Margarita Rivera

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/9439590/publications.pdf

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70	11,283	81743	88477
papers	citations	39 h-index	70 g-index

77	77	77	13550
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Identifying the Common Genetic Basis of Antidepressant Response. Biological Psychiatry Global Open Science, 2022, 2, 115-126.	1.0	31
2	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. Biological Psychiatry, 2022, 91, 313-327.	0.7	114
3	Body mass index interacts with a genetic-risk score for depression increasing the risk of the disease in high-susceptibility individuals. Translational Psychiatry, 2022, 12, 30.	2.4	4
4	Interaction Effect between Physical Activity and the BDNF Val66Met Polymorphism on Depression in Women from the PISMA-ep Study. International Journal of Environmental Research and Public Health, 2022, 19, 2068.	1.2	9
5	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. Nature, 2022, 604, 502-508.	13.7	929
6	Association of polygenic score for major depression with response to lithium in patients with bipolar disorder. Molecular Psychiatry, 2021, 26, 2457-2470.	4.1	44
7	Bipolar multiplex families have an increased burden of common risk variants for psychiatric disorders. Molecular Psychiatry, 2021, 26, 1286-1298.	4.1	33
8	Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. Nature Genetics, 2021, 53, 817-829.	9.4	629
9	The role of the FTO gene in the relationship between depression and obesity. A systematic review. Neuroscience and Biobehavioral Reviews, 2021, 127, 630-637.	2.9	20
10	The Genetic Architecture of Depression in Individuals of East Asian Ancestry. JAMA Psychiatry, 2021, 78, 1258.	6.0	88
11	Investigating rare pathogenic/likely pathogenic exonic variation in bipolar disorder. Molecular Psychiatry, 2021, 26, 5239-5250.	4.1	15
12	EpidemiologÃa de la fobia social en AndalucÃa. Revista De PsiquiatrÃa Y Salud Mental, 2021, , .	1.0	0
13	Classical Human Leukocyte Antigen Alleles and C4 Haplotypes Are Not Significantly Associated With Depression. Biological Psychiatry, 2020, 87, 419-430.	0.7	27
14	The Genetics of the Mood Disorder Spectrum: Genome-wide Association Analyses of More Than 185,000 Cases and 439,000 Controls. Biological Psychiatry, 2020, 88, 169-184.	0.7	137
15	A phenome-wide association and Mendelian Randomisation study of polygenic risk for depression in UK Biobank. Nature Communications, 2020, 11, 2301.	5.8	81
16	Genome-wide gene-environment analyses of major depressive disorder and reported lifetime traumatic experiences in UK Biobank. Molecular Psychiatry, 2020, 25, 1430-1446.	4.1	116
17	Association of Polygenic Liabilities for Major Depression, Bipolar Disorder, and Schizophrenia With Risk for Depression in the Danish Population. JAMA Psychiatry, 2019, 76, 516.	6.0	78
18	Assessment of Bidirectional Relationships Between Physical Activity and Depression Among Adults. JAMA Psychiatry, 2019, 76, 399.	6.0	399

