Peng Xie

List of Publications by Year in descending order

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158 papers 6,955 citations

71102 41 h-index 79698 73 g-index

166 all docs

166 docs citations

166 times ranked 8264 citing authors

#	Article	IF	Citations
1	Comparative efficacy and tolerability of antidepressants for major depressive disorder in children and adolescents: a network meta-analysis. Lancet, The, 2016, 388, 881-890.	13.7	513
2	The gut microbiome from patients with schizophrenia modulates the glutamate-glutamine-GABA cycle and schizophrenia-relevant behaviors in mice. Science Advances, 2019, 5, eaau8317.	10.3	446
3	Reduced default mode network functional connectivity in patients with recurrent major depressive disorder. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 9078-9083.	7.1	441
4	Comparative efficacy and acceptability of psychotherapies for depression in children and adolescents: A systematic review and network meta-analysis. World Psychiatry, 2015, 14, 207-222.	10.4	234
5	Medial reward and lateral non-reward orbitofrontal cortex circuits change in opposite directions in depression. Brain, 2016, 139, 3296-3309.	7.6	224
6	Plasma Metabonomics as a Novel Diagnostic Approach for Major Depressive Disorder. Journal of Proteome Research, 2012, 11, 1741-1748.	3.7	204
7	Landscapes of bacterial and metabolic signatures and their interaction in major depressive disorders. Science Advances, 2020, 6, .	10.3	178
8	Gut microbiota from NLRP3-deficient mice ameliorates depressive-like behaviors by regulating astrocyte dysfunction via circHIPK2. Microbiome, 2019, 7, 116.	11.1	169
9	Microbiota Modulate Anxiety-Like Behavior and Endocrine Abnormalities in Hypothalamic-Pituitary-Adrenal Axis. Frontiers in Cellular and Infection Microbiology, 2017, 7, 489.	3.9	160
10	Diagnosis of major depressive disorder based on changes in multiple plasma neurotransmitters: a targeted metabolomics study. Translational Psychiatry, 2018, 8, 130.	4.8	152
11	Integrated Metabolomics and Proteomics Analysis of Hippocampus in a Rat Model of Depression. Neuroscience, 2018, 371, 207-220.	2.3	132
12	The gut microbiome modulates gut–brain axis glycerophospholipid metabolism in a region-specific manner in a nonhuman primate model of depression. Molecular Psychiatry, 2021, 26, 2380-2392.	7.9	102
13	Gut Microbial Signatures Can Discriminate Unipolar from Bipolar Depression. Advanced Science, 2020, 7, 1902862.	11.2	99
14	The identification of metabolic disturbances in the prefrontal cortex of the chronic restraint stress rat model of depression. Behavioural Brain Research, 2016, 305, 148-156.	2.2	97
15	Comparative efficacy and acceptability of electroconvulsive therapy versus repetitive transcranial magnetic stimulation for major depression: A systematic review and multiple-treatments meta-analysis. Behavioural Brain Research, 2017, 320, 30-36.	2.2	91
16	Meta-Analysis of Infectious Agents and Depression. Scientific Reports, 2014, 4, 4530.	3. 3	83
17	Disrupted intrinsic functional brain topology in patients with major depressive disorder. Molecular Psychiatry, 2021, 26, 7363-7371.	7.9	82
18	Functional connectivity of the right inferior frontal gyrus and orbitofrontal cortex in depression. Social Cognitive and Affective Neuroscience, 2020, 15, 75-86.	3.0	81

