Yong-Yong Shi

List of Publications by Year
in descending order

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		57719	23514
150	13,599	44	111
papers	citations	h-index	g-index

157	157	157	20913
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	SHEsis, a powerful software platform for analyses of linkage disequilibrium, haplotype construction, and genetic association at polymorphism loci. Cell Research, 2005, 15, 97-98.	5.7	2,010
2	The sequence and de novo assembly of the giant panda genome. Nature, 2010, 463, 311-317.	13.7	1,058
3	Identification of loci associated with schizophrenia by genome-wide association and follow-up. Nature Genetics, 2008, 40, 1053-1055.	9.4	977
4	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. Nature, 2022, 604, 502-508.	13.7	929
5	A partition-ligation-combination-subdivision EM algorithm for haplotype inference with multiallelic markers: update of the SHEsis (http://analysis.bio-x.cn). Cell Research, 2009, 19, 519-523.	5.7	706
6	Genome-wide association study identifies susceptibility loci for polycystic ovary syndrome on chromosome 2p16.3, 2p21 and 9q33.3. Nature Genetics, 2011, 43, 55-59.	9.4	604
7	Genome-wide association study identifies eight new risk loci for polycystic ovary syndrome. Nature Genetics, 2012, 44, 1020-1025.	9.4	505
8	Comparative genetic architectures of schizophrenia in East Asian and European populations. Nature Genetics, 2019, 51, 1670-1678.	9.4	440
9	Genome-wide association analysis identifies 30 new susceptibility loci for schizophrenia. Nature Genetics, 2017, 49, 1576-1583.	9.4	395
10	A genome-wide association study identifies two new lung cancer susceptibility loci at 13q12.12 and 22q12.2 in Han Chinese. Nature Genetics, 2011, 43, 792-796.	9.4	340
11	Genome-wide association study in Han Chinese identifies four new susceptibility loci for coronary artery disease. Nature Genetics, 2012, 44, 890-894.	9.4	295
12	The OncoArray Consortium: A Network for Understanding the Genetic Architecture of Common Cancers. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 126-135.	1.1	278
13	Recurrent gain-of-function USP8 mutations in Cushing's disease. Cell Research, 2015, 25, 306-317.	5.7	263
14	Genetic Variants Associated With Phenytoin-Related Severe Cutaneous Adverse Reactions. JAMA - Journal of the American Medical Association, 2014, 312, 525.	3.8	256
15	A genome-wide association study identifies new susceptibility loci for non-cardia gastric cancer at 3q13.31 and 5p13.1. Nature Genetics, 2011, 43, 1215-1218.	9.4	250
16	Common variants on 8p12 and 1q24.2 confer risk of schizophrenia. Nature Genetics, 2011, 43, 1224-1227.	9.4	224
17	GWAS Identifies Novel Susceptibility Loci on 6p21.32 and 21q21.3 for Hepatocellular Carcinoma in Chronic Hepatitis B Virus Carriers. PLoS Genetics, 2012, 8, e1002791.	1.5	177
18	New loci associated with chronic hepatitis B virus infection in Han Chinese. Nature Genetics, 2013, 45, 1499-1503.	9.4	140

