Yong-Yong Shi

List of Publications by Year
in descending order

150 papers	13,599 citations	66250 44 h-index	26792 111 g-index
157 all docs	157 docs citations	157 times ranked	22935 citing authors

#	Article	IF	Citations
1	Structural Comparison and Drug Screening of Spike Proteins of Ten SARS-CoV-2 Variants. Research, 2022, 2022, 9781758.	2.8	15
2	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
3	The Relationship between Alcohol Consumption and Gout: A Mendelian Randomization Study. Genes, 2022, 13, 557.	1.0	5
4	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. Nature, 2022, 604, 502-508.	13.7	929
5	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
6	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
7	Cigarette smoking and schizophrenia: Mendelian randomisation study. British Journal of Psychiatry, 2021, 218, 98-103.	1.7	6
8	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
9	Identification of SHANK2 Pathogenic Variants in a Chinese Uygur Population with Schizophrenia. Journal of Molecular Neuroscience, 2021, 71, 1-8.	1.1	6
10	Genetic risk of clozapine-induced leukopenia and neutropenia: a genome-wide association study. Translational Psychiatry, 2021, 11, 343.	2.4	5
11	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
12	Scrutinizing the causal relationship between schizophrenia and vitamin supplementation. Journal of Bio-X Research, 2021, Publish Ahead of Print, .	0.3	0
13	Structural Analysis of the SARS-CoV-2 Omicron Variant Proteins. Research, 2021, 2021, 9769586.	2.8	27
14	Prediction of functional regulatory elements of bipolar disorder via data integration analysis. Journal of Affective Disorders Reports, 2020, 1, 100015.	0.9	1
15	Four Loci Are Associated with Cardiorespiratory Fitness and Endurance Performance in Young Chinese Females. Scientific Reports, 2020, 10, 10117.	1.6	7
16	Elevated levels of IL â€18 associated with schizophrenia and first episode psychosis: A systematic review and metaâ€anâ€alysis. Microbial Biotechnology, 2020, 15, 896-905.	0.9	9
17	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
18	Identification of rare and common variants in BNIP3L: a schizophrenia susceptibility gene. Human Genomics, 2020, 14, 16.	1.4	6

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19	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
20	Fineâ€mapping of ZDHHC2 identifies risk variants for schizophrenia in the Han Chinese population. Molecular Genetics & Genomic Medicine, 2020, 8, e1190.	0.6	7
21	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
22	Rare and common variants analysis of the EMB gene in patients with schizophrenia. BMC Psychiatry, 2020, 20, 135.	1.1	4
23	The Potential Effect of Aberrant Testosterone Levels on Common Diseases: A Mendelian Randomization Study. Genes, 2020, 11, 721.	1.0	14
24	Functional annotation of regulatory single nucleotide polymorphisms associated with schizophrenia. Schizophrenia Research, 2020, 218, 326-328.	1.1	4
25	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
26	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
27	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
28	The pathogenicity of duck hepatitis A virus types 1 and 3 on ducklings. Poultry Science, 2019, 98, 6333-6339.	1.5	11
29	The schizophrenia genetics knowledgebase: a comprehensive update of findings from candidate gene studies. Translational Psychiatry, 2019, 9, 205.	2.4	19
30	SLC39A8 is a risk factor for schizophrenia in Uygur Chinese: a case-control study. BMC Psychiatry, 2019, 19, 293.	1.1	7
31	Amplicon targeted resequencing for SLC2A9 and SLC22A12 identified novel mutations in hypouricemia subjects. Molecular Genetics & Genomic Medicine, 2019, 7, e00722.	0.6	4
32	Polymorphism of the PPARD Gene and Dynamic Balance Performance in Han Chinese Children. Hereditas, 2019, 156, 15.	0.5	3
33	Immunosuppressive potential of fowl adenovirus serotype 4. Poultry Science, 2019, 98, 3514-3522.	1.5	24
34	Genome-wide association studies identify susceptibility loci for epithelial ovarian cancer in east Asian women. Gynecologic Oncology, 2019, 153, 343-355.	0.6	28
35	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
36	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