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19	Functional Connectivity of the Anterior Cingulate Cortex in Depression and in Health. Cerebral Cortex, 2019, 29, 3617-3630.	2.9	79
20	Effects of gut microbiota on the microRNA and mRNA expression in the hippocampus of mice. Behavioural Brain Research, 2017, 322, 34-41.	2.2	77
21	Altered resting-state dynamic functional brain networks in major depressive disorder: Findings from the REST-meta-MDD consortium. Neurolmage: Clinical, 2020, 26, 102163.	2.7	76
22	Reproducibility of functional brain alterations in major depressive disorder: Evidence from a multisite resting-state functional MRI study with 1,434 individuals. NeuroImage, 2019, 189, 700-714.	4.2	72
23	Divergent Urinary Metabolic Phenotypes between Major Depressive Disorder and Bipolar Disorder Identified by a Combined GC–MS and NMR Spectroscopic Metabonomic Approach. Journal of Proteome Research, 2015, 14, 3382-3389.	3.7	71
24	Amino acid metabolic dysfunction revealed in the prefrontal cortex of a rat model of depression. Behavioural Brain Research, 2015, 278, 286-292.	2.2	70
25	A systematic review and meta-analysis of deep brain stimulation in treatment-resistant depression. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2018, 82, 224-232.	4.8	68
26	Combined Application of NMR- and GC-MS-Based Metabonomics Yields a Superior Urinary Biomarker Panel for Bipolar Disorder. Scientific Reports, 2014, 4, 5855.	3.3	65
27	Serotonin-1A receptor alterations in depression: a meta-analysis of molecular imaging studies. BMC Psychiatry, 2016, 16, 319.	2.6	65
28	Hippocampal proteomic changes of susceptibility and resilience to depression or anxiety in a rat model of chronic mild stress. Translational Psychiatry, 2019, 9, 260.	4.8	65
29	Identification of suitable plasma-based reference genes for miRNAome analysis of major depressive disorder. Journal of Affective Disorders, 2014, 163, 133-139.	4.1	61
30	Burnout, psychological morbidity, job stress, and job satisfaction in Chinese neurologists. Neurology, 2017, 88, 1727-1735.	1.1	61
31	Differential urinary metabolites related with the severity of major depressive disorder. Behavioural Brain Research, 2017, 332, 280-287.	2.2	59
32	Metabolomic identification of molecular changes associated with stress resilience in the chronic mild stress rat model of depression. Metabolomics, 2013, 9, 433-443.	3.0	58
33	Perturbed Microbial Ecology in Myasthenia Gravis: Evidence from the Gut Microbiome and Fecal Metabolome. Advanced Science, 2019, 6, 1901441.	11.2	55
34	Bilateral vs. unilateral repetitive transcranial magnetic stimulation in treating major depression: A meta-analysis of randomized controlled trials. Psychiatry Research, 2014, 219, 51-57.	3.3	53
35	Functional connectivity of the human amygdala in health and in depression. Social Cognitive and Affective Neuroscience, 2018, 13, 557-568.	3.0	51
36	Clostridium butyricum miyairi 588 has preventive effects on chronic social defeat stress-induced depressive-like behaviour and modulates microglial activation in mice. Biochemical and Biophysical Research Communications, 2019, 516, 430-436.	2.1	51

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37	Biotypes of major depressive disorder: Neuroimaging evidence from resting-state default mode network patterns. Neurolmage: Clinical, 2020, 28, 102514.	2.7	51
38	Predictive diagnosis of major depression using NMR-based metabolomics and least-squares support vector machine. Clinica Chimica Acta, 2017, 464, 223-227.	1.1	49
39	Reconfiguration of Cortical Networks in MDD Uncovered by Multiscale Community Detection with fMRI. Cerebral Cortex, 2018, 28, 1383-1395.	2.9	49
40	Metabolomics identifies perturbations in amino acid metabolism in the prefrontal cortex of the learned helplessness rat model of depression. Neuroscience, 2017, 343, 1-9.	2.3	48
41	Characterization of gut microbiome in mice model of depression with divergent response to escitalopram treatment. Translational Psychiatry, 2021, 11, 303.	4.8	48
42	Functional Connectivity of the Precuneus in Unmedicated Patients With Depression. Biological Psychiatry: Cognitive Neuroscience and Neuroimaging, 2018, 3, 1040-1049.	1.5	46
43	Quantitative proteomics analysis of the liver reveals immune regulation and lipid metabolism dysregulation in a mouse model of depression. Behavioural Brain Research, 2016, 311, 330-339.	2.2	45
44	Metabolite signature for diagnosing major depressive disorder in peripheral blood mononuclear cells. Journal of Affective Disorders, 2016, 195, 75-81.	4.1	45
45	Hypothalamic Proteomic Analysis Reveals Dysregulation of Glutamate Balance and Energy Metabolism in a Mouse Model of Chronic Mild Stress-Induced Depression. Neurochemical Research, 2016, 41, 2443-2456.	3. 3	44
46	Metabolomic analysis reveals metabolic disturbances in the prefrontal cortex of the lipopolysaccharide-induced mouse model of depression. Behavioural Brain Research, 2016, 308, 115-127.	2.2	43
47	Selective Serotonin Reuptake Inhibitors Versus Tricyclic Antidepressants in Young Patients: A Meta-analysis of Efficacy and Acceptability. Clinical Therapeutics, 2014, 36, 1087-1095.e4.	2.5	42
48	Effective Connectivity in Depression. Biological Psychiatry: Cognitive Neuroscience and Neuroimaging, 2018, 3, 187-197.	1.5	42
49	2,4-Dihydroxypyrimidine is a potential urinary metabolite biomarker for diagnosing bipolar disorder. Molecular BioSystems, 2014, 10, 813.	2.9	41
50	<p>Metabonomics reveals peripheral and central short-chain fatty acid and amino acid dysfunction in a naturally occurring depressive model of macaques</p> . Neuropsychiatric Disease and Treatment, 2019, Volume 15, 1077-1088.	2.2	41
51	Peripheral blood mononuclear cell-based metabolomic profiling of a chronic unpredictable mild stress rat model of depression. Molecular BioSystems, 2014, 10, 2994-3001.	2.9	40
52	Structural Asymmetry of Dorsolateral Prefrontal Cortex Correlates with Depressive Symptoms: Evidence from Healthy Individuals and Patients with Major Depressive Disorder. Neuroscience Bulletin, 2016, 32, 217-226.	2.9	39
53	Proteomic analysis of olfactory bulb suggests CACNA1E as a promoter of CREB signaling in microbiota-induced depression. Journal of Proteomics, 2019, 194, 132-147.	2.4	39
54	Up-regulation of SIRT6 in the hippocampus induced rats with depression-like behavior via the block Akt/GSK3β signaling pathway. Behavioural Brain Research, 2017, 323, 38-46.	2.2	37

