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List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	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
2	Large recurrent microdeletions associated with schizophrenia. Nature, 2008, 455, 232-236.	27.8	1,619
3	Common variants conferring risk of schizophrenia. Nature, 2009, 460, 744-747.	27.8	1,572
4	A high-resolution recombination map of the human genome. Nature Genetics, 2002, 31, 241-247.	21.4	1,571
5	A variant associated with nicotine dependence, lung cancer and peripheral arterial disease. Nature, 2008, 452, 638-642.	27.8	1,399
6	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
7	Genome-wide association study identifies 30 loci associated with bipolar disorder. Nature Genetics, 2019, 51, 793-803.	21.4	1,191
8	A mega-analysis of genome-wide association studies for major depressive disorder. Molecular Psychiatry, 2013, 18, 497-511.	7.9	1,002
9	A common inversion under selection in Europeans. Nature Genetics, 2005, 37, 129-137.	21.4	747
10	The nature of nurture: Effects of parental genotypes. Science, 2018, 359, 424-428.	12.6	720
11	Sequence variants at CHRNB3–CHRNA6 and CYP2A6 affect smoking behavior. Nature Genetics, 2010, 42, 448-453.	21.4	649
12	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.	21.4	629
13	A Genetic Risk Factor for Periodic Limb Movements in Sleep. New England Journal of Medicine, 2007, 357, 639-647.	27.0	582
14	Sequence variants at the TERT-CLPTM1L locus associate with many cancer types. Nature Genetics, 2009, 41, 221-227.	21.4	572
15	Large-scale association analysis identifies new lung cancer susceptibility loci and heterogeneity in genetic susceptibility across histological subtypes. Nature Genetics, 2017, 49, 1126-1132.	21.4	472
16	Sequence variant on 8q24 confers susceptibility to urinary bladder cancer. Nature Genetics, 2008, 40, 1307-1312.	21.4	377
17	Polygenic risk scores for schizophrenia and bipolar disorder predict creativity. Nature Neuroscience, 2015, 18, 953-955.	14.8	351
18	Genome-wide analyses for personality traits identify six genomic loci and show correlations with	21.4	350

psychiatric disorders. Nature Genetics, 2017, 49, 152-156.

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19	Genome-wide association and genetic functional studies identify <i>autism susceptibility candidate 2</i> gene (<i>AUTS2</i>) in the regulation of alcohol consumption. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 7119-7124.	7.1	258
20	Brain age prediction using deep learning uncovers associated sequence variants. Nature Communications, 2019, 10, 5409.	12.8	238
21	A large-scale genome-wide association study meta-analysis of cannabis use disorder. Lancet Psychiatry,the, 2020, 7, 1032-1045.	7.4	200
22	Recombination rate and reproductive success in humans. Nature Genetics, 2004, 36, 1203-1206.	21.4	176
23	Genome-wide Association for Major Depression Through Age at Onset Stratification: Major Depressive Disorder Working Group of the Psychiatric Genomics Consortium. Biological Psychiatry, 2017, 81, 325-335.	1.3	175
24	A sequence variant at 4p16.3 confers susceptibility to urinary bladder cancer. Nature Genetics, 2010, 42, 415-419.	21.4	169
25	Nanosecond photolysis of rhodopsin: evidence for a new blue-shifted intermediate. Biochemistry, 1990, 29, 1475-1485.	2.5	143
26	Expanding the range of ZNF804A variants conferring risk of psychosis. Molecular Psychiatry, 2011, 16, 59-66.	7.9	140
27	The Genetics of the Mood Disorder Spectrum: Genome-wide Association Analyses of More Than 185,000 Cases and 439,000 Controls. Biological Psychiatry, 2020, 88, 169-184.	1.3	137
28	Transient channel-opening in bacteriorhodopsin: an EPR study 1 1Edited by D. Ress. Journal of Molecular Biology, 1997, 273, 951-957.	4.2	119
29	Neuregulin 1 and schizophrenia. Annals of Medicine, 2004, 36, 62-71.	3.8	119
30	Sequence variants at CYP1A1–CYP1A2 and AHR associate with coffee consumption. Human Molecular Genetics, 2011, 20, 2071-2077.	2.9	114
31	Genetic Differences in the Immediate Transcriptome Response to Stress Predict Risk-Related Brain Function and Psychiatric Disorders. Neuron, 2015, 86, 1189-1202.	8.1	102
32	Genome-wide association study implicates CHRNA2 in cannabis use disorder. Nature Neuroscience, 2019, 22, 1066-1074.	14.8	94
33	Anxiety with Panic Disorder Linked to Chromosome 9q in Iceland. American Journal of Human Genetics, 2003, 72, 1221-1230.	6.2	93
34	Polygenic risk scores for schizophrenia and bipolar disorder associate with addiction. Addiction Biology, 2018, 23, 485-492.	2.6	90
35	The Genetic Architecture of Depression in Individuals of East Asian Ancestry. JAMA Psychiatry, 2021, 78, 1258.	11.0	88
36	Does Childhood Trauma Moderate Polygenic Risk for Depression? A Meta-analysis of 5765 Subjects From the Psychiatric Genomics Consortium. Biological Psychiatry, 2018, 84, 138-147.	1.3	87

