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Genome-wide association with select biomarker traits in the Framingham Heart Study

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#	Paper	IF	Citations
104	The Framingham Heart Study, on its way to becoming the gold standard for Cardiovascular Genetic Epidemiology?. <i>BMC Medical Genetics</i> , 2007 , 8, 63	2.1	25
103	The Framingham Heart Study 100K SNP genome-wide association study resource: overview of 17 phenotype working group reports. <i>BMC Medical Genetics</i> , 2007 , 8 Suppl 1, S1	2.1	152
102	Quantification of the genetic component of basal C-reactive protein expression in SLE nuclear families. 2008 , 72, 611-20		8
101	Novel association of ABO histo-blood group antigen with soluble ICAM-1: results of a genome-wide association study of 6,578 women. 2008 , 4, e1000118		230
100	Tilting at quixotic trait loci (QTL): an evolutionary perspective on genetic causation. 2008 , 179, 1741-56		57
99	Blood and cardiovascular disease: the promise and limitations of gene expression analysis. 2008 , 1, 7-9		4
98	Family study designs in the age of genome-wide association studies: experience from the Framingham Heart Study. 2008 , 19, 144-50		11
97	Family-based genome-wide association studies. 2009 , 10, 181-90		53
96	Population-based genomewide genetic analysis of common clinical chemistry analytes. 2009 , 55, 39-51		9
95	C-reactive protein and coronary heart disease: predictive test or therapeutic target?. 2009 , 55, 239-55		55
94	The role of C-reactive protein polymorphisms in inflammation and cardiovascular risk. 2009 , 11, 124-30		24
93	Tests of association for quantitative traits in nuclear families using principal components to correct for population stratification. 2009 , 73, 601-13		27
92	Cross-sectional relations of multiple inflammatory biomarkers to peripheral arterial disease: The Framingham Offspring Study. 2009 , 203, 509-14		52
91	Pooled versus individual genotyping in a breast cancer genome-wide association study. 2010 , 34, 603-12		10
90	Relation of platelet and leukocyte inflammatory transcripts to body mass index in the Framingham heart study. 2010 , 122, 119-29		105
89	Association between a literature-based genetic risk score and cardiovascular events in women. 2010 , 303, 631-7		256
88	Exhaled carbon monoxide and risk of metabolic syndrome and cardiovascular disease in the community. 2010 , 122, 1470-7		33

87	Disease-associated mutations that alter the RNA structural ensemble. 2010 , 6, e1001074		221
86	PheWAS: demonstrating the feasibility of a phenome-wide scan to discover gene-disease associations. 2010 , 26, 1205-10		668
85	A systematic review of the association between common single nucleotide polymorphisms and 25-hydroxyvitamin D concentrations. 2010 , 121, 471-7		175
84	Genome-wide association study of vitamin D concentrations in Hispanic Americans: the IRAS family study. 2010 , 122, 186-92		52
83	Joint effects of common genetic variants from multiple genes and pathways on the risk of premature coronary artery disease. 2010 , 160, 250-256.e3		26
82	Genetic variants of connexin37 are associated with carotid intima-medial thickness and future onset of ischemic stroke. 2011 , 214, 101-6		37
81	Determinants of plasma vitamin D levels in patients with acute coronary syndromes. 2011 , 41, 1299-309		6
80	Maternal C-reactive protein levels in pregnancy are associated with wheezing and lower respiratory tract infections in the offspring. 2011 , 204, 164.e1-9		22
79	Genetic polymorphisms of innate immunity-related inflammatory pathways and their association with factors related to type 2 diabetes. <i>BMC Medical Genetics</i> , 2011 , 12, 95	2.1	47
78	Genome-wide association analysis and fine mapping of NT-proBNP level provide novel insight into the role of the MTHFR-CLCN6-NPPA-NPPB gene cluster. 2011 , 20, 1660-71		38
77	Epidemiologic considerations in complex disease genetics. 2012 , 2012, 62-70		
76	Chapter 13: Mining electronic health records in the genomics era. 2012 , 8, e1002823		122
75	A genome-wide association scan on the levels of markers of inflammation in Sardinians reveals associations that underpin its complex regulation. 2012 , 8, e1002480		112
74	Structural effects of linkage disequilibrium on the transcriptome. 2012 , 18, 77-87		34
73	Effects of genetic variation on chromatin structure and the transcriptional machinery: analysis of the IL6 gene locus. 2012 , 13, 583-6		7
72	Vitamin D: Genetics and Genomic Effects. 2012 , 151-178		2
71	Genome-wide association replicates the association of Duffy antigen receptor for chemokines (DARC) polymorphisms with serum monocyte chemoattractant protein-1 (MCP-1) levels in Hispanic children. 2012 , 60, 634-8		17
70	A genome-wide association study of total bilirubin and cholelithiasis risk in sickle cell anemia. 2012 , 7, e34741		43