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55	Efficacy and acceptability of interpersonal psychotherapy for depression in adolescents: A meta-analysis of randomized controlled trials. Psychiatry Research, 2017, 253, 226-232.	3.3	37
56	Urinary peptidomics identifies potential biomarkers for major depressive disorder. Psychiatry Research, 2014, 217, 25-33.	3.3	36
57	Novel urinary biomarkers for diagnosing bipolar disorder. Metabolomics, 2013, 9, 800-808.	3.0	33
58	Absence of gut microbiota during early life affects anxiolytic Behaviors and monoamine neurotransmitters system in the hippocampal of mice. Journal of the Neurological Sciences, 2019, 400, 160-168.	0.6	33
59	Frequency Dependant Topological Alterations of Intrinsic Functional Connectome in Major Depressive Disorder. Scientific Reports, 2015, 5, 9710.	3.3	32
60	Potential antidepressant and resilience mechanism revealed by metabolomic study on peripheral blood mononuclear cells of stress resilient rats. Behavioural Brain Research, 2017, 320, 12-20.	2.2	30
61	Early brain changes associated with psychotherapy in major depressive disorder revealed by resting-state fMRI: Evidence for the top-down regulation theory. International Journal of Psychophysiology, 2014, 94, 437-444.	1.0	29
62	Comparative efficacy and acceptability of bibliotherapy for depression and anxiety disorders in children and adolescents: a meta-analysis of randomized clinical trials. Neuropsychiatric Disease and Treatment, 2018, Volume 14, 353-365.	2.2	29
63	Quantitative Proteomic Analysis Reveals Molecular Adaptations in the Hippocampal Synaptic Active Zone of Chronic Mild Stress-Unsusceptible Rats. International Journal of Neuropsychopharmacology, 2016, 19, pyv100.	2.1	27
64	Sex-Specific Urinary Biomarkers for Diagnosing Bipolar Disorder. PLoS ONE, 2014, 9, e115221.	2.5	27
65	Age-specific urinary metabolite signatures and functions in patients with major depressive disorder. Aging, 2019, 11, 6626-6637.	3.1	27
66	Association between processed meat and red meat consumption and risk for glioma: A meta-analysis from 14 articles. Nutrition, 2015, 31, 45-50.	2.4	26
67	Quantitative Proteomic Analysis Reveals Synaptic Dysfunction in the Amygdala of Rats Susceptible to Chronic Mild Stress. Neuroscience, 2018, 376, 24-39.	2.3	26
68	Peripheral metabolic abnormalities of lipids and amino acids implicated in increased risk of suicidal behavior in major depressive disorder. Metabolomics, 2013, 9, 688-696.	3.0	25
69	Chronic mild stress leads to aberrant glucose energy metabolism in depressed Macaca fascicularis models. Psychoneuroendocrinology, 2019, 107, 59-69.	2.7	25
70	Age-related changes in microbial composition and function in cynomolgus macaques. Aging, 2019, 11, 12080-12096.	3.1	25
71	GC–MS-based metabolomic study on the antidepressant-like effects of diterpene ginkgolides in mouse hippocampus. Behavioural Brain Research, 2016, 314, 116-124.	2.2	24
72	Absence of gut microbiota affects lipid metabolism in the prefrontal cortex of mice. Neurological Research, 2019, 41, 1104-1112.	1.3	24

