

Subramaniam Jayanthi

List of Publications by Year in descending order

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67
papers

3,685
citations

147801

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133252

59
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67
all docs

67
docs citations

67
times ranked

2906
citing authors

#	ARTICLE	IF	CITATIONS
1	Epigenetics of Addiction. , 2022, , 383-389.		0
2	Footshock-Induced Abstinence from Compulsive Methamphetamine Self-administration in Rat Model Is Accompanied by Increased Hippocampal Expression of Cannabinoid Receptors (CB1 and CB2). <i>Molecular Neurobiology</i> , 2022, 59, 1238-1248.	4.0	4
3	Sex-Specific Alterations in Dopamine Metabolism in the Brain after Methamphetamine Self-Administration. <i>International Journal of Molecular Sciences</i> , 2022, 23, 4353.	4.1	6
4	Sex differences in methamphetamine use disorder perused from pre-clinical and clinical studies: Potential therapeutic impacts. <i>Neuroscience and Biobehavioral Reviews</i> , 2022, 137, 104674.	6.1	27
5	Sex- and Brain Region-specific Changes in Gene Expression in Male and Female Rats as Consequences of Methamphetamine Self-administration and Abstinence. <i>Neuroscience</i> , 2021, 452, 265-279.	2.3	19
6	Epigenetic Landscape of Methamphetamine Use Disorder. <i>Current Neuropharmacology</i> , 2021, 19, 2060-2066.	2.9	7
7	Footshock-induced abstinence from compulsive methamphetamine self-administration is associated with increased expression of cannabinoid receptors (CB1 and CB2) in the rat hippocampus. <i>FASEB Journal</i> , 2021, 35, .	0.5	0
8	Epigenetics of addiction. <i>Neurochemistry International</i> , 2021, 147, 105069.	3.8	18
9	Neurotoxicity of methamphetamine: Main effects and mechanisms. <i>Experimental Neurology</i> , 2021, 344, 113795.	4.1	88
10	Potassium Channels and Their Potential Roles in Substance Use Disorders. <i>International Journal of Molecular Sciences</i> , 2021, 22, 1249.	4.1	14
11	Epigenetic Regulatory Dynamics in Models of Methamphetamine-Use Disorder. <i>Genes</i> , 2021, 12, 1614.	2.4	12
12	HDAC superfamily promoters acetylation is differentially regulated by modafinil and methamphetamine in the mouse medial prefrontal cortex. <i>Addiction Biology</i> , 2020, 25, e12737.	2.6	15
13	A Single Prior Injection of Methamphetamine Enhances Methamphetamine Self-Administration (SA) and Blocks SA-Induced Changes in DNA Methylation and mRNA Expression of Potassium Channels in the Rat Nucleus Accumbens. <i>Molecular Neurobiology</i> , 2020, 57, 1459-1472.	4.0	24
14	Compulsive methamphetamine taking induces autophagic and apoptotic markers in the rat dorsal striatum. <i>Archives of Toxicology</i> , 2020, 94, 3515-3526.	4.2	14
15	Methamphetamine pre-exposure induces steeper escalation of methamphetamine self-administration with consequent alterations in hippocampal glutamate AMPA receptor mRNAs. <i>European Journal of Pharmacology</i> , 2020, 889, 173732.	3.5	2
16	Neurochemical and behavioral comparisons of contingent and non-contingent methamphetamine exposure following binge or yoked long-access self-administration paradigms. <i>Psychopharmacology</i> , 2020, 237, 1989-2005.	3.1	19
17	The effects of single-dose injections of modafinil and methamphetamine on epigenetic and functional markers in the mouse medial prefrontal cortex: potential role of dopamine receptors. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2019, 88, 222-234.	4.8	26
18	Sex Differences in Escalated Methamphetamine Self-Administration and Altered Gene Expression Associated With Incubation of Methamphetamine Seeking. <i>International Journal of Neuropsychopharmacology</i> , 2019, 22, 710-723.	2.1	38

