Fuquan Zhang

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#	Paper	IF	Citations
113	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. <i>Cell</i> , 2019 , 179, 1469-1482.e11	56.2	402
112	MirSNP, a database of polymorphisms altering miRNA target sites, identifies miRNA-related SNPs in GWAS SNPs and eQTLs. <i>BMC Genomics</i> , 2012 , 13, 661	4.5	207
111	Depression in college: depressive symptoms and personality factors in Beijing and Hong Kong college freshmen. <i>Comprehensive Psychiatry</i> , 2008 , 49, 496-502	7.3	92
110	MicroRNAs and target site screening reveals a pre-microRNA-30e variant associated with schizophrenia. <i>Schizophrenia Research</i> , 2010 , 119, 219-27	3.6	76
109	Five novel loci associated with antipsychotic treatment response in patients with schizophrenia: a genome-wide association study. <i>Lancet Psychiatry,the</i> , 2018 , 5, 327-338	23.3	66
108	Multi-trait analysis for genome-wide association study of five psychiatric disorders. <i>Translational Psychiatry</i> , 2020 , 10, 209	8.6	59
107	Genetic regulatory subnetworks and key regulating genes in rat hippocampus perturbed by prenatal malnutrition: implications for major brain disorders. <i>Aging</i> , 2020 , 12, 8434-8458	5.6	56
106	Implications of Newly Identified Brain eQTL Genes and Their Interactors in Schizophrenia. <i>Molecular Therapy - Nucleic Acids</i> , 2018 , 12, 433-442	10.7	52
105	Co-expression network analysis identified hub genes critical to triglyceride and free fatty acid metabolism as key regulators of age-related vascular dysfunction in mice. <i>Aging</i> , 2019 , 11, 7620-7638	5.6	48
104	Abnormal functional connectivity of brain network hubs associated with symptom severity in treatment-naive patients with obsessive-compulsive disorder: A resting-state functional MRI study. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2016 , 66, 104-111	5.5	43
103	The early growth response protein 1-miR-30a-5p-neurogenic differentiation factor 1 axis as a novel biomarker for schizophrenia diagnosis and treatment monitoring. <i>Translational Psychiatry</i> , 2017 , 7, e998	3 ^{8.6}	37
102	Diagnostic value of blood-derived microRNAs for schizophrenia: results of a meta-analysis and validation. <i>Scientific Reports</i> , 2017 , 7, 15328	4.9	35
101	An association study of DRD2 gene polymorphisms with schizophrenia in a Chinese Han population. <i>Neuroscience Letters</i> , 2010 , 477, 53-6	3.3	35
100	Genetic evidence suggests posttraumatic stress disorder as a subtype of major depressive disorder. Journal of Clinical Investigation, 2021,	15.9	35
99	Causal influences of neuroticism on mental health and cardiovascular disease. <i>Human Genetics</i> , 2021 , 140, 1267-1281	6.3	30
98	Exploring Transcription Factors-microRNAs Co-regulation Networks in Schizophrenia. <i>Schizophrenia Bulletin</i> , 2016 , 42, 1037-45	1.3	29
97	Common variants in CACNA1C and MDD susceptibility: A comprehensive meta-analysis. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2016 , 171, 896-903	3.5	27

(2019-2015)

96	Converging evidence implicates the abnormal microRNA system in schizophrenia. <i>Schizophrenia Bulletin</i> , 2015 , 41, 728-35	1.3	26	
95	Abnormal regional spontaneous neuronal activity associated with symptom severity in treatment-naive patients with obsessive-compulsive disorder revealed by resting-state functional MRI. <i>Neuroscience Letters</i> , 2017 , 640, 99-104	3.3	23	
94	Efficacy of Acupuncture Therapy for Chemotherapy-Related Cognitive Impairment in Breast Cancer Patients. <i>Medical Science Monitor</i> , 2018 , 24, 2919-2927	3.2	22	
93	Replication study confirms link between TSPAN18 mutation and schizophrenia in Han Chinese. <i>PLoS ONE</i> , 2013 , 8, e58785	3.7	20	
92	A Schizophrenia-Related Genetic-Brain-Cognition Pathway Revealed in a Large Chinese Population. <i>EBioMedicine</i> , 2018 , 37, 471-482	8.8	20	
