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Genome-wide association with diabetes-related traits in the Framingham Heart Study

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#	Paper	IF	Citations
74	A 100K genome-wide association scan for diabetes and related traits in the Framingham Heart Study: replication and integration with other genome-wide datasets. <i>Diabetes</i> , 2007 , 56, 3063-74	0.9	74
73	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
72	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
71	Ordered stratification to reduce heterogeneity in linkage to diabetes-related quantitative traits. <i>Obesity</i> , 2008 , 16, 2314-22	8	3
70	Prevalence in the United States of selected candidate gene variants: Third National Health and Nutrition Examination Survey, 1991-1994. <i>American Journal of Epidemiology</i> , 2009 , 169, 54-66	3.8	72
69	Family study designs in the age of genome-wide association studies: experience from the Framingham Heart Study. <i>Current Opinion in Lipidology</i> , 2008 , 19, 144-50	4.4	11
68	Family-based genome-wide association studies. <i>Pharmacogenomics</i> , 2009 , 10, 181-90	2.6	53
67	Risk loci for type 2 diabetes - quo vadis?. <i>Clinical Chemistry and Laboratory Medicine</i> , 2009 , 47, 383-6	5.9	3
66	A genomics study of type 2 diabetes mellitus in U.S. Air Force personnel. <i>Journal of Diabetes Science and Technology</i> , 2009 , 3, 770-5	4.1	2
65	Epistatic interactions of CDKN2B-TCF7L2 for risk of type 2 diabetes and of CDKN2B-JAZF1 for triglyceride/high-density lipoprotein ratio longitudinal change: evidence from the Framingham Heart Study. <i>BMC Proceedings</i> , 2009 , 3 Suppl 7, S71	2.3	9
64	Estudio del componente genético de la cardiopatía isquémica: de los estudios de ligamiento al genotipado integral del genoma. <i>Revista Espanola De Cardiologia Suplementos</i> , 2009 , 9, 24-38	0.2	2
63	The genetics of autoimmune diseases: a networked perspective. <i>Current Opinion in Immunology</i> , 2009 , 21, 596-605	7.8	110
62	Agreement among type 2 diabetes linkage studies but a poor correlation with results from genome-wide association studies. <i>Diabetologia</i> , 2009 , 52, 1061-74	10.3	17
61	Association analysis of v-AKT murine thymoma viral oncogene homolog 1 (AKT1) polymorphisms and type 2 diabetes mellitus in the Korean population. <i>Genes and Genomics</i> , 2009 , 31, 73-83	2.1	1
60	ATRIUM: testing untyped SNPs in case-control association studies with related individuals. <i>American Journal of Human Genetics</i> , 2009 , 85, 667-78	11	6
59	Genetic risk factors for type 2 diabetes with pharmacologic intervention in African-American patients with schizophrenia or schizoaffective disorder. <i>Schizophrenia Research</i> , 2009 , 114, 50-6	3.6	12
58	Antihypertensive pharmacogenetics: missed opportunity. <i>Journal of Hypertension</i> , 2010 , 28, 2007-9	1.9	1

