

# Yong-Yong Shi

## List of Publications by Citations

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148  
papers

9,845  
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39  
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157  
ext. papers

11,912  
ext. citations

9.6  
avg, IF

5.55  
L-index

#	Paper	IF	Citations
148	SHEsis, a powerful software platform for analyses of linkage disequilibrium, haplotype construction, and genetic association at polymorphism loci. <i>Cell Research</i> , <b>2005</b> , 15, 97-8	24.7	1736
147	Identification of loci associated with schizophrenia by genome-wide association and follow-up. <i>Nature Genetics</i> , <b>2008</b> , 40, 1053-5	36.3	877
146	The sequence and de novo assembly of the giant panda genome. <i>Nature</i> , <b>2010</b> , 463, 311-7	50.4	864
145	A partition-ligation-combination-subdivision EM algorithm for haplotype inference with multiallelic markers: update of the SHEsis ( <a href="http://analysis.bio-x.cn">http://analysis.bio-x.cn</a> ). <i>Cell Research</i> , <b>2009</b> , 19, 519-23	24.7	610
144	Genome-wide association study identifies susceptibility loci for polycystic ovary syndrome on chromosome 2p16.3, 2p21 and 9q33.3. <i>Nature Genetics</i> , <b>2011</b> , 43, 55-9	36.3	462
143	Genome-wide association study identifies eight new risk loci for polycystic ovary syndrome. <i>Nature Genetics</i> , <b>2012</b> , 44, 1020-5	36.3	380
142	A genome-wide association study identifies two new lung cancer susceptibility loci at 13q12.12 and 22q12.2 in Han Chinese. <i>Nature Genetics</i> , <b>2011</b> , 43, 792-6	36.3	288
141	Genome-wide association analysis identifies 30 new susceptibility loci for schizophrenia. <i>Nature Genetics</i> , <b>2017</b> , 49, 1576-1583	36.3	272
140	Genome-wide association study in Han Chinese identifies four new susceptibility loci for coronary artery disease. <i>Nature Genetics</i> , <b>2012</b> , 44, 890-4	36.3	243
139	A genome-wide association study identifies new susceptibility loci for non-cardia gastric cancer at 3q13.31 and 5p13.1. <i>Nature Genetics</i> , <b>2011</b> , 43, 1215-8	36.3	215
138	Genetic variants associated with phenytoin-related severe cutaneous adverse reactions. <i>JAMA - Journal of the American Medical Association</i> , <b>2014</b> , 312, 525-34	27.4	209
137	Common variants on 8p12 and 1q24.2 confer risk of schizophrenia. <i>Nature Genetics</i> , <b>2011</b> , 43, 1224-7	36.3	201
136	Recurrent gain-of-function USP8 mutations in Cushing's disease. <i>Cell Research</i> , <b>2015</b> , 25, 306-17	24.7	189
135	Comparative genetic architectures of schizophrenia in East Asian and European populations. <i>Nature Genetics</i> , <b>2019</b> , 51, 1670-1678	36.3	185
134	The OncoArray Consortium: A Network for Understanding the Genetic Architecture of Common Cancers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2017</b> , 26, 126-135	4	183
133	GWAS identifies novel susceptibility loci on 6p21.32 and 21q21.3 for hepatocellular carcinoma in chronic hepatitis B virus carriers. <i>PLoS Genetics</i> , <b>2012</b> , 8, e1002791	6	142
132	New loci associated with chronic hepatitis B virus infection in Han Chinese. <i>Nature Genetics</i> , <b>2013</b> , 45, 1499-503	36.3	108