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19	The genome-wide mutational landscape of pituitary adenomas. Cell Research, 2016, 26, 1255-1259.	5.7	137
20	Association analyses identify multiple new lung cancer susceptibility loci and their interactions with smoking in the Chinese population. Nature Genetics, 2012, 44, 895-899.	9.4	129
21	A genome-wide association study identifies two new cervical cancer susceptibility loci at 4q12 and 17q12. Nature Genetics, 2013, 45, 918-922.	9.4	108
22	Cell Culture System for Analysis of Genetic Heterogeneity WithinÂHepatocellular Carcinomas and Response to Pharmacologic Agents. Gastroenterology, 2017, 152, 232-242.e4.	0.6	107
23	Gold nanoparticles for high-throughput genotyping of long-range haplotypes. Nature Nanotechnology, 2011, 6, 639-644.	15.6	106
24	Identification of recurrent USP48 and BRAF mutations in Cushingâ€™s disease. Nature Communications, 2018, 9, 3171.	5.8	106
25	A genome-wide association study identifies two risk loci for congenital heart malformations in Han Chinese populations. Nature Genetics, 2013, 45, 818-821.	9.4	88
26	Genome-wide association analysis identifies three new risk loci for gout arthritis in Han Chinese. Nature Communications, 2015, 6, 7041.	5.8	88
27	Recurrent deletions of ULK4 in schizophrenia: a novel gene crucial for neuritogenesis and neuronal motility. Journal of Cell Science, 2014, 127, 630-40.	1.2	78
28	SHEsisPlus, a toolset for genetic studies on polyploid species. Scientific Reports, 2016, 6, 24095.	1.6	77
29	Genetic Variants at 6p21.1 and 7p15.3 Are Associated with Risk of Multiple Cancers in Han Chinese. American Journal of Human Genetics, 2012, 91, 928-934.	2.6	76
30	Effect of aerobic exercise and diet on liver fat in pre-diabetic patients with non-alcoholic-fatty-liver-disease: A randomized controlled trial. Scientific Reports, 2017, 7, 15952.	1.6	74
31	DNA origami-based shape IDs for single-molecule nanomechanical genotyping. Nature Communications, 2017, 8, 14738.	5.8	73
32	Common Variants in Major Histocompatibility Complex Region and TCF4 Gene Are Significantly Associated with Schizophrenia in Han Chinese. Biological Psychiatry, 2010, 68, 671-673.	0.7	69
33	Identification of new susceptibility loci for gastric non-cardia adenocarcinoma: pooled results from two Chinese genome-wide association studies. Gut, 2017, 66, 581-587.	6.1	68
34	A Case-control association study between the GRID1 gene and schizophrenia in the Chinese Northern Han population. Schizophrenia Research, 2007, 93, 385-390.	1.1	67
35	SHEsisEpi, a GPU-enhanced genome-wide SNP-SNP interaction scanning algorithm, efficiently reveals the risk genetic epistasis in bipolar disorder. Cell Research, 2010, 20, 854-857.	5.7	63
36	Systematic polymorphism analysis of the CYP2D6 gene in four different geographical Han populations in mainland China. Genomics, 2008, 92, 152-158.	1.3	59

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37	Exome Array Analysis Identifies Variants in SPOCD1 and BTN3A2 That Affect Risk for Gastric Cancer. Gastroenterology, 2017, 152, 2011-2021.	0.6	58
38	Germline Mutations in CDH23, Encoding Cadherin-Related 23, Are Associated with Both Familial and Sporadic Pituitary Adenomas. American Journal of Human Genetics, 2017, 100, 817-823.	2.6	57
39	Susceptibility loci for metabolic syndrome and metabolic components identified in Han Chinese: a multiâ€stage genomeâ€wide association study. Journal of Cellular and Molecular Medicine, 2017, 21, 1106-1116.	1.6	56
40	Genome-wide Analysis of the Role of Copy Number Variation in Schizophrenia Risk in Chinese. Biological Psychiatry, 2016, 80, 331-337.	0.7	55
41	CACNA1C, schizophrenia and major depressive disorder in the Han Chinese population. British Journal of Psychiatry, 2014, 204, 36-39.	1.7	53
42	Rare CNVs and Tag SNPs at 15q11.2 Are Associated With Schizophrenia in the Han Chinese Population. Schizophrenia Bulletin, 2013, 39, 712-719.	2.3	52
43	Genome-wide association study identifies two risk loci for tuberculosis in Han Chinese. Nature Communications, 2018, 9, 4072.	5.8	51
44	CNTNAP2 is significantly associated with schizophrenia and major depression in the Han Chinese population. Psychiatry Research, 2013, 207, 225-228.	1.7	50
45	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
46	Family-based analysis of susceptibility loci for polycystic ovary syndrome on chromosome 2p16.3, 2p21 and 9q33.3. Human Reproduction, 2012, 27, 294-298.	0.4	47
47	A genome-wide gene-environment interaction analysis for tobacco smoke and lung cancer susceptibility. Carcinogenesis, 2014, 35, 1528-1535.	1.3	47
48	A Genome-Wide Association Study Identifies a Locus on TERT for Mean Telomere Length in Han Chinese. PLoS ONE, 2014, 9, e85043.	1.1	46
49	Common SNPs and haplotypes in DGKH are associated with bipolar disorder and schizophrenia in the Chinese Han population. Molecular Psychiatry, 2011, 16, 473-475.	4.1	43
50	Common variants at 10p12.31, 10q21.1 and 13q12.13 are associated with sporadic pituitary adenoma. Nature Genetics, 2015, 47, 793-797.	9.4	43
51	A modifier screen identifies DNAJB6 as a cardiomyopathy susceptibility gene. JCI Insight, 2016, 1, .	2.3	42
52	Genome-Wide Association Study Identifies a Novel Susceptibility Locus at 12q23.1 for Lung Squamous Cell Carcinoma in Han Chinese. PLoS Genetics, 2013, 9, e1003190.	1.5	41
53	Low-Frequency Coding Variants at 6p21.33 and 20q11.21 Are Associated with Lung Cancer Risk in Chinese Populations. American Journal of Human Genetics, 2015, 96, 832-840.	2.6	41
54	MicroRNA-137 Inhibits EFNB2 Expression Affected by a Genetic Variant and Is Expressed Aberrantly in Peripheral Blood of Schizophrenia Patients. EBioMedicine, 2016, 12, 133-142.	2.7	41

