Markus M NÃ¶then

List of Publications by Year
in descending order

609 papers	66,812 citations	1294 109 h-index	1250 226 g-index
641 all docs	641 docs citations	641 times ranked	61113 citing authors

#	Article	IF	Citations
1	Elucidation of the genetic causes of bicuspid aortic valve disease. Cardiovascular Research, 2023, 119, 857-866.	1.8	11
2	Polygenic risk for schizophrenia and schizotypal traits in non-clinical subjects. Psychological Medicine, 2022, 52, 1069-1079.	2.7	10
3	Sex-Dependent Shared and Nonshared Genetic Architecture Across Mood and Psychotic Disorders. Biological Psychiatry, 2022, 91, 102-117.	0.7	61
4	Cis-epistasis at the LPA locus and risk of cardiovascular diseases. Cardiovascular Research, 2022, 118, 1088-1102.	1.8	14
5	Ventral Striatalâ€“Hippocampus Coupling During Reward Processing as a Stratification Biomarker for Psychotic Disorders. Biological Psychiatry, 2022, 91, 216-225.	0.7	10
6	Genetic risk for psychiatric illness is associated with the number of hospitalizations of bipolar disorder patients. Journal of Affective Disorders, 2022, 296, 532-540.	2.0	6
7	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. Biological Psychiatry, 2022, 91, 313-327.	0.7	114
8	Genome-wide interaction study with major depression identifies novel variants associated with cognitive function. Molecular Psychiatry, 2022, 27, 1111-1119.	4.1	24
9	Epigenome-wide association study of alcohol use disorder in five brain regions. Neuropsychopharmacology, 2022, 47, 832-839.	2.8	16
10	Breast and prostate cancer risk: The interplay of polygenic risk, rare pathogenic germline variants, and family history. Genetics in Medicine, 2022, 24, 576-585.	1.1	22
11	First genome-wide association study of esophageal atresia identifies three genetic risk loci at CTNNA3, FOXF1/FOXC2/FOXL1, and HNF1B. Human Genetics and Genomics Advances, 2022, 3, 100093.	1.0	4
12	Cross-tissue transcriptome-wide association studies identify susceptibility genes shared between schizophrenia and inflammatory bowel disease. Communications Biology, 2022, 5, 80.	2.0	12
13	OUP accepted manuscript. Cerebral Cortex, 2022, , .	1.6	0
14	GestaltMatcher facilitates rare disease matching using facial phenotype descriptors. Nature Genetics, 2022, 54, 349-357.	9.4	73
15	Observations that suggest a contribution of altered dermal papilla mitochondrial function to androgenetic alopecia. Experimental Dermatology, 2022, 31, 906-917.	1.4	11
16	Using polygenic scores and clinical data for bipolar disorder patient stratification and lithium response prediction: machine learning approach. British Journal of Psychiatry, 2022, 220, 219-228.	1.7	11
17	Colocalization analysis of pancreas eQTLs with risk loci from alcoholic and novel non-alcoholic chronic pancreatitis GWAS suggests potential disease causing mechanisms. Pancreatology, 2022, 22, 449-456.	0.5	3
18	Associations of common genetic risk variants of the muscarinic acetylcholine receptor M2 with cardiac autonomic dysfunction in patients with schizophrenia. World Journal of Biological Psychiatry, 2022, , 1-11.	1.3	1

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19	A genetic risk score of alleles related to MGUS interacts with socioeconomic position in a population-based cohort. Scientific Reports, 2022, 12, 4409.	1.6	0
20	Genetic variants associated with longitudinal changes in brain structure across the lifespan. Nature Neuroscience, 2022, 25, 421-432.	7.1	75
21	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. Nature, 2022, 604, 502-508.	13.7	929
22	ExomeChip-based rare variant association study in restless legs syndrome. Sleep Medicine, 2022, 94, 26-30.	0.8	0
23	Epigenetic Signatures of Smoking in Five Brain Regions. Journal of Personalized Medicine, 2022, 12, 566.	1.1	4
24	New insights into the genetic etiology of Alzheimerâ€™s disease and related dementias. Nature Genetics, 2022, 54, 412-436.	9.4	700
25	Wie wichtig ist die Kenntnis des genetischen Populationshintergrundes in der Medizin? Ein humangenetischer Beitrag vor dem Hintergrund der aktuellen Diskussion um die Verwendung des Begriffs â€žRasseâ€œ. Medizinische Genetik, 2022, 33, 337-341.	0.1	0
