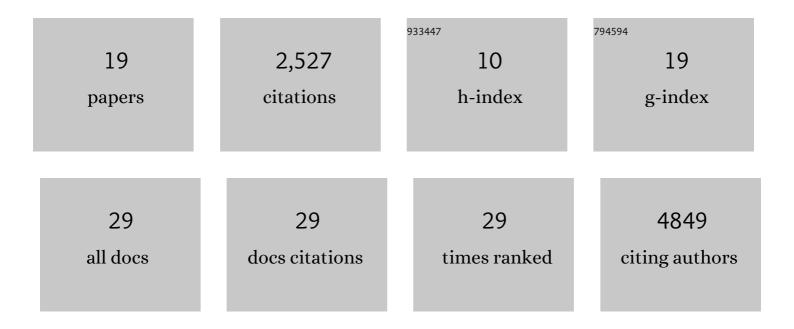
## Miruna C Barbu

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1901001/publications.pdf Version: 2024-02-01



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#	Article	IF	CITATIONS
1	Methylome-wide association study of early life stressors and adult mental health. Human Molecular Genetics, 2022, 31, 651-664.	2.9	7
2	Epigenome-wide association study of global cortical volumes in generation Scotland: Scottish family health study. Epigenetics, 2022, 17, 1143-1158.	2.7	3
3	DNA methylome-wide association study of genetic risk for depression implicates antigen processing and immune responses. Genome Medicine, 2022, 14, 36.	8.2	16
4	Methylome-wide association study of antidepressant use in Generation Scotland and the Netherlands Twin Register implicates the innate immune system. Molecular Psychiatry, 2022, 27, 1647-1657.	7.9	10
5	Structural neuroimaging measures and lifetime depression across levels of phenotyping in UK biobank. Translational Psychiatry, 2022, 12, 157.	4.8	7
6	Complex trait methylation scores in the prediction of major depressive disorder. EBioMedicine, 2022, 79, 104000.	6.1	4
7	Epigenetic prediction of major depressive disorder. Molecular Psychiatry, 2021, 26, 5112-5123.	7.9	44
8	Structural brain correlates of serum and epigenetic markers of inflammation in major depressive disorder. Brain, Behavior, and Immunity, 2021, 92, 39-48.	4.1	53
9	Grey and white matter associations of psychotic-like experiences in a general population sample (UK) Tj ETQq1 1	0.784314	၊ rggT /Overio
10	Hair glucocorticoids are associated with childhood adversity, depressive symptoms and reduced global and lobar grey matter in Generation Scotland. Translational Psychiatry, 2021, 11, 523.	4.8	13
11	Brain structural associations with depression in a large early adolescent sample (the ABCD study®). EClinicalMedicine, 2021, 42, 101204.	7.1	16
12	Automated classification of depression from structural brain measures across two independent communityâ€based cohorts. Human Brain Mapping, 2020, 41, 3922-3937.	3.6	27
13	Expression quantitative trait loci-derived scores and white matter microstructure in UK Biobank: a novel approach to integrating genetics and neuroimaging. Translational Psychiatry, 2020, 10, 55.	4.8	8
14	Associations between vascular risk factors and brain MRI indices in UK Biobank. European Heart Journal, 2019, 40, 2290-2300.	2.2	204
15	Association of Whole-Genome and NETRIN1 Signaling Pathway–Derived Polygenic Risk Scores for Major Depressive Disorder and White Matter Microstructure in the UK Biobank. Biological Psychiatry: Cognitive Neuroscience and Neuroimaging, 2019, 4, 91-100.	1.5	16
16	Reversal of proliferation deficits caused by chromosome 16p13.11 microduplication through targeting NFήB signaling: an integrated study of patient-derived neuronal precursor cells, cerebral organoids and in vivo brain imaging. Molecular Psychiatry, 2019, 24, 294-311.	7.9	36
17	Genome-wide meta-analysis of depression identifies 102 independent variants and highlights the importance of the prefrontal brain regions. Nature Neuroscience, 2019, 22, 343-352.	14.8	1,589
18	Genome-wide association study of depression phenotypes in UK Biobank identifies variants in excitatory synaptic pathways. Nature Communications, 2018, 9, 1470.	12.8	415

#	Article	IF	CITATIONS
19	Addendum: Genome-wide association study of depression phenotypes in UK Biobank identifies variants in excitatory synaptic pathways. Nature Communications, 2018, 9, 3578.	12.8	16