57	The contribution of enteroinsular hormones to the pathogenesis of type 2 diabetes mellitus. <i>Current Diabetes Reports</i> , 2010 , 10, 192-8	5.6	2
56	Cardiovascular disease risk factors, type 2 diabetes mellitus, and the Framingham Heart Study. <i>Trends in Cardiovascular Medicine</i> , 2010 , 20, 90-5	6.9	108
55	Large effects on body mass index and insulin resistance of fat mass and obesity associated gene (FTO) variants in patients with polycystic ovary syndrome (PCOS). <i>BMC Medical Genetics</i> , 2010 , 11, 12	2.1	56
54	Considerations for designing a prototype genetic test for use in translational research. <i>Public Health Genomics</i> , 2010 , 13, 155-65	1.9	22
53	The genetics of obesity and the metabolic syndrome. <i>Endocrine, Metabolic and Immune Disorders - Drug Targets</i> , 2010 , 10, 86-108	2.2	47
52	A genome-wide association study identifies a novel major locus for glycemic control in type 1 diabetes, as measured by both A1C and glucose. <i>Diabetes</i> , 2010 , 59, 539-49	0.9	81
51	Genome-wide significant associations for variants with minor allele frequency of 5% or less--an overview: A HuGE review. <i>American Journal of Epidemiology</i> , 2010 , 172, 869-89	3.8	38
50	Genomics of heart failure. <i>Heart Failure Clinics</i> , 2010 , 6, 115-24	3.3	9
49	A survey of the genetics of stomach, liver, and adipose gene expression from a morbidly obese cohort. <i>Genome Research</i> , 2011 , 21, 1008-16	9.7	141
48	Genetic and clinical risk factors of new-onset diabetes after transplantation in Hispanic kidney transplant recipients. <i>Transplantation</i> , 2011 , 91, 1114-9	1.8	34
47	The "thrifty" gene encoding Ahsg/Fetuin-A meets the insulin receptor: Insights into the mechanism of insulin resistance. <i>Cellular Signalling</i> , 2011 , 23, 980-90	4.9	61
46	Pharmacogenomics of oral antidiabetic medications: current data and pharmacoepigenomic perspective. <i>Pharmacogenomics</i> , 2011 , 12, 1161-91	2.6	53
45	Evaluation of four novel genetic variants affecting hemoglobin A1c levels in a population-based type 2 diabetes cohort (the HUNT2 study). <i>BMC Medical Genetics</i> , 2011 , 12, 20	2.1	10
44	Typ-2-Diabetes-assozierte Gene. <i>Diabetologe</i> , 2012 , 8, 26-34	0.2	3
43	Does familial clustering of risk factors for long-term diabetic complications leave any place for genes that act independently?. <i>Journal of Cardiovascular Translational Research</i> , 2012 , 5, 388-98	3.3	6
42	Expression quantitative trait loci analysis identifies associations between genotype and gene expression in human intestine. <i>Gastroenterology</i> , 2013 , 144, 1488-96, 1496.e1-3	13.3	51
41	Genetic analysis of 16 NMR-lipoprotein fractions in humans, the GOLDN study. <i>Lipids</i> , 2013 , 48, 155-65	1.6	29
40	Common genetic variants in peroxisome proliferator-activated receptor- γ (PPARG) and type 2 diabetes risk among Women@Health Initiative postmenopausal women. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2013 , 98, E600-4	5.6	15

39	Combined genome-wide linkage and association analyses of fasting glucose level in healthy twins and families of Korea. <i>Journal of Korean Medical Science</i> , 2013 , 28, 415-23	4.7	1
38	Autoimmune Disorders. 2013 , 822-838		3
37	Genome-wide association studies and heritability estimates of body mass index related phenotypes in Bangladeshi adults. <i>PLoS ONE</i> , 2014 , 9, e105062	3.7	10
36	Genomic structure of nucleotide diversity among Lyon rat models of metabolic syndrome. <i>BMC Genomics</i> , 2014 , 15, 197	4.5	7
35	Insights into the genetic susceptibility to type 2 diabetes from genome-wide association studies of glycaemic traits. <i>Current Diabetes Reports</i> , 2014 , 14, 551	5.6	34
34	Sex- and age-interacting eQTLs in human complex diseases. <i>Human Molecular Genetics</i> , 2014 , 23, 1947-56	6.6	48
33	APOE and BCHE as modulators of cerebral amyloid deposition: a florbetapir PET genome-wide association study. <i>Molecular Psychiatry</i> , 2014 , 19, 351-7	15.1	145
32	No Evidence of a Causal Relationship between Plasma Homocysteine and Type 2 Diabetes: A Mendelian Randomization Study. <i>Frontiers in Cardiovascular Medicine</i> , 2015 , 2, 11	5.4	17
31	Parent-of-Origin Effects of the APOB Gene on Adiposity in Young Adults. <i>PLoS Genetics</i> , 2015 , 11, e1005673		9
30	Understanding multicellular function and disease with human tissue-specific networks. <i>Nature Genetics</i> , 2015 , 47, 569-76	36.3	473
29	The Impact of Genetic Variants for Different Physiological Characterization of Type 2 Diabetes Loci on Gestational Insulin Signaling in Nondiabetic Pregnant Chinese Women. <i>Reproductive Sciences</i> , 2015 , 22, 1421-8	3	8
28	Characteristics of Bipolar I patients grouped by externalizing disorders. <i>Journal of Affective Disorders</i> , 2015 , 178, 206-14	6.6	7
27	The uniform-score gene set analysis for identifying common pathways associated with different diabetes traits. <i>BMC Genomics</i> , 2015 , 16, 336	4.5	4
26	Genome-wide interaction study of gene-by-occupational exposure and effects on FEV1 levels. <i>Journal of Allergy and Clinical Immunology</i> , 2015 , 136, 1664-1672.e14	11.5	27
25	Bayesian Variable Selection in Multilevel Item Response Theory Models with Application in Genomics. <i>Genetic Epidemiology</i> , 2016 , 40, 253-63	2.6	2
24	The Framingham Heart Study--67 years of discovery in metabolic disease. <i>Nature Reviews Endocrinology</i> , 2016 , 12, 177-83	15.2	37
23	PHLPP phosphatases as a therapeutic target in insulin resistance-related diseases. <i>Expert Opinion on Therapeutic Targets</i> , 2016 , 20, 663-75	6.4	9
22	Genetic variants in , and are associated with type 2 diabetes, BMI and dyslipidemia in families of Northeastern Mexico: A pilot study. <i>Experimental and Therapeutic Medicine</i> , 2017 , 13, 523-529	2.1	6