26	Borderline personality disorder and the big five: molecular genetic analyses indicate shared genetic architecture with neuroticism and openness. Translational Psychiatry, 2022, 12, 153.	2.4	7
27	Genome-wide meta-analysis of monoclonal gammopathy of undetermined significance (MGUS) identifies risk loci impacting IRF-6. Blood Cancer Journal, 2022, 12, 60.	2.8	2
28	Multi-omics signatures of alcohol use disorder in the dorsal and ventral striatum. Translational Psychiatry, 2022, 12, 190.	2.4	11
29	Chemokine receptor 4 expression on blood T lymphocytes predicts severity of major depressive disorder. Journal of Affective Disorders, 2022, 310, 343-353.	2.0	5
30	GWAS of thyroid dysgenesis identifies a risk locus at 2q33.3 linked to regulation of Wnt signaling. Human Molecular Genetics, 2022, 31, 3967-3974.	1.4	2
31	A novel longitudinal clustering approach to psychopathology across diagnostic entities in the hospital-based PsyCourse study. Schizophrenia Research, 2022, 244, 29-38.	1.1	2
32	Hepatic Expression of the Na+-Taurocholate Cotransporting Polypeptide Is Independent from Genetic Variation. International Journal of Molecular Sciences, 2022, 23, 7468.	1.8	6
33	Genome-wide association study of panic disorder reveals genetic overlap with neuroticism and depression. Molecular Psychiatry, 2021, 26, 4179-4190.	4.1	58
34	Genome-wide association study reveals new insights into the heritability and genetic correlates of developmental dyslexia. Molecular Psychiatry, 2021, 26, 3004-3017.	4.1	56
35	Association of polygenic score for major depression with response to lithium in patients with bipolar disorder. Molecular Psychiatry, 2021, 26, 2457-2470.	4.1	44
36	A common variation in HCN1 is associated with heart rate variability in schizophrenia. Schizophrenia Research, 2021, 229, 73-79.	1.1	13

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37	Generative network models of altered structural brain connectivity in schizophrenia. NeuroImage, 2021, 225, 117510.	2.1	24
38	Germline variation in the insulin-like growth factor pathway and risk of Barrettâ€™s esophagus and esophageal adenocarcinoma. Carcinogenesis, 2021, 42, 369-377.	1.3	11
39	Shared Genetics of Multiple System Atrophy and Inflammatory Bowel Disease. Movement Disorders, 2021, 36, 449-459.	2.2	16
40	Identifying multimodal signatures underlying the somatic comorbidity of psychosis: the COMMITMENT roadmap. Molecular Psychiatry, 2021, 26, 722-724.	4.1	7
41	Childhood maltreatment and cognitive functioning: the role of depression, parental education, and polygenic predisposition. Neuropsychopharmacology, 2021, 46, 891-899.	2.8	17
42	Bipolar multiplex families have an increased burden of common risk variants for psychiatric disorders. Molecular Psychiatry, 2021, 26, 1286-1298.	4.1	33
43	Exemplar scoring identifies genetically separable phenotypes of lithium responsive bipolar disorder. Translational Psychiatry, 2021, 11, 36.	2.4	16
44	Prediction of lithium response using genomic data. Scientific Reports, 2021, 11, 1155.	1.6	11
45	Synaptic processes and immune-related pathways implicated in Tourette syndrome. Translational Psychiatry, 2021, 11, 56.	2.4	31
46	Interaction of developmental factors and ordinary stressful life events on brain structure in adults. NeuroImage: Clinical, 2021, 30, 102683.	1.4	5
47	Hyper-Coordinated DNA Methylation is Altered in Schizophrenia and Associated with Brain Function. Schizophrenia Bulletin Open, 2021, 2, .	0.9	0
48	Exome-Wide Association Study Identifies FN3KRP and PGP as New Candidate Longevity Genes. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2021, 76, 786-795.	1.7	14
49	Clinical and genetic differences between bipolar disorder type 1 and 2 in multiplex families. Translational Psychiatry, 2021, 11, 31.	2.4	22
50	â€œThe Heidelberg Fiveâ€•personality dimensions: Genomeâ€wide associations, polygenic risk for neuroticism, and psychopathology 20â€‰years after assessment. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2021, 186, 77-89.	1.1	6
