

# Bing Liu

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8152765/publications.pdf>

Version: 2024-02-01

90  
papers

2,986  
citations

172457

29  
h-index

189892

50  
g-index

91  
all docs

91  
docs citations

91  
times ranked

5381  
citing authors

#	ARTICLE	IF	CITATIONS
1	Four Distinct Subtypes of Alzheimer's Disease Based on Resting-State Connectivity Biomarkers. <i>Biological Psychiatry</i> , 2023, 93, 759-769.	1.3	20
2	Orbitofrontal cortex volume links polygenic risk for smoking with tobacco use in healthy adolescents. <i>Psychological Medicine</i> , 2022, 52, 1175-1182.	4.5	3
3	Dynamic reconfiguration of human brain networks across altered states of consciousness. <i>Behavioural Brain Research</i> , 2022, 419, 113685.	2.2	6
4	Regional Radiomics Similarity Networks Reveal Distinct Subtypes and Abnormality Patterns in Mild Cognitive Impairment. <i>Advanced Science</i> , 2022, 9, e2104538.	11.2	21
5	Predicting Treatment Response in Schizophrenia With Magnetic Resonance Imaging and Polygenic Risk Score. <i>Frontiers in Genetics</i> , 2022, 13, 848205.	2.3	4
6	Consistent brain structural abnormalities and multisite individualised classification of schizophrenia using deep neural networks. <i>British Journal of Psychiatry</i> , 2022, 221, 732-739.	2.8	9
7	A Pathway-Specific Polygenic Risk Score Is Associated with Tau Pathology and Cognitive Decline. <i>Journal of Alzheimer's Disease</i> , 2022, 85, 1745-1754.	2.6	4
8	Uncovering the genetic profiles underlying the intrinsic organization of the human cerebellum. <i>Molecular Psychiatry</i> , 2022, 27, 2619-2634.	7.9	3
9	Structural Brain Atrophy Predict Symptom Severity in Schizophrenia Based on Generalized Additive Models. , 2022, , .		0
10	Structural and functional connectivity abnormalities of the default mode network in patients with Alzheimer's disease and mild cognitive impairment within two independent datasets. <i>Methods</i> , 2022, 205, 29-38.	3.8	14
11	Predicting Conversion to Mild Cognitive Impairment in Cognitively Normal with Incomplete Multi-modal Neuroimages. , 2022, , .		0
12	AI4AD: Artificial intelligence analysis for Alzheimer's disease classification based on a multisite DTI database. <i>Brain Disorders</i> , 2021, 1, 100005.	1.7	14
13	Quantitative Radiomic Features as New Biomarkers for Alzheimer's Disease: An Amyloid PET Study. <i>Cerebral Cortex</i> , 2021, 31, 3950-3961.	2.9	18
14	Multi-template Neuroimaging Feature Selection Using Weight-constrained Low-rank Learning for Alzheimer's Disease Classification. , 2021, , .		1
15	Predicting brain age during typical and atypical development based on structural and functional neuroimaging. <i>Human Brain Mapping</i> , 2021, 42, 5943-5955.	3.6	10
16	Multisite schizophrenia classification by integrating structural magnetic resonance imaging data with polygenic risk score. <i>NeuroImage: Clinical</i> , 2021, 32, 102860.	2.7	8
17	Altered Connection and Diagnosis Utility of White Matter in Alzheimer's Disease: A Multi-site Automated Fiber Quantification Study. , 2021, 2021, 2923-2927.		0
18	<i>MIR137</i> polygenic risk is associated with schizophrenia and affects functional connectivity of the dorsolateral prefrontal cortex. <i>Psychological Medicine</i> , 2020, 50, 1510-1518.	4.5	9