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73	iTRAQ-based proteomics suggests LRP6, NPY and NPY2R perturbation in the hippocampus involved in CSDS may induce resilience and susceptibility. Life Sciences, 2018, 211, 102-117.	4.3	23
74	Relationship between burnout and career choice regret among Chinese neurology postgraduates. BMC Medical Education, 2019, 19, 162.	2.4	23
75	CD36 deficiency affects depressive-like behaviors possibly by modifying gut microbiota and the inflammasome pathway in mice. Translational Psychiatry, 2021, 11, 16.	4.8	23
76	Disrupted hemispheric connectivity specialization in patients with major depressive disorder: Evidence from the REST-meta-MDD Project. Journal of Affective Disorders, 2021, 284, 217-228.	4.1	23
77	Borna Disease Virus Infection Perturbs Energy Metabolites and Amino Acids in Cultured Human Oligodendroglia Cells. PLoS ONE, 2012, 7, e44665.	2.5	22
78	2D-gel based proteomics unravels neurogenesis and energetic metabolism dysfunction of the olfactory bulb in CUMS rat model. Behavioural Brain Research, 2016, 313, 302-309.	2,2	22
79	Venlafaxine exerts antidepressant effects possibly by activating MAPK–ERK1/2 and P13K–AKT pathways in the hippocampus. Behavioural Brain Research, 2017, 335, 63-70.	2.2	22
80	Knock-Down of Endogenous Bornavirus-Like Nucleoprotein 1 Inhibits Cell Growth and Induces Apoptosis in Human Oligodendroglia Cells. International Journal of Molecular Sciences, 2016, 17, 435.	4.1	21
81	Major depression accompanied with inflammation and multiple cytokines alterations: Evidences from clinical patients to macaca fascicularis and LPS-induced depressive mice model. Journal of Affective Disorders, 2020, 271, 262-271.	4.1	21
82	Enhanced Detection of Low-Abundance Human Plasma Proteins by Integrating Polyethylene Glycol Fractionation and Immunoaffinity Depletion. PLoS ONE, 2016, 11, e0166306.	2.5	21
83	Real-Time qPCR Identifies Suitable Reference Genes for Borna Disease Virus-Infected Rat Cortical Neurons. International Journal of Molecular Sciences, 2014, 15, 21825-21839.	4.1	20
84	Differential co-expression and regulation analyses reveal different mechanisms underlying major depressive disorder and subsyndromal symptomatic depression. BMC Bioinformatics, 2015, 16, 112.	2.6	20
85	Chronic d-ribose and d-mannose overload induce depressive/anxiety-like behavior and spatial memory impairment in mice. Translational Psychiatry, 2021, 11, 90.	4.8	20
86	Insight into the metabolic mechanism of Diterpene Ginkgolides on antidepressant effects for attenuating behavioural deficits compared with venlafaxine. Scientific Reports, 2017, 7, 9591.	3.3	19
87	Sema3A - mediated modulation of NR1D1 expression may be involved in the regulation of axonal guidance signaling by the microbiota. Life Sciences, 2019, 223, 54-61.	4.3	19
88	Differential Gut Microbiota and Fecal Metabolites Related With the Clinical Subtypes of Myasthenia Gravis. Frontiers in Microbiology, 2020, 11, 564579.	3.5	19
89	<p>Chronic Stress in a Rat Model of Depression Disturbs the Glutamine–Glutamate–GABA Cycle in the Striatum, Hippocampus, and Cerebellum</p> . Neuropsychiatric Disease and Treatment, 2020, Volume 16, 557-570.	2.2	19
90	Elevated host lipid metabolism revealed by iTRAQ-based quantitative proteomic analysis of cerebrospinal fluid of tuberculous meningitis patients. Biochemical and Biophysical Research Communications, 2015, 466, 689-695.	2.1	18