51	Effects of polygenic risk for major mental disorders and cross-disorder on cortical complexity. Psychological Medicine, 2021, , 1-12.	2.7	7
52	Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. Nature Genetics, 2021, 53, 817-829.	9.4	629
53	Analysis of genetic impact on smell impairment in patients with hereditary angioedema typeÂ1 and 2. JDDG - Journal of the German Society of Dermatology, 2021, 19, 1060-1062.	0.4	0
54	Pharmacogenetic association of diabetes-associated genetic risk score with rapid progression of coronary artery calcification following treatment with HMG-CoA-reductase inhibitors â€”results of the Heinz Nixdorf Recall Study. Naunyn-Schmiedeberg's Archives of Pharmacology, 2021, 394, 1713-1725.	1.4	4

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55	Apolipoprotein E homozygous Îµ4 allele status: Effects on cortical structure and white matter integrity in a young to mid-age sample. European Neuropsychopharmacology, 2021, 46, 93-104.	0.3	2
56	Common variants in Alzheimerâ€™s disease and risk stratification by polygenic risk scores. Nature Communications, 2021, 12, 3417.	5.8	140
57	Identification of transdiagnostic psychiatric disorder subtypes using unsupervised learning. Neuropsychopharmacology, 2021, 46, 1895-1905.	2.8	24
58	DNA Repair Gene Polymorphisms and Chromosomal Aberrations in Exposed Populations. Frontiers in Genetics, 2021, 12, 691947.	1.1	3
59	LAMP-Seq enables sensitive, multiplexed COVID-19 diagnostics using molecular barcoding. Nature Biotechnology, 2021, 39, 1556-1562.	9.4	46
60	Search for AL amyloidosis risk factors using Mendelian randomization. Blood Advances, 2021, 5, 2725-2731.	2.5	5
61	TBK1 and TNFRSF13B mutations and an autoinflammatory disease in a child with lethal COVID-19. Npj Genomic Medicine, 2021, 6, 55.	1.7	38
62	Identification of pleiotropy at the gene level between psychiatric disorders and related traits. Translational Psychiatry, 2021, 11, 410.	2.4	7
63	Interaction of Alzheimerâ€™s Disease-Associated Genetic Risk with Indicators of Socioeconomic Position on Mild Cognitive Impairment in the Heinz Nixdorf Recall Study. Journal of Alzheimer's Disease, 2021, 82, 1715-1725.	1.2	6
64	Characterisation of age and polarity at onset in bipolar disorder. British Journal of Psychiatry, 2021, 219, 659-669.	1.7	20
65	A genetic sum score of effect alleles associated with serum lipid concentrations interacts with educational attainment. Scientific Reports, 2021, 11, 16541.	1.6	1
66	HLA-DRB1 and HLA-DQB1 genetic diversity modulates response to lithium in bipolar affective disorders. Scientific Reports, 2021, 11, 17823.	1.6	10
67	Evidence for a functional interaction of WNT10A and EBF1 in male-pattern baldness. PLoS ONE, 2021, 16, e0256846.	1.1	6
68	Systematic investigation of a potential epidemiological and genetic association between male androgenetic alopecia and COVIDâ€19. Skin Health and Disease, 2021, 1, e72.	0.7	3
69	Association between genetic variants of the cholinergic system and postoperative delirium and cognitive dysfunction in elderly patients. BMC Medical Genomics, 2021, 14, 248.	0.7	8
70	Rare variant analysis in eczema identifies exonic variants in DUSP1, NOTCH4 and SLC9A4. Nature Communications, 2021, 12, 6618.	5.8	17
71	Combining schizophrenia and depression polygenic risk scores improves the genetic prediction of lithium response in bipolar disorder patients. Translational Psychiatry, 2021, 11, 606.	2.4	25
72	Polygenic risk scores across the extended psychosis spectrum. Translational Psychiatry, 2021, 11, 600.	2.4	11

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73	Common and rare variant association analyses in amyotrophic lateral sclerosis identify 15 risk loci with distinct genetic architectures and neuron-specific biology. Nature Genetics, 2021, 53, 1636-1648.	9.4	223
74	Cortical surface area alterations shaped by genetic load for neuroticism. Molecular Psychiatry, 2020, 25, 3422-3431.	4.1	20
75	Protein-coding variants contribute to the risk of atopic dermatitis and skin-specific gene expression. Journal of Allergy and Clinical Immunology, 2020, 145, 1208-1218.	1.5	29