91	Peripheral blood nerve growth factor levels in major psychiatric disorders. <i>Journal of Psychiatric Research</i> , 2017 , 86, 39-45	5.2	19	
90	Altered expression of mRNA profiles in blood of early-onset schizophrenia. <i>Scientific Reports</i> , 2016 , 6, 16767	4.9	19	
89	An updated meta-analysis of the association between SORL1 variants and the risk for sporadic Alzheimer disease. <i>Journal of Alzheimer Disease</i> , 2013 , 37, 429-37	4.3	19	
88	Effect of Damaging Rare Mutations in Synapse-Related Gene Sets on Response to Short-term Antipsychotic Medication in Chinese Patients With Schizophrenia: A Randomized Clinical Trial. <i>JAMA Psychiatry</i> , 2018 , 75, 1261-1269	14.5	19	
87	Lack of association between microRNA-137 SNP rs1625579 and schizophrenia in a replication study of Han Chinese. <i>Molecular Genetics and Genomics</i> , 2015 , 290, 297-301	3.1	17	
86	Converging evidence implicates the dopamine D3 receptor gene in vulnerability to schizophrenia. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2011 , 156B, 613-9	3.5	17	
85	Identifying common genome-wide risk genes for major psychiatric traits. <i>Human Genetics</i> , 2020 , 139, 185-198	6.3	17	
84	Rumination mediates the relationship between overgeneral autobiographical memory and depression in patients with major depressive disorder. <i>BMC Psychiatry</i> , 2017 , 17, 103	4.2	16	
83	Overgeneral autobiographical memory at baseline predicts depressive symptoms at follow-up in patients with first-episode depression. <i>Psychiatry Research</i> , 2016 , 243, 123-7	9.9	15	
82	The SORL1 polymorphism rs985421 may confer the risk for amnestic mild cognitive impairment and Alzheimer disease in the Han Chinese population. <i>Neuroscience Letters</i> , 2014 , 563, 80-4	3.3	15	
81	A competitive PCR assay confirms the association of a copy number variation in the VIPR2 gene with schizophrenia in Han Chinese. <i>Schizophrenia Research</i> , 2014 , 156, 66-70	3.6	14	
80	Blood-Derived Plasma Protein Biomarkers for Alzheimer\\Disease in Han Chinese. <i>Frontiers in Aging Neuroscience</i> , 2018 , 10, 414	5.3	14	
79	Altered expression of the CSMD1 gene in the peripheral blood of schizophrenia patients. <i>BMC Psychiatry</i> , 2019 , 19, 113	4.2	13	

78	Increased Variability of Genomic Transcription in Schizophrenia. Scientific Reports, 2015, 5, 17995	4.9	13
77	The norepinephrine transporter gene modulates the relationship between urban/rural residency and major depressive disorder in a Chinese population. <i>Psychiatry Research</i> , 2009 , 168, 213-7	9.9	13
76	Metabolic Effects of 7 Antipsychotics on Patients With Schizophrenia: A Short-Term, Randomized, Open-Label, Multicenter, Pharmacologic Trial. <i>Journal of Clinical Psychiatry</i> , 2020 , 81,	4.6	13
75	The Shared and Distinct White Matter Networks Between Drug-Naive Patients With Obsessive-Compulsive Disorder and Schizophrenia. <i>Frontiers in Neuroscience</i> , 2019 , 13, 96	5.1	12
74	Potential involvement of the interleukin-18 pathway in schizophrenia. <i>Journal of Psychiatric Research</i> , 2016 , 74, 10-6	5.2	12
73	Lack of association between MPC2 variants and schizophrenia in a replication study of Han Chinese. <i>Neuroscience Letters</i> , 2013 , 552, 120-3	3.3	12
72	Genetic association of rs1344706 in ZNF804A with bipolar disorder and schizophrenia susceptibility in Chinese populations. <i>Scientific Reports</i> , 2017 , 7, 41140	4.9	11
71	Regional homogeneity associated with overgeneral autobiographical memory of first-episode treatment-naive patients with major depressive disorder in the orbitofrontal cortex: A resting-state fMRI study. <i>Journal of Affective Disorders</i> , 2017 , 209, 163-168	6.6	11
