Sean P Farris

List of Publications by Year
in descending order

22 papers	800 citations	567281 15 h-index	713466 21 g-index
23 all docs	23 docs citations	23 times ranked	980 citing authors

#	Article	IF	Citations
1	Alcohol Dependence in Rats Is Associated with Global Changes in Gene Expression in the Central Amygdala. Brain Sciences, 2021, 11, 1149.	2.3	7
2	Epigenetic and non-coding regulation of alcohol abuse and addiction. International Review of Neurobiology, 2021, 156, 63-86.	2.0	8
3	Transcriptome Analysis of Alcohol Drinking in Non-Dependent and Dependent Mice Following Repeated Cycles of Forced Swim Stress Exposure. Brain Sciences, 2020, 10, 275.	2.3	11
4	Bioinformatic and biological avenues for understanding alcohol use disorder. Alcohol, 2019, 74, 65-71.	1.7	3
5	Cross-Species Co-analysis of Prefrontal Cortex Chronic Ethanol Transcriptome Responses in Mice and Monkeys. Frontiers in Molecular Neuroscience, 2019, 12, 197.	2.9	21
6	Analysis of whole genome-transcriptomic organization in brain to identify genes associated with alcoholism. Translational Psychiatry, 2019, 9, 89.	4.8	66
7	Ethanol and a rapid-acting antidepressant produce overlapping changes in exon expression in the synaptic transcriptome. Neuropharmacology, 2019, 146, 289-299.	4.1	9
8	Beyond genomeâ€wide significance: integrative approaches to the interpretation and extension of GWAS findings for alcohol use disorder. Addiction Biology, 2019, 24, 275-289.	2.6	15
9	Astrocyte-specific transcriptome responses to chronic ethanol consumption. Pharmacogenomics Journal, 2018, 18, 578-589.	2.0	35
10	Microglial-specific transcriptome changes following chronic alcohol consumption. Neuropharmacology, 2018, 128, 416-424.	4.1	37
11	Glycogen synthase kinase 3 beta regulates ethanol consumption and is a risk factor for alcohol dependence. Neuropsychopharmacology, 2018, 43, 2521-2531.	5.4	25
12	FMRP regulates an ethanol-dependent shift in GABABR function and expression with rapid antidepressant properties. Nature Communications, 2016, 7, 12867.	12.8	48
13	Moving Toward Understanding the Proteome Involved in Substance Abuse. Biological Psychiatry, 2016, 79, 422-424.	1.3	0
14	Epigenetic modulation of brain gene networks for cocaine and alcohol abuse. Frontiers in Neuroscience, 2015, 9, 176.	2.8	69
15	Chronic Ethanol Exposure Produces Time- and Brain Region-Dependent Changes in Gene Coexpression Networks. PLoS ONE, 2015, 10, e0121522.	2.5	92
16	Applying the new genomics to alcohol dependence. Alcohol, 2015, 49, 825-836.	1.7	15
17	Transcriptome organization for chronic alcohol abuse in human brain. Molecular Psychiatry, 2015, 20, 1438-1447.	7.9	111
18	Effects of chronic HIV-1 Tat exposure in the CNS: heightened vulnerability of males versus females to changes in cell numbers, synaptic integrity, and behavior. Brain Structure and Function, 2015, 220, 605-623.	2.3	74

#	Article	IF	Citations
19	RNA-Seq Reveals Novel Transcriptional Reorganization in Human Alcoholic Brain. International Review of Neurobiology, 2014, 116, 275-300.	2.0	50
20	Fyn-Dependent Gene Networks in Acute Ethanol Sensitivity. PLoS ONE, 2013, 8, e82435.	2.5	40
21	Ethanol modulation of gene networks: Implications for alcoholism. Neurobiology of Disease, 2012, 45, 115-121.	4.4	23
22	Using Expression Genetics to Study the Neurobiology of Ethanol and Alcoholism. International Review of Neurobiology, 2010, 91, 95-128.	2.0	41