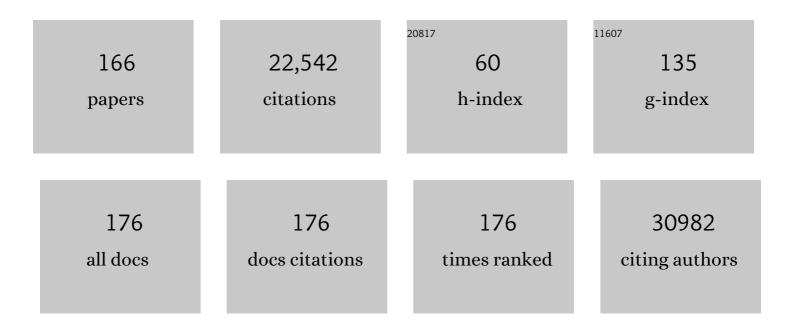
Gonneke Willemsen

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Association analyses of 249,796 individuals reveal 18 new loci associated with body mass index. Nature Genetics, 2010, 42, 937-948.	21.4	2,634
2	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. Nature Genetics, 2018, 50, 668-681.	21.4	2,224
3	Association studies of up to 1.2 million individuals yield new insights into the genetic etiology of tobacco and alcohol use. Nature Genetics, 2019, 51, 237-244.	21.4	1,307
4	Genome-wide association study identifies 74 loci associated with educational attainment. Nature, 2016, 533, 539-542.	27.8	1,204
5	Genetic analysis of over 1 million people identifies 535 new loci associated with blood pressure traits. Nature Genetics, 2018, 50, 1412-1425.	21.4	924
6	Genome-wide study for circulating metabolites identifies 62 loci and reveals novel systemic effects of LPA. Nature Communications, 2016, 7, 11122.	12.8	576
7	Parent-of-origin-specific allelic associations among 106 genomic loci for age at menarche. Nature, 2014, 514, 92-97.	27.8	548
8	Shared genetic origin of asthma, hay fever and eczema elucidates allergic disease biology. Nature Genetics, 2017, 49, 1752-1757.	21.4	432
9	Genomic analyses identify hundreds of variants associated with age at menarche and support a role for puberty timing in cancer risk. Nature Genetics, 2017, 49, 834-841.	21.4	426
10	Genome-wide associations for birth weight and correlations with adult disease. Nature, 2016, 538, 248-252.	27.8	406
11	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
12	Heritability and genomics of gene expression in peripheral blood. Nature Genetics, 2014, 46, 430-437.	21.4	370
13	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. Nature Genetics, 2015, 47, 1294-1303.	21.4	357
14	Netherlands Twin Register: From Twins to Twin Families. Twin Research and Human Genetics, 2006, 9, 849-857.	0.6	356
15	The power of genetic diversity in genome-wide association studies of lipids. Nature, 2021, 600, 675-679.	27.8	353
16	Polygenic risk scores for schizophrenia and bipolar disorder predict creativity. Nature Neuroscience, 2015, 18, 953-955.	14.8	351
17	The trans-ancestral genomic architecture of glycemic traits. Nature Genetics, 2021, 53, 840-860.	21.4	341
18	The Influence of Age and Sex on Genetic Associations with Adult Body Size and Shape: A Large-Scale Genome-Wide Interaction Study. PLoS Genetics, 2015, 11, e1005378.	3.5	331

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19	Genome Analyses of >200,000 Individuals Identify 58 Loci for Chronic Inflammation and Highlight Pathways that Link Inflammation and Complex Disorders. American Journal of Human Genetics, 2018, 103, 691-706.	6.2	326
20	Genetic and environmental influences interact with age and sex in shaping the human methylome. Nature Communications, 2016, 7, 11115.	12.8	299
21	Genome-wide analysis identifies 12 loci influencing human reproductive behavior. Nature Genetics, 2016, 48, 1462-1472.	21.4	284
22	Selective maternal seeding and environment shape the human gut microbiome. Genome Research, 2018, 28, 561-568.	5.5	247
23	The Genome of the Netherlands: design, and project goals. European Journal of Human Genetics, 2014, 22, 221-227.	2.8	246
24	Multivariate genome-wide analyses of the well-being spectrum. Nature Genetics, 2019, 51, 445-451.	21.4	228
25	Genome-wide association meta-analysis of human longevity identifies a novel locus conferring survival beyond 90 years of age. Human Molecular Genetics, 2014, 23, 4420-4432.	2.9	227
26	Genetic Evidence for Causal Relationships Between Maternal Obesity-Related Traits and Birth Weight. JAMA - Journal of the American Medical Association, 2016, 315, 1129.	7.4	220