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55	Genomic dissection of 43 serum urate-associated loci provides multiple insights into molecular mechanisms of urate control. Human Molecular Genetics, 2020, 29, 923-943.	1.4	40
56	Common Variants in the BCL9 Gene Conferring Risk of Schizophrenia. Archives of General Psychiatry, 2011, 68, 232.	13.8	39
57	A panel of ancestry informative markers to estimate and correct potential effects of population stratification in Han Chinese. European Journal of Human Genetics, 2014, 22, 248-253.	1.4	39
58	Body Mass Index and Polycystic Ovary Syndrome: A 2-Sample Bidirectional Mendelian Randomization Study. Journal of Clinical Endocrinology and Metabolism, 2020, 105, 1778-1784.	1.8	39
59	Genetic Structure Adds Power to Detect Schizophrenia Susceptibility at SLIT3 in the Chinese Han Population. Genome Research, 2004, 14, 1345-1349.	2.4	36
60	Apoptotic Engulfment Pathway and Schizophrenia. PLoS ONE, 2009, 4, e6875.	1.1	35
61	The MDGA1 gene confers risk to schizophrenia and bipolar disorder. Schizophrenia Research, 2011, 125, 194-200.	1.1	35
62	Glucose and Insulin-Related Traits, Type 2 Diabetes and Risk of Schizophrenia: A Mendelian Randomization Study. EBioMedicine, 2018, 34, 182-188.	2.7	34
63	Fine mapping the MHC region identified four independent variants modifying susceptibility to chronic hepatitis B in Han Chinese. Human Molecular Genetics, 2016, 25, 1225-1232.	1.4	33
64	Genome-wide association study of cervical cancer suggests a role forARRDC3gene in human papillomavirus infection. Human Molecular Genetics, 2019, 28, 341-348.	1.4	33
65	ZNF804A and schizophrenia susceptibility in Asian populations. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2012, 159B, 794-802.	1.1	30
66	A genome-wide geneâ€“gene interaction analysis identifies an epistatic gene pair for lung cancer susceptibility in Han Chinese. Carcinogenesis, 2014, 35, 572-577.	1.3	29
67	Loci with genome-wide associations with schizophrenia in the Han Chinese population. British Journal of Psychiatry, 2015, 207, 490-494.	1.7	29
68	A meta-analysis of three polymorphisms in the endothelial nitric oxide synthase gene (NOS3) and their effect on the risk of diabetic nephropathy. Human Genetics, 2010, 127, 373-381.	1.8	28
69	Genome-wide association studies identify susceptibility loci for epithelial ovarian cancer in east Asian women. Gynecologic Oncology, 2019, 153, 343-355.	0.6	28
70	Structural Analysis of the SARS-CoV-2 Omicron Variant Proteins. Research, 2021, 2021, 9769586.	2.8	27
71	Genetic association between NRG1 and schizophrenia, major depressive disorder, bipolar disorder in Han Chinese population. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2016, 171, 468-478.	1.1	26
72	CTLA-4 confers a risk of recurrent schizophrenia, major depressive disorder and bipolar disorder in the Chinese Han population. Brain, Behavior, and Immunity, 2011, 25, 429-433.	2.0	24