131	Association analyses identify multiple new lung cancer susceptibility loci and their interactions with smoking in the Chinese population. <i>Nature Genetics</i> , <b>2012</b> , 44, 895-9	36.3	108
130	Gold nanoparticles for high-throughput genotyping of long-range haplotypes. <i>Nature Nanotechnology</i> , <b>2011</b> , 6, 639-44	28.7	97
129	Cell Culture System for Analysis of Genetic Heterogeneity Within Hepatocellular Carcinomas and Response to Pharmacologic Agents. <i>Gastroenterology</i> , <b>2017</b> , 152, 232-242.e4	13.3	81
128	The genome-wide mutational landscape of pituitary adenomas. <i>Cell Research</i> , <b>2016</b> , 26, 1255-1259	24.7	81
127	A genome-wide association study identifies two new cervical cancer susceptibility loci at 4q12 and 17q12. <i>Nature Genetics</i> , <b>2013</b> , 45, 918-22	36.3	80
126	Genome-wide association analysis identifies three new risk loci for gout arthritis in Han Chinese. <i>Nature Communications</i> , <b>2015</b> , 6, 7041	17.4	68
125	DNA origami-based shape IDs for single-molecule nanomechanical genotyping. <i>Nature Communications</i> , <b>2017</b> , 8, 14738	17.4	62
124	Common variants in major histocompatibility complex region and TCF4 gene are significantly associated with schizophrenia in Han Chinese. <i>Biological Psychiatry</i> , <b>2010</b> , 68, 671-3	7.9	62
123	A genome-wide association study identifies two risk loci for congenital heart malformations in Han Chinese populations. <i>Nature Genetics</i> , <b>2013</b> , 45, 818-21	36.3	60
122	Genetic variants at 6p21.1 and 7p15.3 are associated with risk of multiple cancers in Han Chinese. <i>American Journal of Human Genetics</i> , <b>2012</b> , 91, 928-34	11	59
121	SHESisEpi, a GPU-enhanced genome-wide SNP-SNP interaction scanning algorithm, efficiently reveals the risk genetic epistasis in bipolar disorder. <i>Cell Research</i> , <b>2010</b> , 20, 854-7	24.7	57
120	Identification of recurrent USP48 and BRAF mutations in Cushing's disease. <i>Nature Communications</i> , <b>2018</b> , 9, 3171	17.4	56
119	SHESisPlus, a toolset for genetic studies on polyploid species. <i>Scientific Reports</i> , <b>2016</b> , 6, 24095	4.9	55
118	A case-control association study between the GRID1 gene and schizophrenia in the Chinese Northern Han population. <i>Schizophrenia Research</i> , <b>2007</b> , 93, 385-90	3.6	53
117	Recurrent deletions of ULK4 in schizophrenia: a gene crucial for neuritogenesis and neuronal motility. <i>Journal of Cell Science</i> , <b>2014</b> , 127, 630-40	5.3	52
116	Systematic polymorphism analysis of the CYP2D6 gene in four different geographical Han populations in mainland China. <i>Genomics</i> , <b>2008</b> , 92, 152-8	4.3	52
115	Identification of new susceptibility loci for gastric non-cardia adenocarcinoma: pooled results from two Chinese genome-wide association studies. <i>Gut</i> , <b>2017</b> , 66, 581-587	19.2	51
114	CACNA1C, schizophrenia and major depressive disorder in the Han Chinese population. <i>British Journal of Psychiatry</i> , <b>2014</b> , 204, 36-9	5.4	44

