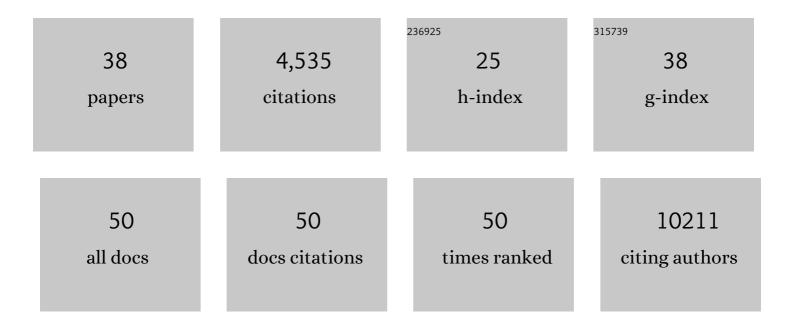
Marcel den Hoed

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Multiancestry genome-wide association study of 520,000 subjects identifies 32 loci associated with stroke subtypes. Nature Genetics, 2018, 50, 524-537.	21.4	1,124
2	Variability in the Heritability of Body Mass Index: A Systematic Review and Meta-Regression. Frontiers in Endocrinology, 2012, 3, 29.	3.5	489
3	New loci associated with birth weight identify genetic links between intrauterine growth and adult height and metabolism. Nature Genetics, 2013, 45, 76-82.	21.4	293
4	Genome-wide analysis identifies 12 loci influencing human reproductive behavior. Nature Genetics, 2016, 48, 1462-1472.	21.4	284
5	Identification of heart rate–associated loci and their effects on cardiac conduction and rhythm disorders. Nature Genetics, 2013, 45, 621-631.	21.4	282
6	Genetic association study of QT interval highlights role for calcium signaling pathways in myocardial repolarization. Nature Genetics, 2014, 46, 826-836.	21.4	281
7	Genome-wide physical activity interactions in adiposity ― A meta-analysis of 200,452 adults. PLoS Genetics, 2017, 13, e1006528.	3.5	158
8	Assessing Causality in the Association between Child Adiposity and Physical Activity Levels: A Mendelian Randomization Analysis. PLoS Medicine, 2014, 11, e1001618.	8.4	147
9	Identification of additional risk loci for stroke and small vessel disease: a meta-analysis of genome-wide association studies. Lancet Neurology, The, 2016, 15, 695-707.	10.2	130
10	Heritability of objectively assessed daily physical activity and sedentary behavior. American Journal of Clinical Nutrition, 2013, 98, 1317-1325.	4.7	121
11	Genetic Susceptibility to Obesity and Related Traits in Childhood and Adolescence. Diabetes, 2010, 59, 2980-2988.	0.6	120
12	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.	12.4	100
13	Genetic loci associated with heart rate variability and their effects on cardiac disease risk. Nature Communications, 2017, 8, 15805.	12.8	95
14	Postprandial responses in hunger and satiety are associated with the rs9939609 single nucleotide polymorphism in FTO. American Journal of Clinical Nutrition, 2009, 90, 1426-1432.	4.7	93
15	Biological/Genetic Regulation of Physical Activity Level. Medicine and Science in Sports and Exercise, 2018, 50, 863-873.	0.4	80
16	PR interval genome-wide association meta-analysis identifies 50 loci associated with atrial and atrioventricular electrical activity. Nature Communications, 2018, 9, 2904.	12.8	71
17	Genome-wide discovery of genetic loci that uncouple excess adiposity from its comorbidities. Nature Metabolism, 2021, 3, 228-243.	11.9	70
18	CRISPR-Cas9 induces large structural variants at on-target and off-target sites in vivo that segregate across generations. Nature Communications, 2022, 13, 627.	12.8	65

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#	Article	IF	CITATIONS
19	Identification of 371 genetic variants for age at first sex and birth linked to externalising behaviour. Nature Human Behaviour, 2021, 5, 1717-1730.	12.0	62
20	Obesity-susceptibility loci have a limited influence on birth weight: a meta-analysis of up to 28,219 individuals. American Journal of Clinical Nutrition, 2011, 93, 851-860.	4.7	58
21	Contribution of common non-synonymous variants in PCSK1 to body mass index variation and risk of obesity: a systematic review and meta-analysis with evidence from up to 331 175 individuals. Human Molecular Genetics, 2015, 24, 3582-3594.	2.9	53
22	Relationship between perilipin gene polymorphisms and body weight and body composition during weight loss and weight maintenance. Physiology and Behavior, 2009, 96, 723-728.	2.1	37
23	Amplification-free long-read sequencing reveals unforeseen CRISPR-Cas9 off-target activity. Genome Biology, 2020, 21, 290.	8.8	35
24	Habitual physical activity in daily life correlates positively with markers for mitochondrial capacity. Journal of Applied Physiology, 2008, 105, 561-568.	2.5	33
25	Pleiotropic effects of obesity-susceptibility loci on metabolic traits: a meta-analysis of up to 37,874 individuals. Diabetologia, 2013, 56, 2134-2146.	6.3	32
26	GWAS-identified loci for coronary heart disease are associated with intima-media thickness and plaque presence at the carotid artery bulb. Atherosclerosis, 2015, 239, 304-310.	0.8	31
27	Body composition is associated with physical activity in daily life as measured using a triaxial accelerometer in both men and women. International Journal of Obesity, 2008, 32, 1264-1270.	3.4	20
28	Testâ€retest variability of VO _{2max} using totalâ€capture indirect calorimetry reveals linear relationship of VO ₂ and Power. Scandinavian Journal of Medicine and Science in Sports, 2019, 29, 213-222.	2.9	18
29	SNP analyses of postprandial responses in (an)orexigenic hormones and feelings of hunger reveal long-term physiological adaptations to facilitate homeostasis. International Journal of Obesity, 2008, 32, 1790-1798.	3.4	17
30	Evaluation of common genetic variants identified by GWAS for early onset and morbid obesity in population-based samples. International Journal of Obesity, 2013, 37, 191-196.	3.4	16
31	Dysregulation of macrophage PEPD in obesity determines adipose tissue fibro-inflammation and insulin resistance. Nature Metabolism, 2022, 4, 476-494.	11.9	16
32	Maternal diabetes and incidence of childhood cancer – a nationwide cohort study and exploratory genetic analysis. Clinical Epidemiology, 2017, Volume 9, 633-642.	3.0	12
33	Translating GWAS-identified loci for cardiac rhythm and rate using an in vivo image- and CRISPR/Cas9-based approach. Scientific Reports, 2020, 10, 11831.	3.3	12
34	Multiâ€phenotype analyses of hemostatic traits with cardiovascular events reveal novel genetic associations. Journal of Thrombosis and Haemostasis, 2022, 20, 1331-1349.	3.8	12
35	Body composition in 10–13-year-old children: A comparison between air displacement plethysmography and deuterium dilution. Pediatric Obesity, 2009, 4, 397-404.	3.2	7
36	Skeletal muscle fiberâ€ŧype distribution and habitual physical activity in daily life. Scandinavian Journal of Medicine and Science in Sports, 2009, 19, 373-380.	2.9	5

#	Article	IF	CITATIONS
37	Identification of a novel proinsulin-associated SNP and demonstration that proinsulin is unlikely to be a causal factor in subclinical vascular remodelling using Mendelian randomisation. Atherosclerosis, 2017, 266, 196-204.	0.8	3

Gender, Ethnic, and Geographic Variation in Adiposity. , 2014, , 97-112.