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37	Neuropathic pain phenotyping by international consensus (NeuroPPIC) for genetic studies. Pain, 2015, 156, 2337-2353.	4.2	86
38	Genome-wide meta-analysis reveals common splice site acceptor variant in CHRNA4 associated with nicotine dependence. Translational Psychiatry, 2015, 5, e651-e651.	4.8	86
39	Bacteriorhodopsin D85N: Three spectroscopic species in equilibrium. Biochemistry, 1993, 32, 1332-1337.	2.5	84
40	Effects of temperature on rhodopsin photointermediates from lumirhodopsin to metarhodopsin II. Biochemistry, 1993, 32, 13861-13872.	2.5	82
41	Effects of detergent environments on the photocycle of purified monomeric bacteriorhodopsin. Biochemistry, 1991, 30, 1751-1761.	2.5	81
42	Genome-wide association study across European and African American ancestries identifies a SNP in DNMT3B contributing to nicotine dependence. Molecular Psychiatry, 2018, 23, 1911-1919.	7.9	80
43	Direct Determination of the Membrane Affinities of Individual Amino Acidsâ€. Biochemistry, 1996, 35, 1803-1809.	2.5	78
44	Increased Genetic Vulnerability to Smoking at CHRNA5 in Early-Onset Smokers. Archives of General Psychiatry, 2012, 69, 854.	12.3	71
45	Effects of Asp-96 .fwdarw. Asn, Asp-85 .fwdarw. Asn, and Arg-82 .fwdarw. Gln single-site substitutions on the photocycle of bacteriorhodopsin. Biochemistry, 1991, 30, 9133-9142.	2.5	70
46	Sequence variants in ARHGAP15, COLQ and FAM155A associate with diverticular disease and diverticulitis. Nature Communications, 2017, 8, 15789.	12.8	67
47	MgATP Binding and Hydrolysis Determinants of NtrC, a Bacterial Enhancer-Binding Protein. Journal of Bacteriology, 1999, 181, 4628-4638.	2.2	64
48	Truncating mutations in RBM12 are associated with psychosis. Nature Genetics, 2017, 49, 1251-1254.	21.4	63
49	Photoinduced electron transfer and enhanced triplet yields in benzo[a]pyrene derivative-nucleic acid complexes and covalent adducts. Journal of the American Chemical Society, 1994, 116, 63-72.	13.7	60
50	The association between lower educational attainment and depression owing to shared genetic effects? Results in ~25 000 subjects. Molecular Psychiatry, 2015, 20, 735-743.	7.9	59
51	Two Modes of Ligand Binding in Maltose-binding Protein ofEscherichia coli. Journal of Biological Chemistry, 1997, 272, 17610-17614.	3.4	58
52	Genome-wide association study of panic disorder reveals genetic overlap with neuroticism and depression. Molecular Psychiatry, 2021, 26, 4179-4190.	7.9	58
53	Topology of an Amphiphilic Mitochondrial Signal Sequence in the Membrane-Inserted State: A Spin Labeling Study. Biochemistry, 1994, 33, 14221-14226.	2.5	51
54	A common biological basis of obesity and nicotine addiction. Translational Psychiatry, 2013, 3, e308.e308.	4.8	51

