Alexandre A Lussier

List of Publications by Year
in descending order

25	757	932766 10	642321 23
papers	citations	h-index	g-index

35	35	35	1177
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Sensitive Periods for the Effect of Childhood Adversity on DNA Methylation: Updated Results From a Prospective, Longitudinal Study. Biological Psychiatry Global Open Science, 2023, 3, 567-571.	1.0	3
2	Sensitive period-regulating genetic pathways and exposure to adversity shape risk for depression. Neuropsychopharmacology, 2022, 47, 497-506.	2.8	8
3	Updates to data versions and analytic methods influence the reproducibility of results from epigenome-wide association studies. Epigenetics, 2022, 17, 1373-1388.	1.3	9
4	Examining the epigenetic mechanisms of childhood adversity and sensitive periods: A gene set-based approach. Psychoneuroendocrinology, 2022, 144, 105854.	1.3	2
5	Genetic susceptibility for major depressive disorder associates with trajectories of depressive symptoms across childhood and adolescence. Journal of Child Psychology and Psychiatry and Allied Disciplines, 2021, 62, 895-904.	3.1	9
6	A Structured Approach to Evaluating Life-Course Hypotheses: Moving Beyond Analyses of Exposed Versus Unexposed in the -Omics Context. American Journal of Epidemiology, 2021, 190, 1101-1112.	1.6	11
7	OUP accepted manuscript. Nucleic Acids Research, 2021, 49, 9097-9116.	6.5	19
8	Childhood Emotional Neglect and Adolescent Depression: Assessing the Protective Role of Peer Social Support in a Longitudinal Birth Cohort. Frontiers in Psychiatry, 2021, 12, 681176.	1.3	8
9	Prenatal Adversity Alters the Epigenetic Profile of the Prefrontal Cortex: Sexually Dimorphic Effects of Prenatal Alcohol Exposure and Food-Related Stress. Genes, 2021, 12, 1773.	1.0	10
10	Association of Maternal Stress and Social Support During Pregnancy With Growth Marks in Childrenâ€™s Primary Tooth Enamel. JAMA Network Open, 2021, 4, e2129129.	2.8	10
11	Associations between indicators of socioeconomic position and DNA methylation: a scoping review. Clinical Epigenetics, 2021, 13, 221.	1.8	23
12	Intersection of Epigenetic and Immune Alterations: Implications for Fetal Alcohol Spectrum Disorder and Mental Health. Frontiers in Neuroscience, 2021, 15, 788630.	1.4	10
13	Neonatal Alcohol Exposure in Mice Induces Select Differentiation- and Apoptosis-Related Chromatin Changes Both Independent of and Dependent on Sex. Frontiers in Genetics, 2020, 11, 35.	1.1	15
14	Crowdsourced genealogies and genomes. Science, 2018, 360, 153-154.	6.0	4
15	Epigenetics and Genetics of Development. , 2018, , 153-210.		2
16	Prenatal Alcohol Exposure: Profiling Developmental DNA Methylation Patterns in Central and Peripheral Tissues. Frontiers in Genetics, 2018, 9, 610.	1.1	27
17	Epigenetic analysis of human postmortem brain tissue. Handbook of Clinical Neurology / Edited By P J Vinken and G W Bruyn, 2018, 150, 237-261.	1.0	3
18	DNA methylation as a predictor of fetal alcohol spectrum disorder. Clinical Epigenetics, 2018, 10, 5.	1.8	89

#	Article	IF	Citations
19	Epigenetics studies of fetal alcohol spectrum disorder: where are we now?. Epigenomics, 2017, 9, 291-311.	1.0	84
20	DNA methylation signature of human fetal alcohol spectrum disorder. Epigenetics and Chromatin, 2016, 9, 25.	1.8	129
21	La-related Protein 1 (LARP1) Represses Terminal Oligopyrimidine (TOP) mRNA Translation Downstream of mTOR Complex 1 (mTORC1). Journal of Biological Chemistry, 2015, 290, 15996-16020.	1.6	198
22	Prenatal Alcohol Exposure Alters Steadyâ€State and Activated Gene Expression in the Adult Rat Brain. Alcoholism: Clinical and Experimental Research, 2015, 39, 251-261.	1.4	41
23	ISDN2014_0378: Prenatal alcohol exposure alters the developmental methylation profile of the rat hypothalamus. International Journal of Developmental Neuroscience, 2015, 47, 109-109.	0.7	0
24	Prenatal alcohol exposure alters gene expression in the rat brain: Experimental design and bioinformatic analysis of microarray data. Data in Brief, 2015, 4, 239-252.	0.5	2
25	Molecular pathways underpinning ethanol-induced neurodegeneration. Frontiers in Genetics, 2014, 5, 203.	1.1	30