113	Rare CNVs and tag SNPs at 15q11.2 are associated with schizophrenia in the Han Chinese population. <i>Schizophrenia Bulletin</i> , <b>2013</b> , 39, 712-9	1.3	42
112	Genome-wide Analysis of the Role of Copy Number Variation in Schizophrenia Risk in Chinese. <i>Biological Psychiatry</i> , <b>2016</b> , 80, 331-337	7.9	40
111	Effect of aerobic exercise and diet on liver fat in pre-diabetic patients with non-alcoholic-fatty-liver-disease: A randomized controlled trial. <i>Scientific Reports</i> , <b>2017</b> , 7, 15952	4.9	40
110	Common SNPs and haplotypes in DGKH are associated with bipolar disorder and schizophrenia in the Chinese Han population. <i>Molecular Psychiatry</i> , <b>2011</b> , 16, 473-5	15.1	40
109	Susceptibility loci for metabolic syndrome and metabolic components identified in Han Chinese: a multi-stage genome-wide association study. <i>Journal of Cellular and Molecular Medicine</i> , <b>2017</b> , 21, 1106-1116	5.6	39
108	Family-based analysis of susceptibility loci for polycystic ovary syndrome on chromosome 2p16.3, 2p21 and 9q33.3. <i>Human Reproduction</i> , <b>2012</b> , 27, 294-8	5.7	39
107	Germline Mutations in CDH23, Encoding Cadherin-Related 23, Are Associated with Both Familial and Sporadic Pituitary Adenomas. <i>American Journal of Human Genetics</i> , <b>2017</b> , 100, 817-823	11	37
106	CNTNAP2 is significantly associated with schizophrenia and major depression in the Han Chinese population. <i>Psychiatry Research</i> , <b>2013</b> , 207, 225-8	9.9	37
105	MicroRNA-137 Inhibits EFNB2 Expression Affected by a Genetic Variant and Is Expressed Aberrantly in Peripheral Blood of Schizophrenia Patients. <i>EBioMedicine</i> , <b>2016</b> , 12, 133-142	8.8	35
104	A genome-wide gene-environment interaction analysis for tobacco smoke and lung cancer susceptibility. <i>Carcinogenesis</i> , <b>2014</b> , 35, 1528-35	4.6	35
103	Mapping genomic loci implicates genes and synaptic biology in schizophrenia.. <i>Nature</i> , <b>2022</b> ,	50.4	35
102	A genome-wide association study identifies a locus on TERT for mean telomere length in Han Chinese. <i>PLoS ONE</i> , <b>2014</b> , 9, e85043	3.7	34
101	Genome-wide association study identifies a novel susceptibility locus at 12q23.1 for lung squamous cell carcinoma in han chinese. <i>PLoS Genetics</i> , <b>2013</b> , 9, e1003190	6	34
100	Genetics of schizophrenia: A consensus paper of the WFSBP Task Force on Genetics. <i>World Journal of Biological Psychiatry</i> , <b>2017</b> , 18, 492-505	3.8	33
99	Genetic structure adds power to detect schizophrenia susceptibility at SLIT3 in the Chinese Han population. <i>Genome Research</i> , <b>2004</b> , 14, 1345-9	9.7	33
98	Exome Array Analysis Identifies Variants in SPOCD1 and BTN3A2 That Affect Risk for Gastric Cancer. <i>Gastroenterology</i> , <b>2017</b> , 152, 2011-2021	13.3	32
97	Common variants at 10p12.31, 10q21.1 and 13q12.13 are associated with sporadic pituitary adenoma. <i>Nature Genetics</i> , <b>2015</b> , 47, 793-7	36.3	32
96	Common variants in the BCL9 gene conferring risk of schizophrenia. <i>Archives of General Psychiatry</i> , <b>2011</b> , 68, 232-40		32