27	Genomic and phenotypic insights from an atlas of genetic effects on DNA methylation. Nature Genetics, 2021, 53, 1311-1321.	21.4	218
28	The Adult Netherlands Twin Register: Twenty-Five Years of Survey and Biological Data Collection. Twin Research and Human Genetics, 2013, 16, 271-281.	0.6	186
29	Cenetic insights into biological mechanisms governing human ovarian ageing. Nature, 2021, 596, 393-397.	27.8	183
30	Genome-wide association study identifies novel genetic variants contributing to variation in blood metabolite levels. Nature Communications, 2015, 6, 7208.	12.8	178
31	Genetic and environmental effects on body mass index from infancy to the onset of adulthood: an individual-based pooled analysis of 45 twin cohorts participating in the COllaborative project of Development of Anthropometrical measures in Twins (CODATwins) study. American Journal of Clinical Nutrition, 2016, 104, 371-379.	4.7	175
32	Directional dominance on stature and cognition inÂdiverse human populations. Nature, 2015, 523, 459-462.	27.8	173
33	Genome-wide meta-analysis of 241,258 adults accounting for smoking behaviour identifies novel loci for obesity traits. Nature Communications, 2017, 8, 14977.	12.8	169
34	Genome-wide physical activity interactions in adiposity ― A meta-analysis of 200,452 adults. PLoS Genetics, 2017, 13, e1006528.	3.5	158
35	Blood lipids influence DNA methylation in circulating cells. Genome Biology, 2016, 17, 138.	8.8	154
36	Investigating the possible causal association of smoking with depression and anxiety using Mendelian randomisation meta-analysis: the CARTA consortium. BMJ Open, 2014, 4, e006141.	1.9	150

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37	Circulating metabolites and general cognitive ability and dementia: Evidence from 11 cohort studies. Alzheimer's and Dementia, 2018, 14, 707-722.	0.8	143
38	The Netherlands Twin Register Biobank: A Resource for Genetic Epidemiological Studies. Twin Research and Human Genetics, 2010, 13, 231-245.	0.6	141
39	Meta-analysis of epigenome-wide association studies in neonates reveals widespread differential DNA methylation associated with birthweight. Nature Communications, 2019, 10, 1893.	12.8	140
40	Population structure, migration, and diversifying selection in the Netherlands. European Journal of Human Genetics, 2013, 21, 1277-1285.	2.8	137
41	Genetic and environmental influences on height from infancy to early adulthood: An individual-based pooled analysis of 45 twin cohorts. Scientific Reports, 2016, 6, 28496.	3.3	133
42	Heritability and Stability of Resting Blood Pressure. Twin Research and Human Genetics, 2005, 8, 499-508.	0.6	129
43	The Concordance and Heritability of Type 2 Diabetes in 34,166 Twin Pairs From International Twin Registers: The Discordant Twin (DISCOTWIN) Consortium. Twin Research and Human Genetics, 2015, 18, 762-771.	0.6	125
44	The Netherlands Twin Register: Longitudinal Research Based on Twin and Twin-Family Designs. Twin Research and Human Genetics, 2019, 22, 623-636.	0.6	112
45	Genetic variants linked to education predict longevity. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 13366-13371.	7.1	110
46	Differences in genetic and environmental variation in adult BMI by sex, age, time period, and region: an individual-based pooled analysis of 40 twin cohorts. American Journal of Clinical Nutrition, 2017, 106, 457-466.	4.7	107
47	Genome-wide association and HLA fine-mapping studies identify risk loci and genetic pathways underlying allergic rhinitis. Nature Genetics, 2018, 50, 1072-1080.	21.4	106
48	Effects of Metformin on Metabolite Profiles and LDL Cholesterol in Patients With Type 2 Diabetes. Diabetes Care, 2015, 38, 1858-1867.	8.6	97
49	Genetic loci associated with heart rate variability and their effects on cardiac disease risk. Nature Communications, 2017, 8, 15805.	12.8	95