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91	Memory Impairment Induced by Borna Disease Virus 1 Infection is Associated with Reduced H3K9 Acetylation. Cellular Physiology and Biochemistry, 2018, 49, 381-394.	1.6	18
92	Depressive symptoms and quality of life among Chinese medical postgraduates: a national cross-sectional study. Psychology, Health and Medicine, 2019, 24, 1015-1027.	2.4	18
93	The potential for metabolomics in the study and treatment of major depressive disorder and related conditions. Expert Review of Proteomics, 2020, 17, 309-322.	3.0	18
94	Transition and Dynamic Reconfiguration of Whole-Brain Network in Major Depressive Disorder. Molecular Neurobiology, 2020, 57, 4031-4044.	4.0	18
95	Brain structural alterations in MDD patients with gastrointestinal symptoms: Evidence from the REST-meta-MDD project. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2021, 111, 110386.	4.8	18
96	Normal Thoracic Radiographic Appearance of the Cynomolgus Monkey (Macaca fascicularis). PLoS ONE, 2014, 9, e84599.	2.5	17
97	GC–MS-Based Metabonomic Profiling Displayed Differing Effects of Borna Disease Virus Natural Strain Hu-H1 and Laboratory Strain V Infection in Rat Cortical Neurons. International Journal of Molecular Sciences, 2015, 16, 19347-19368.	4.1	17
98	Recombinant tissue plasminogen activator induces long-term anxiety-like behaviors via the ERK1/2-GAD1-GABA cascade in the hippocampus of a rat model. Neuropharmacology, 2018, 128, 119-131.	4.1	17
99	Biogeography of the large intestinal mucosal and luminal microbiome in cynomolgus macaques with depressive-like behavior. Molecular Psychiatry, 2022, 27, 1059-1067.	7.9	17
100	An entorhinal-visual cortical circuit regulates depression-like behaviors. Molecular Psychiatry, 2022, 27, 3807-3820.	7.9	17
101	Human borna disease virus infection impacts host proteome and histone lysine acetylation in human oligodendroglia cells. Virology, 2014, 464-465, 196-205.	2.4	16
102	Proteomic analysis reveals energy metabolic dysfunction and neurogenesis in the prefrontal cortex of a lipopolysaccharide-induced mouse model of depression. Molecular Medicine Reports, 2016, 13, 1813-1820.	2.4	16
103	Chronic mild stress-induced protein dysregulations correlated with susceptibility and resiliency to depression or anxiety revealed by quantitative proteomics of the rat prefrontal cortex. Translational Psychiatry, 2021, 11, 143.	4.8	16
104	1H NMR-Based Metabolic Profiling Reveals the Effects of Fluoxetine on Lipid and Amino Acid Metabolism in Astrocytes. International Journal of Molecular Sciences, 2015, 16, 8490-8504.	4.1	15
105	Behavioral characterization of CD36 knockout mice with SHIRPA primary screen. Behavioural Brain Research, 2016, 299, 90-96.	2.2	15
106	miR-146a promotes Borna disease virus 1 replication through IRAK1/TRAF6/NF-κB signaling pathway. Virus Research, 2019, 271, 197671.	2.2	15
107	Free Language Selection in the Bilingual Brain: An Event-Related fMRI Study. Scientific Reports, 2015, 5, 11704.	3.3	14
108	Changes in gut viral and bacterial species correlate with altered 1,2-diacylglyceride levels and structure in the prefrontal cortex in a depression-like non-human primate model. Translational Psychiatry, 2022, 12, 74.	4.8	14