69	Heritability, genetic correlation and linkage to the 9p21.3 region of mixed platelet-leukocyte conjugates in families with and without early myocardial infarction. 2013 , 23, 684-92	8
68	Genetic analysis of 16 NMR-lipoprotein fractions in humans, the GOLDN study. 2013 , 48, 155-65	29
67	Genetic regulation of vitamin D levels. 2013 , 92, 106-17	58
66	Relationship among circulating inflammatory proteins, platelet gene expression, and cardiovascular risk. 2013 , 33, 2666-73	49
65	A novel, privacy-preserving cryptographic approach for sharing sequencing data. 2013 , 20, 69-76	8
64	Predictors of hemoglobin variability in a population of weaning age (3- to 4-month old) rhesus monkeys. 2013 , 75, 1139-46	
63	Two variants of the C-reactive protein gene are associated with risk of pre-eclampsia in an American Indian population. 2013 , 8, e71231	14
62	Genome-wide linkage analysis of cardiovascular disease biomarkers in a large, multigenerational family. 2013 , 8, e71779	10
61	Single-Nucleotide Polymorphisms Within MicroRNAs Sequences and Their 3QTR Target Sites May Regulate Gene Expression in Gastrointestinal Tract Cancers. 2014 , 16, e16659	18
60	Mendelian randomization studies in coronary artery disease. 2014 , 35, 1917-24	122
59	Links between allergy and cardiovascular or hemostatic system. 2014 , 170, 278-85	38
58	Developmental plasticity of red blood cell homeostasis. 2014 , 89, 459-66	13
57	Identification of novel loci for bipolar I disorder in a multi-stage genome-wide association study. 2014 , 51, 58-64	22
56	The potential of the riboSNitch in personalized medicine. 2015 , 6, 517-32	31
55	Association of the C-Reactive Protein Gene (CRP) rs1205 C>T Polymorphism with Aortic Valve Calcification in Patients with Aortic Stenosis. 2015 , 16, 23745-59	10
54	Prediction of breast cancer survival using clinical and genetic markers by tumor subtypes. 2015 , 10, e0122413	8
53	Systems Genetic Validation of the SNP-Metabolite Association in Rice Via Metabolite-Pathway-Based Phenome-Wide Association Scans. 2015 , 6, 1027	4
52	Association of exome sequences with plasma C-reactive protein levels in >9000 participants. 2015 , 24, 559-71	31

51	Multiple conformations are a conserved and regulatory feature of the RB1 5'UTR. 2015 , 21, 1274-85	45
50	Association of CRP genetic variants with blood concentrations of C-reactive protein and colorectal cancer risk. 2015 , 136, 1181-92	53
49	Common variants in the CRP promoter are associated with a high C-reactive protein level in Kawasaki disease. 2015 , 36, 438-44	12
48	Genome-wide association analysis of plasma B-type natriuretic peptide in blacks: the Jackson Heart Study. 2015 , 8, 122-30	26
47	Potential relationship between single nucleotide polymorphisms used in forensic genetics and diseases or other traits in European population. 2015 , 129, 435-43	0
46	C-reactive protein genetic variant is associated with diabetic retinopathy in Chinese patients with type 2 diabetes. 2015 , 15, 8	10
45	A common variant in RAB27A gene is associated with fractional exhaled nitric oxide levels in adults. 2015 , 45, 797-806	9
44	Serum IgE levels are associated with coronary artery disease severity. 2016 , 251, 355-360	24
43	Mendelian Randomization for the Identification of Causal Pathways in Atherosclerotic Vascular Disease. 2016 , 30, 41-9	8
42	Shared genetic susceptibility of vascular-related biomarkers with ischemic and recurrent stroke. 2016 , 86, 351-9	26
41	Genome-Wide Association Analysis Reveals Genetic Heterogeneity of Sjögren's Syndrome According to Ancestry. 2017 , 69, 1294-1305	56
40	Enhanced Missing Proteins Detection in NCI60 Cell Lines Using an Integrative Search Engine Approach. 2017 , 16, 4374-4390	6
39	Coronary collateralization shows sex and racial-ethnic differences in obstructive artery disease patients. 2017 , 12, e0183836	11
38	From genome-wide association studies to Mendelian randomization: novel opportunities for understanding cardiovascular disease causality, pathogenesis, prevention, and treatment. 2018 , 114, 1192-1208	36
37	Incidence and Risk of Gallstone Disease in Gilbert's Syndrome Patients in Indian Population. 2018 , 8, 362-366	9
36	Genetic and epigenetic factors influencing vitamin D status. 2018 , 233, 4033-4043	58
35	Genetic variation in 9p21 is associated with fasting insulin in women but not men. 2018 , 13, e0202365	3
34	Genome-Wide Association Study of Serum 25-Hydroxyvitamin D in US Women. 2018 , 9, 67	21