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37	VariFAST: a variant filter by automated scoring based on tagged-signatures. BMC Bioinformatics, 2019, 20, 713.	1.2	3
38	Comparative genetic architectures of schizophrenia in East Asian and European populations. Nature Genetics, 2019, 51, 1670-1678.	9.4	440
39	Common variants in SATB2 are associated with schizophrenia in Uygur Chinese population. Psychiatric Genetics, 2019, 29, 120-126.	0.6	2
40	The association between rs12807809 polymorphism in neurogranin gene and risk of schizophrenia. Medicine (United States), 2019, 98, e18518.	0.4	4
41	ACTN3 is associated with childrenâ€™s physical fitness in Han Chinese. Molecular Genetics and Genomics, 2019, 294, 47-56.	1.0	8
42	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
43	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
44	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
45	Identifying the Genotypes of Hepatitis B Virus (HBV) with DNA Origami Label. Small, 2018, 14, 1701718.	5.2	23
46	Genetic analysis of common variants in the ZNF804A gene with schizophrenia and major depressive disorder. Psychiatric Genetics, 2018, 28, 1-7.	0.6	8
47	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
48	Genome-wide association study identifies two risk loci for tuberculosis in Han Chinese. Nature Communications, 2018, 9, 4072.	5.8	51
49	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
50	Genetic risk between the CACNA1I gene and schizophrenia in Chinese Uygur population. Hereditas, 2018, 155, 5.	0.5	12
51	Glucose and Insulin-Related Traits, Type 2 Diabetes and Risk of Schizophrenia: A Mendelian Randomization Study. EBioMedicine, 2018, 34, 182-188.	2.7	34
52	Identification of recurrent USP48 and BRAF mutations in Cushingâ€™s disease. Nature Communications, 2018, 9, 3171.	5.8	106
53	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
54	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

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55	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
56	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
57	Exome Array Analysis Identifies Variants in SPOCD1 and BTN3A2 That Affect Risk for Gastric Cancer. Gastroenterology, 2017, 152, 2011-2021.	0.6	58
58	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
59	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
60	Common variants in ZMIZ1 and near NGF confer risk for primary dysmenorrhoea. Nature Communications, 2017, 8, 14900.	5.8	9
61	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
62	DNA origami-based shape IDs for single-molecule nanomechanical genotyping. Nature Communications, 2017, 8, 14738.	5.8	73
63	Genome-wide association analysis identifies 30 new susceptibility loci for schizophrenia. Nature Genetics, 2017, 49, 1576-1583.	9.4	395
64	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
65	Replication of Gout/Urate Concentrations GWAS Susceptibility Loci Associated with Gout in a Han Chinese Population. Scientific Reports, 2017, 7, 4094.	1.6	15
66	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
67	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
68	Cell Culture System for Analysis of Genetic Heterogeneity WithinÂHepatocellular Carcinomas and Response to Pharmacologic Agents. Gastroenterology, 2017, 152, 232-242.e4.	0.6	107
69	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
70	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
71	SHEsisPlus, a toolset for genetic studies on polyploid species. Scientific Reports, 2016, 6, 24095.	1.6	77
72	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

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73	The genome-wide mutational landscape of pituitary adenomas. Cell Research, 2016, 26, 1255-1259.	5.7	137
74	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
75	eRFSVM: a hybrid classifier to predict enhancers-integrating random forests with support vector machines. Hereditas, 2016, 153, 6.	0.5	5
76	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
77	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
78	Genome-wide Analysis of the Role of Copy Number Variation in Schizophrenia Risk in Chinese. Biological Psychiatry, 2016, 80, 331-337.	0.7	55
79	Noninvasive fetal trisomy detection by multiplexed semiconductor sequencing: a barcoding analysis strategy. Journal of Human Genetics, 2016, 61, 247-252.	1.1	1
80	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
81	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
82	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
83	Psychiatric genetics in China: achievements and challenges. Molecular Psychiatry, 2016, 21, 4-9.	4.1	6
84	A modifier screen identifies DNAJB6 as a cardiomyopathy susceptibility gene. JCI Insight, 2016, 1, .	2.3	42
85	Polymorphisms of Renin-Angiotensin-Aldosterone System Gene in Chinese Han Patients with Nonfamilial Atrial Fibrillation. PLoS ONE, 2015, 10, e0117489.	1.1	12
86	Systematic Evaluation of Genetic Variants for Polycystic Ovary Syndrome in a Chinese Population. PLoS ONE, 2015, 10, e0140695.	1.1	17
87	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
88	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
89	Genome-wide association analysis identifies three new risk loci for gout arthritis in Han Chinese. Nature Communications, 2015, 6, 7041.	5.8	88
90	Recurrent gain-of-function USP8 mutations in Cushing's disease. Cell Research, 2015, 25, 306-317.	5.7	263