50	Identification of Common Genetic Variants Influencing Spontaneous Dizygotic Twinning and Female Fertility. American Journal of Human Genetics, 2016, 98, 898-908.	6.2	89
51	Sex-dimorphic genetic effects and novel loci for fasting glucose and insulin variability. Nature Communications, 2021, 12, 24.	12.8	87
52	Genome-wide association meta-analysis of individuals of European ancestry identifies new loci explaining a substantial fraction of hair color variation and heritability. Nature Genetics, 2018, 50, 652-656.	21.4	86
53	An Analysis of Two Genome-wide Association Meta-analyses Identifies a New Locus for Broad Depression Phenotype. Biological Psychiatry, 2017, 82, 322-329.	1.3	84
54	Associations between smoking and caffeine consumption in two European cohorts. Addiction, 2016, 111, 1059-1068.	3.3	80

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55	Genome-wide association study identifies 48 common genetic variants associated with handedness. Nature Human Behaviour, 2021, 5, 59-70.	12.0	79
56	The Genetic Architecture of Neuroticism in 3301 Dutch Adolescent Twins as a Function of Age and Sex: A Study From the Dutch Twin Register. Twin Research and Human Genetics, 2006, 9, 24-29.	0.6	77
57	Discovery and Fine-Mapping of Glycaemic and Obesity-Related Trait Loci Using High-Density Imputation. PLoS Genetics, 2015, 11, e1005230.	3.5	77
58	A principal component meta-analysis on multiple anthropometric traits identifies novel loci for body shape. Nature Communications, 2016, 7, 13357.	12.8	74
59	A meta-analysis of 120 246 individuals identifies 18 new loci for fibrinogen concentration. Human Molecular Genetics, 2016, 25, 358-370.	2.9	73
60	Associations between loneliness and personality are mostly driven by a genetic association with Neuroticism. Journal of Personality, 2019, 87, 386-397.	3.2	66
61	Environmental Factors Determine Where the Dutch Live: Results From the Netherlands Twin Register. Twin Research and Human Genetics, 2005, 8, 312-317.	0.6	65
62	Heritability estimates for 361 blood metabolites across 40 genome-wide association studies. Nature Communications, 2020, 11, 39.	12.8	64
63	Differential gene expression patterns between smokers and nonâ€smokers: cause or consequence?. Addiction Biology, 2017, 22, 550-560.	2.6	62
64	Phenome-wide investigation of health outcomes associated with genetic predisposition to loneliness. Human Molecular Genetics, 2019, 28, 3853-3865.	2.9	62
65	An integrative cross-omics analysis of DNA methylation sites of glucose and insulin homeostasis. Nature Communications, 2019, 10, 2581.	12.8	62
66	Heritability of Self-Reported Asthma and Allergy: A Study in Adult Dutch Twins, Siblings and Parents. Twin Research and Human Genetics, 2008, 11, 132-142.	0.6	61
67	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. Biological Psychiatry, 2022, 91, 102-117.	1.3	61
68	Familial Clustering of Major Depression and Anxiety Disorders in Australian and Dutch Twins and Siblings. Twin Research and Human Genetics, 2005, 8, 609-615.	0.6	60
69	CNV Concordance in 1,097 MZ Twin Pairs. Twin Research and Human Genetics, 2015, 18, 1-12.	0.6	59
70	Genetic and environmental variation in educational attainment: an individual-based analysis of 28 twin cohorts. Scientific Reports, 2020, 10, 12681.	3.3	59
71	The genetics of alcohol dependence: Twin and SNPâ€based heritability, and genomeâ€wide association study based on AUDIT scores. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2015, 168, 739-748.	1.7	56
72	The CODATwins Project: The Cohort Description of Collaborative Project of Development of Anthropometrical Measures in Twins to Study Macro-Environmental Variation in Genetic and Environmental Effects on Anthropometric Traits. Twin Research and Human Genetics, 2015, 18, 348-360.	0.6	55