76	Eight novel loci implicate shared genetic etiology in multiple myeloma, AL amyloidosis, and monoclonal gammopathy of unknown significance. Leukemia, 2020, 34, 1187-1191.	3.3	13
77	Genome-wide study on uveal melanoma patients finds association to DNA repair gene TDP1. Melanoma Research, 2020, 30, 166-172.	0.6	6
78	Predictive power of the ADHD GWAS 2019 polygenic risk scores in independent samples of bipolar patients with childhood ADHD. Journal of Affective Disorders, 2020, 265, 651-659.	2.0	15
79	A large-scale genome-wide association study meta-analysis of cannabis use disorder. Lancet Psychiatry,the, 2020, 7, 1032-1045.	3.7	200
80	Infection fatality rate of SARS-CoV2 in a super-spreading event in Germany. Nature Communications, 2020, 11, 5829.	5.8	207
81	Mapping of cis-acting expression quantitative trait loci in human scalp hair follicles. BMC Dermatology, 2020, 20, 16.	2.1	6
82	Insights Into the Biology of Persistent Chemotherapy-Induced Alopecia via Genomic Approachesâ€”An Avenue to Clinical Translation?. JAMA Dermatology, 2020, 156, 947.	2.0	2
83	Shared Genetic Etiology of Obesity-Related Traits and Barrett's Esophagus/Adenocarcinoma: Insights from Genome-Wide Association Studies. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 427-433.	1.1	7
84	Replication of a hippocampus specific effect of the tescalcin regulating variant rs7294919 on gray matter structure. European Neuropsychopharmacology, 2020, 36, 10-17.	0.3	2
85	Hormonal regulation in male androgenetic alopeciaâ€”Sex hormones and beyond: Evidence from recent genetic studies. Experimental Dermatology, 2020, 29, 814-827.	1.4	27
86	The genetic architecture of the human cerebral cortex. Science, 2020, 367, .	6.0	450
87	Acute alcohol withdrawal and recovery in men lead to profound changes in DNA methylation profiles: a longitudinal clinical study. Addiction, 2020, 115, 2034-2044.	1.7	21
88	Whole-exome sequencing of 81 individuals from 27 multiply affected bipolar disorder families. Translational Psychiatry, 2020, 10, 57.	2.4	23
89	Association of a Reproducible Epigenetic Risk Profile for Schizophrenia With Brain Methylation and Function. JAMA Psychiatry, 2020, 77, 628.	6.0	46
90	Genome-wide gene-environment analyses of major depressive disorder and reported lifetime traumatic experiences in UK Biobank. Molecular Psychiatry, 2020, 25, 1430-1446.	4.1	116

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91	Association between lipoprotein(a) (Lp(a)) levels and Lp(a) genetic variants with coronary artery calcification. BMC Medical Genetics, 2020, 21, 62.	2.1	23
92	An Investigation of Psychosis Subgroups With Prognostic Validation and Exploration of Genetic Underpinnings. JAMA Psychiatry, 2020, 77, 523.	6.0	39
93	A Novel Locus and Candidate Gene for Familial Developmental Dyslexia on Chromosome 4q. Zeitschrift FÃœr Kinder- Und Jugendpsychiatrie Und Psychotherapie, 2020, 48, 478-489.	0.4	5
94	The role of environmental stress and DNA methylation in the longitudinal course of bipolar disorder. International Journal of Bipolar Disorders, 2020, 8, 9.	0.8	13
95	Insights into the genomics of affective disorders. Medizinische Genetik, 2020, 32, 9-18.	0.1	2
96	Out of the lab and into the clinic: steps to a pragmatic new era in psychiatric genetics. Medizinische Genetik, 2020, 32, 5-7.	0.1	0
97	Polygenic scores for psychiatric disease: from research tool to clinical application. Medizinische Genetik, 2020, 32, 39-45.	0.1	14
98	Investigating polygenic burden in age at disease onset in bipolar disorder: Findings from an international multicentric study. Bipolar Disorders, 2019, 21, 68-75.	1.1	20
99	Pathway-Specific Genetic Risk for Alzheimerâ€™s Disease Differentiates Regional Patterns of Cortical Atrophy in Older Adults. Cerebral Cortex, 2019, 30, 801-811.	1.6	11
100	Effects of a neurodevelopmental genes based polygenic risk score for schizophrenia and single gene variants on brain structure in non-clinical subjects: A preliminary report. Schizophrenia Research, 2019, 212, 225-228.	1.1	7
