## Cunyou Zhao

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	GABAA receptor subtype selectivity underlying anxiolytic effect of 6-hydroxyflavone. Biochemical Pharmacology, 2010, 79, 1337-1344.	4.4	73
2	Two isoforms of GABAA receptor β2 subunit with different electrophysiological properties: differential expression and genotypical correlations in schizophrenia. Molecular Psychiatry, 2006, 11, 1092-1105.	7.9	50
3	Imprinting in the schizophrenia candidate gene GABRB2 encoding GABAA receptor β2 subunit. Molecular Psychiatry, 2011, 16, 557-568.	7.9	50
4	GABRB2 Association with Schizophrenia: Commonalities and Differences Between Ethnic Groups and Clinical Subtypes. Biological Psychiatry, 2007, 61, 653-660.	1.3	45
5	Hypomethylation of LINE-1 elements in schizophrenia and bipolar disorder. Journal of Psychiatric Research, 2018, 107, 68-72.	3.1	41
6	Schizophrenia-associated rs4702 G allele-specific downregulation of FURIN expression by miR-338-3p reduces BDNF production. Schizophrenia Research, 2018, 199, 176-180.	2.0	39
7	Positive Selection within the Schizophrenia-Associated GABAA Receptor Î <sup>2</sup> 2 Gene. PLoS ONE, 2007, 2, e462.	2.5	38
8	Alternative-Splicing in the Exon-10 Region of GABAA Receptor β2 Subunit Gene: Relationships between Novel Isoforms and Psychotic Disorders. PLoS ONE, 2009, 4, e6977.	2.5	35
9	<i>GABRB2</i> in schizophrenia and bipolar disorder: disease association, gene expression and clinical correlations. Biochemical Society Transactions, 2009, 37, 1415-1418.	3.4	34
10	Epigenetic regulation on GABRB2 isoforms expression: Developmental variations and disruptions in psychotic disorders. Schizophrenia Research, 2012, 134, 260-266.	2.0	31
11	A Genetic Variant Ameliorates β-Thalassemia Severity by Epigenetic-Mediated Elevation of Human Fetal Hemoglobin Expression. American Journal of Human Genetics, 2017, 101, 130-138.	6.2	31
12	Genetic and epigenetic regulation on the transcription of GABRB2 : Genotype-dependent hydroxymethylation and methylation alterations in schizophrenia. Journal of Psychiatric Research, 2017, 88, 9-17.	3.1	29
13	Social Cognitive Role of Schizophrenia Candidate Gene GABRB2. PLoS ONE, 2013, 8, e62322.	2.5	29
14	A Recombination Hotspot in a Schizophrenia-Associated Region of GABRB2. PLoS ONE, 2010, 5, e9547.	2.5	28
15	LncRNA-AC006129.1 reactivates a SOCS3-mediated anti-inflammatory response through DNA methylation-mediated CIC downregulation in schizophrenia. Molecular Psychiatry, 2021, 26, 4511-4528.	7.9	26
16	Simultaneous detection of influenza virus type B and influenza A virus subtypes H1N1, H3N2, and H5N1 using multiplex real-time RT-PCR. Applied Microbiology and Biotechnology, 2011, 90, 1463-1470.	3.6	24
17	Variation of global DNA methylation levels with age and in autistic children. Human Genomics, 2016, 10, 31.	2.9	21
18	<scp>DNA</scp> methylation regulates <i>gabrb2</i> <scp>mRNA</scp> expression: developmental variations and disruptions in <scp>l</scp> â€methionineâ€induced zebrafish with schizophreniaâ€ike symptoms. Genes, Brain and Behavior, 2016, 15, 702-710.	2.2	21