70	Association analyses of the interaction between the ADSS and ATM genes with schizophrenia in a Chinese population. <i>BMC Medical Genetics</i> , 2008 , 9, 119	2.1	11
69	The schizophrenia genetics knowledgebase: a comprehensive update of findings from candidate gene studies. <i>Translational Psychiatry</i> , 2019 , 9, 205	8.6	10
68	No association between ZNF804A rs1344706 and schizophrenia in a case-control study of Han Chinese. <i>Neuroscience Letters</i> , 2016 , 618, 14-18	3.3	10
67	Systematic association analysis of microRNA machinery genes with schizophrenia informs further study. <i>Neuroscience Letters</i> , 2012 , 520, 47-50	3.3	10
66	Further evidence supporting the association of NKAPL with schizophrenia. <i>Neuroscience Letters</i> , 2015 , 605, 49-52	3.3	9
65	No association between FOXP2 rs10447760 and schizophrenia in a replication study of the Chinese Han population. <i>Psychiatric Genetics</i> , 2018 , 28, 19-23	2.9	9
64	Meta-analysis of the association of brain-derived neurotrophic factor Val66Met polymorphism with obsessive-compulsive disorder. <i>Acta Neuropsychiatrica</i> , 2015 , 27, 327-35	3.9	9
63	Myosin Vb gene is associated with schizophrenia in Chinese Han population. <i>Psychiatry Research</i> , 2013 , 207, 13-8	9.9	9
62	Gene expression profiling in peripheral blood mononuclear cells of early-onset schizophrenia. <i>Genomics Data</i> , 2015 , 5, 169-70		9
61	A two-stage association study suggests BRAP as a susceptibility gene for schizophrenia. <i>PLoS ONE</i> , 2014 , 9, e86037	3.7	9

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60	Integrative analysis of shared genetic pathogenesis by autism spectrum disorder and obsessive-compulsive disorder. <i>Bioscience Reports</i> , 2019 , 39,	4.1	9	
59	Changes in the level of Long Non-Coding RNA Gomafu gene expression in schizophrenia patients before and after antipsychotic medication. <i>Schizophrenia Research</i> , 2018 , 195, 318-319	3.6	8	
58	Association analysis of a functional variant in ATXN2 with schizophrenia. <i>Neuroscience Letters</i> , 2014 , 562, 24-7	3.3	8	
57	Current antipsychotic agent use and risk of venous thromboembolism and pulmonary embolism: a systematic review and meta-analysis of observational studies. <i>Therapeutic Advances in Psychopharmacology</i> , 2021 , 11, 2045125320982720	4.9	8	
56	No association of catechol-O-methyltransferase polymorphisms with schizophrenia in the Han Chinese population. <i>Genetic Testing and Molecular Biomarkers</i> , 2012 , 16, 1138-41	1.6	7	
55	Testing the role of genetic variation of the MC4R gene in Chinese population in antipsychotic-induced metabolic disturbance. <i>Science China Life Sciences</i> , 2019 , 62, 535-543	8.5	6	
54	New findings support the association of DISC1 genetic variants with susceptibility to schizophrenia in the Han Chinese population. <i>Psychiatry Research</i> , 2015 , 228, 966-8	9.9	6	
53	Association study of DISC1 genetic variants with the risk of schizophrenia. <i>Psychiatric Genetics</i> , 2016 , 26, 132-5	2.9	6	
52	Increased Insular Cortical Thickness Associated With Symptom Severity in Male Youths With Internet Gaming Disorder: A Surface-Based Morphometric Study. <i>Frontiers in Psychiatry</i> , 2018 , 9, 99	5	6	
51	An Integrative Computational Approach to Evaluate Genetic Markers for Bipolar Disorder. <i>Scientific Reports</i> , 2017 , 7, 6745	4.9	6	
50	rTMS modulates precuneus-hippocampal subregion circuit in patients with subjective cognitive decline. <i>Aging</i> , 2020 , 13, 1314-1331	5.6	6	
49	Genetic mechanisms of COVID-19 and its association with smoking and alcohol consumption. <i>Briefings in Bioinformatics</i> , 2021 , 22,	13.4	6	
48	No association between the rs10503253 polymorphism in the CSMD1 gene and schizophrenia in a Han Chinese population. <i>BMC Psychiatry</i> , 2016 , 16, 206	4.2	6	