101	HDAC9 is implicated in atherosclerotic aortic calcification and affects vascular smooth muscle cell phenotype. Nature Genetics, 2019, 51, 1580-1587.	9.4	92
102	Transcriptome-wide association study of multiple myeloma identifies candidate susceptibility genes. Human Genomics, 2019, 13, 37.	1.4	14
103	A genetic sum score of risk alleles associated with body mass index interacts with socioeconomic position in the Heinz Nixdorf Recall Study. PLoS ONE, 2019, 14, e0221252.	1.1	11
104	Distinct pathways associated with chromosomal aberration frequency in a cohort exposed to genotoxic compounds compared to general population. Mutagenesis, 2019, 34, 323-330.	1.0	6
105	Reproducible grey matter patterns index a multivariate, global alteration of brain structure in schizophrenia and bipolar disorder. Translational Psychiatry, 2019, 9, 12.	2.4	35
106	PEDIA: prioritization of exome data by image analysis. Genetics in Medicine, 2019, 21, 2807-2814.	1.1	58
107	Apolipoprotein E Homozygous Îµ4 Allele Status: A Deteriorating Effect on Visuospatial Working Memory and Global Brain Structure. Frontiers in Neurology, 2019, 10, 552.	1.1	10
108	Associations of schizophrenia risk genes ZNF804A and CACNA1C with schizotypy and modulation of attention in healthy subjects. Schizophrenia Research, 2019, 208, 67-75.	1.1	20

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109	Genome-wide association study identifies 30 loci associated with bipolar disorder. Nature Genetics, 2019, 51, 793-803.	9.4	1,191
110	Interrogating the Genetic Determinants of Touretteâ€™s Syndrome and Other Tic Disorders Through Genome-Wide Association Studies. American Journal of Psychiatry, 2019, 176, 217-227.	4.0	242
111	Transcriptome-wide analysis of filarial extract-primed human monocytes reveal changes in LPS-induced PTX3 expression levels. Scientific Reports, 2019, 9, 2562.	1.6	3
112	Genome-wide interaction and pathway-based identification of key regulators in multiple myeloma. Communications Biology, 2019, 2, 89.	2.0	14
113	Cover Image, Volume 180B, Number 2, March 2019. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2019, 180, i.	1.1	0
114	No Association Between Vitamin D Status and Risk of Barrett's Esophagus or Esophageal Adenocarcinoma: A Mendelian Randomization Study. Clinical Gastroenterology and Hepatology, 2019, 17, 2227-2235.e1.	2.4	16
115	First genotypeâ€“phenotype study reveals HLAâ€ÐQÎ²1 insertion heterogeneity in highâ€resolution manometry achalasia subtypes. United European Gastroenterology Journal, 2019, 7, 45-51.	1.6	5
116	Genome-wide association study of monoclonal gammopathy of unknown significance (MGUS): comparison with multiple myeloma. Leukemia, 2019, 33, 1817-1821.	3.3	14
117	Genome-wide association scan identifies new variants associated with a cognitive predictor of dyslexia. Translational Psychiatry, 2019, 9, 77.	2.4	82
118	Genetic meta-analysis of diagnosed Alzheimerâ€™s disease identifies new risk loci and implicates AÎ², tau, immunity and lipid processing. Nature Genetics, 2019, 51, 414-430.	9.4	1,962
119	Male-pattern baldness and incident coronary heart disease and risk factors in the Heinz Nixdorf Recall Study. PLoS ONE, 2019, 14, e0225521.	1.1	6
120	Identification of loci of functional relevance to Barrettâ€™s esophagus and esophageal adenocarcinoma: Cross-referencing of expression quantitative trait loci data from disease-relevant tissues with genetic association data. PLoS ONE, 2019, 14, e0227072.	1.1	5
121	Genetic architecture of subcortical brain structures in 38,851 individuals. Nature Genetics, 2019, 51, 1624-1636.	9.4	192
122	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. Cell, 2019, 179, 1469-1482.e11.	13.5	935
123	Genetic correlation between multiple myeloma and chronic lymphocytic leukaemia provides evidence for shared aetiology. Blood Cancer Journal, 2019, 9, 1.	2.8	40
124	The influence of religious activity and polygenic schizophrenia risk on religious delusions in schizophrenia. Schizophrenia Research, 2019, 210, 255-261.	1.1	9
125	Effects of BDNF Val66Met genotype and schizophrenia familial risk on a neural functional network for cognitive control in humans. Neuropsychopharmacology, 2019, 44, 590-597.	2.8	19