95	A panel of ancestry informative markers to estimate and correct potential effects of population stratification in Han Chinese. <i>European Journal of Human Genetics</i> , <b>2014</b> , 22, 248-53	5.3	31
94	Low-frequency coding variants at 6p21.33 and 20q11.21 are associated with lung cancer risk in Chinese populations. <i>American Journal of Human Genetics</i> , <b>2015</b> , 96, 832-40	11	30
93	Apoptotic engulfment pathway and schizophrenia. <i>PLoS ONE</i> , <b>2009</b> , 4, e6875	3.7	30
92	ZNF804A and schizophrenia susceptibility in Asian populations. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2012</b> , 159B, 794-802	3.5	28
91	A modifier screen identifies as a cardiomyopathy susceptibility gene. <i>JCI Insight</i> , <b>2016</b> , 1,	9.9	28
90	The MDGA1 gene confers risk to schizophrenia and bipolar disorder. <i>Schizophrenia Research</i> , <b>2011</b> , 125, 194-200	3.6	27
89	A meta-analysis of three polymorphisms in the endothelial nitric oxide synthase gene (NOS3) and their effect on the risk of diabetic nephropathy. <i>Human Genetics</i> , <b>2010</b> , 127, 373-81	6.3	25
88	Loci with genome-wide associations with schizophrenia in the Han Chinese population. <i>British Journal of Psychiatry</i> , <b>2015</b> , 207, 490-4	5.4	24
87	Fine mapping the MHC region identified four independent variants modifying susceptibility to chronic hepatitis B in Han Chinese. <i>Human Molecular Genetics</i> , <b>2016</b> , 25, 1225-32	5.6	24
86	Genome-wide association study identifies two risk loci for tuberculosis in Han Chinese. <i>Nature Communications</i> , <b>2018</b> , 9, 4072	17.4	23
85	A genome-wide gene-gene interaction analysis identifies an epistatic gene pair for lung cancer susceptibility in Han Chinese. <i>Carcinogenesis</i> , <b>2014</b> , 35, 572-7	4.6	22
84	CTLA-4 confers a risk of recurrent schizophrenia, major depressive disorder and bipolar disorder in the Chinese Han population. <i>Brain, Behavior, and Immunity</i> , <b>2011</b> , 25, 429-33	16.6	22
83	Genomic dissection of 43 serum urate-associated loci provides multiple insights into molecular mechanisms of urate control. <i>Human Molecular Genetics</i> , <b>2020</b> , 29, 923-943	5.6	20
82	Body Mass Index and Polycystic Ovary Syndrome: A 2-Sample Bidirectional Mendelian Randomization Study. <i>Journal of Clinical Endocrinology and Metabolism</i> , <b>2020</b> , 105,	5.6	18
81	Genetic association between NRG1 and schizophrenia, major depressive disorder, bipolar disorder in Han Chinese population. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2016</b> , 171B, 468-78	3.5	18
80	Identifying the Genotypes of Hepatitis B Virus (HBV) with DNA Origami Label. <i>Small</i> , <b>2018</b> , 14, 1701718	11	17
79	Glucose and Insulin-Related Traits, Type 2 Diabetes and Risk of Schizophrenia: A Mendelian Randomization Study. <i>EBioMedicine</i> , <b>2018</b> , 34, 182-188	8.8	17
78	Genome-wide association study in Han Chinese identifies three novel loci for human height. <i>Human Genetics</i> , <b>2013</b> , 132, 681-9	6.3	17

77	Genome-wide association studies identify susceptibility loci for epithelial ovarian cancer in east Asian women. <i>Gynecologic Oncology</i> , <b>2019</b> , 153, 343-355	4.9	16
76	Association between SREBF2 gene polymorphisms and metabolic syndrome in clozapine-treated patients with schizophrenia. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , <b>2015</b> , 56, 136-41	5.5	16
75	Genome-wide association study of cervical cancer suggests a role for ARRDC3 gene in human papillomavirus infection. <i>Human Molecular Genetics</i> , <b>2019</b> , 28, 341-348	5.6	16
74	The NVL gene confers risk for both major depressive disorder and schizophrenia in the Han Chinese population. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , <b>2015</b> , 62, 7-13	5.5	15
73	Immunosuppressive potential of fowl adenovirus serotype 4. <i>Poultry Science</i> , <b>2019</b> , 98, 3514-3522	3.9	14
72	Comparison of the performance of Ion Torrent chips in noninvasive prenatal trisomy detection. <i>Journal of Human Genetics</i> , <b>2014</b> , 59, 393-6	4.3	14
71	ITIH family genes confer risk to schizophrenia and major depressive disorder in the Han Chinese population. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , <b>2014</b> , 51, 34-8	5.5	13
70	Mapping of hepatic expression quantitative trait loci (eQTLs) in a Han Chinese population. <i>Journal of Medical Genetics</i> , <b>2014</b> , 51, 319-26	5.8	13
69	Polymorphisms in GCKR, SLC17A1 and SLC22A12 were associated with phenotype gout in Han Chinese males: a case-control study. <i>BMC Medical Genetics</i> , <b>2015</b> , 16, 66	2.1	12
68	Association study between polymorphisms of PRMT6, PEX10, SOX5, and nonobstructive azoospermia in the Han Chinese population. <i>Biology of Reproduction</i> , <b>2014</b> , 90, 96	3.9	12
67	Replication of Gout/Urate Concentrations GWAS Susceptibility Loci Associated with Gout in a Han Chinese Population. <i>Scientific Reports</i> , <b>2017</b> , 7, 4094	4.9	12
66	Systematic Evaluation of Genetic Variants for Polycystic Ovary Syndrome in a Chinese Population. <i>PLoS ONE</i> , <b>2015</b> , 10, e0140695	3.7	12
65	Analysis of association between common SNPs in ErbB4 and bipolar affective disorder, major depressive disorder and schizophrenia in the Han Chinese population. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , <b>2012</b> , 36, 17-21	5.5	12
64	The schizophrenia genetics knowledgebase: a comprehensive update of findings from candidate gene studies. <i>Translational Psychiatry</i> , <b>2019</b> , 9, 205	8.6	10
63	Both class I and II regions identified as genome-wide significant susceptibility loci for adult-onset Still's disease in Chinese individuals. <i>Annals of the Rheumatic Diseases</i> , <b>2020</b> , 79, 161-163	2.4	10
62	Polymorphisms of renin-angiotensin-aldosterone system gene in chinese han patients with nonfamilial atrial fibrillation. <i>PLoS ONE</i> , <b>2015</b> , 10, e0117489	3.7	10
61	Genetic association of ACSM1 variation with schizophrenia and major depressive disorder in the Han Chinese population. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2015</b> , 168B, 144-9	3.5	10
60	Genetic risk between the gene and schizophrenia in Chinese Uygur population. <i>Hereditas</i> , <b>2018</b> , 155, 5	2.4	9