#	ARTICLE	IF	CITATIONS
19	Polygenic effects of schizophrenia on hippocampal grey matter volume and hippocampusâ€‘medial prefrontal cortex functional connectivity. <i>British Journal of Psychiatry</i> , 2020, 216, 267-274.	2.8	30
20	A common variant in OXTR rs53576 impacts topological patterns of brain functional networks. <i>European Child and Adolescent Psychiatry</i> , 2020, 29, 993-1002.	4.7	8
21	<scp>Grabâ€‘AD</scp>: Generalizability and reproducibility of altered brain activity and diagnostic classification in Alzheimer's Disease. <i>Human Brain Mapping</i> , 2020, 41, 3379-3391.	3.6	38
22	Generalizable, Reproducible, and Neuroscientifically Interpretable Imaging Biomarkers for Alzheimer's Disease. <i>Advanced Science</i> , 2020, 7, 2000675.	11.2	53
23	Characterizing white matter connectivity in Alzheimer's disease and mild cognitive impairment: An automated fiber quantification analysis with two independent datasets. <i>Cortex</i> , 2020, 129, 390-405.	2.4	30
24	Association of DTNBP1 With Schizophrenia: Findings From Two Independent Samples of Han Chinese Population. <i>Frontiers in Psychiatry</i> , 2020, 11, 446.	2.6	7
25	A neuroimaging biomarker for striatal dysfunction in schizophrenia. <i>Nature Medicine</i> , 2020, 26, 558-565.	30.7	152
26	Independent and reproducible hippocampal radiomic biomarkers for multisite Alzheimerâ€™s disease: diagnosis, longitudinal progress and biological basis. <i>Science Bulletin</i> , 2020, 65, 1103-1113.	9.0	70
27	Characterizing White Matter Connectivity in Alzheimerâ€™s Disease and Mild Cognitive Impairment: Automated Fiber Quantification. , 2019, , .		2
28	Impact of COMT haplotypes on functional connectivity density and its association with the gene expression of dopamine receptors. <i>Brain Structure and Function</i> , 2019, 224, 2619-2630.	2.3	5
29	Attention-based 3D Convolutional Network for Alzheimerâ€™s Disease Diagnosis and Biomarkers Exploration. , 2019, , .		33
30	ASAF: altered spontaneous activity fingerprinting in Alzheimerâ€™s disease based on multisite fMRI. <i>Science Bulletin</i> , 2019, 64, 998-1010.	9.0	24
31	Characterization of white matter changes along fibers by automated fiber quantification in the early stages of Alzheimer's disease. <i>NeuroImage: Clinical</i> , 2019, 22, 101723.	2.7	37
32	Common and Specific Functional Activity Features in Schizophrenia, Major Depressive Disorder, and Bipolar Disorder. <i>Frontiers in Psychiatry</i> , 2019, 10, 52.	2.6	45
33	Polygenic risk for Alzheimer's disease influences precuneal volume in two independent general populations. <i>Neurobiology of Aging</i> , 2018, 64, 116-122.	3.1	35
34	Multilocus genetic profile in dopaminergic pathway modulates the striatum and working memory. <i>Scientific Reports</i> , 2018, 8, 5372.	3.3	11
35	Left Parietal Functional Connectivity Mediates the Association Between COMT rs4633 and Verbal Intelligence in Healthy Adults. <i>Frontiers in Neuroscience</i> , 2018, 12, 233.	2.8	3
36	Association between cerebral dopamine neurotrophic factor (CDNF) 2 polymorphisms and schizophrenia susceptibility and symptoms in the Han Chinese population. <i>Behavioral and Brain Functions</i> , 2018, 14, 1.	3.3	21