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109	Persistent human Borna disease virus infection modifies the acetylome of human oligodendroglia cells towards higher energy and transporter levels. Virology, 2015, 485, 58-78.	2.4	13
110	Proteomic analysis of the intestine reveals SNARE-mediated immunoregulatory and amino acid absorption perturbations in a rat model of depression. Life Sciences, 2019, 234, 116778.	4.3	13
111	The 25(OH)D/VDR signaling may play a role in major depression. Biochemical and Biophysical Research Communications, 2020, 523, 405-410.	2.1	13
112	Vascular endothelial growth factor in major depressive disorder, schizophrenia, and bipolar disorder: A network meta-analysis. Psychiatry Research, 2020, 292, 113319.	3.3	13
113	Borna disease virus infection impacts microRNAs associated with nervous system development, cell differentiation, proliferation and apoptosis in the hippocampi of neonatal rats. Molecular Medicine Reports, 2015, 12, 3697-3703.	2.4	12
114	Prolonged chronic social defeat stress promotes less resilience and higher uniformity in depression-like behaviors in adult male mice. Biochemical and Biophysical Research Communications, 2021, 553, 107-113.	2.1	12
115	The C825T Polymorphism of the G-Protein \hat{I}^23 Gene as a Risk Factor for Depression: A Meta-Analysis. PLoS ONE, 2015, 10, e0132274.	2.5	12
116	Differential Gut Microbiota Compositions Related With the Severity of Major Depressive Disorder. Frontiers in Cellular and Infection Microbiology, 0, 12, .	3.9	12
117	Proteomic and network analysis of human serum albuminome by integrated use of quick crosslinking and two-step precipitation. Scientific Reports, 2017, 7, 9856.	3.3	11
118	Effects of chronic stress on intestinal amino acid pathways. Physiology and Behavior, 2019, 204, 199-209.	2.1	11
119	Metabolomic analysis of animal models of depression. Metabolic Brain Disease, 2020, 35, 979-990.	2.9	11
120	Identification and validation of argininosuccinate synthase as a candidate urinary biomarker for major depressive disorder. Clinica Chimica Acta, 2015, 451, 142-148.	1.1	10
121	Brain region-specific metabolite networks regulate antidepressant effects of venlafaxine. RSC Advances, 2017, 7, 46358-46369.	3.6	10
122	Ginkgo biloba extract and its diterpene ginkgolide constituents ameliorate the metabolic disturbances caused by recombinant tissue plasminogen activator in rat prefrontal cortex. Neuropsychiatric Disease and Treatment, 2018, Volume 14, 1755-1772.	2,2	10
123	Pigment epithelium-derived factor alleviates depressive-like behaviors in mice by modulating adult hippocampal synaptic growth and Wnt pathway. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2020, 98, 109792.	4.8	10
124	Neural correlates of causal attribution in negative events of depressed patients: Evidence from an fMRI study. Clinical Neurophysiology, 2015, 126, 1331-1337.	1.5	9
125	Preparation and application of a partially degradable gel in mass spectrometry-based proteomic analysis. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2011, 879, 2957-2962.	2.3	8
126	In-context language control with production tasks in bilinguals: An fMRI study. Brain Research, 2014, 1585, 131-140.	2.2	8

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127	Effective lock-in strategy for proteomic analysis of corona complexes bound to amino-free ligands of gold nanoparticles. Nanoscale, 2018, 10, 12413-12423.	5.6	8
128	The reductions in the subcallosal region cortical volume and surface area in major depressive disorder across the adult life span. Psychological Medicine, 2020, 50, 422-430.	4.5	8
129	Are human Borna disease virus 1 infections zoonotic and fatal?. Lancet Infectious Diseases, The, 2020, 20, 650-651.	9.1	8
130	Overlap of burnout-depression symptoms among Chinese neurology graduate students in a national cross-sectional study. BMC Medical Education, 2021, 21, 83.	2.4	8
131	Glutamate and Lipid Metabolic Perturbation in the Hippocampi of Asymptomatic Borna Disease Virus-Infected Horses. PLoS ONE, 2014, 9, e99752.	2.5	8
132	Impaired robust interhemispheric function integration of depressive brain from RESTâ€metaâ€MDD database in China. Bipolar Disorders, 2022, 24, 400-411.	1.9	8
133	Gut microbiota: a new insight into neurological diseases. Chinese Medical Journal, 2023, 136, 1261-1277.	2.3	8
134	Identification of suitable reference genes for BDV-infected primary rat hippocampal neurons. Molecular Medicine Reports, 2016, 14, 5587-5594.	2.4	7
135	Biochemical effects of venlafaxine on astrocytes as revealed by ¹ H NMR-based metabolic profiling. Molecular BioSystems, 2017, 13, 338-349.	2.9	7
136	Dysfunction of the anterior and intermediate hippocampal functional network in major depressive disorders across the adult lifespan. Biological Psychology, 2021, 165, 108192.	2.2	6
137	Molecular epidemiology of human Borna disease virus 1 infection revisited. Emerging Microbes and Infections, 2022, , 1 -36.	6.5	6
138	Non-targeted Metabolomics Profiling of Plasma Samples From Patients With Major Depressive Disorder. Frontiers in Psychiatry, 2021, 12, 810302.	2.6	5
139	Identification and bioinformatic analysis of dysregulated microRNAs in human oligodendroglial cells infected with borna disease virus. Molecular Medicine Reports, 2016, 14, 4715-4722.	2.4	4
140	Modulatory interactions of resting-state brain functional connectivity in major depressive disorder. Neuropsychiatric Disease and Treatment, 2018, Volume 14, 2461-2472.	2.2	4
141	Changed PGA and POSTN levels in choroid plexus are associated with depressive-like behaviors in mice. Biochemical and Biophysical Research Communications, 2020, 524, 231-235.	2.1	4
142	Dynamic changes occur in the DNA gut virome of female cynomolgus macaques during aging. MicrobiologyOpen, 2021, 10, e1186.	3.0	4
143	Depressed female cynomolgus monkeys (Macaca fascicularis) display a higher second-to-fourth (2D:4D) digit ratio. Zoological Research, 2019, 40, 219-225.	2.1	4
144	A multiple-reaction-monitoring mass spectrometric method for simultaneous quantitative analysis of five plasma apolipoproteins. Science China Chemistry, 2014, 57, 723-731.	8.2	3

