Thrombosis and Haemostasis, 2015, 13, S281-S289.	1.9	10
161	Association Between Shortened Leukocyte Telomere Length and Cardiometabolic Outcomes. Circulation: Cardiovascular Genetics, 2015, 8, 82-90.	5.1	277
162	Testosterone suppression in opioid users: A systematic review and meta-analysis. Drug and Alcohol Dependence, 2015, 149, 1-9.	1.6	93

#	ARTICLE	IF	CITATIONS
163	Genome-wide studies to identify risk factors for kidney disease with a focus on patients with diabetes. <i>Nephrology Dialysis Transplantation</i> , 2015, 30, iv26-iv34.	0.4	41
164	Contribution of Large Region Joint Associations to Complex Traits Genetics. <i>PLoS Genetics</i> , 2015, 11, e1005103.	1.5	10
165	Effect of Bile Acid Sequestrants on the Risk of Cardiovascular Events. <i>Circulation: Cardiovascular Genetics</i> , 2015, 8, 618-627.	5.1	61
166	Mendelian randomization analysis supports the causal role of dysglycaemia and diabetes in the risk of coronary artery disease. <i>European Heart Journal</i> , 2015, 36, 1454-1462.	1.0	106
167	Methadone induces testosterone suppression in patients with opioid addiction. <i>Scientific Reports</i> , 2015, 4, 6189.	1.6	37
168	The impact of chronic pain on opioid addiction treatment: a systematic review protocol. <i>Systematic Reviews</i> , 2015, 4, 49.	2.5	9
169	Active metabolite concentration of clopidogrel in patients taking different doses of aspirin: results of the interaction trial. <i>Journal of Thrombosis and Haemostasis</i> , 2015, 13, 347-352.	1.9	7
170	Identifying Novel Biomarkers for Cardiovascular Events or Death in People With Dysglycemia. <i>Circulation</i> , 2015, 132, 2297-2304.	1.6	64
171	Multiple daily doses of acetylsalicylic acid (ASA) overcome reduced platelet response to once-daily ASA after coronary artery bypass graft surgery: a pilot randomized controlled trial. <i>Journal of Thrombosis and Haemostasis</i> , 2015, 13, 448-456.	1.9	30
172	Recommendations From the International Stroke Genetics Consortium, Part 2. <i>Stroke</i> , 2015, 46, 285-290.	1.0	8
173	Biological Rational for the Use of Heparin in Septic Shock: Translational Data from the Halo Pilot RCT. <i>Blood</i> , 2015, 126, 2336-2336.	0.6	4
174	A Gene Variant in CERS2 Is Associated with Rate of Increase in Albuminuria in Patients with Diabetes from ONTARGET and TRANSCEND. <i>PLoS ONE</i> , 2014, 9, e106631.	1.1	31
175	Evaluation of clinical and inflammatory profile in opioid addiction patients with comorbid pain: results from a multicenter investigation. <i>Neuropsychiatric Disease and Treatment</i> , 2014, 10, 2239.	1.0	16
176	Genetic influence on methadone treatment outcomes in patients undergoing methadone maintenance treatment for opioid addiction: a pilot study. <i>Neuropsychiatric Disease and Treatment</i> , 2014, 10, 1503.	1.0	26
177	Meta-analysis of SNPs involved in variance heterogeneity using Levene's test for equal variances. <i>European Journal of Human Genetics</i> , 2014, 22, 427-430.	1.4	7
178	The effectiveness of opioid substitution treatments for patients with opioid dependence: a systematic review and multiple treatment comparison protocol. <i>Systematic Reviews</i> , 2014, 3, 105.	2.5	49
179	The pharmacogenetics of carboxylesterases: <i>CES1</i> and <i>CES2</i> genetic variants and their clinical effect. <i>Drug Metabolism and Drug Interactions</i> , 2014, 29, 143-151.	0.3	79
180	Variation at the DPP4 locus influences apolipoprotein B levels in South Asians and exhibits heterogeneity in Europeans related to BMI. <i>Diabetologia</i> , 2014, 57, 738-745.	2.9	9

#	ARTICLE	IF	CITATIONS
181	Association of cyclooxygenase-2 genetic variant with cardiovascular disease. <i>European Heart Journal</i> , 2014, 35, 2242-2248.	1.0	42
182	Statistical genetics with application to population-based study design: a primer for clinicians. <i>European Heart Journal</i> , 2014, 35, 495-500.	1.0	7