59	Genetic variants at 5p15 are associated with risk and early onset of gastric cancer in Chinese populations. <i>Carcinogenesis</i> , <b>2013</b> , 34, 2539-42	4.6	9
58	No association of the YWHAE gene with schizophrenia, major depressive disorder or bipolar disorder in the Han Chinese population. <i>Behavior Genetics</i> , <b>2011</b> , 41, 557-64	3.2	9
57	Structural Analysis of the SARS-CoV-2 Omicron Variant Proteins.. <i>Research</i> , <b>2021</b> , 2021, 9769586	7.8	8
56	Comparative genetic architectures of schizophrenia in East Asian and European populations		8
55	Association between SCAP and SREBF1 gene polymorphisms and metabolic syndrome in schizophrenia patients treated with atypical antipsychotics. <i>World Journal of Biological Psychiatry</i> , <b>2016</b> , 17, 467-74	3.8	8
54	Genome-wide two-locus interaction analysis identifies multiple epistatic SNP pairs that confer risk of prostate cancer: A cross-population study. <i>International Journal of Cancer</i> , <b>2017</b> , 140, 2075-2084	7.5	7
53	Prediction of causal genes and gene expression analysis of attention-deficit hyperactivity disorder in the different brain region, a comprehensive integrative analysis of ADHD. <i>Behavioural Brain Research</i> , <b>2019</b> , 364, 183-192	3.4	7
52	A new risk locus in the ZEB2 gene for schizophrenia in the Han Chinese population. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , <b>2016</b> , 66, 97-103	5.5	7
51	Polymorphisms in NRG1 are associated with schizophrenia, major depressive disorder and bipolar disorder in the Han Chinese population. <i>Journal of Affective Disorders</i> , <b>2016</b> , 194, 180-7	6.6	7
50	The CMYA5 gene confers risk for both schizophrenia and major depressive disorder in the Han Chinese population. <i>World Journal of Biological Psychiatry</i> , <b>2014</b> , 15, 553-60	3.8	7
49	Association analysis of the GRM8 gene with schizophrenia in the Uygur Chinese population. <i>Hereditas</i> , <b>2014</b> , 151, 140-4	2.4	7
48	Association of SCN10A Polymorphisms with the Recurrence of Atrial Fibrillation after Catheter Ablation in a Chinese Han Population. <i>Scientific Reports</i> , <b>2017</b> , 7, 44003	4.9	6
47	The pathogenicity of duck hepatitis A virus types 1 and 3 on ducklings. <i>Poultry Science</i> , <b>2019</b> , 98, 6333-6339	3.9	6
46	Genetic analysis of common variants in the ZNF804A gene with schizophrenia and major depressive disorder. <i>Psychiatric Genetics</i> , <b>2018</b> , 28, 1-7	2.9	6
45	Association between the variability of the ABCA13 gene and the risk of major depressive disorder and schizophrenia in the Han Chinese population. <i>World Journal of Biological Psychiatry</i> , <b>2017</b> , 18, 550-556	3.8	5
44	Meta-analysis of GABRB2 polymorphisms and the risk of schizophrenia combined with GWAS data of the Han Chinese population and psychiatric genomics consortium. <i>PLoS ONE</i> , <b>2018</b> , 13, e0198690	3.7	5
43	The YWHAE gene confers risk to major depressive disorder in the male group of Chinese Han population. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , <b>2017</b> , 77, 172-177	5.5	4
42	Common variants in ZMIZ1 and near NGF confer risk for primary dysmenorrhoea. <i>Nature Communications</i> , <b>2017</b> , 8, 14900	17.4	4