126	Insights into Male Androgenetic Alopecia: Differential Gene Expression Profiling of PluckedÂHair Follicles and Integration with Genetic Data. Journal of Investigative Dermatology, 2019, 139, 235-238.	0.3	10

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127	Genetic variation associated with chromosomal aberration frequency: A genomeâ€wide association study. Environmental and Molecular Mutagenesis, 2019, 60, 17-28.	0.9	9
128	Efficient region-based test strategy uncovers genetic risk factors for functional outcome in bipolar disorder. European Neuropsychopharmacology, 2019, 29, 156-170.	0.3	7
129	Neurobiology of the major psychoses: a translational perspective on brain structure and functionâ€"the FOR2107 consortium. European Archives of Psychiatry and Clinical Neuroscience, 2019, 269, 949-962.	1.8	103
130	Association of Polygenic Score for Schizophrenia and HLA Antigen and Inflammation Genes With Response to Lithium in Bipolar Affective Disorder. JAMA Psychiatry, 2018, 75, 65-74.	6.0	102
131	Polygenic risk for schizophrenia affects working memory and its neural correlates in healthy subjects. Schizophrenia Research, 2018, 197, 315-320.	1.1	11
132	Genome-wide association study identifies susceptibility loci for B-cell childhood acute lymphoblastic leukemia. Nature Communications, 2018, 9, 1340.	5.8	58
133	Impact on birthÂweight of maternal smoking throughout pregnancy mediated by DNA methylation. BMC Genomics, 2018, 19, 290.	1.2	41
134	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. Nature Genetics, 2018, 50, 668-681.	9.4	2,224
135	The protocadherin 17 gene affects cognition, personality, amygdala structure and function, synapse development and risk of major mood disorders. Molecular Psychiatry, 2018, 23, 400-412.	4.1	60
136	El estudio Andalusian Bipolar Family (ABiF): protocolo y descripciÃ³n de la muestra. Revista De PsiquiatrÃa Y Salud Mental, 2018, 11, 199-207.	1.0	5
137	The influence of MIR137 on white matter fractional anisotropy and cortical surface area in individuals with familial risk for psychosis. Schizophrenia Research, 2018, 195, 190-196.	1.1	6
138	Genome-Wide MicroRNA Analysis Implicates miR-30b/d in the Etiology ofÂAlopecia Areata. Journal of Investigative Dermatology, 2018, 138, 549-556.	0.3	21
139	Investigation of dominant and recessive inheritance models in genomeâ€wide association studies data of nonsyndromic cleft lip with or without cleft palate. Birth Defects Research, 2018, 110, 336-341.	0.8	8
140	Gene set enrichment analysis and expression pattern exploration implicate an involvement of neurodevelopmental processes in bipolar disorder. Journal of Affective Disorders, 2018, 228, 20-25.	2.0	14
141	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. Nature Neuroscience, 2018, 21, 1656-1669.	7.1	490
142	Exploring Genetic Associations of Alzheimerâ€™s Disease Loci With Mild Cognitive Impairment Neurocognitive Endophenotypes. Frontiers in Aging Neuroscience, 2018, 10, 340.	1.7	12
143	Elevated expression of a minor isoform of ANK3 is a risk factor for bipolar disorder. Translational Psychiatry, 2018, 8, 210.	2.4	24
144	Identification of multiple risk loci and regulatory mechanisms influencing susceptibility to multiple myeloma. Nature Communications, 2018, 9, 3707.	5.8	86

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145	Genome-wide association study implicates immune dysfunction in the development of Hodgkin lymphoma. Blood, 2018, 132, 2040-2052.	0.6	17
146	Enrichment of B cell receptor signaling and epidermal growth factor receptor pathways in monoclonal gammopathy of undetermined significance: a genome-wide genetic interaction study. Molecular Medicine, 2018, 24, 30.	1.9	9
147	Evidence for <scp>PTGER</scp>4,<scp>PSCA</scp>, and <scp>MBOAT</scp>7 as risk genes for gastric cancer on the genome and transcriptome level. Cancer Medicine, 2018, 7, 5057-5065.	1.3	22
148	Analysis of the Influence of microRNAs in Lithium Response in Bipolar Disorder. Frontiers in Psychiatry, 2018, 9, 207.	1.3	28