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91	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
92	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
93	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
94	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
95	Loci with genome-wide associations with schizophrenia in the Han Chinese population. British Journal of Psychiatry, 2015, 207, 490-494.	1.7	29
96	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
97	USP8 mutation in Cushing's disease. Oncotarget, 2015, 6, 18240-18241.	0.8	6
98	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
99	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
100	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
101	Mapping of hepatic expression quantitative trait loci (eQTLs) in a Han Chinese population. Journal of Medical Genetics, 2014, 51, 319-326.	1.5	16
102	Genetic Variants Associated With Phenytoin-Related Severe Cutaneous Adverse Reactions. JAMA - Journal of the American Medical Association, 2014, 312, 525.	3.8	256
103	Association analysis of theGRM8gene with schizophrenia in the Uygur Chinese population. Hereditas, 2014, 151, 140-144.	0.5	9
104	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
105	CACNA1C, schizophrenia and major depressive disorder in the Han Chinese population. British Journal of Psychiatry, 2014, 204, 36-39.	1.7	53
106	A genome-wide gene-environment interaction analysis for tobacco smoke and lung cancer susceptibility. Carcinogenesis, 2014, 35, 1528-1535.	1.3	47
107	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
108	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

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109	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
110	Comparison of the performance of Ion Torrent chips in noninvasive prenatal trisomy detection. Journal of Human Genetics, 2014, 59, 393-396.	1.1	15
111	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
112	Genome-wide association study in Han Chinese identifies three novel loci for human height. Human Genetics, 2013, 132, 681-689.	1.8	21
113	CNTNAP2 is significantly associated with schizophrenia and major depression in the Han Chinese population. Psychiatry Research, 2013, 207, 225-228.	1.7	50
114	New loci associated with chronic hepatitis B virus infection in Han Chinese. Nature Genetics, 2013, 45, 1499-1503.	9.4	140
115	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
116	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
117	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
118	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
119	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
120	An Evaluation of Association between a Novel Hippocampal Biology Related SNP (rs7294919) and Schizophrenia. PLoS ONE, 2013, 8, e80696.	1.1	1
121	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
122	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
123	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
124	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
125	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
126	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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127	ZNF804A and schizophrenia susceptibility in Asian populations. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2012, 159B, 794-802.	1.1	30
128	Genome-wide association study identifies eight new risk loci for polycystic ovary syndrome. Nature Genetics, 2012, 44, 1020-1025.	9.4	505
129	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
130	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
131	Common variants on 8p12 and 1q24.2 confer risk of schizophrenia. Nature Genetics, 2011, 43, 1224-1227.	9.4	224
132	The MDGA1 gene confers risk to schizophrenia and bipolar disorder. Schizophrenia Research, 2011, 125, 194-200.	1.1	35
133	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
134	Gold nanoparticles for high-throughput genotyping of long-range haplotypes. Nature Nanotechnology, 2011, 6, 639-644.	15.6	106
135	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
136	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
137	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
138	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
139	Common Variants in the BCL9 Gene Conferring Risk of Schizophrenia. Archives of General Psychiatry, 2011, 68, 232.	13.8	39
140	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
141	The sequence and de novo assembly of the giant panda genome. Nature, 2010, 463, 311-317.	13.7	1,058
142	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
143	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
144	Apoptotic Engulfment Pathway and Schizophrenia. PLoS ONE, 2009, 4, e6875.	1.1	35

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145	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
146	Identification of loci associated with schizophrenia by genome-wide association and follow-up. Nature Genetics, 2008, 40, 1053-1055.	9.4	977
147	Systematic polymorphism analysis of the CYP2D6 gene in four different geographical Han populations in mainland China. Genomics, 2008, 92, 152-158.	1.3	59
148	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
149	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
150	Genetic Structure Adds Power to Detect Schizophrenia Susceptibility at SLIT3 in the Chinese Han Population. Genome Research, 2004, 14, 1345-1349.	2.4	36