41	Common variants in the gene are associated with serum uric acid level and hyperuricemia and gout in Han Chinese. <i>Hereditas</i> , <b>2019</b> , 156, 4	2.4	4
40	A novel variant associated with HDL-C levels by modifying DAGLB expression levels: An annotation-based genome-wide association study. <i>European Journal of Human Genetics</i> , <b>2018</b> , 26, 838-847	5.3	4
39	ACTN3 is associated with children's physical fitness in Han Chinese. <i>Molecular Genetics and Genomics</i> , <b>2019</b> , 294, 47-56	3.1	4
38	The Potential Effect of Aberrant Testosterone Levels on Common Diseases: A Mendelian Randomization Study. <i>Genes</i> , <b>2020</b> , 11,	4.2	3
37	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. <i>Endocrine Journal</i> , <b>2018</b> , 65, 783-791	2.9	3
36	eRFSVM: a hybrid classifier to predict enhancers-integrating random forests with support vector machines. <i>Hereditas</i> , <b>2016</b> , 153, 6	2.4	3
35	Association study of NDST3 gene for schizophrenia, bipolar disorder, major depressive disorder in the Han Chinese population. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2018</b> , 177, 3-9	3.5	3
34	SLC39A8 is a risk factor for schizophrenia in Uygur Chinese: a case-control study. <i>BMC Psychiatry</i> , <b>2019</b> , 19, 293	4.2	2
33	Polymorphism of the Gene and Dynamic Balance Performance in Han Chinese Children. <i>Hereditas</i> , <b>2019</b> , 156, 15	2.4	2
32	Common variants in FAN1, located in 15q13.3, confer risk for schizophrenia and bipolar disorder in Han Chinese. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , <b>2020</b> , 103, 109973	5.5	2
31	Functional annotation of regulatory single nucleotide polymorphisms associated with schizophrenia. <i>Schizophrenia Research</i> , <b>2020</b> , 218, 326-328	3.6	2
30	Role played by the SP4 gene in schizophrenia and major depressive disorder in the Han Chinese population. <i>British Journal of Psychiatry</i> , <b>2016</b> , 208, 441-5	5.4	2
29	Analysis of association between common variants in the SLC6A1 gene with schizophrenia, bipolar disorder and major depressive disorder in the Han Chinese population. <i>World Journal of Biological Psychiatry</i> , <b>2016</b> , 17, 140-6	3.8	2
28	Common variants in the CDH7 gene are associated with major depressive disorder in the Han Chinese population. <i>Behavior Genetics</i> , <b>2014</b> , 44, 97-101	3.2	2
27	Structural Comparison and Drug Screening of Spike Proteins of Ten SARS-CoV-2 Variants.. <i>Research</i> , <b>2022</b> , 2022, 9781758	7.8	2
26	Genomic dissection of 43 serum urate-associated loci provides multiple insights into molecular mechanisms of urate control		2
25	Four Loci Are Associated with Cardiorespiratory Fitness and Endurance Performance in Young Chinese Females. <i>Scientific Reports</i> , <b>2020</b> , 10, 10117	4.9	2
24	Elevated levels of IL-18 associated with schizophrenia and first episode psychosis: A systematic review and meta-analysis. <i>Microbial Biotechnology</i> , <b>2021</b> , 15, 896-905	3.3	2

