

# Ilja M Nolte

## List of Publications by Year in descending order

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Version: 2024-02-01

161  
papers

27,460  
citations

20759

60  
h-index

7496

151  
g-index

173  
all docs

173  
docs citations

173  
times ranked

35160  
citing authors

#	ARTICLE	IF	CITATIONS
1	Polygenic risk for aggressive behavior from late childhood through early adulthood. <i>European Child and Adolescent Psychiatry</i> , 2023, 32, 651-660.	2.8	6
2	Plasma Lead Concentration and Risk of Late Kidney Allograft Failure: Findings From the TransplantLines Biobank and Cohort Studies. <i>American Journal of Kidney Diseases</i> , 2022, 80, 87-97.e1.	2.1	6
3	Killer Cell Immunoglobulin-Like Receptor Haplotype B Modulates Susceptibility to EBV-Associated Classic Hodgkin Lymphoma. <i>Frontiers in Immunology</i> , 2022, 13, 829943.	2.2	4
4	Early increase in single-kidney glomerular filtration rate after living kidney donation predicts long-term kidney function. <i>Kidney International</i> , 2022, 101, 1251-1259.	2.6	8
5	Genome-wide Association Meta-analysis of Childhood and Adolescent Internalizing Symptoms. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2022, 61, 934-945.	0.3	26
6	Influence of Receptor Polymorphisms on the Response to $\alpha_1$ -Adrenergic Receptor Blockers in Pheochromocytoma Patients. <i>Biomedicines</i> , 2022, 10, 896.	1.4	1
7	Plasma phosphate and all-cause mortality in individuals with and without type 2 diabetes: the Dutch population-based lifelines cohort study. <i>Cardiovascular Diabetology</i> , 2022, 21, 61.	2.7	2
8	MO944: Persistent Microscopic Hematuria At Kidney Donor Screening and Long-Term Post-Donation Kidney Outcomes. <i>Nephrology Dialysis Transplantation</i> , 2022, 37, .	0.4	0
9	Genetic loci and prioritization of genes for kidney function decline derived from a meta-analysis of 62 longitudinal genome-wide association studies. <i>Kidney International</i> , 2022, 102, 624-639.	2.6	18
10	Differential and shared genetic effects on kidney function between diabetic and non-diabetic individuals. <i>Communications Biology</i> , 2022, 5, .	2.0	17
11	Gene-environment interplay in externalizing behavior from childhood through adulthood. <i>Journal of Child Psychology and Psychiatry and Allied Disciplines</i> , 2022, 63, 1206-1213.	3.1	7
12	Thyroid function and risk of all-cause and cardiovascular mortality: a prospective population-based cohort study. <i>Endocrine</i> , 2021, 71, 385-396.	1.1	10
13	Serum uric acid is associated with increased risk of posttransplantation diabetes in kidney transplant recipients: a prospective cohort study. <i>Metabolism: Clinical and Experimental</i> , 2021, 116, 154465.	1.5	4
14	Multi-ancestry genome-wide association study accounting for gene-psychosocial factor interactions identifies novel loci for blood pressure traits. <i>Human Genetics and Genomics Advances</i> , 2021, 2, 100013.	1.0	2
15	Meta-analysis uncovers genome-wide significant variants for rapid kidney function decline. <i>Kidney International</i> , 2021, 99, 926-939.	2.6	42
16	Plasma cadmium is associated with increased risk of long-term kidney graft failure. <i>Kidney International</i> , 2021, 99, 1213-1224.	2.6	18
17	Sex-dimorphic genetic effects and novel loci for fasting glucose and insulin variability. <i>Nature Communications</i> , 2021, 12, 24.	5.8	87
18	GWASInspector: comprehensive quality control of genome-wide association study results. <i>Bioinformatics</i> , 2021, 37, 129-130.	1.8	11