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73	Immunosuppressive potential of fowl adenovirus serotype 4. Poultry Science, 2019, 98, 3514-3522.	1.5	24
74	Identifying the Genotypes of Hepatitis B Virus (HBV) with DNA Origami Label. Small, 2018, 14, 1701718.	5.2	23
75	Genome-wide association study in Han Chinese identifies three novel loci for human height. Human Genetics, 2013, 132, 681-689.	1.8	21
76	Association Study Between Polymorphisms of PRMT6, PEX10, SOX5, and Nonobstructive Azoospermia in the Han Chinese Population1. Biology of Reproduction, 2014, 90, 96.	1.2	19
77	Association between SREBF2 gene polymorphisms and metabolic syndrome in clozapine-treated patients with schizophrenia. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2015, 56, 136-141.	2.5	19
78	The schizophrenia genetics knowledgebase: a comprehensive update of findings from candidate gene studies. Translational Psychiatry, 2019, 9, 205.	2.4	19
79	Prediction of causal genes and gene expression analysis of attention-deficit hyperactivity disorder in the different brain region, a comprehensive integrative analysis of ADHD. Behavioural Brain Research, 2019, 364, 183-192.	1.2	18
80	ITIH family genes confer risk to schizophrenia and major depressive disorder in the Han Chinese population. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2014, 51, 34-38.	2.5	17
81	Systematic Evaluation of Genetic Variants for Polycystic Ovary Syndrome in a Chinese Population. PLoS ONE, 2015, 10, e0140695.	1.1	17
82	The NVL gene confers risk for both major depressive disorder and schizophrenia in the Han Chinese population. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2015, 62, 7-13.	2.5	17
83	Both HLA class I and II regions identified as genome-wide significant susceptibility loci for adult-onset Still's disease in Chinese individuals. Annals of the Rheumatic Diseases, 2020, 79, 161-163.	0.5	17
84	Mapping of hepatic expression quantitative trait loci (eQTLs) in a Han Chinese population. Journal of Medical Genetics, 2014, 51, 319-326.	1.5	16
85	Comparison of the performance of Ion Torrent chips in noninvasive prenatal trisomy detection. Journal of Human Genetics, 2014, 59, 393-396.	1.1	15
86	Polymorphisms in GCKR, SLC17A1 and SLC22A12 were associated with phenotype gout in Han Chinese males: a caseâ€“control study. BMC Medical Genetics, 2015, 16, 66.	2.1	15
87	Replication of Gout/Urate Concentrations GWAS Susceptibility Loci Associated with Gout in a Han Chinese Population. Scientific Reports, 2017, 7, 4094.	1.6	15
88	Structural Comparison and Drug Screening of Spike Proteins of Ten SARS-CoV-2 Variants. Research, 2022, 2022, 9781758.	2.8	15
89	The Potential Effect of Aberrant Testosterone Levels on Common Diseases: A Mendelian Randomization Study. Genes, 2020, 11, 721.	1.0	14
90	Analysis of association between common SNPs in ErbB4 and bipolar affective disorder, major depressive disorder and schizophrenia in the Han Chinese population. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2012, 36, 17-21.	2.5	13

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91	Genetic variants at 5p15 are associated with risk and early onset of gastric cancer in Chinese populations. Carcinogenesis, 2013, 34, 2539-2542.	1.3	13
92	TheCMYA5gene confers risk for both schizophrenia and major depressive disorder in the Han Chinese population. World Journal of Biological Psychiatry, 2014, 15, 553-560.	1.3	13
93	Polymorphisms of Renin-Angiotensin-Aldosterone System Gene in Chinese Han Patients with Nonfamilial Atrial Fibrillation. PLoS ONE, 2015, 10, e0117489.	1.1	12
94	Genetic association of ACSM1 variation with schizophrenia and major depressive disorder in the Han Chinese population. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2015, 168, 144-149.	1.1	12
95	Association betweenSCAPandSREBF1gene polymorphisms and metabolic syndrome in schizophrenia patients treated with atypical antipsychotics. World Journal of Biological Psychiatry, 2016, 17, 467-474.	1.3	12
96	Genetic risk between the CACNA1I gene and schizophrenia in Chinese Uygur population. Hereditas, 2018, 155, 5.	0.5	12
97	Association of SCN10A Polymorphisms with the Recurrence of Atrial Fibrillation after Catheter Ablation in a Chinese Han Population. Scientific Reports, 2017, 7, 44003.	1.6	11
98	The pathogenicity of duck hepatitis A virus types 1 and 3 on ducklings. Poultry Science, 2019, 98, 6333-6339.	1.5	11
99	No Association of the YWHAE Gene with Schizophrenia, Major Depressive Disorder or Bipolar Disorder in the Han Chinese Population. Behavior Genetics, 2011, 41, 557-564.	1.4	10
100	A new risk locus in the ZEB2 gene for schizophrenia in the Han Chinese population. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2016, 66, 97-103.	2.5	10
101	Polymorphisms in NRGN are associated with schizophrenia, major depressive disorder and bipolar disorder in the Han Chinese population. Journal of Affective Disorders, 2016, 194, 180-187.	2.0	10
102	Genomeâ€wide twoâ€locus interaction analysis identifies multiple epistatic SNP pairs that confer risk of prostate cancer: A crossâ€population study. International Journal of Cancer, 2017, 140, 2075-2084.	2.3	10
103	SNX29, a new susceptibility gene shared with major mental disorders in Han Chinese population. World Journal of Biological Psychiatry, 2021, 22, 526-534.	1.3	10
104	Association analysis of theGRM8gene with schizophrenia in the Uygur Chinese population. Hereditas, 2014, 151, 140-144.	0.5	9
105	Common variants in ZMIZ1 and near NGF confer risk for primary dysmenorrhoea. Nature Communications, 2017, 8, 14900.	5.8	9
106	Association between the variability of theABCA13gene and the risk of major depressive disorder and schizophrenia in the Han Chinese population. World Journal of Biological Psychiatry, 2017, 18, 550-556.	1.3	9
107	Elevated levels of IL â€18 associated with schizophrenia and first episode psychosis: A systematic review and metaâ€analysis. Microbial Biotechnology, 2020, 15, 896-905.	0.9	9
108	Genetic analysis of common variants in the ZNF804A gene with schizophrenia and major depressive disorder. Psychiatric Genetics, 2018, 28, 1-7.	0.6	8

