Hugoline G De Haan

List of Publications by Year
in descending order

25	2,850	471509 17	580821 25
papers	citations	h-index	g-index

26	26	26	6952
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Genome-wide associations for birth weight and correlations with adult disease. Nature, 2016, 538, 248-252.	27.8	406
2	Maternal and fetal genetic effects on birth weight and their relevance to cardio-metabolic risk factors. Nature Genetics, 2019, 51, 804-814.	21.4	402
3	Refining the accuracy of validated target identification through coding variant fine-mapping in type 2 diabetes. Nature Genetics, 2018, 50, 559-571.	21.4	356
4	The trans-ancestral genomic architecture of glycemic traits. Nature Genetics, 2021, 53, 840-860.	21.4	341
5	Genome-wide analysis identifies 12 loci influencing human reproductive behavior. Nature Genetics, 2016, 48, 1462-1472.	21.4	284
6	Meta-analysis of 65,734 Individuals Identifies TSPAN15 and SLC44A2 as Two Susceptibility Loci for Venous Thromboembolism. American Journal of Human Genetics, 2015, 96, 532-542.	6.2	222
7	Directional dominance on stature and cognition inÂdiverse human populations. Nature, 2015, 523, 459-462.	27.8	173
8	Multiple SNP testing improves risk prediction of first venous thrombosis. Blood, 2012, 120, 656-663.	1.4	132
9	Associations of Mitochondrial and Nuclear Mitochondrial Variants and Genes with Seven Metabolic Traits. American Journal of Human Genetics, 2019, 104, 112-138.	6.2	106
10	Genome-Wide Association Transethnic Meta-Analyses Identifies Novel Associations Regulating Coagulation Factor VIII and von Willebrand Factor Plasma Levels. Circulation, 2019, 139, 620-635.	1.6	102
11	Associations of autozygosity with a broad range of human phenotypes. Nature Communications, 2019, 10, 4957.	12.8	84
12	Assessing the causal relationship between obesity and venous thromboembolism through a Mendelian Randomization study. Human Genetics, 2017, 136, 897-902.	3.8	46
13	Genome-wide association study with additional genetic and post-transcriptional analyses reveals novel regulators of plasma factor XI levels. Human Molecular Genetics, 2017, 26, ddw401.	2.9	35
14	A genome-wide association study identifies new loci for factor VII and implicates factor VII in ischemic stroke etiology. Blood, 2019, 133, 967-977.	1.4	34
15	Discovery of novel heart rate-associated loci using the Exome Chip. Human Molecular Genetics, 2017, 26, 2346-2363.	2.9	29
16	Genetic susceptibility to radiation-induced breast cancer after Hodgkin lymphoma. Blood, 2019, 133, 1130-1139.	1.4	29
17	Targeted sequencing to identify novel genetic risk factors for deep vein thrombosis: a study of 734 genes. Journal of Thrombosis and Haemostasis, 2018, 16, 2432-2441.	3.8	17
18	Identification of coagulation gene 3â€²UTR variants that are potentially regulated by microRNAs. British Journal of Haematology, 2017, 177, 782-790.	2.5	15

#	Article	IF	Citations
19	Single Nucleotide Variant rs2232710 in the Protein Z-Dependent Protease Inhibitor (ZPI, SERPINA10) Gene Is Not Associated with Deep Vein Thrombosis. PLoS ONE, 2016, 11, e0151347.	2.5	9
20	Genetic determinants of activity and antigen levels of contact system factors. Journal of Thrombosis and Haemostasis, 2019, 17, 157-168.	3.8	7
21	Next-Generation Sequencing and In Vitro Expression Study of ADAMTS13 Single Nucleotide Variants in Deep Vein Thrombosis. PLoS ONE, 2016, 11, e0165665.	2.5	7
22	Male-specific risk of first and recurrent venous thrombosis: a phylogenetic analysis of the Y chromosome. Journal of Thrombosis and Haemostasis, 2016, 14, 1971-1977.	3.8	5
23	Genetic variants in Cell Adhesion Molecule 1 (CADM1): A validation study of a novel endothelial cell venous thrombosis risk factor. Thrombosis Research, 2014, 134, 1186-1192.	1.7	4
24	Next-generation DNA sequencing to identify novel genetic risk factors for cerebral vein thrombosis. Thrombosis Research, 2018, 169, 76-81.	1.7	4
25	Clinical and laboratory predictors of deep vein thrombosis after acute stroke; does D-dimer really improve predictive power?. Thrombosis Research, 2016, 146, 131-132.	1.7	1