Jessica Connelly

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Neuroepigenetic impact on mentalizing in childhood. Developmental Cognitive Neuroscience, 2022, 54, 101080.	1.9	3
2	Lifetime marijuana use and epigenetic age acceleration: A 17-year prospective examination. Drug and Alcohol Dependence, 2022, 233, 109363.	1.6	14
3	Exercise during pregnancy mitigates negative effects of parental obesity on metabolic function in adult mouse offspring. Journal of Applied Physiology, 2021, 130, 605-616.	1.2	11
4	OXTR DNA methylation moderates the developmental calibration of neural reward sensitivity. Developmental Psychobiology, 2021, 63, 114-124.	0.9	8
5	Genetic, epigenetic, and environmental factors controlling oxytocin receptor gene expression. Clinical Epigenetics, 2021, 13, 23.	1.8	41
6	An epigenetic rheostat of experience: DNA methylation of OXTR as a mechanism of early life allostasis. Comprehensive Psychoneuroendocrinology, 2021, 8, 100098.	0.7	12
7	ls Oxytocin "Nature's Medicine�. Pharmacological Reviews, 2020, 72, 829-861.	7.1	190
8	Epigenetic tuning of brain signal entropy in emergent human social behavior. BMC Medicine, 2020, 18, 244.	2.3	11
9	Epigenetic Dynamics of the Oxytocin Receptor Gene Across the Menstrual Cycle. Biological Psychiatry, 2020, 87, S391.	0.7	1
10	Epigenetic dynamics in infancy and the impact of maternal engagement. Science Advances, 2019, 5, eaay0680.	4.7	48
11	Epigenetic modification of the oxytocin receptor gene is associated with emotion processing in the infant brain. Developmental Cognitive Neuroscience, 2019, 37, 100648.	1.9	55
12	Behavioral and epigenetic consequences of oxytocin treatment at birth. Science Advances, 2019, 5, eaav2244.	4.7	50
13	T56. DNA Methylation of the Oxytocin Receptor Changes During Infancy and is Impacted by Maternal Behavior. Biological Psychiatry, 2019, 85, S150.	0.7	0
14	Early nurture epigenetically tunes the oxytocin receptor. Psychoneuroendocrinology, 2019, 99, 128-136.	1.3	83
15	Oxytocin receptor genotype and low economic privilege reverses ventral striatum-social anxiety association. Social Neuroscience, 2019, 14, 67-79.	0.7	7
16	Associations between oxytocin receptor gene (OXTR) methylation, plasma oxytocin, and attachment across adulthood. International Journal of Psychophysiology, 2019, 136, 22-32.	0.5	55
17	The Role of Endogenous Oxytocin in Anxiolysis: Structural and Functional Correlates. Biological Psychiatry: Cognitive Neuroscience and Neuroimaging, 2018, 3, 618-625.	1.1	16
18	Neuroimaging Epigenetics: Challenges and Recommendations for Best Practices. Neuroscience, 2018, 370, 88-100.	1.1	19