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19	Genetic pre-screening for glaucoma in population-based epidemiology: protocol for a double-blind prospective screening study within Lifelines (EyeLife). <i>BMC Ophthalmology</i> , 2021, 21, 18.	0.6	9
20	Interaction between ERAP Alleles and HLA Class I Types Support a Role of Antigen Presentation in Hodgkin Lymphoma Development. <i>Cancers</i> , 2021, 13, 414.	1.7	6
21	Genome-wide association study of circulating interleukin 6 levels identifies novel loci. <i>Human Molecular Genetics</i> , 2021, 30, 393-409.	1.4	32
22	Multi-ancestry genome-wide gene×sleep interactions identify novel loci for blood pressure. <i>Molecular Psychiatry</i> , 2021, 26, 6293-6304.	4.1	13
23	The trans-ancestral genomic architecture of glycemic traits. <i>Nature Genetics</i> , 2021, 53, 840-860.	9.4	341
24	Meat intake and risk of mortality and graft failure in kidney transplant recipients. <i>American Journal of Clinical Nutrition</i> , 2021, 114, 1505-1517.	2.2	5
25	Genetic association study of childhood aggression across raters, instruments, and age. <i>Translational Psychiatry</i> , 2021, 11, 413.	2.4	31
26	Spontaneous baroreflex sensitivity and its association with age, sex, obesity indices and hypertension: a population study. <i>American Journal of Hypertension</i> , 2021, 34, 1276-1283.	1.0	8
27	The associations of <i>CNR1</i> SNPs and haplotypes with vulnerability and treatment response phenotypes in Han Chinese with major depressive disorder: A case×control association study. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2021, 9, e1752.	0.6	6
28	Continuity of Genetic Risk for Aggressive Behavior Across the Life-Course. <i>Behavior Genetics</i> , 2021, 51, 592-606.	1.4	13
29	Sex and Gender-Related Differences in COVID-19 Diagnoses and SARS-CoV-2 Testing Practices During the First Wave of the Pandemic: The Dutch Lifelines COVID-19 Cohort Study. <i>Journal of Women's Health</i> , 2021, 30, 1686-1692.	1.5	20
30	Lack of association of <i>FKBP5</i> SNPs and haplotypes with susceptibility and treatment response phenotypes in Han Chinese with major depressive disorder. <i>Medicine (United States)</i> , 2021, 100, e26983.	0.4	2
31	Galectin-3 and Risk of Late Graft Failure in Kidney Transplant Recipients: A 10-year Prospective Cohort Study. <i>Transplantation</i> , 2021, 105, 1106-1115.	0.5	8
32	HLA Expression in Relation to HLA Type in Classic Hodgkin Lymphoma Patients. <i>Cancers</i> , 2021, 13, 5833.	1.7	1
33	Mitochondrial Genome Study Identifies Association Between Primary Open-Angle Glaucoma and Variants in <i>MT-CYB</i> , <i>MT-ND4</i> Genes and Haplogroups. <i>Frontiers in Genetics</i> , 2021, 12, 781189.	1.1	13
34	Consumption of fruits and vegetables and cardiovascular mortality in renal transplant recipients: a prospective cohort study. <i>Nephrology Dialysis Transplantation</i> , 2020, 35, 357-365.	0.4	25
35	Metasubtract: an R-package to analytically produce leave-one-out meta-analysis GWAS summary statistics. <i>Bioinformatics</i> , 2020, 36, 4521-4522.	1.8	17
36	Heritability and the Genetic Correlation of Heart Rate Variability and Blood Pressure in >29,000 Families. <i>Hypertension</i> , 2020, 76, 1256-1262.	1.3	13