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19	Genetic and Environmental Risk Factors for Cannabis Use: Preliminary Results for the Role of Parental Care Perception. <i>Substance Use and Misuse</i> , 2019, 54, 670-680.	1.4	18
20	Compulsive methamphetamine taking and abstinence in the presence of adverse consequences: Epigenetic and transcriptional consequences in the rat brain. <i>Pharmacology Biochemistry and Behavior</i> , 2019, 179, 98-108.	2.9	29
21	Molecular Adaptations in the Rat Dorsal Striatum and Hippocampus Following Abstinence-Induced Incubation of Drug Seeking After Escalated Oxycodone Self-Administration. <i>Molecular Neurobiology</i> , 2019, 56, 3603-3615.	4.0	39
22	Selective Activation of Striatal NGF-TrkA/p75NTR/MAPK Intracellular Signaling in Rats That Show Suppression of Methamphetamine Intake 30 Days following Drug Abstinence. <i>International Journal of Neuropsychopharmacology</i> , 2018, 21, 281-290.	2.1	15
23	Gene variants and educational attainment in cannabis use: mediating role of DNA methylation. <i>Translational Psychiatry</i> , 2018, 8, 23.	4.8	32
24	Methamphetamine Induces TET1- and TET3-Dependent DNA Hydroxymethylation of Crh and Avp Genes in the Rat Nucleus Accumbens. <i>Molecular Neurobiology</i> , 2018, 55, 5154-5166.	4.0	38
25	Repeated methamphetamine and modafinil induce differential cognitive effects and specific histone acetylation and DNA methylation profiles in the mouse medial prefrontal cortex. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2018, 82, 1-11.	4.8	39
26	Compulsive methamphetamine taking under punishment is associated with greater cue-induced drug seeking in rats. <i>Behavioural Brain Research</i> , 2017, 326, 265-271.	2.2	31
27	Compulsive methamphetamine taking in the presence of punishment is associated with increased oxytocin expression in the nucleus accumbens of rats. <i>Scientific Reports</i> , 2017, 7, 8331.	3.3	26
28	Genome-wide DNA hydroxymethylation identifies potassium channels in the nucleus accumbens as discriminators of methamphetamine addiction and abstinence. <i>Molecular Psychiatry</i> , 2017, 22, 1196-1204.	7.9	65
29	Increased expression of proenkephalin and prodynorphin mRNAs in the nucleus accumbens of compulsive methamphetamine taking rats. <i>Scientific Reports</i> , 2016, 6, 37002.	3.3	22
30	An Acute Methamphetamine Injection Downregulates the Expression of Several Histone Deacetylases (HDACs) in the Mouse Nucleus Accumbens: Potential Regulatory Role of HDAC2 Expression. <i>Neurotoxicity Research</i> , 2016, 30, 32-40.	2.7	19
31	Differential Expression of mRNAs Coding for Histone Deacetylases (HDACs) in the Nucleus Accumbens of Compulsive Methamphetamine Takers and Abstinent Rats. <i>Journal of Drug and Alcohol Research</i> , 2016, 5, 1-9.	0.9	3
32	CAMKII-conditional deletion of histone deacetylase 2 potentiates acute methamphetamine-induced expression of immediate early genes in the mouse nucleus accumbens. <i>Scientific Reports</i> , 2015, 5, 13396.	3.3	16
33	Epigenetic landscape of amphetamine and methamphetamine addiction in rodents. <i>Epigenetics</i> , 2015, 10, 574-580.	2.7	101
34	Transcriptional and Epigenetic Substrates of Methamphetamine Addiction and Withdrawal: Evidence from a Long-Access Self-Administration Model in the Rat. <i>Molecular Neurobiology</i> , 2015, 51, 696-717.	4.0	64
35	Methamphetamine Downregulates Striatal Glutamate Receptors via Diverse Epigenetic Mechanisms. <i>Biological Psychiatry</i> , 2014, 76, 47-56.	1.3	109
36	Differential effects of binge methamphetamine injections on the mRNA expression of histone deacetylases (HDACs) in the rat striatum. <i>NeuroToxicology</i> , 2014, 45, 178-184.	3.0	27