183	Does genetic heterogeneity account for the divergent risk of type 2 diabetes in South Asian and white European populations?. <i>Diabetologia</i> , 2014, 57, 2270-2281.	2.9	29
184	Enabling the genomic revolution in Africa. <i>Science</i> , 2014, 344, 1346-1348.	6.0	361
185	Meta-analysis in more than 17,900 cases of ischemic stroke reveals a novel association at 12q24.12. <i>Neurology</i> , 2014, 83, 678-685.	1.5	89
186	Effect of Genetic Variants Associated With Plasma Homocysteine Levels on Stroke Risk. <i>Stroke</i> , 2014, 45, 1920-1924.	1.0	30
187	Genomics and epigenomics in pediatric oncology and clinical laboratory genetics. <i>Clinical Biochemistry</i> , 2014, 47, 731-732.	0.8	3
188	Myocardial Injury after Noncardiac Surgery. <i>Anesthesiology</i> , 2014, 120, 564-578.	1.3	740
189	Of stroke and biomarkers: The elusive quest for a clinical biomarker panel. <i>Clinical Biochemistry</i> , 2013, 46, 705-706.	0.8	6
190	Aboriginal birth cohort (ABC): a prospective cohort study of early life determinants of adiposity and associated risk factors among Aboriginal people in Canada. <i>BMC Public Health</i> , 2013, 13, 608.	1.2	17
191	Rationale and design of South Asian Birth Cohort (START): a Canada-India collaborative study. <i>BMC Public Health</i> , 2013, 13, 79.	1.2	49
192	Genetic Information and the Prediction of Incident Type 2 Diabetes in a High-Risk Multiethnic Population. <i>Diabetes Care</i> , 2013, 36, 2836-2842.	4.3	22
193	Pollen Count and Presentation of Angiotensin-Converting Enzyme Inhibitor-Associated Angioedema. <i>Journal of Allergy and Clinical Immunology: in Practice</i> , 2013, 1, 468-473.e4.	2.0	9
194	Pharmacogenetics of antiplatelets and anticoagulants: a report on clopidogrel, warfarin and dabigatran. <i>Pharmacogenomics</i> , 2013, 14, 1565-1572.	0.6	19
195	Secretory Phospholipase A2-IIA and Cardiovascular Disease. <i>Journal of the American College of Cardiology</i> , 2013, 62, 1966-1976.	1.2	115
196	A Genome-Wide Association Study for Venous Thromboembolism: The Extended Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) Consortium. <i>Genetic Epidemiology</i> , 2013, 37, 512-521.	0.6	99
197	Genetic Determinants of Dabigatran Plasma Levels and Their Relation to Bleeding. <i>Circulation</i> , 2013, 127, 1404-1412.	1.6	222
198	The protective effect of the obesity-associated rs9939609 A variant in fat mass- and obesity-associated gene on depression. <i>Molecular Psychiatry</i> , 2013, 18, 1281-1286.	4.1	115

#	ARTICLE	IF	CITATIONS
199	Common genetic loci influencing plasma homocysteine concentrations and their effect on risk of coronary artery disease. <i>American Journal of Clinical Nutrition</i> , 2013, 98, 668-676.	2.2	161
200	Genetic Markers of Inflammation and Their Role in Cardiovascular Disease. <i>Canadian Journal of Cardiology</i> , 2013, 29, 67-74.	0.8	47
201	Exploring Gene-Environment Relationships in Cardiovascular Disease. <i>Canadian Journal of Cardiology</i> , 2013, 29, 37-45.	0.8	46
202	Genetic Dissection of Diabetes: Facing the Giant. <i>Diabetes</i> , 2013, 62, 3338-3340.	0.3	3
203	Causal Relationship between Obesity and Vitamin D Status: Bi-Directional Mendelian Randomization Analysis of Multiple Cohorts. <i>PLoS Medicine</i> , 2013, 10, e1001383.	3.9	753
204	All SNPs Are Not Created Equal: Genome-Wide Association Studies Reveal a Consistent Pattern of Enrichment among Functionally Annotated SNPs. <i>PLoS Genetics</i> , 2013, 9, e1003449.	1.5	268
205	Gene × Physical Activity Interactions in Obesity: Combined Analysis of 111,421 Individuals of European Ancestry. <i>PLoS Genetics</i> , 2013, 9, e1003607.	1.5	168
206	Genetic variants associated with angiotensin-converting enzyme inhibitor-associated angioedema. <i>Pharmacogenetics and Genomics</i> , 2013, 23, 470-478.	0.7	68
207	A common biological basis of obesity and nicotine addiction. <i>Translational Psychiatry</i> , 2013, 3, e308-e308.	2.4	51