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37	Genetic Risk Scores for Complex Disease Traits in Youth. <i>Circulation Genomic and Precision Medicine</i> , 2020, 13, e002775.	1.6	17
38	Gene-educational attainment interactions in a multi-ancestry genome-wide meta-analysis identify novel blood pressure loci. <i>Molecular Psychiatry</i> , 2020, 26, 2111-2125.	4.1	17
39	Erythropoietin, Fibroblast Growth Factor 23, and Death After Kidney Transplantation. <i>Journal of Clinical Medicine</i> , 2020, 9, 1737.	1.0	0
40	The miR-26b-5p/KPNA2 Axis Is an Important Regulator of Burkitt Lymphoma Cell Growth. <i>Cancers</i> , 2020, 12, 1464.	1.7	19
41	Validating the doubly weighted genetic risk score for the prediction of type 2 diabetes in the Lifelines and Estonian Biobank cohorts. <i>Genetic Epidemiology</i> , 2020, 44, 589-600.	0.6	6
42	Altered Gut Microbial Fermentation and Colonization with <i>Methanobrevibacter smithii</i> in Renal Transplant Recipients. <i>Journal of Clinical Medicine</i> , 2020, 9, 518.	1.0	7
43	Heritability and genetic correlations of obesity indices with ambulatory and office beat-to-beat blood pressure in the Oman Family Study. <i>Journal of Hypertension</i> , 2020, 38, 1474-1480.	0.3	10
44	Duality of Tocopherol Isoforms and Novel Associations with Vitamins Involved in One-Carbon Metabolism: Results from an Elderly Sample of the LifeLines Cohort Study. <i>Nutrients</i> , 2020, 12, 580.	1.7	0
45	Ethnic differences in prevalence of Dupuytren disease can partly be explained by known genetic risk variants. <i>European Journal of Human Genetics</i> , 2019, 27, 1876-1884.	1.4	13
46	New alcohol-related genes suggest shared genetic mechanisms with neuropsychiatric disorders. <i>Nature Human Behaviour</i> , 2019, 3, 950-961.	6.2	75
47	Associations of autozygosity with a broad range of human phenotypes. <i>Nature Communications</i> , 2019, 10, 4957.	5.8	84
48	Using Polygenic Scores in Social Science Research: Unraveling Childlessness. <i>Frontiers in Sociology</i> , 2019, 4, 74.	1.0	4
49	Multi-ancestry sleep-by-SNP interaction analysis in 126,926 individuals reveals lipid loci stratified by sleep duration. <i>Nature Communications</i> , 2019, 10, 5121.	5.8	62
50	The association of depression and anxiety with cardiac autonomic activity: The role of confounding effects of antidepressants. <i>Depression and Anxiety</i> , 2019, 36, 1163-1172.	2.0	36
51	Target genes, variants, tissues and transcriptional pathways influencing human serum urate levels. <i>Nature Genetics</i> , 2019, 51, 1459-1474.	9.4	251
52	Multiancestry Genome-Wide Association Study of Lipid Levels Incorporating Gene-Alcohol Interactions. <i>American Journal of Epidemiology</i> , 2019, 188, 1033-1054.	1.6	85
53	Multi-ancestry study of blood lipid levels identifies four loci interacting with physical activity. <i>Nature Communications</i> , 2019, 10, 376.	5.8	64
54	Iron deficiency, elevated erythropoietin, fibroblast growth factor 23, and mortality in the general population of the Netherlands: A cohort study. <i>PLoS Medicine</i> , 2019, 16, e1002818.	3.9	16