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37	Enhanced Upregulation of CRH mRNA Expression in the Nucleus Accumbens of Male Rats after a Second Injection of Methamphetamine Given Thirty Days Later. <i>PLoS ONE</i> , 2014, 9, e84665.	2.5	35
38	Genome-wide profiling identifies a subset of methamphetamine (METH)-induced genes associated with METH-induced increased H4K5Ac binding in the rat striatum. <i>BMC Genomics</i> , 2013, 14, 545.	2.8	43
39	Epigenetics of Methamphetamine-Induced Changes in Glutamate Function. <i>Neuropsychopharmacology</i> , 2013, 38, 248-249.	5.4	27
40	CREB phosphorylation regulates striatal transcriptional responses in the self-administration model of methamphetamine addiction in the rat. <i>Neurobiology of Disease</i> , 2013, 58, 132-143.	4.4	115
41	Methamphetamine Causes Differential Alterations in Gene Expression and Patterns of Histone Acetylation/Hypoacetylation in the Rat Nucleus Accumbens. <i>PLoS ONE</i> , 2012, 7, e34236.	2.5	111
42	Involvement of Dopamine Receptors in Binge Methamphetamine-Induced Activation of Endoplasmic Reticulum and Mitochondrial Stress Pathways. <i>PLoS ONE</i> , 2011, 6, e28946.	2.5	78
43	Chronic methamphetamine exposure suppresses the striatal expression of members of multiple families of immediate early genes (IEGs) in the rat: normalization by an acute methamphetamine injection. <i>Psychopharmacology</i> , 2011, 215, 353-365.	3.1	47
44	Differential histone modifications induced by chronic methamphetamine exposure in the rat striatum. <i>FASEB Journal</i> , 2011, 25, 896.6.	0.5	0
45	Differential effects of methamphetamine and SCH23390 on the expression of members of IEG families of transcription factors in the rat striatum. <i>Brain Research</i> , 2010, 1318, 1-10.	2.2	36
46	Methamphetamine Self-Administration Is Associated with Persistent Biochemical Alterations in Striatal and Cortical Dopaminergic Terminals in the Rat. <i>PLoS ONE</i> , 2010, 5, e8790.	2.5	119
47	Dopamine D1 Receptors, Regulation of Gene Expression in the Brain, and Neurodegeneration. <i>CNS and Neurological Disorders - Drug Targets</i> , 2010, 9, 526-538.	1.4	90
48	Methamphetamine Induces Dopamine D1 Receptor-Dependent Endoplasmic Reticulum Stress-Related Molecular Events in the Rat Striatum. <i>PLoS ONE</i> , 2009, 4, e6092.	2.5	76
49	Methamphetamine Administration Causes Death of Dopaminergic Neurons in the Mouse Olfactory Bulb. <i>Biological Psychiatry</i> , 2007, 61, 1235-1243.	1.3	62
50	Neurotoxicity of substituted amphetamines: Molecular and cellular mechanisms. <i>Neurotoxicity Research</i> , 2007, 11, 183-202.	2.7	252
51	Identification of Putative Biomarkers in the Serum of Marijuana Users by Surface-Enhanced Laser Desorption/Ionization Time of Flight Mass Spectrometry (SELDI-TOF-MS). <i>FASEB Journal</i> , 2007, 21, A421.	0.5	0
52	Methamphetamine-induced neuronal apoptosis involves the activation of multiple death pathways. Review. <i>Neurotoxicity Research</i> , 2005, 8, 199-206.	2.7	114
53	Calcineurin/NFAT-induced up-regulation of the Fas ligand/Fas death pathway is involved in methamphetamine-induced neuronal apoptosis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 868-873.	7.1	208
54	Methamphetamine induces neuronal apoptosis via cross-talks between endoplasmic reticulum and mitochondria-dependent death cascades. <i>FASEB Journal</i> , 2004, 18, 238-251.	0.5	255

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55	Substituted Amphetamines That Produce Long-Term Serotonin Depletion in Rat Brain (â€œNeurotoxicityâ€) Do Not Decrease Serotonin Transporter Protein Expression. <i>Annals of the New York Academy of Sciences</i> , 2004, 1025, 151-161.	3.8	12
56	High-dose fenfluramine administration decreases serotonin transporter binding, but not serotonin transporter protein levels, in rat forebrain. <i>Synapse</i> , 2003, 50, 233-239.	1.2	56
57	Speed kills: cellular and molecular bases of methamphetamineâ€induced nerve terminal degeneration and neuronal apoptosis. <i>FASEB Journal</i> , 2003, 17, 1775-1788.	0.5	265
58	Methamphetamine Causes Coordinate Regulation of Src, Cas, Crk, and the Jun N-Terminal Kinaseâ€Jun Pathway. <i>Molecular Pharmacology</i> , 2002, 61, 1124-1131.	2.3	63
59	Mice with Partial Deficiency of c-Jun Show Attenuation of Methamphetamine-Induced Neuronal Apoptosis. <i>Molecular Pharmacology</i> , 2002, 62, 993-1000.	2.3	49
60	Amphetamine-induced toxicity in dopamine terminals in CD-1 and C57BL/6J mice: complex roles for oxygen-based species and temperature regulation. <i>Neuroscience</i> , 2001, 107, 265-274.	2.3	43
61	Methamphetamine increases expression of the apoptotic c-myc and l-myc genes in the mouse brain. <i>Molecular Brain Research</i> , 2001, 90, 202-204.	2.3	14
62	Fas-induced apoptosis of glioma cells is associated with down-regulation of the hSCO1 protein, a subunit of complex IV. <i>Molecular Brain Research</i> , 2001, 91, 131-136.	2.3	9
63	Temporal profiling of methamphetamine-induced changes in gene expression in the mouse brain: Evidence from cDNA array. <i>Synapse</i> , 2001, 41, 40-48.	1.2	99
64	Methamphetamine causes differential regulation of proâ€death and antiâ€death Bclâ€2 genes in the mouse neocortex. <i>FASEB Journal</i> , 2001, 15, 1745-1752.	0.5	149
65	Overexpression of human copper/zinc superoxide dismutase in transgenic mice attenuates oxidative stress caused by methylenedioxymethamphetamine (Ecstasy). <i>Neuroscience</i> , 1999, 91, 1379-1387.	2.3	74
66	Dual mechanism of Fas-induced cell death in neuroglioma cells: a role for reactive oxygen species. <i>Molecular Brain Research</i> , 1999, 72, 158-165.	2.3	38
67	Methamphetamine-Induced Changes in Antioxidant Enzymes and Lipid Peroxidation in Copper/Zinc-Superoxide Dismutase Transgenic Mice. <i>Annals of the New York Academy of Sciences</i> , 1998, 844, 92-102.	3.8	120