47	Brain function, structure and genomic data are linked but show different sensitivity to duration of illness and disease stage in schizophrenia. <i>NeuroImage: Clinical</i> , 2019 , 23, 101887	5.3	5	
46	Genome-wide mRNA expression analysis of peripheral blood from patients with obsessive-compulsive disorder. <i>Scientific Reports</i> , 2018 , 8, 12583	4.9	5	
45	Mapsnp: an R package to plot a genomic map for single nucleotide polymorphisms. <i>PLoS ONE</i> , 2015 , 10, e0123609	3.7	5	
44	GAB2 polymorphism rs2373115 confers susceptibility to sporadic Alzheimer disease. <i>Neuroscience Letters</i> , 2013 , 556, 216-20	3.3	4	
43	A replication study of schizophrenia-related rare copy number variations in a Han Southern Chinese population. <i>Hereditas</i> , 2017 , 154, 2	2.4	4	

42	Impairments in Negative Facial Emotion Recognition in Chinese Schizophrenia Patients Detected With a Newly Designed Task. <i>Journal of Nervous and Mental Disease</i> , 2015 , 203, 718-24	1.8	4
41	Analysis of the association of VIPR2 polymorphisms with susceptibility to schizophrenia. <i>Psychiatry Research</i> , 2016 , 241, 104-7	9.9	4
40	Efficacy of anticonvulsant ethosuximide for major depressive disorder: a randomized, placebo-control clinical trial. <i>European Archives of Psychiatry and Clinical Neuroscience</i> , 2021 , 271, 487-49.	9 5 .1	4
39	Altered Frequency-Dependent Brain Activation and White Matter Integrity Associated With Cognition in Characterizing Preclinical Alzheimer Widisease Stages. Frontiers in Human Neuroscience, 2021, 15, 625232	3.3	4
38	Unraveling Risk Genes of COVID-19 by Multi-Omics Integrative Analyses. <i>Frontiers in Medicine</i> , 2021 , 8, 738687	4.9	4
37	miRNA-Coordinated Schizophrenia Risk Network Cross-Talk With Cardiovascular Repair and Opposed Gliomagenesis. <i>Frontiers in Genetics</i> , 2020 , 11, 149	4.5	3
36	Shared Genetic Liability and Causal Associations Between Major Depressive Disorder and Cardiovascular Diseases. <i>Frontiers in Cardiovascular Medicine</i> , 2021 , 8, 735136	5.4	3
35	BDNF Gene W Role in Schizophrenia: From Risk Allele to Methylation Implications. <i>Frontiers in Psychiatry</i> , 2020 , 11, 564277	5	3
34	Smoking quantitatively increases risk for COVID-19. European Respiratory Journal, 2021,	13.6	3
33	Association between MKL1 rs6001946 and schizophrenia in a Han Chinese population. <i>Neuroscience Letters</i> , 2016 , 631, 36-39	3.3	3
32	Effect of the gene on regional cortical grey matter density in the Han Chinese population. <i>British Journal of Psychiatry</i> , 2020 , 216, 254-258	5.4	3
31	Causal links between major depressive disorder and insomnia: A Mendelian randomisation study. <i>Gene</i> , 2021 , 768, 145271	3.8	3
30	Comprehensive literature data-mining analysis reveals a broad genetic network functionally associated with autism spectrum disorder. <i>International Journal of Molecular Medicine</i> , 2018 , 42, 2353-2	36 2	3
29	Classifying major mental disorders genetically. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2022 , 112, 110410	5.5	3
28	Association of the angiotensin-converting enzyme gene insertion/deletion polymorphism with schizophrenia: a meta-analysis. <i>Psychiatry Research</i> , 2014 , 220, 1169-71	9.9	2
27	. Tsinghua Science and Technology, 2009 , 14, 534-540	3.4	2
26	Shared genetic liability and causal effects between major depressive disorder and insomnia. <i>Human Molecular Genetics</i> , 2021 ,	5.6	2
25	Gene Set and Pathway Enrichment Analyses Highlight Involvement of Ion Transport in Cholinergic Pathways in Autism: Rationale for Nutritional Intervention. <i>Frontiers in Neuroscience</i> , 2021 , 15, 648410	5.1	2

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24	Transcriptome-Wide Identification of G-to-A RNA Editing in Chronic Social Defeat Stress Mouse Models. <i>Frontiers in Genetics</i> , 2021 , 12, 680548	4.5	2