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55	A catalog of genetic loci associated with kidney function from analyses of a million individuals. <i>Nature Genetics</i> , 2019, 51, 957-972.	9.4	549
56	The effect of high compared with low dairy consumption on glucose metabolism, insulin sensitivity, and metabolic flexibility in overweight adults: a randomized crossover trial. <i>American Journal of Clinical Nutrition</i> , 2019, 109, 1555-1568.	2.2	17
57	Genome-Wide Association Scan of Serum Urea in European Populations Identifies Two Novel Loci. <i>American Journal of Nephrology</i> , 2019, 49, 193-202.	1.4	5
58	Multi-ancestry genome-wide gene-smoking interaction study of 387,272 individuals identifies new loci associated with serum lipids. <i>Nature Genetics</i> , 2019, 51, 636-648.	9.4	112
59	A Weighted Genetic Risk Score Predicts Surgical Recurrence Independent of High-Risk Clinical Features in Dupuytren's Disease. <i>Plastic and Reconstructive Surgery</i> , 2019, 143, 512-518.	0.7	8
60	Heritability and genetic and environmental correlations of heart rate variability and baroreceptor reflex sensitivity with ambulatory and beat-to-beat blood pressure. <i>Scientific Reports</i> , 2019, 9, 1664.	1.6	8
61	Plasma versus Erythrocyte Vitamin E in Renal Transplant Recipients, and Duality of Tocopherol Species. <i>Nutrients</i> , 2019, 11, 2821.	1.7	2
62	The narrow-sense and common single nucleotide polymorphism heritability of early repolarization. <i>International Journal of Cardiology</i> , 2019, 279, 135-140.	0.8	7
63	KCND3 potassium channel gene variant confers susceptibility to electrocardiographic early repolarization pattern. <i>JCI Insight</i> , 2019, 4, .	2.3	15
64	Evaluation of a genetic risk score based on creatinine-estimated glomerular filtration rate and its association with kidney outcomes. <i>Nephrology Dialysis Transplantation</i> , 2018, 33, 1757-1764.	0.4	11
65	Active Smoking and Hematocrit and Fasting Circulating Erythropoietin Concentrations in the General Population. <i>Mayo Clinic Proceedings</i> , 2018, 93, 337-343.	1.4	16
66	Heritability and genetic correlations of heart rate variability at rest and during stress in the Oman Family Study. <i>Journal of Hypertension</i> , 2018, 36, 1477-1485.	0.3	13
67	Genetic regulation of <i>IL1RL1</i> methylation and <i>IL1RL1</i> -a protein levels in asthma. <i>European Respiratory Journal</i> , 2018, 51, 1701377.	3.1	24
68	GWAS and colocalization analyses implicate carotid intima-media thickness and carotid plaque loci in cardiovascular outcomes. <i>Nature Communications</i> , 2018, 9, 5141.	5.8	119
69	Switching iron sucrose to ferric carboxymaltose associates to better control of iron status in hemodialysis patients. <i>BMC Nephrology</i> , 2018, 19, 242.	0.8	11
70	Genome-wide analyses identify a role for <i>SLC17A4</i> and <i>AADAT</i> in thyroid hormone regulation. <i>Nature Communications</i> , 2018, 9, 4455.	5.8	181
71	Association of different iron deficiency cutoffs with adverse outcomes in chronic kidney disease. <i>BMC Nephrology</i> , 2018, 19, 225.	0.8	35
72	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

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73	PR interval genome-wide association meta-analysis identifies 50 loci associated with atrial and atrioventricular electrical activity. <i>Nature Communications</i> , 2018, 9, 2904.	5.8	71
74	Genome-wide association meta-analysis of age at first cannabis use. <i>Addiction</i> , 2018, 113, 2073-2086.	1.7	24
75	Novel genetic associations for blood pressure identified via gene-alcohol interaction in up to 570K individuals across multiple ancestries. <i>PLoS ONE</i> , 2018, 13, e0198166.	1.1	94
76	Genetic evidence of assortative mating in humans. <i>Nature Human Behaviour</i> , 2017, 1, .	6.2	242
77	Missing heritability: is the gap closing? An analysis of 32 complex traits in the Lifelines Cohort Study. <i>European Journal of Human Genetics</i> , 2017, 25, 877-885.	1.4	67
78	miR-24-3p Is Overexpressed in Hodgkin Lymphoma and Protects Hodgkin and Reed-Sternberg Cells from Apoptosis. <i>American Journal of Pathology</i> , 2017, 187, 1343-1355.	1.9	46
79	Genomic analyses identify hundreds of variants associated with age at menarche and support a role for puberty timing in cancer risk. <i>Nature Genetics</i> , 2017, 49, 834-841.	9.4	426
80	Genome-wide meta-analysis of 241,258 adults accounting for smoking behaviour identifies novel loci for obesity traits. <i>Nature Communications</i> , 2017, 8, 14977.	5.8	169
81	1000 Genomes-based meta-analysis identifies 10 novel loci for kidney function. <i>Scientific Reports</i> , 2017, 7, 45040.	1.6	98
82	Genetic loci associated with heart rate variability and their effects on cardiac disease risk. <i>Nature Communications</i> , 2017, 8, 15805.	5.8	95
83	CNV-association meta-analysis in 191,161 European adults reveals new loci associated with anthropometric traits. <i>Nature Communications</i> , 2017, 8, 744.	5.8	64
84	A Comparison of Heritability Estimates by Classical Twin Modeling and Based on Genome-Wide Genetic Relatedness for Cardiac Conduction Traits. <i>Twin Research and Human Genetics</i> , 2017, 20, 489-498.	0.3	14
85	Genetic and environmental influences on stability and change in baseline levels of C-reactive protein: A longitudinal twin study. <i>Atherosclerosis</i> , 2017, 265, 172-178.	0.4	13
86	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
87	Lipid and lipoprotein reference values from 133,450 Dutch Lifelines participants: Age- and gender-specific baseline lipid values and percentiles. <i>Journal of Clinical Lipidology</i> , 2017, 11, 1055-1064.e6.	0.6	67
88	C-Terminal Fibroblast Growth Factor 23, Iron Deficiency, and Mortality in Renal Transplant Recipients. <i>Journal of the American Society of Nephrology: JASN</i> , 2017, 28, 3639-3646.	3.0	46
89	Genotype-covariate interaction effects and the heritability of adult body mass index. <i>Nature Genetics</i> , 2017, 49, 1174-1181.	9.4	119
90	Associations between genetic risk, functional brain network organization and neuroticism. <i>Brain Imaging and Behavior</i> , 2017, 11, 1581-1591.	1.1	19