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19	DNA methylation of <i>OXTR</i> is associated with parasympathetic nervous system activity and amygdala morphology. Social Cognitive and Affective Neuroscience, 2018, 13, 1155-1162.	1.5	18
20	S38. Epigenetic Modification of the Oxytocin Receptor Gene Impacts Infant Neural Response to Emotional Faces. Biological Psychiatry, 2018, 83, S361-S362.	0.7	0
21	Epigenetic regulation of the oxytocin receptor is associated with neural response during selective social attention. Translational Psychiatry, 2018, 8, 116.	2.4	46
22	Smooth muscle cell-specific deletion of <i>Col15a1</i> unexpectedly leads to impaired development of advanced atherosclerotic lesions. American Journal of Physiology - Heart and Circulatory Physiology, 2017, 312, H943-H958.	1.5	34
23	Activation of the pluripotency factor OCT4 in smooth muscle cells is atheroprotective. Nature Medicine, 2016, 22, 657-665.	15.2	165
24	Sex and Diagnosis-Specific Associations Between DNA Methylation of the Oxytocin Receptor Gene With Emotion Processing and Temporal-Limbic and Prefrontal Brain Volumes in Psychotic Disorders. Biological Psychiatry: Cognitive Neuroscience and Neuroimaging, 2016, 1, 141-151.	1.1	45
25	Childbirth and symptoms of postpartum depression and anxiety: a prospective birth cohort study. Archives of Women's Mental Health, 2016, 19, 219-227.	1.2	67
26	Interaction between oxytocin receptor DNA methylation and genotype is associated with risk of postpartum depression in women without depression in pregnancy. Frontiers in Genetics, 2015, 6, 243.	1.1	82
27	Plasma oxytocin explains individual differences in neural substrates of social perception. Frontiers in Human Neuroscience, 2015, 9, 132.	1.0	41
28	Epigenetic modification of the oxytocin receptor gene influences the perception of anger and fear in the human brain. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 3308-3313.	3.3	170
29	ZNF277 microdeletions, specific language impairment and the meiotic mismatch methylation (3M) hypothesis. European Journal of Human Genetics, 2015, 23, 1113-1113.	1.4	4
30	A Functionally Significant Polymorphism in ID3 Is Associated with Human Coronary Pathology. PLoS ONE, 2014, 9, e90222.	1.1	18
31	Personality, Behavior and Environmental Features Associated with OXTR Genetic Variants in British Mothers. PLoS ONE, 2014, 9, e90465.	1.1	29
32	Response to Comment on Laker et al. Exercise Prevents Maternal High-Fat Diet–Induced Hypermethylation of thePgc-1αGene and Age-Dependent Metabolic Dysfunction in the Offspring. Diabetes 2014;63:1605â°'1611. Diabetes, 2014, 63, e6-e7.	0.3	5
33	Exercise Prevents Maternal High-Fat Diet–Induced Hypermethylation of the <i>Pgc-1α</i> Gene and Age-Dependent Metabolic Dysfunction in the Offspring. Diabetes, 2014, 63, 1605-1611.	0.3	184
34	Oxytocin receptor gene variation predicts empathic concern and autonomic arousal while perceiving harm to others. Social Neuroscience, 2014, 9, 1-9.	0.7	123
35	Epigenetic origins of metabolic disease: The impact of the maternal condition to the offspring epigenome and later health consequences. Food Science and Human Wellness, 2013, 2, 1-11.	2.2	61
36	Epigenetic regulation of COL15A1 in smooth muscle cell replicative aging and atherosclerosis. Human Molecular Genetics, 2013, 22, 5107-5120.	1.4	66

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37	Gestational Exposure to Bisphenol A Produces Transgenerational Changes in Behaviors and Gene Expression. Endocrinology, 2012, 153, 3828-3838.	1.4	276
38	DNA methylation of the oxytocin receptor gene predicts neural response to ambiguous social stimuli. Frontiers in Human Neuroscience, 2012, 6, 280.	1.0	155
39	The role of Bisphenol A in shaping the brain, epigenome and behavior. Hormones and Behavior, 2011, 59, 296-305.	1.0	256
40	Gestational Exposure to Low Dose Bisphenol A Alters Social Behavior in Juvenile Mice. PLoS ONE, 2011, 6, e25448.	1.1	144
41	Polymorphic variants in tenascin-C (TNC) are associated with atherosclerosis and coronary artery disease. Human Genetics, 2011, 129, 641-654.	1.8	25
42	Cigarette smoking status has a modifying effect on the association between polymorphisms in KALRN and measures of cardiovascular risk in the diabetes heart study. Genes and Genomics, 2011, 33, 483-490.	0.5	2
43	Aging-related atherosclerosis is exacerbated by arterial expression of tumor necrosis factor receptor-1: evidence from mouse models and human association studies. Human Molecular Genetics, 2010, 19, 2754-2766.	1.4	32
44	Neuropeptide Y Gene Polymorphisms Confer Risk of Early-Onset Atherosclerosis. PLoS Genetics, 2009, 5, e1000318.	1.5	87
45	Genomic and epigenetic evidence for oxytocin receptor deficiency in autism. BMC Medicine, 2009, 7, 62.	2.3	497
46	Genetic effects in the leukotriene biosynthesis pathway and association with atherosclerosis. Human Genetics, 2009, 125, 217-229.	1.8	51
47	Refinement of 2q and 7p loci in a large multiplex NTD family. Birth Defects Research Part A: Clinical and Molecular Teratology, 2008, 82, 441-452.	1.6	12
48	ALOX5AP variants are associated with in-stent restenosis after percutaneous coronary intervention. Atherosclerosis, 2008, 201, 148-154.	0.4	22
49	The Crystal Structure of Cdc42 in Complex with Collybistin II, a Gephyrin-interacting Guanine Nucleotide Exchange Factor. Journal of Molecular Biology, 2006, 359, 35-46.	2.0	63
50	GATA2 Is Associated with Familial Early-Onset Coronary Artery Disease. PLoS Genetics, 2006, 2, e139.	1.5	82
51	Structure and Function of the Saccharomyces cerevisiae Sir3 BAH Domain. Molecular and Cellular Biology, 2006, 26, 3256-3265.	1.1	56
52	Importance of the Sir3 N Terminus and Its Acetylation for Yeast Transcriptional Silencing. Genetics, 2004, 168, 547-551.	1.2	68