33	SNPs related to vitamin D and breast cancer risk: a case-control study. 2018 , 20, 1	18
32	CYP27A1, CYP24A1, and RXR- β Polymorphisms, Vitamin D, and Multiple Sclerosis: a Pilot Study. 2018 , 66, 77-84	19
31	Genetics of the Vitamin D Endocrine System. 2018 , 151-165	
30	Hypoxia Signaling in Vascular Homeostasis. 2018 , 33, 328-337	10
29	The Relationship of Serum Antigen-Specific and Total Immunoglobulin E with Adult Cardiovascular Diseases. 2018 , 15, 1098-1104	2
28	Identification of Novel Genetic Polymorphisms Associated with Type 2 Diabetes Mellitus: A Chinese Sib-Pair Study. 2019 , 23, 435-441	2
27	A critical evaluation of results from genome-wide association studies of micronutrient status and their utility in the practice of precision nutrition. 2019 , 122, 121-130	3
26	Genetic analysis of hsCRP in American Indians: The Strong Heart Family Study. 2019 , 14, e0223574	2
25	Vitamin D-Related Genes, Blood Vitamin D Levels and Colorectal Cancer Risk in Western European Populations. 2019 , 11,	11
24	Genomewide association study for C-reactive protein in Indians replicates known associations of common variants. 2019 , 98, 1	7
23	The genetics of vitamin D. 2019 , 126, 59-77	27
22	LRP1 and APOA1 Polymorphisms: Impact on Warfarin International Normalized Ratio-Related Phenotypes. 2020 , 76, 71-76	0
21	polymorphisms associated with warfarin stable dose in Chinese patients: a stepwise conditional analysis. 2020 , 21, 1169-1178	0
20	DNA methylation loci in placenta associated with birthweight and expression of genes relevant for early development and adult diseases. 2020 , 12, 78	14
19	Genetic Contributions to Maternal and Neonatal Vitamin D Levels. 2020 , 214, 1091-1102	3
18	The Role of Electronic Health Records in Advancing Genomic Medicine. 2021 , 22, 219-238	2
17	Genetic Link Determining the Maternal-Fetal Circulation of Vitamin D. 2021 , 12, 721488	1
16	Plasma Metabolomic Signatures of Chronic Obstructive Pulmonary Disease and the Impact of Genetic Variants on Phenotype-Driven Modules. 2020 , 3, 159-181	8

15	Identifying Pleiotropic Effects: A Two-Stage Approach Using Genome-Wide Association Meta-Analysis Data.	2
14	Placental genome and maternal-placental genetic interactions: a genome-wide and candidate gene association study of placental abruption. 2014 , 9, e116346	22
13	[Current status of genome-wide association study]. 2011 , 33, 25-35	2
12	Obesity, Metabolic Syndrome and Inflammation. 2022 , 133-149	
11	"Diabetes Associated Genes from the Dark Matter of the Human Proteome". 2014 , 1,	1
10	Quantile-specific heritability of monocyte chemoattractant protein-1, and relevance to rs1024611-disease interactions. 2022 , 149, 155722	1
9	Evaluating the Risk Factors of Development and Progression of Diabetic Retinopathy: A Review Study. 2022 , In Press,	
8	Identification of microsatellite markers and their application in yellow catfish (Richardson, 1846) population genetics of Korea. 2019 , 98,	
7	DataSheet1.docx. 2018 ,	
6	Pre-diagnostic C-reactive protein concentrations, CRP genetic variation and mortality among individuals with colorectal cancer in Western European populations. 2022 , 22,	
5	Vitamin D Metabolism Gene Polymorphisms and Their Associated Disorders: A Literature Review. 2022 , 23,	0
4	Features of gallstones in adult sickle cell patients. 2022 , 25, 1-18	
3	Genetic Determinants of 25-Hydroxyvitamin D Concentrations and Their Relevance to Public Health. 2022 , 14, 4408	3
2	The genetic and epigenetic contributions to the development of nutritional rickets. 13,	0
1	Vitamin D Determinants, Status, and Antioxidant/Antiinflammatory-Related Effects in Cardiovascular Risk and Disease: Not the Last Word in the Controversy. 2023 , 12, 948	0