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19	DNA hypermethylation and X chromosome inactivation are major determinants of phenotypic variation in women heterozygous for G6PD mutations. Blood Cells, Molecules, and Diseases, 2014, 53, 241-245.	1.4	18
20	Epigenetic inactivation of ERF reactivates γ-globin expression in β-thalassemia. American Journal of Human Genetics, 2021, 108, 709-721.	6.2	18
21	AluScan: a method for genome-wide scanning of sequence and structure variations in the human genome. BMC Genomics, 2011, 12, 564.	2.8	17
22	Variation in global DNA hydroxymethylation with age associated with schizophrenia. Psychiatry Research, 2017, 257, 497-500.	3.3	16
23	Altered DNA methylation of the <i>Alu</i> Y subfamily in schizophrenia and bipolar disorder. Epigenomics, 2019, 11, 581-586.	2.1	15
24	Epigenetic Age Acceleration Was Delayed in Schizophrenia. Schizophrenia Bulletin, 2021, 47, 803-811.	4.3	14
25	Applicability of a sensitive duplex real-time PCR assay for identifying B/Yamagata and B/Victoria lineages of influenza virus from clinical specimens. Applied Microbiology and Biotechnology, 2012, 93, 797-805.	3.6	13
26	MicroRNAâ€2355â€5p regulates γâ€globin expression in human erythroid cells by inhibiting KLF6. British Journal of Haematology, 2021, 193, 401-405.	2.5	11
27	A simple method for high-throughput quantification of genome-wide DNA methylation by fluorescence polarization. Epigenetics, 2012, 7, 335-339.	2.7	10
28	Clustering Pattern and Functional Effect of SNPs in Human miRNA Seed Regions. International Journal of Genomics, 2018, 2018, 1-4.	1.6	9
29	Effects of flavone 6-substitutions on GABAA receptors efficacy. European Journal of Pharmacology, 2011, 670, 121-129.	3.5	7
30	Allele-specific DNA methylation maps in monozygotic twins discordant for psychiatric disorders reveal that disease-associated switching at the EIPR1 regulatory loci modulates neural function. Molecular Psychiatry, 2021, 26, 6630-6642.	7.9	7
31	Glioma Association and Balancing Selection of ZFPM2. PLoS ONE, 2015, 10, e0133003.	2.5	6
32	LncRNA RP5-998N21.4 promotes immune defense through upregulation of IFIT2 and IFIT3 in schizophrenia. NPJ Schizophrenia, 2022, 8, 11.	3.6	6
33	A rapid fluorescence polarization-based method for genotypic detection of drug resistance in Mycobacterium tuberculosis. Applied Microbiology and Biotechnology, 2014, 98, 4095-4105.	3.6	5
34	DNA methylation patterns of β-globin cluster in β-thalassemia patients. Clinical Epigenetics, 2020, 12, 187.	4.1	5
35	Mediating effect of neurocognition between severity of symptoms and social-occupational function in anxious depression. Journal of Affective Disorders, 2019, 246, 667-673.	4.1	4
36	Fluorescence polarization-based method with bisulfite conversion-specific one-label extension for quantification of single CpG dinucleotide methylation. Genome, 2015, 58, 357-363.	2.0	3

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37	Downregulation by CNNM2 of ATP5MD expression in the 10q24.32 schizophrenia-associated locus involved in impaired ATP production and neurodevelopment. NPJ Schizophrenia, 2021, 7, 27.	3.6	3
38	Tat-dependent repression of human immunodeficiency virus type 1 long terminal repeat promoter activity by fusion of cellular transcription factors. Biochemical and Biophysical Research Communications, 2004, 322, 614-622.	2.1	1
39	an Alternative-splicing Hot-spot in <i>gabrb2</i> : Novel Splicing Variants Associated with Major Psychotic Disorders. European Psychiatry, 2009, 24, .	0.2	0
40	Imprinting in the schizophrenia candidate gene GABRB2. European Psychiatry, 2011, 26, 823-823.	0.2	0
41	Association of SNPs and haplotypes in <i>GABRB2</i> with schizophrenia in Japanese and German aucasians. FASEB Journal, 2006, 20, A909.	0.5	0
42	Downregulation by <i>CNNM2</i> of <i>ATP5MD</i> Expression in the 10q24.32 Schizophrenia-Associated Locus Involved in Impaired ATP Production and Neurodevelopment. SSRN Electronic Journal, 0, , .	0.4	0