

Natalie Beveridge

List of Publications by Year in descending order

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Version: 2024-02-01

18
papers

2,315
citations

516561

16
h-index

839398

18
g-index

18
all docs

18
docs citations

18
times ranked

3569
citing authors

#	ARTICLE	IF	CITATIONS
1	Transcriptome-wide mega-analyses reveal joint dysregulation of immunologic genes and transcription regulators in brain and blood in schizophrenia. <i>Schizophrenia Research</i> , 2016, 176, 114-124.	1.1	74
2	Repair of UVB-induced DNA damage is reduced in melanoma due to low XPC and global genome repair. <i>Oncotarget</i> , 2016, 7, 60940-60953.	0.8	28
3	Understanding Xeroderma Pigmentosum Complementation Groups Using Gene Expression Profiling after UV-Light Exposure. <i>International Journal of Molecular Sciences</i> , 2015, 16, 15985-15996.	1.8	16
4	Maturation of the Human Dorsolateral Prefrontal Cortex Coincides With a Dynamic Shift in MicroRNA Expression. <i>Schizophrenia Bulletin</i> , 2014, 40, 399-409.	2.3	44
5	The long non-coding RNA Gomafu is acutely regulated in response to neuronal activation and involved in schizophrenia-associated alternative splicing. <i>Molecular Psychiatry</i> , 2014, 19, 486-494.	4.1	356
6	Gene-microRNA interactions associated with antipsychotic mechanisms and the metabolic side effects of olanzapine. <i>Psychopharmacology</i> , 2013, 227, 67-78.	1.5	39
7	Gene expression analysis reveals schizophrenia-associated dysregulation of immune pathways in peripheral blood mononuclear cells. <i>Journal of Psychiatric Research</i> , 2013, 47, 425-437.	1.5	83
8	Gene expression profiling in treatment-naive schizophrenia patients identifies abnormalities in biological pathways involving AKT1 that are corrected by antipsychotic medication. <i>International Journal of Neuropsychopharmacology</i> , 2013, 16, 1483-1503.	1.0	59
9	Decreased cortical muscarinic M1 receptors in schizophrenia are associated with changes in gene promoter methylation, mRNA and gene targeting microRNA. <i>Translational Psychiatry</i> , 2013, 3, e230-e230.	2.4	59
10	Imprinted DLK1-DIO3 region of 14q32 defines a schizophrenia-associated miRNA signature in peripheral blood mononuclear cells. <i>Molecular Psychiatry</i> , 2012, 17, 827-840.	4.1	210
11	The 3rd Schizophrenia International Research Society Conference, 14-18 April 2012, Florence, Italy: Summaries of oral sessions. <i>Schizophrenia Research</i> , 2012, 141, e1-e24.	1.1	8
12	MicroRNA dysregulation in schizophrenia. <i>Neurobiology of Disease</i> , 2012, 46, 263-271.	2.1	180
13	Transcriptome Sequencing Revealed Significant Alteration of Cortical Promoter Usage and Splicing in Schizophrenia. <i>PLoS ONE</i> , 2012, 7, e36351.	1.1	89
14	Upregulation of Dicer and MicroRNA Expression in the Dorsolateral Prefrontal Cortex Brodmann Area 46 in Schizophrenia. <i>Biological Psychiatry</i> , 2011, 69, 180-187.	0.7	236
15	Schizophrenia is associated with an increase in cortical microRNA biogenesis. <i>Molecular Psychiatry</i> , 2010, 15, 1176-1189.	4.1	396
16	Down-regulation of miR-17 family expression in response to retinoic acid induced neuronal differentiation. <i>Cellular Signalling</i> , 2009, 21, 1837-1845.	1.7	98
17	Different Forms of Glycine- and GABA _A -Receptor Mediated Inhibitory Synaptic Transmission in Mouse Superficial and Deep Dorsal Horn Neurons. <i>Molecular Pain</i> , 2009, 5, 1744-8069-5-65.	1.0	28
18	Dysregulation of miRNA 181b in the temporal cortex in schizophrenia. <i>Human Molecular Genetics</i> , 2008, 17, 1156-1168.	1.4	312