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109	ACTN3 is associated with childrenâ€™s physical fitness in Han Chinese. Molecular Genetics and Genomics, 2019, 294, 47-56.	1.0	8
110	Common Variants in the CDH7 Gene are Associated with Major Depressive Disorder in the Han Chinese Population. Behavior Genetics, 2014, 44, 97-101.	1.4	7
111	A novel variant associated with HDL-C levels by modifying DAGLB expression levels: An annotation-based genome-wide association study. European Journal of Human Genetics, 2018, 26, 838-847.	1.4	7
112	SLC39A8 is a risk factor for schizophrenia in Uygur Chinese: a case-control study. BMC Psychiatry, 2019, 19, 293.	1.1	7
113	Four Loci Are Associated with Cardiorespiratory Fitness and Endurance Performance in Young Chinese Females. Scientific Reports, 2020, 10, 10117.	1.6	7
114	Fineâ€mapping of ZDHHC2 identifies risk variants for schizophrenia in the Han Chinese population. Molecular Genetics & Genomic Medicine, 2020, 8, e1190.	0.6	7
115	Psychiatric genetics in China: achievements and challenges. Molecular Psychiatry, 2016, 21, 4-9.	4.1	6
116	Association study of NDST3 gene for schizophrenia, bipolar disorder, major depressive disorder in the Han Chinese population. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2018, 177, 3-9.	1.1	6
117	Meta-analysis of GABRB2 polymorphisms and the risk of schizophrenia combined with GWAS data of the Han Chinese population and psychiatric genomics consortium. PLoS ONE, 2018, 13, e0198690.	1.1	6
118	Identification of rare and common variants in BNIP3L: a schizophrenia susceptibility gene. Human Genomics, 2020, 14, 16.	1.4	6
119	Cigarette smoking and schizophrenia: Mendelian randomisation study. British Journal of Psychiatry, 2021, 218, 98-103.	1.7	6
120	Trans-ancestral dissection of urate- and gout-associated major loci SLC2A9 and ABCG2 reveals primate-specific regulatory effects. Journal of Human Genetics, 2021, 66, 161-169.	1.1	6
121	Identification of SHANK2 Pathogenic Variants in a Chinese Uygur Population with Schizophrenia. Journal of Molecular Neuroscience, 2021, 71, 1-8.	1.1	6
122	USP8 mutation in Cushing's disease. Oncotarget, 2015, 6, 18240-18241.	0.8	6
123	eRFSVM: a hybrid classifier to predict enhancers-integrating random forests with support vector machines. Hereditas, 2016, 153, 6.	0.5	5
124	The YWHAE gene confers risk to major depressive disorder in the male group of Chinese Han population. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2017, 77, 172-177.	2.5	5
125	Common variants in the SLC28A2 gene are associated with serum uric acid level and hyperuricemia and gout in Han Chinese. Hereditas, 2019, 156, 4.	0.5	5
126	Common variants in FAN1, located in 15q13.3, confer risk for schizophrenia and bipolar disorder in Han Chinese. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2020, 103, 109973.	2.5	5