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19	Associations of major depressive disorder with chronic physical conditions, obesity and medication use: Results from the PISMA-ep study. European Psychiatry, 2019, 60, 20-27.	0.1	19
20	Genome-wide association study identifies 30 loci associated with bipolar disorder. Nature Genetics, 2019, 51, 793-803.	9.4	1,191
21	Genome-wide Burden of Rare Short Deletions Is Enriched in Major Depressive Disorder in Four Cohorts. Biological Psychiatry, 2019, 85, 1065-1073.	0.7	25
22	Physical exercise and body mass index as correlates of major depressive disorder in community-dwelling adults: Results from the PISMA-ep study. Journal of Affective Disorders, 2019, 251, 263-269.	2.0	14
23	Evidence of causal effect of major depression on alcohol dependence: findings from the psychiatric genomics consortium. Psychological Medicine, 2019, 49, 1218-1226.	2.7	74
24	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. Cell, 2019, 179, 1469-1482.e11.	13.5	935
25	Association of Whole-Genome and NETRIN1 Signaling Pathwayâ€"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.	1.1	16
26	Evidence for increased genetic risk load for major depression in patients assigned to electroconvulsive therapy. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2019, 180, 35-45.	1.1	18
27	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. Nature Genetics, 2018, 50, 668-681.	9.4	2,224
28	Does Childhood Trauma Moderate Polygenic Risk for Depression? A Meta-analysis of 5765 Subjects From the Psychiatric Genomics Consortium. Biological Psychiatry, 2018, 84, 138-147.	0.7	87
29	Genome-wide interaction study of a proxy for stress-sensitivity and its prediction of major depressive disorder. PLoS ONE, 2018, 13, e0209160.	1.1	14
30	Reduction in the levels of CoQ biosynthetic proteins is related to an increase in lifespan without evidence of hepatic mitohormesis. Scientific Reports, 2018, 8, 14013.	1.6	9
31	A Cross-Sectional Study on the Prevalence and Risk Correlates of Mental Disorders: The GRANADÎ£P Study. Journal of Nervous and Mental Disease, 2018, 206, 716-725.	0.5	8
32	Analysis of shared heritability in common disorders of the brain. Science, 2018, 360, .	6.0	1,085
33	Genome-wide Association for Major Depression Through Age at Onset Stratification: Major Depressive Disorder Working Group of the Psychiatric Genomics Consortium. Biological Psychiatry, 2017, 81, 325-335.	0.7	175
34	Genetic effects influencing risk for major depressive disorder in China and Europe. Translational Psychiatry, 2017, 7, e1074-e1074.	2.4	64
35	An Analysis of Two Genome-wide Association Meta-analyses Identifies a New Locus for Broad Depression Phenotype. Biological Psychiatry, 2017, 82, 322-329.	0.7	84
36	Genetic Association of Major Depression With Atypical Features and Obesity-Related Immunometabolic Dysregulations. JAMA Psychiatry, 2017, 74, 1214.	6.0	174

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37	Hair Cortisol in Twins: Heritability and Genetic Overlap with Psychological Variables and Stress-System Genes. Scientific Reports, 2017, 7, 15351.	1.6	50
38	Interaction between the FTO gene, body mass index and depression: meta-analysis of 13701 individuals. British Journal of Psychiatry, 2017, 211, 70-76.	1.7	49
39	Polygenic interactions with environmental adversity in the aetiology of major depressive disorder. Psychological Medicine, 2016, 46, 759-770.	2.7	176
40	Association of CRTC1 polymorphisms with obesity markers in subjects from the general population with lifetime depression. Journal of Affective Disorders, 2016, 198, 43-49.	2.0	18
41	Protocolo y metodologÃa del estudio epidemiolÃ³gico de la salud mental en AndalucÃa: PISMA-ep. Revista De PsiquiatrÃa Y Salud Mental, 2016, 9, 185-194.	1.0	17
42	Immune signatures and disorder-specific patterns in a cross-disorder gene expression analysis. British Journal of Psychiatry, 2016, 209, 202-208.	1.7	31
43	Genome-wide assessment of Parkinson's disease in a Southern Spanish population. Neurobiology of Aging, 2016, 45, 213.e3-213.e9.	1.5	35
44	Phenotypic Association Analyses With Copy Number Variation in Recurrent Depressive Disorder. Biological Psychiatry, 2016, 79, 329-336.	0.7	21
45	Familiality and SNP heritability of age at onset and episodicity in major depressive disorder. Psychological Medicine, 2015, 45, 2215-2225.	2.7	21
46	The interaction between stress and genetic factors in the etiopathogenesis of depression. World Psychiatry, 2015, 14, 161-163.	4.8	51
47	Epidemiological support for genetic variability at hypothalamic–pituitary–adrenal axis and serotonergic system as risk factors for major depression. Neuropsychiatric Disease and Treatment, 2015, 11, 2743.	1.0	21
48	A genetic risk score combining 32 SNPs is associated with body mass index and improves obesity prediction in people with major depressive disorder. BMC Medicine, 2015, 13, 86.	2.3	56
49	Molecular Signatures of Major Depression. Current Biology, 2015, 25, 1146-1156.	1.8	224
50	The successful search for genetic loci associated with depression. Genome Medicine, 2015, 7, 92.	3.6	3
51	The risk for major depression conferred by childhood maltreatment is multiplied by BDNF and SERT genetic vulnerability: a replication study. Journal of Psychiatry and Neuroscience, 2015, 40, 187-196.	1.4	41
52	Training and capacity building evaluation: Maximizing resources and results with Success Case Method. Evaluation and Program Planning, 2015, 52, 126-132.	0.9	15
53	Comorbid medical illness in bipolar disorder. British Journal of Psychiatry, 2014, 205, 465-472.	1.7	113
54	Genetic Studies of Major Depressive Disorder: Why Are There No Genome-wide Association Study Findings and What Can We Do About It?. Biological Psychiatry, 2014, 76, 510-512.	0.7	161