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91	Impact of common genetic determinants of Hemoglobin A1c on type 2 diabetes risk and diagnosis in ancestrally diverse populations: A transethnic genome-wide meta-analysis. <i>PLoS Medicine</i> , 2017, 14, e1002383.	3.9	341
92	Genome-wide physical activity interactions in adiposity • A meta-analysis of 200,452 adults. <i>PLoS Genetics</i> , 2017, 13, e1006528.	1.5	158
93	HLA expression and HLA type associations in relation to EBV status in Hispanic Hodgkin lymphoma patients. <i>PLoS ONE</i> , 2017, 12, e0174457.	1.1	7
94	Personality Polygenes, Positive Affect, and Life Satisfaction. <i>Twin Research and Human Genetics</i> , 2016, 19, 407-417.	0.3	16
95	Meta-analysis of 49â€¦549 individuals imputed with the 1000 Genomes Project reveals an exonic damaging variant in <i>ANGPTL4</i> determining fasting TG levels. <i>Journal of Medical Genetics</i> , 2016, 53, 441-449.	1.5	34
96	Genetic variants associated with subjective well-being, depressive symptoms, and neuroticism identified through genome-wide analyses. <i>Nature Genetics</i> , 2016, 48, 624-633.	9.4	870
97	A genome-wide approach to children's aggressive behavior: <i>The EAGLE consortium</i> . <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2016, 171, 562-572.	1.1	153
98	A principal component meta-analysis on multiple anthropometric traits identifies novel loci for body shape. <i>Nature Communications</i> , 2016, 7, 13357.	5.8	74
99	52 Genetic Loci Influencing Myocardial Mass. <i>Journal of the American College of Cardiology</i> , 2016, 68, 1435-1448.	1.2	113
100	The genetics of blood pressure regulation and its target organs from association studies in 342,415 individuals. <i>Nature Genetics</i> , 2016, 48, 1171-1184.	9.4	362
101	Genome-wide analysis identifies 12 loci influencing human reproductive behavior. <i>Nature Genetics</i> , 2016, 48, 1462-1472.	9.4	284
102	Does refining the phenotype improve replication rates? A review and replication of candidate gene studies on Major Depressive Disorder and Chronic Major Depressive Disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2016, 171, 215-236.	1.1	13
103	The association of single nucleotide polymorphisms of the maternal cystathionine-Î²-synthase gene with early-onset preeclampsia. <i>Pregnancy Hypertension</i> , 2016, 6, 60-65.	0.6	12
104	lodGWAS: a software package for genome-wide association analysis of biomarkers with a limit of detection. <i>Bioinformatics</i> , 2016, 32, 1552-1554.	1.8	5
105	Genetic associations at 53 loci highlight cell types and biological pathways relevant for kidney function. <i>Nature Communications</i> , 2016, 7, 10023.	5.8	412
106	Connecting the dots, genome-wide association studies in substance use. <i>Molecular Psychiatry</i> , 2016, 21, 733-735.	4.1	31
107	Investigating the Causal Relationship of C-Reactive Protein with 32 Complex Somatic and Psychiatric Outcomes: A Large-Scale Cross-Consortium Mendelian Randomization Study. <i>PLoS Medicine</i> , 2016, 13, e1001976.	3.9	150
108	Fine mapping the CETP region reveals a common intronic insertion associated to HDL-C. <i>Npj Aging and Mechanisms of Disease</i> , 2015, 1, 15011.	4.5	8



