

Christine SÃ¸holm Hansen

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5993225/publications.pdf>

Version: 2024-02-01

15
papers

10,231
citations

758635

12
h-index

940134

16
g-index

21
all docs

21
docs citations

21
times ranked

13567
citing authors

#	ARTICLE	IF	CITATIONS
1	Identifying the Common Genetic Basis of Antidepressant Response. <i>Biological Psychiatry Global Open Science</i> , 2022, 2, 115-126.	1.0	31
2	Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. <i>Nature Genetics</i> , 2021, 53, 817-829.	9.4	629
3	Genetic regulation of spermine oxidase activity and cancer risk: a Mendelian randomization study. <i>Scientific Reports</i> , 2021, 11, 17463.	1.6	1
4	The Genetics of the Mood Disorder Spectrum: Genome-wide Association Analyses of More Than 185,000 Cases and 439,000 Controls. <i>Biological Psychiatry</i> , 2020, 88, 169-184.	0.7	137
5	Large-Scale Exome Sequencing Study Implicates Both Developmental and Functional Changes in the Neurobiology of Autism. <i>Cell</i> , 2020, 180, 568-584.e23.	13.5	1,422
6	Identification of common genetic risk variants for autism spectrum disorder. <i>Nature Genetics</i> , 2019, 51, 431-444.	9.4	1,538
7	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. <i>Cell</i> , 2019, 179, 1469-1482.e11.	13.5	935
8	Discovery of the first genome-wide significant risk loci for attention deficit/hyperactivity disorder. <i>Nature Genetics</i> , 2019, 51, 63-75.	9.4	1,594
9	Common schizophrenia alleles are enriched in mutation-intolerant genes and in regions under strong background selection. <i>Nature Genetics</i> , 2018, 50, 381-389.	9.4	1,332
10	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. <i>Nature Genetics</i> , 2018, 50, 668-681.	9.4	2,224
11	RNA sequencing of archived neonatal dried blood spots. <i>Molecular Genetics and Metabolism Reports</i> , 2017, 10, 33-37.	0.4	8
12	Evaluation of whole genome amplified DNA to decrease material expenditure and increase quality. <i>Molecular Genetics and Metabolism Reports</i> , 2017, 11, 36-45.	0.4	9
13	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.	1.3	30
14	An epigenetic clock for gestational age at birth based on blood methylation data. <i>Genome Biology</i> , 2016, 17, 206.	3.8	193
15	Gene expression profiling of archived dried blood spot samples from the Danish Neonatal Screening Biobank. <i>Molecular Genetics and Metabolism</i> , 2015, 116, 119-124.	0.5	19