23	Exploring different impaired speed of genetic-related brain function and structures in schizophrenic progress using multimodal analysis. Annual International Conference of the IEEE Engineering in Medicine and Biology Society IEEE Engineering in Medicine and Biology Society Annual	0.9	2
22	Shared Genetic Liability Between Major Depressive Disorder and Atopic Diseases. <i>Frontiers in Immunology</i> , 2021 , 12, 665160	8.4	2
21	Altered Functional Connectivity of the Nucleus Accumbens Network Between Deficit and Non-deficit Schizophrenia. <i>Frontiers in Psychiatry</i> , 2021 , 12, 704631	5	2
20	Interaction Between Variations in Dopamine D2 and Serotonin 2A Receptor is Associated with Short-Term Response to Antipsychotics in Schizophrenia. <i>Neuroscience Bulletin</i> , 2019 , 35, 1102-1105	4.3	1
19	Exploring the mRNA expression level of RELN in peripheral blood of schizophrenia patients before and after antipsychotic treatment. <i>Hereditas</i> , 2020 , 157, 43	2.4	1
18	plot2groups: an R package to plot scatter points for two groups of values. <i>Source Code for Biology and Medicine</i> , 2014 , 9, 23	1.9	1
17	Involvement of the long intergenic non-coding RNA LINC00461 in schizophrenia <i>BMC Psychiatry</i> , 2022 , 22, 59	4.2	1
16	Longitudinal trajectory analysis of antipsychotic response in patients with schizophrenia: 6-week, randomised, open-label, multicentre clinical trial. <i>BJPsych Open</i> , 2020 , 6, e126	5	1
15	Preeclampsia Drives Molecular Networks to Shift Toward Greater Vulnerability to the Development of Autism Spectrum Disorder. <i>Frontiers in Neurology</i> , 2020 , 11, 590	4.1	1
14	ATP-binding cassette transporter 13 mRNA expression level in schizophrenia patients. <i>Scientific Reports</i> , 2020 , 10, 21498	4.9	1
13	Association study of KIBRA rs17070145 polymorphism with the risk of schizophrenia in the Han Chinese population. <i>Psychiatry Research</i> , 2016 , 239, 331-2	9.9	1
12	Altered Insular Subregional Connectivity Associated With Cognitions for Distinguishing the Spectrum of Pre-clinical Alzheimer Disease. <i>Frontiers in Aging Neuroscience</i> , 2021 , 13, 597455	5.3	1
11	Causal Association and Shared Genetics Between Asthma and COVID-19 <i>Frontiers in Immunology</i> , 2022 , 13, 705379	8.4	1
10	An association study of ADSS gene polymorphisms with schizophrenia. <i>Behavioral and Brain Functions</i> , 2008 , 4, 39	4.1	0
9	Shared genetic liability between major depressive disorder and osteoarthritis <i>Bone and Joint Research</i> , 2022 , 11, 12-22	4.2	O
8	Eye movement characteristics in male patients with deficit and non-deficit schizophrenia and their relationships with psychiatric symptoms and cognitive function. <i>BMC Neuroscience</i> , 2021 , 22, 70	3.2	0
7	Altered expression of the DISC1 gene in peripheral blood of patients with schizophrenia. <i>BMC Medical Genetics</i> , 2020 , 21, 194	2.1	O

6	Protocol for a pharmacogenomic study on individualised antipsychotic drug treatment for patients with schizophrenia. <i>BJPsych Open</i> , 2021 , 7, e121	5	O
5	Convergent lines of evidence supporting involvement of NFKB1 in schizophrenia <i>Psychiatry Research</i> , 2022 , 312, 114588	9.9	O
4	A flexible tool to plot a genomic map for single nucleotide polymorphisms. <i>Source Code for Biology and Medicine</i> , 2016 , 11, 5	1.9	
3	Allelic frequency differences of variants between Caucasians and Asians and their association with major mood disorders. <i>Signal Transduction and Targeted Therapy</i> , 2019 , 4, 39	21	
2	Association analysis between CAMKK2 rs1063843 and patients with schizophrenia in a Han Chinese population. <i>Asian Journal of Psychiatry</i> , 2020 , 52, 102055	6.7	
1	Further evidence for the association between CMYA5 rs7714250 and schizophrenia in a Han Chinese population. <i>Psychiatry Research</i> , 2018 , 270, 1177-1178	9.9	