23	Common variants in QPCT gene confer risk of schizophrenia in the Han Chinese population. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2016</b> , 171B, 237-42	3.5	2
22	VariFAST: a variant filter by automated scoring based on tagged-signatures. <i>BMC Bioinformatics</i> , <b>2019</b> , 20, 713	3.6	2
21	The association between rs12807809 polymorphism in neurogranin gene and risk of schizophrenia: A meta-analysis. <i>Medicine (United States)</i> , <b>2019</b> , 98, e18518	1.8	2
20	Identification of SHANK2 Pathogenic Variants in a Chinese Uygur Population with Schizophrenia. <i>Journal of Molecular Neuroscience</i> , <b>2021</b> , 71, 1-8	3.3	2
19	Amplicon targeted resequencing for SLC2A9 and SLC22A12 identified novel mutations in hypouricemia subjects. <i>Molecular Genetics &amp; Genomic Medicine</i> , <b>2019</b> , 7, e00722	2.3	1
18	SHesisPCA: a GPU-based software to correct for population stratification that efficiently accelerates the process for handling genome-wide datasets. <i>Journal of Genetics and Genomics</i> , <b>2015</b> , 42, 445-53	4	1
17	Identification of rare and common variants in BNIP3L: a schizophrenia susceptibility gene. <i>Human Genomics</i> , <b>2020</b> , 14, 16	6.8	1
16	Fine-mapping of ZDHHC2 identifies risk variants for schizophrenia in the Han Chinese population. <i>Molecular Genetics &amp; Genomic Medicine</i> , <b>2020</b> , 8, e1190	2.3	1
15	Noninvasive fetal trisomy detection by multiplexed semiconductor sequencing: a barcoding analysis strategy. <i>Journal of Human Genetics</i> , <b>2016</b> , 61, 247-52	4.3	1
14	An evaluation of association between a novel hippocampal biology related SNP (rs7294919) and schizophrenia. <i>PLoS ONE</i> , <b>2013</b> , 8, e80696	3.7	1
13	MicroRNA Microarray Analysis Combined with Interaction Network Analysis to Investigate the Influence of Clozapine to Metabolic Syndrome. <i>International Journal of Pharmacology</i> , <b>2013</b> , 9, 366-372	0.7	1
12	Genetic risk of clozapine-induced leukopenia and neutropenia: a genome-wide association study. <i>Translational Psychiatry</i> , <b>2021</b> , 11, 343	8.6	1
11	Disruption of Gene Function Increases the Risk of Doxorubicin-Induced Cardiomyopathy and Heart Failure. <i>BioMed Research International</i> , <b>2021</b> , 2021, 8569921	3	1
10	Common variants in SATB2 are associated with schizophrenia in Uygur Chinese population. <i>Psychiatric Genetics</i> , <b>2019</b> , 29, 120-126	2.9	1
9	Cigarette smoking and schizophrenia: Mendelian randomisation study. <i>British Journal of Psychiatry</i> , <b>2021</b> , 218, 98-103	5.4	1
8	Trans-ancestral dissection of urate- and gout-associated major loci SLC2A9 and ABCG2 reveals primate-specific regulatory effects. <i>Journal of Human Genetics</i> , <b>2021</b> , 66, 161-169	4.3	1
7	SNX29, a new susceptibility gene shared with major mental disorders in Han Chinese population. <i>World Journal of Biological Psychiatry</i> , <b>2021</b> , 22, 526-534	3.8	0
6	Rare and common variants analysis of the EMB gene in patients with schizophrenia. <i>BMC Psychiatry</i> , <b>2020</b> , 20, 135	4.2	0

5	Polymorphisms and rare variants identified by next-generation sequencing confer risk for lung cancer in han Chinese population. <i>Pathology Research and Practice</i> , <b>2020</b> , 216, 152873	3.4	○
4	Association Analysis Between Common Variants of the Gene and Three Mental Disorders in the Han Chinese Population. <i>Genetic Testing and Molecular Biomarkers</i> , <b>2020</b> , 24, 649-657	1.6	○
3	A machine learning-assisted model for renal urate underexcretion with genetic and clinical variables among Chinese men with gout.. <i>Arthritis Research and Therapy</i> , <b>2022</b> , 24, 67	5.7	○
2	Prediction of functional regulatory elements of bipolar disorder via data integration analysis. <i>Journal of Affective Disorders Reports</i> , <b>2020</b> , 1, 100015	1.4	
1	Association analysis of potentially functional variants within 8p12 with schizophrenia in the Han Chinese population. <i>World Journal of Biological Psychiatry</i> , <b>2021</b> , 22, 27-33	3.8	