Patrick C Mchugh

List of Publications by Year in descending order

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471371 552653 33 710 17 26 citations h-index g-index papers 34 34 34 1247 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	High incidence of single nucleotide substitutions in the mitochondrial genome is associated with poor semen parameters in men. Journal of Developmental and Physical Disabilities, 2001, 24, 175-182.	3.6	99
2	Evolution of a maternal immune activation (mIA) model in rats: Early developmental effects. Brain, Behavior, and Immunity, 2019, 75, 48-59.	2.0	66
3	A dopamine transporter polymorphism is a risk factor for borderline personality disorder in depressed patients. Psychological Medicine, 2006, 36, 807-813.	2.7	57
4	The Structure and Function of the Dopamine Transporter and its Role in CNS Diseases. Vitamins and Hormones, 2015, 98, 339-369.	0.7	54
5	Relationships Between Angry-Impulsive Personality Traits and Genetic Polymorphisms of the Dopamine Transporter. Biological Psychiatry, 2009, 66, 717-721.	0.7	35
6	Proteomic analysis of embryonic stem cell–derived neural cells exposed to the antidepressant paroxetine. Journal of Neuroscience Research, 2008, 86, 306-316.	1.3	34
7	Role of carbohydrate response element-binding protein (ChREBP) in generating an aerobic metabolic phenotype and in breast cancer progression. British Journal of Cancer, 2014, 110, 715-723.	2.9	30
8	Effects of different isoforms of apoE on aggregation of the αâ€synuclein protein implicated in Parkinson's disease. Neuroscience Letters, 2016, 618, 146-151.	1.0	28
9	The presence of both serotonin 1A receptor (HTR1A) and dopamine transporter (DAT1) gene variants increase the risk of borderline personality disorder. Frontiers in Genetics, 2014, 4, 313.	1.1	24
10	Contribution of Individual Histidines to Prion Protein Copper Binding. Biochemistry, 2011, 50, 10781-10791.	1.2	21
11	Proteomic analysis of rat hippocampus exposed to the antidepressant paroxetine. Journal of Psychopharmacology, 2010, 24, 1243-1251.	2.0	20
12	Association of a functional polymorphism in the adrenomedullin gene (ADM) with response to paroxetine. Pharmacogenomics Journal, 2010, 10, 126-133.	0.9	20
13	Cognitive ability in Down syndrome and its relationship to urinary neopterin, a marker of activated cellular immunity. Neuroscience Letters, 2017, 636, 254-257.	1.0	20
14	Molecular analysis of polymerase gamma gene and mitochondrial polymorphism in fertile and subfertile men. Journal of Developmental and Physical Disabilities, 2006, 29, 421-433.	3.6	18
15	Counter-regulation of alpha- and beta-synuclein expression at the transcriptional level. Molecular and Cellular Neurosciences, 2013, 57, 33-41.	1.0	17
16	Memory Decline in Down Syndrome and Its Relationship to iPF2alpha, a Urinary Marker of Oxidative Stress. PLoS ONE, 2014, 9, e97709.	1.1	17
17	Transcriptional Regulation of the Beta-Synuclein 5′-Promoter Metal Response Element by Metal Transcription Factor-1. PLoS ONE, 2011, 6, e17354.	1.1	17
18	Repurposing of Tranilast for Potential Neuropathic Pain Treatment by Inhibition of Sepiapterin Reductase in the BH ₄ Pathway. ACS Omega, 2019, 4, 11960-11972.	1.6	15

#	Article	IF	CITATIONS
19	Polymorphisms of sepiapterin reductase gene alter promoter activity and may influence risk of bipolar disorder. Pharmacogenetics and Genomics, 2009, 19, 330-337.	0.7	14
20	A polymorphism of the GTP-cyclohydrolase I feedback regulator gene alters transcriptional activity and may affect response to SSRI antidepressants. Pharmacogenomics Journal, 2011, 11, 207-213.	0.9	13
21	Prion protein expression alters APP cleavage without interaction with BACE-1. Neurochemistry International, 2012, 61, 672-680.	1.9	12
22	The Development of Translational Biomarkers as a Tool for Improving the Understanding, Diagnosis and Treatment of Chronic Neuropathic Pain. Molecular Neurobiology, 2018, 55, 2420-2430.	1.9	12
23	Activation and repression of prion protein expression by key regions of intron 1. Cellular and Molecular Life Sciences, 2009, 66, 3809-3820.	2.4	11
24	The tetrahydrobiopterin pathway: a novel target for the treatment of depression. Pharmacogenomics, 2011, 12, 1625-1627.	0.6	10
25	The Identification of Blood Biomarkers of Chronic Neuropathic Pain by Comparative Transcriptomics. NeuroMolecular Medicine, 2022, 24, 320-338.	1.8	10
26	Increased Oxidative Stress as a Risk Factor in Chronic Idiopathic Axonal Polyneuropathy. Journal of Molecular Neuroscience, 2018, 66, 547-551.	1.1	7
27	Astragalus and human mesenchymal stem cells promote wound healing by mediating immunomodulatory effects through paracrine signaling. Regenerative Medicine, 2022, 17, 219-232.	0.8	7
28	Research of single mitochondrial nucleotide substitutions in male infertility should consider human mitochondrial haplogroups. Journal of Developmental and Physical Disabilities, 2002, 25, 372-373.	3.6	6
29	Downregulation of <i>Ccnd1 </i> and <i> Hes6 </i> in rat hippocampus after chronic exposure to the antidepressant paroxetine. Acta Neuropsychiatrica, 2008, 20, 307-313.	1.0	6
30	Nonsynonymous Polymorphism in Guanine Monophosphate Synthetase Is a Risk Factor for Unfavorable Thiopurine Metabolite Ratios in Patients With Inflammatory Bowel Disease. Inflammatory Bowel Diseases, 2018, 24, 2606-2612.	0.9	4
31	MicroRNA Quantitation During Dendritic Cell Endocytosis Using Imaging Flow Cytometry: Key Factors and Requirements. Cellular Physiology and Biochemistry, 2018, 51, 793-811.	1.1	3
32	Sphingosine and dihydrosphingosine as biomarkers for multiple sclerosis identified by metabolomic profiling using coupled UPLC-MS. Analytical Methods, 2017, 9, 5929-5934.	1.3	2
33	Tetrahydrobiopterin Pathway may Provide Novel Molecular Targets for Acute and Long Term Efficacy of Mood-Regulating Drugs. Current Pharmacogenomics and Personalized Medicine, 2010, 8, 174-181.	0.2	1