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127	Genetic risk of clozapine-induced leukopenia and neutropenia: a genome-wide association study. Translational Psychiatry, 2021, 11, 343.	2.4	5
128	Disruption of MAP7D1 Gene Function Increases the Risk of Doxorubicin-Induced Cardiomyopathy and Heart Failure. BioMed Research International, 2021, 2021, 1-9.	0.9	5
129	The Relationship between Alcohol Consumption and Gout: A Mendelian Randomization Study. Genes, 2022, 13, 557.	1.0	5
130	Analysis of association between common variants in theSLCO6A1gene with schizophrenia, bipolar disorder and major depressive disorder in the Han Chinese population. World Journal of Biological Psychiatry, 2016, 17, 140-146.	1.3	4
131	Association of fat mass and obesity-associated and retinitis pigmentosa guanosine triphosphatase (GTPase) regulator-interacting protein-1 like polymorphisms with body mass index in Chinese women. Endocrine Journal, 2018, 65, 783-791.	0.7	4
132	Amplicon targeted resequencing for SLC2A9 and SLC22A12 identified novel mutations in hypouricemia subjects. Molecular Genetics & Genomic Medicine, 2019, 7, e00722.	0.6	4
133	The association between rs12807809 polymorphism in neurogranin gene and risk of schizophrenia. Medicine (United States), 2019, 98, e18518.	0.4	4
134	Rare and common variants analysis of the EMB gene in patients with schizophrenia. BMC Psychiatry, 2020, 20, 135.	1.1	4
135	Functional annotation of regulatory single nucleotide polymorphisms associated with schizophrenia. Schizophrenia Research, 2020, 218, 326-328.	1.1	4
136	A machine learning-assisted model for renal urate underexcretion with genetic and clinical variables among Chinese men with gout. Arthritis Research and Therapy, 2022, 24, 67.	1.6	4
137	Common variants in QPCT gene confer risk of schizophrenia in the Han Chinese population. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2016, 171, 237-242.	1.1	3
138	Role played by the SP4 gene in schizophrenia and major depressive disorder in the Han Chinese population. British Journal of Psychiatry, 2016, 208, 441-445.	1.7	3
139	Polymorphism of the PPARD Gene and Dynamic Balance Performance in Han Chinese Children. Hereditas, 2019, 156, 15.	0.5	3
140	VariFAST: a variant filter by automated scoring based on tagged-signatures. BMC Bioinformatics, 2019, 20, 713.	1.2	3
141	Common variants in SATB2 are associated with schizophrenia in Uygur Chinese population. Psychiatric Genetics, 2019, 29, 120-126.	0.6	2
142	Polymorphisms and rare variants identified by next-generation sequencing confer risk for lung cancer in han Chinese population. Pathology Research and Practice, 2020, 216, 152873.	1.0	2
143	SHEsisPCA: A GPU-Based Software to Correct for Population Stratification that Efficiently Accelerates the Process for Handling Genome-Wide Datasets. Journal of Genetics and Genomics, 2015, 42, 445-453.	1.7	1
144	Noninvasive fetal trisomy detection by multiplexed semiconductor sequencing: a barcoding analysis strategy. Journal of Human Genetics, 2016, 61, 247-252.	1.1	1

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145	Prediction of functional regulatory elements of bipolar disorder via data integration analysis. Journal of Affective Disorders Reports, 2020, 1, 100015.	0.9	1
146	An Evaluation of Association between a Novel Hippocampal Biology Related SNP (rs7294919) and Schizophrenia. PLoS ONE, 2013, 8, e80696.	1.1	1
147	MicroRNA Microarray Analysis Combined with Interaction Network Analysis to Investigate the Influence of Clozapine to Metabolic Syndrome. International Journal of Pharmacology, 2013, 9, 366-372.	0.1	1
148	Association Analysis Between Common Variants of the TRPM1 Gene and Three Mental Disorders in the Han Chinese Population. Genetic Testing and Molecular Biomarkers, 2020, 24, 649-657.	0.3	1
149	Association analysis of potentially functional variants within 8p12 with schizophrenia in the Han Chinese population. World Journal of Biological Psychiatry, 2021, 22, 27-33.	1.3	0
150	Scrutinizing the causal relationship between schizophrenia and vitamin supplementation. Journal of Bio-X Research, 2021, Publish Ahead of Print, .	0.3	0