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73	Neighbourhood characteristics and prevalence and severity of depression: pooled analysis of eight Dutch cohort studies. British Journal of Psychiatry, 2019, 215, 468-475.	2.8	54
74	Integration of epidemiologic, pharmacologic, genetic and gut microbiome data in a drug–metabolite atlas. Nature Medicine, 2020, 26, 110-117.	30.7	54
75	Heavier smoking may lead to a relative increase in waist circumference: evidence for a causal relationship from a Mendelian randomisation meta-analysis. The CARTA consortium: TableÂ1. BMJ Open, 2015, 5, e008808.	1.9	53
76	Smoking During Adolescence as a Risk Factor for Attention Problems. Biological Psychiatry, 2015, 78, 656-663.	1.3	52
77	Prevalence of dieting and fear of weight gain across ages: a community sample from adolescents to the elderly. International Journal of Public Health, 2017, 62, 911-919.	2.3	52
78	Stratification by Smoking Status Reveals an Association of CHRNA5-A3-B4 Genotype with Body Mass Index in Never Smokers. PLoS Genetics, 2014, 10, e1004799.	3.5	45
79	Genetic and environmental influences on adult human height across birth cohorts from 1886 to 1994. ELife, 2016, 5, .	6.0	42
80	Genome-Wide Meta-Analysis of Cotinine Levels in Cigarette Smokers Identifies Locus at 4q13.2. Scientific Reports, 2016, 6, 20092.	3.3	42
81	DNA methylation signatures of educational attainment. Npj Science of Learning, 2018, 3, 7.	2.8	42
82	A fluid response: Alpha-amylase reactions to acute laboratory stress are related to sample timing and saliva flow rate. Biological Psychology, 2015, 109, 111-119.	2.2	39
83	Genome-wide analysis of DNA methylation in buccal cells: a study of monozygotic twins and mQTLs. Epigenetics and Chromatin, 2018, 11, 54.	3.9	39
84	Causes of variation in the neutrophil–lymphocyte and platelet–lymphocyte ratios: a twin-family study. Biomarkers in Medicine, 2016, 10, 1061-1072.	1.4	38
85	An Extended Twin-Pedigree Study of Neuroticism in the Netherlands Twin Register. Behavior Genetics, 2018, 48, 1-11.	2.1	36
86	Educational Attainment Influences Levels of Homozygosity through Migration and Assortative Mating. PLoS ONE, 2015, 10, e0118935.	2.5	36
87	Twin Family Registries Worldwide: An Important Resource for Scientific Research. Twin Research and Human Genetics, 2019, 22, 427-437.	0.6	33
88	Contribution of Genetics to the Susceptibility to Hidradenitis Suppurativa in a Large, Cross-sectional Dutch Twin Cohort. JAMA Dermatology, 2020, 156, 1359.	4.1	33
89	Identification, Heritability, and Relation With Gene Expression of Novel DNA Methylation Loci for Blood Pressure. Hypertension, 2020, 76, 195-205.	2.7	33
90	Environmental Factors Determine Where the Dutch Live: Results From the Netherlands Twin Register. Twin Research and Human Genetics, 2005, 8, 312-317.	0.6	33

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91	2SNP heritability and effects of genetic variants for neutrophil-to-lymphocyte and platelet-to-lymphocyte ratio. Journal of Human Genetics, 2017, 62, 979-988.	2.3	32
92	Metabolite ratios as potential biomarkers for type 2 diabetes: a DIRECT study. Diabetologia, 2018, 61, 117-129.	6.3	32
93	Genome-wide association study of circulating interleukin 6 levels identifies novel loci. Human Molecular Genetics, 2021, 30, 393-409.	2.9	32
94	Heritability and Genome-Wide Association Studies for Hair Color in a Dutch Twin Family Based Sample. Genes, 2015, 6, 559-576.	2.4	31
95	Validated inference of smoking habits from blood with a finite DNA methylation marker set. European Journal of Epidemiology, 2019, 34, 1055-1074.	5.7	31
96	Healthy Cotwins Share Gut Microbiome Signatures With Their Inflammatory Bowel Disease Twins and Unrelated Patients. Gastroenterology, 2021, 160, 1970-1985.	1.3	31