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109	The Influence of Age and Sex on Genetic Associations with Adult Body Size and Shape: A Large-Scale Genome-Wide Interaction Study. <i>PLoS Genetics</i> , 2015, 11, e1005378.	1.5	331
110	A Population Based Study of the Genetic Association between Catecholamine Gene Variants and Spontaneous Low-Frequency Fluctuations in Reaction Time. <i>PLoS ONE</i> , 2015, 10, e0126461.	1.1	2
111	Validity of (Ultra-)Short Recordings for Heart Rate Variability Measurements. <i>PLoS ONE</i> , 2015, 10, e0138921.	1.1	225
112	In Silico Post Genome-Wide Association Studies Analysis of C-Reactive Protein Loci Suggests an Important Role for Interferons. <i>Circulation: Cardiovascular Genetics</i> , 2015, 8, 487-497.	5.1	24
113	Interactions between uncoupling protein 2 gene polymorphisms, obesity and alcohol intake on liver function: a large meta-analysed population-based study. <i>European Journal of Endocrinology</i> , 2015, 173, 863-872.	1.9	7
114	New genetic loci link adipose and insulin biology to body fat distribution. <i>Nature</i> , 2015, 518, 187-196.	13.7	1,328
115	Genetic studies of body mass index yield new insights for obesity biology. <i>Nature</i> , 2015, 518, 197-206.	13.7	3,823
116	Dominance Genetic Variation Contributes Little to the Missing Heritability for Human Complex Traits. <i>American Journal of Human Genetics</i> , 2015, 96, 377-385.	2.6	191
117	Directional dominance on stature and cognition in diverse human populations. <i>Nature</i> , 2015, 523, 459-462.	13.7	173
118	Genetic variance estimation with imputed variants finds negligible missing heritability for human height and body mass index. <i>Nature Genetics</i> , 2015, 47, 1114-1120.	9.4	709
119	Genome-wide association study of kidney function decline in individuals of European descent. <i>Kidney International</i> , 2015, 87, 1017-1029.	2.6	113
120	Genome-wide meta-analysis identifies six novel loci associated with habitual coffee consumption. <i>Molecular Psychiatry</i> , 2015, 20, 647-656.	4.1	235
121	Modulation of Genetic Associations with Serum Urate Levels by Body-Mass-Index in Humans. <i>PLoS ONE</i> , 2015, 10, e0119752.	1.1	64
122	Postoperative Vision-Related Quality of Life in Macula-Off Rhegmatogenous Retinal Detachment Patients and Its Relation to Visual Function. <i>PLoS ONE</i> , 2014, 9, e114489.	1.1	14
123	Identifying Genetic Variants for Heart Rate Variability in the Acetylcholine Pathway. <i>PLoS ONE</i> , 2014, 9, e112476.	1.1	13
124	Genetic Associations in Classical Hodgkin Lymphoma: A Systematic Review and Insights into Susceptibility Mechanisms. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 2737-2747.	1.1	52
125	QCCWAS: A flexible R package for automated quality control of genome-wide association results. <i>Bioinformatics</i> , 2014, 30, 1185-1186.	1.8	22
126	Gene-centric Meta-analysis in 87,736 Individuals of European Ancestry Identifies Multiple Blood-Pressure-Related Loci. <i>American Journal of Human Genetics</i> , 2014, 94, 349-360.	2.6	158



