

Anna Starnawska

List of Publications by Year in descending order

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Version: 2024-02-01

28
papers

1,076
citations

623734

14
h-index

642732

23
g-index

36
all docs

36
docs citations

36
times ranked

2097
citing authors

#	ARTICLE	IF	CITATIONS
1	IGHV-associated methylation signatures more accurately predict clinical outcomes of chronic lymphocytic leukemia patients than IGHV mutation load. <i>Haematologica</i> , 2022, 107, 877-886.	3.5	5
2	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. <i>Biological Psychiatry</i> , 2022, 91, 313-327.	1.3	114
3	Comparison of electrostatic and mechanical cell sorting with limited starting material. <i>Cytometry Part A: the Journal of the International Society for Analytical Cytology</i> , 2022, 101, 298-310.	1.5	3
4	Role of DNA methylation in aging-related cognitive functioning. , 2021, , 499-508.		0
5	Role of DNA Methylation in Mediating Genetic Risk of Psychiatric Disorders. <i>Frontiers in Psychiatry</i> , 2021, 12, 596821.	2.6	14
6	DNA methylation of the KLK8 gene in depression symptomatology. <i>Clinical Epigenetics</i> , 2021, 13, 200.	4.1	7
7	Genetics of suicide attempts in individuals with and without mental disorders: a population-based genome-wide association study. <i>Molecular Psychiatry</i> , 2020, 25, 2410-2421.	7.9	124
8	Neurospisin in mental health. <i>Journal of Physiological Sciences</i> , 2020, 70, 26.	2.1	7
9	Epigenome-wide association study of depression symptomatology in elderly monozygotic twins. <i>Translational Psychiatry</i> , 2019, 9, 214.	4.8	48
10	GWAS of Suicide Attempt in Psychiatric Disorders and Association With Major Depression Polygenic Risk Scores. <i>American Journal of Psychiatry</i> , 2019, 176, 651-660.	7.2	186
11	GENOME-WIDE METHYLOMIC ANALYSIS OF NEONATAL BLOOD FROM DANISH TWINS DISCORDANT FOR MENTAL ILLNESS. <i>European Neuropsychopharmacology</i> , 2019, 29, S794-S795.	0.7	0
12	ESTIMATED DNA METHYLATION GESTATIONAL AGE IN NEWBORN MONOZYGOTIC TWINS ASSOCIATE WITH LATER PSYCHIATRIC DISORDERS BETWEEN CON/DISCORDANT PAIRS. <i>European Neuropsychopharmacology</i> , 2019, 29, S795.	0.7	0
13	Evaluating the Feasibility of DNA Methylation Analyses Using Long-Term Archived Brain Formalin-Fixed Paraffin-Embedded Samples. <i>Molecular Neurobiology</i> , 2018, 55, 668-681.	4.0	6
14	Biological age of the endometrium using DNA methylation. <i>Reproduction</i> , 2018, 155, 165-170.	2.6	13
15	Blood DNA methylation age is not associated with cognitive functioning in middle-aged monozygotic twins. <i>Neurobiology of Aging</i> , 2017, 50, 60-63.	3.1	28
16	Differential DNA methylation at birth associated with mental disorder in individuals with 22q11.2 deletion syndrome. <i>Translational Psychiatry</i> , 2017, 7, e1221-e1221.	4.8	21
17	A MBD-seq protocol for large-scale methylome-wide studies with (very) low amounts of DNA. <i>Epigenetics</i> , 2017, 12, 743-750.	2.7	42
18	Saliva as a Blood Alternative for Genome-Wide DNA Methylation Profiling by Methylated DNA Immunoprecipitation (MeDIP) Sequencing. <i>Epigenomes</i> , 2017, 1, 14.	1.8	8

#	ARTICLE	IF	CITATIONS
19	Epigenome-Wide Association Study of Cognitive Functioning in Middle-Aged Monozygotic Twins. <i>Frontiers in Aging Neuroscience</i> , 2017, 9, 413.	3.4	52
20	The Genome-Wide DNA Methylation Profile of Peripheral Blood Is Not Systematically Changed by Short-Time Storage at Room Temperature. <i>Epigenomes</i> , 2017, 1, 23.	1.8	0
21	Lung function discordance in monozygotic twins and associated differences in blood DNA methylation. <i>Clinical Epigenetics</i> , 2017, 9, 132.	4.1	18
22	Lung function discordance in monozygotic twins and associated differences in blood DNA methylation., 2017,, .		0
23	DNA Methylation at the Neonatal State and at the Time of Diagnosis: Preliminary Support for an Association with the Estrogen Receptor 1, Gamma-Aminobutyric Acid B Receptor 1, and Myelin Oligodendrocyte Glycoprotein in Female Adolescent Patients with OCD. <i>Frontiers in Psychiatry</i> , 2016, 7, 35.	2.6	30
24	Hypomethylation of FAM63B in bipolar disorder patients. <i>Clinical Epigenetics</i> , 2016, 8, 52.	4.1	24
25	Identification of active denitrifiers in full-scale nutrient removal wastewater treatment systems. <i>Environmental Microbiology</i> , 2016, 18, 50-64.	3.8	226
26	Genome-wide DNA methylation profiling with MeDIP-seq using archived dried blood spots. <i>Clinical Epigenetics</i> , 2016, 8, 81.	4.1	36
27	CACNA1C hypermethylation is associated with bipolar disorder. <i>Translational Psychiatry</i> , 2016, 6, e831-e831.	4.8	39
28	A Novel Locus Harboring a Functional CD164 Nonsense Mutation Identified in a Large Danish Family with Nonsyndromic Hearing Impairment. <i>PLoS Genetics</i> , 2015, 11, e1005386.	3.5	18