97	Individual Differences in Exercise Behavior: Stability and Change in Genetic and Environmental Determinants From Age 7 to 18. Behavior Genetics, 2016, 46, 665-679.	2.1	30
98	Cohort profile: the Geoscience and Health Cohort Consortium (GECCO) in the Netherlands. BMJ Open, 2018, 8, e021597.	1.9	29
99	Unraveling the Genetic and Environmental Relationship Between Well-Being and Depressive Symptoms Throughout the Lifespan. Frontiers in Psychiatry, 2018, 9, 261.	2.6	29
100	QTLs for height: results of a full genome scan in Dutch sibling pairs. European Journal of Human Genetics, 2004, 12, 820-828.	2.8	28
101	Genome-Wide Significance for <i>PCLO</i> as a Gene for Major Depressive Disorder. Twin Research and Human Genetics, 2017, 20, 267-270.	0.6	28
102	Genetic variants in RBFOX3 are associated with sleep latency. European Journal of Human Genetics, 2016, 24, 1488-1495.	2.8	27
103	Harmonising and linking biomedical and clinical data across disparate data archives to enable integrative cross-biobank research. European Journal of Human Genetics, 2016, 24, 521-528.	2.8	27
104	Parental Education and Genetics of BMI from Infancy to Old Age: A Pooled Analysis of 29 Twin Cohorts. Obesity, 2019, 27, 855-865.	3.0	27
105	The Etiology of Mathematical and Reading (Dis)ability Covariation in a Sample of Dutch Twins. Twin Research and Human Genetics, 2005, 8, 585-593.	0.6	26
106	The association of alcohol intake with gamma-glutamyl transferase (GGT) levels: Evidence for correlated genetic effects. Drug and Alcohol Dependence, 2014, 134, 99-105.	3.2	26
107	Cohabitation is associated with a greater resemblance in gut microbiota which can impact cardiometabolic and inflammatory risk. BMC Microbiology, 2019, 19, 230.	3.3	26
108	Identical twins carry a persistent epigenetic signature of early genome programming. Nature Communications, 2021, 12, 5618.	12.8	26

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109	Zygosity Differences in Height and Body Mass Index of Twins From Infancy to Old Age: A Study of the CODATwins Project. Twin Research and Human Genetics, 2015, 18, 557-570.	0.6	24
110	Establishing a Twin Register: An Invaluable Resource for (Behavior) Genetic, Epidemiological, Biomarker, and †Omics' Studies. Twin Research and Human Genetics, 2018, 21, 239-252.	0.6	24
111	DNA Methylation Changes in the <i>IGF1R</i> Gene in Birth Weight Discordant Adult Monozygotic Twins. Twin Research and Human Genetics, 2015, 18, 635-646.	0.6	23
112	Association between birthweight and later body mass index: an individual-based pooled analysis of 27 twin cohorts participating in the CODATwins project. International Journal of Epidemiology, 2017, 46, 1488-1498.	1.9	22
113	Relative Telomere Repeat Mass in Buccal and Leukocyte-Derived DNA. PLoS ONE, 2017, 12, e0170765.	2.5	22
114	A powerful phenotype for geneâ€finding studies derived from trajectory analyses of symptoms of anxiety and depression between age seven and 18. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2016, 171, 948-957.	1.7	21
115	Twin's Birth-Order Differences in Height and Body Mass Index From Birth to Old Age: A Pooled Study of 26 Twin Cohorts Participating in the CODATwins Project. Twin Research and Human Genetics, 2016, 19, 112-124.	0.6	21
116	Birth size and gestational age in opposite-sex twins as compared to same-sex twins: An individual-based pooled analysis of 21 cohorts. Scientific Reports, 2018, 8, 6300.	3.3	21
117	Predicting Loneliness from Where and What People Do. Social Sciences, 2020, 9, 51.	1.4	21
118	The Dopaminergic Reward System and Leisure Time Exercise Behavior: A Candidate Allele Study. BioMed Research International, 2014, 2014, 1-9.	1.9	20
119	Associations between birth size and later height from infancy through adulthood: An individual based pooled analysis of 28 twin cohorts participating in the CODATwins project. Early Human Development, 2018, 120, 53-60.	1.8	20