#	ARTICLE	IF	CITATIONS
37	Prefrontal Volume Mediates Effect of <i>COMT</i> Polymorphism on Interference Resolution Capacity in Healthy Male Adults. <i>Cerebral Cortex</i> , 2017, 27, 5211-5221.	2.9	6
38	<i>APOE</i> and <i>KIBRA</i> Interactions on Brain Functional Connectivity in Healthy Young Adults. <i>Cerebral Cortex</i> , 2017, 27, 4797-4805.	2.9	10
39	Modulation of <i>APOE</i> and <i>SORL1</i> genes on hippocampal functional connectivity in healthy young adults. <i>Brain Structure and Function</i> , 2017, 222, 2877-2889.	2.3	16
40	Polygenic risk for five psychiatric disorders and cross-disorder and disorder-specific neural connectivity in two independent populations. <i>NeuroImage: Clinical</i> , 2017, 14, 441-449.	2.7	81
41	Interaction effect between 5-HTTLPR and HTR1A rs6295 polymorphisms on the frontoparietal network. <i>Neuroscience</i> , 2017, 362, 239-247.	2.3	5
42	Disrupted functional connectivity between perirhinal and parahippocampal cortices with hippocampal subfields in patients with mild cognitive impairment and Alzheimer's disease. <i>Oncotarget</i> , 2017, 8, 99112-99124.	1.8	5
43	Impaired Parahippocampus Connectivity in Mild Cognitive Impairment and Alzheimer's Disease. <i>Journal of Alzheimer's Disease</i> , 2016, 49, 1051-1064.	2.6	50
44	Interaction of <i>COMT</i> rs4680 and <i>BDNF</i> rs6265 polymorphisms on functional connectivity density of the left frontal eye field in healthy young adults. <i>Human Brain Mapping</i> , 2016, 37, 2468-2478.	3.6	10
45	Polygenic Risk for Schizophrenia Influences Cortical Gyrification in 2 Independent General Populations. <i>Schizophrenia Bulletin</i> , 2016, 43, sbw051.	4.3	40
46	The effects of a genome-wide supported variant in the <i>CACNA1C</i> gene on cortical morphology in schizophrenia patients and healthy subjects. <i>Scientific Reports</i> , 2016, 6, 34298.	3.3	4
47	Impact of <i>PICALM</i> and <i>CLU</i> on hippocampal degeneration. <i>Human Brain Mapping</i> , 2016, 37, 2419-2430.	3.6	19
48	Sex-specific mediation effect of the right fusiform face area volume on the association between variants in repeat length of <i>AVPR1A</i> rs3 and altruistic behavior in healthy adults. <i>Human Brain Mapping</i> , 2016, 37, 2700-2709.	3.6	21
49	Interactions of genetic variants reveal inverse modulation patterns of dopamine system on brain gray matter volume and resting-state functional connectivity in healthy young adults. <i>Brain Structure and Function</i> , 2016, 221, 3891-3901.	2.3	16
50	Genetic Effects on Fine-Grained Human Cortical Regionalization. <i>Cerebral Cortex</i> , 2016, 26, 3732-3743.	2.9	8
51	Modulation effect of the <i>SORL1</i> gene on functional connectivity density in healthy young adults. <i>Brain Structure and Function</i> , 2016, 221, 4103-4110.	2.3	2
52	<i>ALDH2</i> Glu504Lys Confers Susceptibility to Schizophrenia and Impacts Hippocampal-Prefrontal Functional Connectivity. <i>Cerebral Cortex</i> , 2016, 27, bhw056.	2.9	9
53	Multiple Effect of <i>APOE</i> Genotype on Clinical and Neuroimaging Biomarkers Across Alzheimer's Disease Spectrum. <i>Molecular Neurobiology</i> , 2016, 53, 4539-4547.	4.0	46
54	Primate-specific miR-603 is implicated in the risk and pathogenesis of Alzheimer's disease. <i>Aging</i> , 2016, 8, 272-290.	3.1	31

#	ARTICLE	IF	CITATIONS
55	<i>RAB2A</i> Polymorphism impacts prefrontal morphology, functional connectivity, and working memory. <i>Human Brain Mapping</i> , 2015, 36, 4372-4382.	3.6	9
56	The Structural Connectivity Pattern of the Default Mode Network and Its Association with Memory and Anxiety. <i>Frontiers in Neuroanatomy</i> , 2015, 9, 152.	1.7	33
57	APOE Effects on Default Mode Network in Chinese Cognitive Normal Elderly: Relationship with Clinical Cognitive Performance. <i>PLoS ONE</i> , 2015, 10, e0133179.	2.5	22
58	The catechol-o-methyltransferase Val158Met polymorphism modulates the intrinsic functional network centrality of the parahippocampal cortex in healthy subjects. <i>Scientific Reports</i> , 2015, 5, 10105.	3.3	10
59	Scalable and Dil-compatible optical clearance of the mammalian brain. <i>Frontiers in Neuroanatomy</i> , 2015, 9, 19.	1.7	154
60	Bridging Integrator 1 (BIN1) Genotype Effects on Working Memory, Hippocampal Volume, and Functional Connectivity in Young Healthy Individuals. <i>Neuropsychopharmacology</i> , 2015, 40, 1794-1803.	5.4	55
61	Compromised small-world efficiency of structural brain networks in schizophrenic patients and their unaffected parents. <i>Neuroscience Bulletin</i> , 2015, 31, 275-287.	2.9	24
62	The cortical surface area of the insula mediates the effect of DBH rs7040170 on novelty seeking. <i>NeuroImage</i> , 2015, 117, 184-190.	4.2	6
63	DISC1 Ser704Cys impacts thalamic-prefrontal connectivity. <i>Brain Structure and Function</i> , 2015, 220, 91-100.	2.3	21
64	Impacts of PICALM and CLU variants associated with Alzheimer's disease on the functional connectivity of the hippocampus in healthy young adults. <i>Brain Structure and Function</i> , 2015, 220, 1463-1475.	2.3	35
65	The Impact of MIR137 on Dorsolateral Prefrontal-Hippocampal Functional Connectivity in Healthy Subjects. <i>Neuropsychopharmacology</i> , 2014, 39, 2153-2160.	5.4	48
66	Neural mechanisms of oxytocin receptor gene mediating anxiety-related temperament. <i>Brain Structure and Function</i> , 2014, 219, 1543-1554.	2.3	64
67	Disrupted thalamo-cortical connectivity in schizophrenia: A morphometric correlation analysis. <i>Schizophrenia Research</i> , 2014, 153, 129-135.	2.0	18
68	Impaired Long Distance Functional Connectivity and Weighted Network Architecture in Alzheimer's Disease. <i>Cerebral Cortex</i> , 2014, 24, 1422-1435.	2.9	202
69	Dosage Effects of BDNF Val66Met Polymorphism on Cortical Surface Area and Functional Connectivity. <i>Journal of Neuroscience</i> , 2014, 34, 2645-2651.	3.6	37
70	A potential ethnic difference in the association between 5-HTTLPR polymorphisms and the brain default mode network. <i>Science Bulletin</i> , 2014, 59, 1355-1361.	1.7	5
71	Authors' response to "Maternal age as a potential explanation of the role of the L allele of the serotonin transporter gene in anxiety and depression in Asians". <i>Neuroscience Bulletin</i> , 2014, 30, 536-537.	2.9	2
72	Brainnetome-wide association studies in schizophrenia: The advances and future. <i>Neuroscience and Biobehavioral Reviews</i> , 2013, 37, 2818-2835.	6.1	25