149	Analysis of shared heritability in common disorders of the brain. Science, 2018, 360, .	6.0	1,085
150	Evaluation of food allergy candidate loci in the Genetics of Food Allergy study. Journal of Allergy and Clinical Immunology, 2018, 142, 1368-1370.e2.	1.5	1
151	Shared genetic etiology between alcohol dependence and major depressive disorder. Psychiatric Genetics, 2018, 28, 66-70.	0.6	19
152	Genome-wide significant risk factors for Alzheimerâ€™s disease: role in progression to dementia due to Alzheimer's disease among subjects with mild cognitive impairment. Molecular Psychiatry, 2017, 22, 153-160.	4.1	102
153	Genomewide analysis of copy number variants in alopecia areata in a <scp>C</scp>entral <scp>E</scp>uropean cohort reveals association with <scp>MCHR</scp>2. Experimental Dermatology, 2017, 26, 536-541.	1.4	21
154	Novel genetic loci associated with hippocampal volume. Nature Communications, 2017, 8, 13624.	5.8	250
155	Genetics of schizophrenia: A consensus paper of the WFSBP Task Force on Genetics. World Journal of Biological Psychiatry, 2017, 18, 492-505.	1.3	48
156	Influence of age and cognitive performance on resting-state brain networks of older adults in a population-based cohort. Cortex, 2017, 89, 28-44.	1.1	53
157	Expression profiling and bioinformatic analyses suggest new target genes and pathways for human hair follicle related microRNAs. BMC Dermatology, 2017, 17, 3.	2.1	35
158	Meta-analysis identifies novel risk loci and yields systematic insights into the biology of male-pattern baldness. Nature Communications, 2017, 8, 14694.	5.8	58
159	Socioeconomic Status Interacts with the Genetic Effect of a Chromosome 9p21.3 Common Variant to Influence Coronary Artery Calcification and Incident Coronary Events in the Heinz Nixdorf Recall Study (Risk Factors, Evaluation of Coronary Calcium, and Lifestyle). Circulation: Cardiovascular Genetics, 2017, 10, .	5.1	13
160	Assessing the effect of obesity-related traits on multiple myeloma using a Mendelian randomisation approach. Blood Cancer Journal, 2017, 7, e573-e573.	2.8	12
161	Rare Copy Number Variants in NRXN1 and CNTN6 Increase Risk for Tourette Syndrome. Neuron, 2017, 94, 1101-1111.e7.	3.8	137
162	Polygenic risk for depression and the neural correlates of working memory in healthy subjects. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2017, 79, 67-76.	2.5	41

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163	Genome-wide association study of borderline personality disorder reveals genetic overlap with bipolar disorder, major depression and schizophrenia. Translational Psychiatry, 2017, 7, e1155-e1155.	2.4	150
164	An Expanded Genome-Wide Association Study of Type 2 Diabetes in Europeans. Diabetes, 2017, 66, 2888-2902.	0.3	615
165	Genome-wide association study of glioma subtypes identifies specific differences in genetic susceptibility to glioblastoma and non-glioblastoma tumors. Nature Genetics, 2017, 49, 789-794.	9.4	259
166	Genomewide association study on monoclonal gammopathy of unknown significance (MGUS). European Journal of Haematology, 2017, 99, 70-79.	1.1	16
167	Genetic effects influencing risk for major depressive disorder in China and Europe. Translational Psychiatry, 2017, 7, e1074-e1074.	2.4	64
168	Genome-wide association study of immunoglobulin light chain amyloidosis in three patient cohorts: comparison with myeloma. Leukemia, 2017, 31, 1735-1742.	3.3	32
169	An Analysis of Two Genome-wide Association Meta-analyses Identifies a New Locus for Broad Depression Phenotype. Biological Psychiatry, 2017, 82, 322-329.	0.7	84
170	Genome-wide association study identifies the SERPINB gene cluster as a susceptibility locus for food allergy. Nature Communications, 2017, 8, 1056.	5.8	75
171	Genetic regulatory effects modified by immune activation contribute to autoimmune disease associations. Nature Communications, 2017, 8, 266.	5.8	157
172	Rare coding variants in PLCG2, ABI3, and TREM2 implicate microglial-mediated innate immunity in Alzheimer's disease. Nature Genetics, 2017, 49, 1373-1384.	9.4	783
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