120	Heritability and GWAS Studies for Monocyte–Lymphocyte Ratio. Twin Research and Human Genetics, 2017, 20, 97-107.	0.6	19
121	Polygenic risk for alcohol consumption and its association with alcohol-related phenotypes: Do stress and life satisfaction moderate these relationships?. Drug and Alcohol Dependence, 2018, 183, 7-12.	3.2	19
122	Genetic and environmental factors affecting birth size variation: a pooled individual-based analysis of secular trends and global geographical differences using 26 twin cohorts. International Journal of Epidemiology, 2018, 47, 1195-1206.	1.9	19
123	Association of CRTC1 polymorphisms with obesity markers in subjects from the general population with lifetime depression. Journal of Affective Disorders, 2016, 198, 43-49.	4.1	18
124	The genetic architecture of body mass index from infancy to adulthood modified by parental education. Obesity, 2016, 24, 2004-2011.	3.0	18
125	Sex Differences in Heritability of BMI: A Comparative Study of Results from Twin Studies in Eight Countries. Twin Research and Human Genetics, 2003, 6, 409-421.	1.0	18
126	The Genetic Overlap Between Hair and Eye Color. Twin Research and Human Genetics, 2016, 19, 595-599.	0.6	17

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127	Genome-Wide Meta-Analyses of FTND and TTFC Phenotypes. Nicotine and Tobacco Research, 2020, 22, 900-909.	2.6	17
128	Genetic and environmental influences on human height from infancy through adulthood at different levels of parental education. Scientific Reports, 2020, 10, 7974.	3.3	17
129	Substance use: Interplay between polygenic risk and neighborhood environment. Drug and Alcohol Dependence, 2020, 209, 107948.	3.2	17
130	Familial Resemblance for Serum Metabolite Concentrations. Twin Research and Human Genetics, 2013, 16, 948-961.	0.6	14
131	Causes of Variation in Food Preference in the Netherlands. Twin Research and Human Genetics, 2020, 23, 195-203.	0.6	14
132	Predicting Complex Traits and Exposures From Polygenic Scores and Blood and Buccal DNA Methylation Profiles. Frontiers in Psychiatry, 2021, 12, 688464.	2.6	14
133	Religious Upbringing and Neuroticism in Dutch Twin Families. Twin Research and Human Genetics, 2007, 10, 327-333.	0.6	13
134	Effect of Genome and Environment on Metabolic and Inflammatory Profiles. PLoS ONE, 2015, 10, e0120898.	2.5	13
135	Metabolomics reveals a link between homocysteine and lipid metabolism and leukocyte telomere length: the ENGAGE consortium. Scientific Reports, 2019, 9, 11623.	3.3	13
136	A Comparison of the ASEBA Adult Self Report (ASR) and the Brief Problem Monitor (BPM/18-59). Behavior Genetics, 2020, 50, 363-373.	2.1	13
137	Genetic and Environmental Contributions to Stability in Adult Obsessive Compulsive Behavior. Twin Research and Human Genetics, 2015, 18, 52-60.	0.6	12
138	Genetic Vulnerability for Smoking and Cannabis Use: Associations With E-Cigarette and Water Pipe Use. Nicotine and Tobacco Research, 2019, 21, 723-730.	2.6	12
139	DNA methylation in peripheral tissues and left-handedness. Scientific Reports, 2022, 12, 5606.	3.3	12
140	Are Migraine and Tension-Type Headache Genetically Related? An Investigation of Twin Family Data. Twin Research and Human Genetics, 2018, 21, 112-118.	0.6	11
141	Gene-by-Crisis Interaction for Optimism and Meaning in Life: The Effects of the COVID-19 Pandemic. Behavior Genetics, 2021, , 1.	2.1	11
142	Explaining Individual Differences in Alcohol Intake in Adults: Evidence for Genetic and Cultural Transmission?. Journal of Studies on Alcohol and Drugs, 2014, 75, 201-210.	1.0	10
143	Heritability of Working in a Creative Profession. Behavior Genetics, 2017, 47, 298-304.	2.1	10