#	ARTICLE	IF	CITATIONS
73	Variant in OXTR gene and functional connectivity of the hypothalamus in normal subjects. <i>NeuroImage</i> , 2013, 81, 199-204.	4.2	36
74	KIBRA gene variants are associated with synchronization within the default-mode and executive control networks. <i>NeuroImage</i> , 2013, 69, 213-222.	4.2	18
75	The long rather than the short allele of 5-HTTLPR predisposes Han Chinese to anxiety and reduced connectivity between prefrontal cortex and amygdala. <i>Neuroscience Bulletin</i> , 2013, 29, 4-15.	2.9	49
76	Less Efficient Information Transfer in Cys-Allele Carriers of DISC1: A Brain Network Study Based on Diffusion MRI. <i>Cerebral Cortex</i> , 2013, 23, 1715-1723.	2.9	32
77	Functional Connectivity in Healthy Subjects Is Nonlinearly Modulated by the COMT and DRD2 Polymorphisms in a Functional System-Dependent Manner. <i>Journal of Neuroscience</i> , 2013, 33, 17519-17526.	3.6	32
78	Catechol-O-Methyltransferase Val158Met Polymorphism Modulates Gray Matter Volume and Functional Connectivity of the Default Mode Network. <i>PLoS ONE</i> , 2013, 8, e78697.	2.5	22
79	Disrupted Small-World Brain Networks in Moderate Alzheimer's Disease: A Resting-State fMRI Study. <i>PLoS ONE</i> , 2012, 7, e33540.	2.5	192
80	Prefrontal-Related Functional Connectivities within the Default Network Are Modulated by COMT Val158Met in Healthy Young Adults. <i>Journal of Neuroscience</i> , 2010, 30, 64-69.	3.6	88
81	Cortical thickness is associated with different apolipoprotein E genotypes in healthy elderly adults. <i>Neuroscience Letters</i> , 2010, 479, 332-336.	2.1	38
82	Haplotypes of catechol-O-methyltransferase modulate intelligence-related brain white matter integrity. <i>NeuroImage</i> , 2010, 50, 243-249.	4.2	28
83	COMT val158met modulates association between brain white matter architecture and IQ. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2009, 150B, 375-380.	1.7	42
84	Hippocampal volume and asymmetry in mild cognitive impairment and Alzheimer's disease: Meta-analyses of MRI studies. <i>Hippocampus</i> , 2009, 19, 1055-1064.	1.9	390
85	Adenovirus-Mediated Wild-Type p53 Transfer Radiosensitizes H1299 Cells to Subclinical-Dose Carbon-Ion Irradiation Through the Restoration of p53 Function. <i>Cancer Biotherapy and Radiopharmaceuticals</i> , 2009, 24, 57-66.	1.0	4
86	Multimodal Magnetic Resonance Imaging for Brain Disorders: Advances and Perspectives. <i>Brain Imaging and Behavior</i> , 2008, 2, 249-257.	2.1	8
87	Applications of Psychological Principles and Fatigue Feedback-Learning Technique in E-Learning. , 2008, , .		0
88	Systematic benchmarking of microarray data feature extraction and classification. <i>International Journal of Computer Mathematics</i> , 2008, 85, 803-811.	1.8	2
89	A combinational feature selection and ensemble neural network method for classification of gene expression data. <i>BMC Bioinformatics</i> , 2004, 5, 136.	2.6	107
90	Bridging Integrator 1 (BIN1) Genotype Effects on Working Memory, Hippocampal Volume, and Functional Connectivity in Young Healthy Individuals. , 0, .		1