## Nadine Provençal

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3391467/publications.pdf

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37 papers

2,010 citations

331670 21 h-index 395702 33 g-index

41 all docs

41 docs citations

41 times ranked

2897 citing authors

#	Article	IF	CITATIONS
1	Combined effects of genotype and childhood adversity shape variability of DNA methylation across age. Translational Psychiatry, 2021, 11, 88.	4.8	27
2	Glucocorticoid exposure during hippocampal neurogenesis primes future stress response by inducing changes in DNA methylation. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 23280-23285.	7.1	141
3	Investigation of MORC1 DNA methylation as biomarker of early life stress and depressive symptoms. Journal of Psychiatric Research, 2020, 120, 154-162.	3.1	9
4	A polyepigenetic glucocorticoid exposure score at birth and childhood mental and behavioral disorders. Neurobiology of Stress, 2020, 13, 100275.	4.0	8
5	Intergenerational Effects of Maternal Holocaust Exposure on <i>FKBP5</i> Methylation. American Journal of Psychiatry, 2020, 177, 744-753.	7.2	49
6	INTERACTIONS BETWEEN GENOTYPE AND ENVIRONMENT HAVE A STRONG EFFECT ON VARIABILITY IN DNA METHYLATION IN PSYCHIATRIC PATIENTS. European Neuropsychopharmacology, 2019, 29, S984.	0.7	0
7	SU56EPIGENETIC REGULATION OF THE NOVEL EARLY LIFE ADVERSITY RESPONSIVE GENE MORC1 IN MAJOR DEPRESSIVE DISORDER. European Neuropsychopharmacology, 2019, 29, S1297.	0.7	O
8	Identification of dynamic glucocorticoid-induced methylation changes at the FKBP5 locus. Clinical Epigenetics, 2019, 11, 83.	4.1	49
9	A Role of Oxytocin Receptor Gene Brain Tissue Expression Quantitative Trait Locus rs237895 in the Intergenerational Transmission of the Effects of Maternal Childhood Maltreatment. Journal of the American Academy of Child and Adolescent Psychiatry, 2019, 58, 1207-1216.	0.5	15
10	Dynamic DNA methylation changes in the maternal oxytocin gene locus (OXT) during pregnancy predict postpartum maternal intrusiveness. Psychoneuroendocrinology, 2019, 103, 156-162.	2.7	22
11	<i>DRD4</i> methylation as a potential biomarker for physical aggression: An epigenomeâ€wide, crossâ€tissue investigation. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2018, 177, 746-764.	1.7	33
12	HAM-TBS: high-accuracy methylation measurements via targeted bisulfite sequencing. Epigenetics and Chromatin, 2018, 11, 39.	3.9	22
13	F121. Investigating Glucocorticoid Receptor Binding in Lymphoblastoid Cell Lines. Biological Psychiatry, 2018, 83, S284-S285.	1.3	O
14	Central Neuroepigenetic Regulation of the Hypothalamic–Pituitary–Adrenal Axis. Progress in Molecular Biology and Translational Science, 2018, 158, 105-127.	1.7	13
15	From Epigenetic Associations to Biological and Psychosocial Explanations in Mental Health. Progress in Molecular Biology and Translational Science, 2018, 158, 299-323.	1.7	1
16	Early life stress, FK506 binding protein 5 gene ( <i>FKBP5</i> ) methylation, and inhibition-related prefrontal function: A prospective longitudinal study. Development and Psychopathology, 2017, 29, 1895-1903.	2.3	46
17	How Can GxE Research Help Prevent the Development of Chronic Physical Aggression?., 2017, , 177-207.		1
18	Dynamic Changes in DNA Methylation Occur during the First Year of Life in Preterm Infants. Frontiers in Endocrinology, 2016, 7, 158.	3.5	24

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19	The Impact of Environmental Stressors on DNA Methylation, Neurobehavioral Development, and Chronic Physical Aggression: Prospects for Early Protective Interventions. Molecular and Integrative Toxicology, 2016, , 295-319.	0.5	1
20	Alterations in DNA Methylation and Hydroxymethylation Due to Parental Care in Rhesus Macaques. Epigenetics and Human Health, 2016, , 165-190.	0.2	1
21	Epigenetic mechanisms involved in the effects of stress exposure: focus on 5-hydroxymethylcytosine: Table 1:. Environmental Epigenetics, 2016, 2, dvw016.	1.8	16
22	Impact of Early Environment on Children's Mental Health: Lessons From DNA Methylation Studies With Monozygotic Twins. Twin Research and Human Genetics, 2015, 18, 623-634.	0.6	16
23	The developmental origins of chronic physical aggression: biological pathways triggered by early life adversity. Journal of Experimental Biology, 2015, 218, 123-133.	1.7	83
24	Epigenetics of Posttraumatic Stress Disorder: Current Evidence, Challenges, and Future Directions. Biological Psychiatry, 2015, 78, 327-335.	1.3	166
25	The effects of early life stress on the epigenome: From the womb to adulthood and even before. Experimental Neurology, 2015, 268, 10-20.	4.1	190
26	The neurobiological effects of stress as contributors to psychiatric disorders: focus on epigenetics. Current Opinion in Neurobiology, 2015, 30, 31-37.	4.2	55
27	DNA Methylation Signature of Childhood Chronic Physical Aggression in T Cells of Both Men and Women. PLoS ONE, 2014, 9, e86822.	2.5	81
28	Epigenetics in Posttraumatic Stress Disorder. Progress in Molecular Biology and Translational Science, 2014, 128, 29-50.	1.7	23
29	Hydroxymethylation and DNA methylation profiles in the prefrontal cortex of the non-human primate rhesus macaque and the impact of maternal deprivation on hydroxymethylation. Neuroscience, 2014, 268, 139-148.	2.3	52
30	Association of Childhood Chronic Physical Aggression with a DNA Methylation Signature in Adult Human T Cells. PLoS ONE, 2014, 9, e89839.	2.5	76
31	Childhood Chronic Physical Aggression Associates with Adult Cytokine Levels in Plasma. PLoS ONE, 2013, 8, e69481.	2.5	37
32	Differential DNA Methylation Regions in Cytokine and Transcription Factor Genomic Loci Associate with Childhood Physical Aggression. PLoS ONE, 2013, 8, e71691.	2.5	60
33	The Signature of Maternal Rearing in the Methylome in Rhesus Macaque Prefrontal Cortex and T Cells. Journal of Neuroscience, 2012, 32, 15626-15642.	3.6	340
34	Peripheral SLC6A4 DNA Methylation Is Associated with In Vivo Measures of Human Brain Serotonin Synthesis and Childhood Physical Aggression. PLoS ONE, 2012, 7, e39501.	2.5	181
35	Histone deacetylase inhibitor Trichostatin A induces global and gene-specific DNA demethylation in human cancer cell lines. Biochemical Pharmacology, 2007, 73, 1297-1307.	4.4	168
36	HOW DOES EARLY LIFE SOCIAL ENVIRONMENT SCULPT OUR GENES?. Biology of Reproduction, 2007, 77, 64-64.	2.7	1

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37	Analysis of a variable number tandem repeat polymorphism in the huntingtin interacting protein-1 related gene for anticipation in bipolar affective disorder. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2004, 28, 1299-1303.	4.8	2