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55	Genetic relationships between suicide attempts, suicidal ideation and major psychiatric disorders: A genomeâ€wide association and polygenic scoring study. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2014, 165, 428-437.	1.1	99
56	Investigating the genetic variation underlying episodicity in major depressive disorder: Suggestive evidence for a bipolar contribution. Journal of Affective Disorders, 2014, 155, 81-89.	2.0	15
57	Relationship between obesity and the risk of clinically significant depression: Mendelian randomisation study. British Journal of Psychiatry, 2014, 205, 24-28.	1.7	62
58	Genome-wide association analysis of copy number variation in recurrent depressive disorder. Molecular Psychiatry, 2013, 18, 183-189.	4.1	45
59	Body mass index, but not FTO genotype or major depressive disorder, influences brain structure. Neuroscience, 2013, 252, 109-117.	1.1	40
60	Molecular genetic geneâ€“environment studies using candidate genes in schizophrenia: A systematic review. Schizophrenia Research, 2013, 150, 356-365.	1.1	80
61	The protective effect of the obesity-associated rs9939609 A variant in fat mass- and obesity-associated gene on depression. Molecular Psychiatry, 2013, 18, 1281-1286.	4.1	115
62	Depressive disorder moderates the effect of the FTO gene on body mass index. Molecular Psychiatry, 2012, 17, 604-611.	4.1	72
63	Pharmacogenetics of Response to Antipsychotics in Patients with Schizophrenia. CNS Drugs, 2011, 25, 933-969.	2.7	90
64	Polymorphic variation at the serotonin 1-A receptor gene is associated with comorbid depression and generalized anxiety. Psychiatric Genetics, 2011, 21, 195-201.	0.6	48
65	Genome-Wide Searches for Bipolar Disorder Genes. Current Psychiatry Reports, 2011, 13, 522-527.	2.1	12
66	Genome-Wide Association Study of Major Recurrent Depression in the U.K. Population. American Journal of Psychiatry, 2010, 167, 949-957.	4.0	221
67	Highâ€activity variants of the uMAOA polymorphism increase the risk for depression in a large primary care sample. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2009, 150B, 395-402.	1.1	44
68	Variabilidad en el gen COMT y modificaciÃ³n del riesgo de esquizofrenia conferido por consumo de cannabis. Revista De PsiquiatrÃa Y Salud Mental, 2009, 2, 89-94.	1.0	8
69	The risk for depression conferred by stressful life events is modified by variation at the serotonin transporter 5HTTLPR genotype: evidence from the Spanish PREDICT-Gene cohort. Molecular Psychiatry, 2007, 12, 748-755.	4.1	118
70	The 5-HTTLPR s/s genotype at the serotonin transporter gene (SLC6A4) increases the risk for depression in a large cohort of primary care attendees: The PREDICT-gene study. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2006, 141B, 912-917.	1.1	83