208	Genome-Wide Association Studies (GWAS)., 2013, , 25-50.		4
209	South Asian Heart Risk Assessment (SAHARA): Randomized Controlled Trial Design and Pilot Study. <i>JMIR Research Protocols</i> , 2013, 2, e33.	0.5	17
210	Genetic and phenotypic determinants of blood pressure and other cardiovascular risk factors (GAPP). <i>Swiss Medical Weekly</i> , 2013, 143, w13728.	0.8	30
211	Novel Loci for Adiponectin Levels and Their Influence on Type 2 Diabetes and Metabolic Traits: A Multi-Ethnic Meta-Analysis of 45,891 Individuals. <i>PLoS Genetics</i> , 2012, 8, e1002607.	1.5	419
212	Effect of <i>PON1</i> Q192R Genetic Polymorphism on Clopidogrel Efficacy and Cardiovascular Events in the Clopidogrel in the Unstable Angina to Prevent Recurrent Events Trial and the Atrial Fibrillation Clopidogrel Trial With Irbesartan for Prevention of Vascular Events. <i>Circulation: Cardiovascular Genetics</i> , 2012, 5, 250-256.	5.1	25
213	Combined vaccination and immunostimulatory antibodies provides durable cure of murine melanoma and induces transcriptional changes associated with positive outcome in human melanoma patients. <i>Oncolmmunology</i> , 2012, 1, 419-431.	2.1	25
214	Promises and challenges of pharmacogenetics: an overview of study design, methodological and statistical issues. <i>JRSM Cardiovascular Disease</i> , 2012, 1, 1-13.	0.4	41
215	The relationship between CYP2C19 polymorphisms and ischaemic and bleeding outcomes in stable outpatients: the CHARISMA genetics study. <i>European Heart Journal</i> , 2012, 33, 2143-2150.	1.0	86
216	Relation between clopidogrel active metabolite levels and different platelet aggregation methods in patients receiving clopidogrel and aspirin. <i>Journal of Thrombosis and Thrombolysis</i> , 2012, 34, 429-436.	1.0	13

#	ARTICLE	IF	CITATIONS
217	BRCA2 Variants and cardiovascular disease in a multi-ethnic study. <i>BMC Medical Genetics</i> , 2012, 13, 56.	2.1	13
218	Meta-analyses identify 13 loci associated with age at menopause and highlight DNA repair and immune pathways. <i>Nature Genetics</i> , 2012, 44, 260-268.	9.4	303
219	Genetic risk factors for ischaemic stroke and its subtypes (the METASTROKE Collaboration): a meta-analysis of genome-wide association studies. <i>Lancet Neurology</i> , The, 2012, 11, 951-962.	4.9	445
220	Meta-analysis identifies multiple loci associated with kidney functionâ€related traits in east Asian populations. <i>Nature Genetics</i> , 2012, 44, 904-909.	9.4	254
221	Meta-Analysis of Genome-Wide Association Studies in >80 000 Subjects Identifies Multiple Loci for C-Reactive Protein Levels. <i>Circulation</i> , 2011, 123, 731-738.	1.6	461
222	Genome-Wide Association Identifies Nine Common Variants Associated With Fasting Proinsulin Levels and Provides New Insights Into the Pathophysiology of Type 2 Diabetes. <i>Diabetes</i> , 2011, 60, 2624-2634.	0.3	335
223	Genome-Wide Association Analysis Identifies Variants Associated with Nonalcoholic Fatty Liver Disease That Have Distinct Effects on Metabolic Traits. <i>PLoS Genetics</i> , 2011, 7, e1001324.	1.5	796
224	A fast algorithm to optimize SNP prioritization for gene-gene and gene-environment interactions. <i>Genetic Epidemiology</i> , 2011, 35, 729-738.	0.6	17
225	CUBN Is a Gene Locus for Albuminuria. <i>Journal of the American Society of Nephrology: JASN</i> , 2011, 22, 555-570.	3.0	208
226	Associations of the <i>FTO</i> rs9939609 variant with discrete body fat depots and dietary intake in a multi-ethnic cohort. <i>Genetical Research</i> , 2011, 93, 419-426.	0.3	17
227	Association of Variation at the <i>ABO</i> Locus With Circulating Levels of Soluble Intercellular Adhesion Molecule-1, Soluble P-selectin, and Soluble E-selectin. <i>Circulation: Cardiovascular Genetics</i> , 2011, 4, 681-686.	5.1	77