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145	iTRAQ-based proteomics implies inflammasome pathway activation in the prefrontal cortex of CSDS mice may influence resilience and susceptibility. Life Sciences, 2020, 262, 118501.	4.3	3
146	The Mutual Influences between Depressed Macaca fascicularis Mothers and Their Infants. PLoS ONE, 2014, 9, e89931.	2.5	3
147	A Non-Invasive Method to Assess Cerebral Perfusion Pressure in Geriatric Patients with Suspected Cerebrovascular Disease. PLoS ONE, 2015, 10, e0120146.	2.5	3
148	Neuroinflammatory transcriptional signatures in the entorhinal cortex based on lipopolysaccharide-induced depression model in mice. Biochemical and Biophysical Research Communications, 2022, 590, 109-116.	2.1	3
149	Altered Metabolism of the Microbiota–Gut–Brain Axis Is Linked With Comorbid Anxiety in Fecal Recipient Mice of Myasthenia Gravis. Frontiers in Microbiology, 2022, 13, 804537.	3.5	3
150	Primary testicular NK/T-cell lymphoma: A study of two cases and review of literature. Chinese Journal of Cancer Research: Official Journal of China Anti-Cancer Association, Beijing Institute for Cancer Research, 2010, 22, 239-244.	2.2	2
151	Different inhibitory effects on the proliferation and apoptosis of human and laboratory Borna disease virus‑infected human neuroblastoma SH‑SY5Y cells inïÂį½vitro. Molecular Medicine Reports, 2017, 17, 925-931.	2.4	2
152	Angioplasty and stenting for severe vertebral artery orifice stenosis: effects on cerebellar function remodeling verified by blood oxygen level-dependent functional magnetic resonance imaging. Neural Regeneration Research, 2014, 9, 2095.	3.0	2
153	A diathesis-stress rat model induced suicide-implicated endophenotypes and prefrontal cortex abnormalities in the PKA and GABA receptor signaling pathways. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2022, 116, 110538.	4.8	2
154	Amplification refractory mutation system polymerase chain reaction versus optimized polymerase chain reaction restriction-fragment length polymorphism for apolipoprotein E genotyping of majorly depressed patients. Molecular Medicine Reports, 2015, 12, 6829-6834.	2.4	1
155	Application of antibodies against Borna disease virus phosphoprotein and nucleoprotein on paraffin sections. Molecular Medicine Reports, 2018, 17, 5416-5422.	2.4	1
156	Proteomic Profiling of Lysine Acetylation Indicates Mitochondrial Dysfunction in the Hippocampus of Gut Microbiota-Absent Mice. Frontiers in Molecular Neuroscience, 2021, 14, 594332.	2.9	1
157	Notice of Retraction: Primary Breast Diffuse Large B-Cell Lymphomas. , 2011, , .		0
158	è,é⁵å³¼®ç"Ÿç‰©ï¹¼šè§£æžç¥žç»ç²³¼ç¥žç—¾ç—…的新视窗. Scientia Sinica Vitae, 2022, , .	0.3	0