144	Genetic and environmental influences on quality of life: The <scp>COVID</scp> â€19 pandemic as a natural experiment. Genes, Brain and Behavior, 2022, 21, e12796.	2.2	10

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145	Integrative network analysis highlights biological processes underlying GLP-1 stimulated insulin secretion: A DIRECT study. PLoS ONE, 2018, 13, e0189886.	2.5	9
146	Education in Twins and Their Parents Across Birth Cohorts Over 100 years: An Individual-Level Pooled Analysis of 42-Twin Cohorts. Twin Research and Human Genetics, 2017, 20, 395-405.	0.6	8
147	Does the sex of one's co-twin affect height and BMI in adulthood? A study of dizygotic adult twins from 31 cohorts. Biology of Sex Differences, 2017, 8, 14.	4.1	8
148	Fat metabolism is associated with telomere length in six population-based studies. Human Molecular Genetics, 2022, 31, 1159-1170.	2.9	7
149	Genetics and Not Shared Environment Explains Familial Resemblance in Adult Metabolomics Data. Twin Research and Human Genetics, 2020, 23, 145-155.	0.6	6
150	Genetic meta-analysis of twin birth weight shows high genetic correlation with singleton birth weight. Human Molecular Genetics, 2021, 30, 1894-1905.	2.9	6
151	Testing Familial Transmission of Smoking With Two Different Research Designs. Nicotine and Tobacco Research, 2018, 20, 836-842.	2.6	5
152	Expanding the environmental scope: an environment-wide association study for mental well-being. Journal of Exposure Science and Environmental Epidemiology, 2022, 32, 195-204.	3.9	5
153	Evidence for Gender-Dependent Genotype by Environment Interaction in Adult Depression. Behavior Genetics, 2016, 46, 59-71.	2.1	4
154	Comparing ecstasy users and non-users in a population-based and co-twin control design across multiple traits. Addictive Behaviors, 2020, 108, 106421.	3.0	4
155	Establishment of the Avera Twin Register in the Midwest USA. Twin Research and Human Genetics, 2017, 20, 414-418.	0.6	3
156	Using a multivariate model to assess the interactive effects of demographics and lifestyle on the hematological profile. Biomarkers in Medicine, 2017, 11, 427-438.	1.4	3
157	Genetic factors explain a significant part of associations between adolescent well-being and the social environment. European Child and Adolescent Psychiatry, 2022, 31, 1611-1622.	4.7	3
158	Heritability of lifetime ecstasy use. Drug and Alcohol Dependence, 2017, 178, 66-69.	3.2	2
159	Association Between rs1051730 and Smoking During Pregnancy in Dutch Women. Nicotine and Tobacco Research, 2019, 21, 835-840.	2.6	2
160	Educational attainment of same-sex and opposite-sex dizygotic twins: An individual-level pooled study of 19 twin cohorts. Hormones and Behavior, 2021, 136, 105054.	2.1	1
161	Smoking Status of Parents, Siblings and Friends: Predictors of Regular Smoking? Findings from a Longitudinal Twin-family Study. Twin Research and Human Genetics, 2003, 6, 209-217.	1.0	1
162	Ethical Issues and GenomEUtwin. Twin Research and Human Genetics, 2003, 6, 455-463.	1.0	1

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163	Further Evidence for a QTL Influencing Body Mass Index on Chromosome 7p from a Genome-wide Scan in Dutch Families. Twin Research and Human Genetics, 2004, 7, 192-196.	1.0	1
	Socioemotional Development and Health from Adolescence to Adulthood (Cambridge Studies on) Tj ETQq0 0 0	rgBT /Over	lock 10 Tf 50
164	York: Cambridge University Press, 1st edition, 440 pp, US\$85.00, ISBN 0-521-84631-5 Twin Research and Human Genetics, 2007, 10, 234-234.	0.6	0
165	Plasma biomarkers predict amyloid pathology in cognitively unimpaired individuals. Alzheimer's and Dementia, 2020, 16, e045470.	0.8	Ο
166	Plasma Pâ€ŧau181 levels predict amyloid pathology in cognitively unimpaired individuals after 10 years. Alzheimer's and Dementia, 2021, 17, .	0.8	0