228	Lifestyle Interaction With Fat Mass and Obesity-Associated ( <i>FTO</i> ) Genotype and Risk of Obesity in Apparently Healthy U.S. Women. <i>Diabetes Care</i> , 2011, 34, 675-680.	4.3	84
229	Current Evidence for Genetic Testing in Clopidogrel-Treated Patients Undergoing Coronary Stenting. <i>Circulation: Cardiovascular Interventions</i> , 2011, 4, 505-513.	1.4	16
230	Genome-Wide Association Analysis of Soluble ICAM-1 Concentration Reveals Novel Associations at the NFKB1, PNPLA3, RELA, and SH2B3 Loci. <i>PLoS Genetics</i> , 2011, 7, e1001374.	1.5	76
231	CYP2C19 Genetic Testing Should Not Be Done in All Patients Treated With Clopidogrel Who Are Undergoing Percutaneous Coronary Intervention. <i>Circulation: Cardiovascular Interventions</i> , 2011, 4, 514-521.	1.4	19
232	Association Between a Literature-Based Genetic Risk Score and Cardiovascular Events in Women. <i>Obstetrical and Gynecological Survey</i> , 2010, 65, 368-369.	0.2	1
233	A large-scale candidate gene association study of age at menarche and age at natural menopause. <i>Human Genetics</i> , 2010, 128, 515-527.	1.8	114
234	Genome-Wide Association Studiesâ€Data Generation, Storage, Interpretation, and Bioinformatics. <i>Journal of Cardiovascular Translational Research</i> , 2010, 3, 183-188.	1.1	14

#	ARTICLE	IF	CITATIONS
235	Biological, clinical and population relevance of 95 loci for blood lipids. <i>Nature</i> , 2010, 466, 707-713.	13.7	3,249
236	Hundreds of variants clustered in genomic loci and biological pathways affect human height. <i>Nature</i> , 2010, 467, 832-838.	13.7	1,789
237	New loci associated with kidney function and chronic kidney disease. <i>Nature Genetics</i> , 2010, 42, 376-384.	9.4	710
238	Genome-wide meta-analyses identify multiple loci associated with smoking behavior. <i>Nature Genetics</i> , 2010, 42, 441-447.	9.4	1,083
239	Meta-analysis identifies 13 new loci associated with waist-hip ratio and reveals sexual dimorphism in the genetic basis of fat distribution. <i>Nature Genetics</i> , 2010, 42, 949-960.	9.4	836
240	Association analyses of 249,796 individuals reveal 18 new loci associated with body mass index. <i>Nature Genetics</i> , 2010, 42, 937-948.	9.4	2,634
241	Thirty new loci for age at menarche identified by a meta-analysis of genome-wide association studies. <i>Nature Genetics</i> , 2010, 42, 1077-1085.	9.4	445
242	Fine Mapping of the Insulin-Induced Gene 2 Identifies a Variant Associated With LDL Cholesterol and Total Apolipoprotein B Levels. <i>Circulation: Cardiovascular Genetics</i> , 2010, 3, 454-461.	5.1	7
243	A new method for measurement of total plasma PCSK9: clinical applications. <i>Journal of Lipid Research</i> , 2010, 51, 140-149.	2.0	197
244	Association Between a Literature-Based Genetic Risk Score and Cardiovascular Events in Women. <i>JAMA - Journal of the American Medical Association</i> , 2010, 303, 631.	3.8	320
245	Genetic variants at 2q24 are associated with susceptibility to type 2 diabetes. <i>Human Molecular Genetics</i> , 2010, 19, 2706-2715.	1.4	178
246	Interpreting Metabolomic Profiles using Unbiased Pathway Models. <i>PLoS Computational Biology</i> , 2010, 6, e1000692.	1.5	52
247	Multiple Genetic Loci Influence Serum Urate Levels and Their Relationship With Gout and Cardiovascular Disease Risk Factors. <i>Circulation: Cardiovascular Genetics</i> , 2010, 3, 523-530.	5.1	285
248	On the Use of Variance per Genotype as a Tool to Identify Quantitative Trait Interaction Effects: A Report from the Women's Genome Health Study. <i>PLoS Genetics</i> , 2010, 6, e1000981.	1.5	175
249	Effects of <i>CYP2C19</i> Genotype on Outcomes of Clopidogrel Treatment. <i>New England Journal of Medicine</i> , 2010, 363, 1704-1714.	13.9	497
250	Failure to Validate Association between 12p13 Variants and Ischemic Stroke. <i>New England Journal of Medicine</i> , 2010, 362, 1547-1550.	13.9	75