21	Linking Alzheimer [®] disease and type 2 diabetes: Novel shared susceptibility genes detected by cFDR approach. <i>Journal of the Neurological Sciences</i> , 2017 , 380, 262-272	3.2	23
20	Convergent and divergent genetic changes in the genome of Chinese and European pigs. <i>Scientific Reports</i> , 2017 , 7, 8662	4.9	12
19	Shared genetic regulatory networks for cardiovascular disease and type 2 diabetes in multiple populations of diverse ethnicities in the United States. <i>PLoS Genetics</i> , 2017 , 13, e1007040	6	48
18	Revealing pathway cross-talk related to diabetes mellitus by Monte Carlo Cross-Validation analysis. <i>Open Life Sciences</i> , 2017 , 12, 473-480	1.2	
17	Brief Overview of a Decade of Genome-Wide Association Studies on Primary Hypertension. <i>International Journal of Endocrinology</i> , 2018 , 2018, 7259704	2.7	6
16	Genome-wide association study: Exploring the genetic basis for responsiveness to ketogenic dietary therapies for drug-resistant epilepsy. <i>Epilepsia</i> , 2018 , 59, 1557-1566	6.4	6
15	Heritability and genome-wide association analyses of fasting plasma glucose in Chinese adult twins. <i>BMC Genomics</i> , 2020 , 21, 491	4.5	1
14	Genome-Wide Selection Scan in an Arabian Peninsula Population Identifies a TNKS Haplotype Linked to Metabolic Traits and Hypertension. <i>Genome Biology and Evolution</i> , 2020 , 12, 77-87	3.9	4
13	Identification of transcriptome alterations in the prefrontal cortex, hippocampus, amygdala and hippocampus of suicide victims. <i>Scientific Reports</i> , 2021 , 11, 18853	4.9	5
12	Identification of ALK in Thinness. <i>Cell</i> , 2020 , 181, 1246-1262.e22	56.2	21
11	Identifying tissues implicated in Anorexia Nervosa using Transcriptomic Imputation.		1
10	Placental genome and maternal-placental genetic interactions: a genome-wide and candidate gene association study of placental abruption. <i>PLoS ONE</i> , 2014 , 9, e116346	3.7	22
9	Polymorphisms associated with type 2 diabetes in familial longevity: The Leiden Longevity Study. <i>Aging</i> , 2011 , 3, 55-62	5.6	17
8	No Association of Obesity and Type 2 Diabetes Mellitus Related Genetic Variants With Colon Cancer. <i>Gastroenterology Research</i> , 2009 , 2, 311-316	1.8	
7	Mapping of Susceptibility Genes for Obesity, Type 2 Diabetes, and the Metabolic Syndrome in Human Populations. 2015 , 181-245		
6	Pathways to neurodegeneration: mechanistic insights from GWAS in Alzheimer [®] disease, Parkinson [®] disease, and related disorders. <i>American Journal of Neurodegenerative Disease</i> , 2013 , 2, 145-75	2.5	106
5	Promoter DNA Methylation in GWAS-Identified Genes as Potential Functional Elements for Blood Pressure: An Observational and Mendelian Randomization Study.. <i>Frontiers in Genetics</i> , 2021 , 12, 791146	4.5	0
4	Linking Variants of Hemoglobin A1C and Glycemic Status. <i>Biomarkers in Disease</i> , 2022 , 1-17		

- 3 Graph pangenome reveals functional, evolutionary, and phenotypic significance of human nonreference sequences.
- 2 Linking Variants of Hemoglobin A1C and Glycemic Status. **2023**, 3-19
- 1 PHLPP Signaling in Immune Cells. **2022**, 117-143