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127	Pleiotropic genes for metabolic syndrome and inflammation. <i>Molecular Genetics and Metabolism</i> , 2014, 112, 317-338.	0.5	107
128	Gene-Age Interactions in Blood Pressure Regulation: A Large-Scale Investigation with the CHARGE, Global BPgen, and ICBP Consortia. <i>American Journal of Human Genetics</i> , 2014, 95, 24-38.	2.6	109
129	Parent-of-origin-specific allelic associations among 106 genomic loci for age at menarche. <i>Nature</i> , 2014, 514, 92-97.	13.7	548
130	Filling the Gap: Relationship Between the Serotonin-Transporter-Linked Polymorphic Region and Amygdala Activation. <i>Psychological Science</i> , 2014, 25, 2058-2066.	1.8	52
131	Defining the role of common variation in the genomic and biological architecture of adult human height. <i>Nature Genetics</i> , 2014, 46, 1173-1186.	9.4	1,818
132	Association of vitamin D status with arterial blood pressure and hypertension risk: a mendelian randomisation study. <i>Lancet Diabetes and Endocrinology</i> , 2014, 2, 719-729.	5.5	319
133	Genetic association study of QT interval highlights role for calcium signaling pathways in myocardial repolarization. <i>Nature Genetics</i> , 2014, 46, 826-836.	9.4	281
134	Novel childhood asthma genes interact with in utero and early-life tobacco smoke exposure. <i>Journal of Allergy and Clinical Immunology</i> , 2014, 133, 885-888.	1.5	47
135	Postoperative Recovery of Visual Function after Macula-Off Rhegmatogenous Retinal Detachment. <i>PLoS ONE</i> , 2014, 9, e99787.	1.1	19
136	Discovery and refinement of loci associated with lipid levels. <i>Nature Genetics</i> , 2013, 45, 1274-1283.	9.4	2,641
137	Common variants associated with plasma triglycerides and risk for coronary artery disease. <i>Nature Genetics</i> , 2013, 45, 1345-1352.	9.4	754
138	Loci influencing blood pressure identified using a cardiovascular gene-centric array. <i>Human Molecular Genetics</i> , 2013, 22, 1663-1678.	1.4	141
139	Genome-wide association analyses identify 18 new loci associated with serum urate concentrations. <i>Nature Genetics</i> , 2013, 45, 145-154.	9.4	675
140	Identification of heart rate-associated loci and their effects on cardiac conduction and rhythm disorders. <i>Nature Genetics</i> , 2013, 45, 621-631.	9.4	282
141	Sex-stratified Genome-wide Association Studies Including 270,000 Individuals Show Sexual Dimorphism in Genetic Loci for Anthropometric Traits. <i>PLoS Genetics</i> , 2013, 9, e1003500.	1.5	371
142	A Meta-Analysis of Thyroid-Related Traits Reveals Novel Loci and Gender-Specific Differences in the Regulation of Thyroid Function. <i>PLoS Genetics</i> , 2013, 9, e1003266.	1.5	194
143	Genome-Wide Association Study of Classical Hodgkin Lymphoma and Epstein-Barr Virus Status-Defined Subgroups. <i>Journal of the National Cancer Institute</i> , 2012, 104, 240-253.	3.0	141
144	Seventy-five genetic loci influencing the human red blood cell. <i>Nature</i> , 2012, 492, 369-375.	13.7	320

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145	HLA Associations in Classical Hodgkin Lymphoma: EBV Status Matters. PLoS ONE, 2012, 7, e39986.	1.1	52
146	Meta-Analysis of Genome-Wide Association Studies in >80 000 Subjects Identifies Multiple Loci for C-Reactive Protein Levels. Circulation, 2011, 123, 731-738.	1.6	461
147	New gene functions in megakaryopoiesis and platelet formation. Nature, 2011, 480, 201-208.	13.7	401
148	Genome-wide association study identifies six new loci influencing pulse pressure and mean arterial pressure. Nature Genetics, 2011, 43, 1005-1011.	9.4	403
149	A genome-wide association study of Hodgkin's lymphoma identifies new susceptibility loci at 2p16.1 (REL), 8q24.21 and 10p14 (GATA3). Nature Genetics, 2010, 42, 1126-1130.	9.4	177
150	Common variants in 22 loci are associated with QRS duration and cardiac ventricular conduction. Nature Genetics, 2010, 42, 1068-1076.	9.4	308
151	Genome-Wide Association Study of Blood Pressure Extremes Identifies Variant near UMOD Associated with Hypertension. PLoS Genetics, 2010, 6, e1001177.	1.5	312
152	Genome-wide association analysis identifies multiple loci related to resting heart rate. Human Molecular Genetics, 2010, 19, 3885-3894.	1.4	133
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