251	The Fat-Mass and Obesity-Associated (FTO) gene, physical activity, and risk of incident cardiovascular events in white women. <i>American Heart Journal</i> , 2010, 160, 1163-1169.	1.2	51
252	Mendelian randomisation, triglycerides, and CHD. <i>Lancet</i> , The, 2010, 375, 1584-1586.	6.3	15

#	ARTICLE	IF	CITATIONS
253	Population-Based Genomewide Genetic Analysis of Common Clinical Chemistry Analytes. <i>Clinical Chemistry</i> , 2009, 55, 39-51.	1.5	13
254	Getting Closer to P-Selectin. <i>Clinical Chemistry</i> , 2009, 55, 1051-1052.	1.5	2
255	Genetic Variants Associated With Myocardial Infarction Risk Factors in Over 8000 Individuals From Five Ethnic Groups. <i>Circulation: Cardiovascular Genetics</i> , 2009, 2, 16-25.	5.1	67
256	Polymorphism in the <i>CETP</i> Gene Region, HDL Cholesterol, and Risk of Future Myocardial Infarction. <i>Circulation: Cardiovascular Genetics</i> , 2009, 2, 26-33.	5.1	186
257	Novel Loci, Including Those Related to Crohn Disease, Psoriasis, and Inflammation, Identified in a Genome-Wide Association Study of Fibrinogen in 17 686 Women. <i>Circulation: Cardiovascular Genetics</i> , 2009, 2, 134-141.	5.1	76
258	Multiple loci associated with indices of renal function and chronic kidney disease. <i>Nature Genetics</i> , 2009, 41, 712-717.	9.4	553
259	Genome-wide association studies identify loci associated with age at menarche and age at natural menopause. <i>Nature Genetics</i> , 2009, 41, 724-728.	9.4	348
260	Candidate genetic variants in the fibrinogen, methylenetetrahydrofolate reductase, and intercellular adhesion molecule-1 genes and plasma levels of fibrinogen, homocysteine, and intercellular adhesion molecule-1 among various race/ethnic groups: Data from the Women's Genome Health Study. <i>American Heart Journal</i> , 2009, 157, 777-783.e1.	1.2	21
261	Novel Associations of CPS1, MUT, NOX4, and DPEP1 With Plasma Homocysteine in a Healthy Population. <i>Circulation: Cardiovascular Genetics</i> , 2009, 2, 142-150.	5.1	96
262	Forty-Three Loci Associated with Plasma Lipoprotein Size, Concentration, and Cholesterol Content in Genome-Wide Analysis. <i>PLoS Genetics</i> , 2009, 5, e1000730.	1.5	300
263	K45R variant of squalene synthase increases total cholesterol levels in two study samples from a French Canadian population. <i>Human Mutation</i> , 2008, 29, 689-694.	1.1	5
264	Loci Related to Metabolic-Syndrome Pathways Including LEPR, HNF1A, IL6R, and GCKR Associate with Plasma C-Reactive Protein: The Women's Genome Health Study. <i>American Journal of Human Genetics</i> , 2008, 82, 1185-1192.	2.6	299
265	Novel Association of ABO Histo-Blood Group Antigen with Soluble ICAM-1: Results of a Genome-Wide Association Study of 6,578 Women. <i>PLoS Genetics</i> , 2008, 4, e1000118.	1.5	289
266	Novel Association of HK1 with Glycated Hemoglobin in a Non-Diabetic Population: A Genome-Wide Evaluation of 14,618 Participants in the Women's Genome Health Study. <i>PLoS Genetics</i> , 2008, 4, e1000312.	1.5	87
267	Genetic Loci Associated With Plasma Concentration of Low-Density Lipoprotein Cholesterol, High-Density Lipoprotein Cholesterol, Triglycerides, Apolipoprotein A1, and Apolipoprotein B Among 6382 White Women in Genome-Wide Analysis With Replication. <i>Circulation: Cardiovascular Genetics</i> , 2008, 1, 21-30.	5.1	117
268	Correction of Population Stratification in Large Multi-Ethnic Association Studies. <i>PLoS ONE</i> , 2008, 3, e1382.	1.1	60
269	Genetic Analysis of 103 Candidate Genes for Coronary Artery Disease and Associated Phenotypes in a Founder Population Reveals a New Association between Endothelin-1 and High-Density Lipoprotein Cholesterol. <i>American Journal of Human Genetics</i> , 2007, 80, 673-682.	2.6	79
270	The AIDS Disease of CD4C/HIV Transgenic Mice Shows Impaired Germinal Centers and Autoantibodies and Develops in the Absence of IFN- $\beta$ and IL-6. <i>Immunity</i> , 2001, 15, 173-185.	6.6	59