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55	Direct Measurement of Small Ligand-Induced Conformational Changes in the Aspartate Chemoreceptor Using EPRâ€. Biochemistry, 1998, 37, 7062-7069.	2.5	45
56	PHOTOLYSIS OF RHODOPSIN RESULTS IN DEPROTONATION OF ITS RETINAL SCHIFF'S BASE PRIOR TO FORMATION OF METARHODOPSIN II. Photochemistry and Photobiology, 1992, 56, 1135-1144.	2.5	41
57	Physical evidence for a phosphorylation-dependent conformational change in the enhancer-binding protein NtrC. Proceedings of the National Academy of Sciences of the United States of America, 1999, 96, 4880-4885.	7.1	41
58	Neuregulin 1 in schizophrenia: out of Iceland. Molecular Psychiatry, 2003, 8, 639-640.	7.9	36
59	Temperature Dependence of Polypeptide Partitioning between Water and Phospholipid Bilayersâ€. Biochemistry, 1996, 35, 9526-9532.	2.5	33
60	Classical Human Leukocyte Antigen Alleles and C4 Haplotypes Are Not Significantly Associated With Depression. Biological Psychiatry, 2020, 87, 419-430.	1.3	27
61	A Limiting Law for the Electrostatics of the Binding of Polypeptides to Phospholipid Bilayers. Biochemistry, 1995, 34, 5518-5522.	2.5	26
62	Commentary: Gene-environment interactions and smoking-related cancers. International Journal of Epidemiology, 2010, 39, 577-579.	1.9	26
63	Genome-Wide Significant Association Between a Sequence Variant at 15q15.2 and Lung Cancer Risk. Cancer Research, 2011, 71, 1356-1361.	0.9	26
64	A rare missense mutation in CHRNA4 associates with smoking behavior and its consequences. Molecular Psychiatry, 2016, 21, 594-600.	7.9	26
65	Genetics of Smoking Behavior and Its Consequences: The Role of Nicotinic Acetylcholine Receptors. Biological Psychiatry, 2008, 64, 919-921.	1.3	24
66	Addictions and their familiality in Iceland. Annals of the New York Academy of Sciences, 2010, 1187, 208-217.	3.8	22
67	The Membrane Affinities of the Aliphatic Amino Acid Side Chains in an α-Helical Context Are Independent of Membrane Immersion Depthâ€. Biochemistry, 1999, 38, 337-346.	2.5	17
68	Photolysis intermediates of human rhodopsin. Biochemistry, 1991, 30, 11372-11376.	2.5	16
69	Association of Whole-Genome and NETRIN1 Signaling Pathway–Derived Polygenic Risk Scores for Major Depressive Disorder and White Matter Microstructure in the UK Biobank. Biological Psychiatry: Cognitive Neuroscience and Neuroimaging, 2019, 4, 91-100.	1.5	16
70	GPC5 rs2352028 variant and risk of lung cancer in never smokers. Lancet Oncology, The, 2010, 11, 714-716.	10.7	15
71	Integration of evidence across human and model organism studies: A meeting report. Genes, Brain and Behavior, 2021, 20, e12738.	2.2	12
72	Common and rare sequence variants influencing tumor biomarkers in blood. Cancer Epidemiology Biomarkers and Prevention, 2019, 29, cebp.1060.2018.	2.5	9

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73	De novo design of a peptide which partitions between water and phospholipid bilayers as a monomeric alpha-helix. Protein Engineering, Design and Selection, 1998, 11, 539-547.	2.1	8
74	A genome-wide meta-analysis identifies 50 genetic loci associated with carpal tunnel syndrome. Nature Communications, 2022, 13, 1598.	12.8	8
75	From paper to web: Mode equivalence of the ARHQ and NEO-FFI. Computers in Human Behavior, 2014, 41, 384-392.	8.5	7
76	Psychometric properties of the Icelandic NEO-FFI in a general population sample compared to a sample recruited for a study on the genetics of addiction. Personality and Individual Differences, 2014, 58, 71-75.	2.9	6
77	Genetic propensities for verbal and spatial ability have opposite effects on body mass index and risk of schizophrenia. Intelligence, 2021, 88, 101565.	3.0	2