

Margarita Rivera

List of Publications by Citations

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71
papers

5,851
citations

33
h-index

76
g-index

77
ext. papers

8,819
ext. citations

9.8
avg. IF

4.51
L-index

#	Paper	IF	Citations
71	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. <i>Nature Genetics</i> , 2018 , 50, 668-681	36.3	1301
70	Analysis of shared heritability in common disorders of the brain. <i>Science</i> , 2018 , 360,	33.3	666
69	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , 2019 , 51, 793-803	36.3	662
68	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. <i>Cell</i> , 2019 , 179, 1469-1482.e11	56.2	402
67	Genome-wide association study of major recurrent depression in the U.K. population. <i>American Journal of Psychiatry</i> , 2010 , 167, 949-57	11.9	194
66	Assessment of Bidirectional Relationships Between Physical Activity and Depression Among Adults: A 2-Sample Mendelian Randomization Study. <i>JAMA Psychiatry</i> , 2019 , 76, 399-408	14.5	165
65	Molecular signatures of major depression. <i>Current Biology</i> , 2015 , 25, 1146-56	6.3	162
64	Genome-wide Association for Major Depression Through Age at Onset Stratification: Major Depressive Disorder Working Group of the Psychiatric Genomics Consortium. <i>Biological Psychiatry</i> , 2017 , 81, 325-335	7.9	129
63	Genetic studies of major depressive disorder: why are there no genome-wide association study findings and what can we do about it?. <i>Biological Psychiatry</i> , 2014 , 76, 510-2	7.9	125
62	Polygenic interactions with environmental adversity in the aetiology of major depressive disorder. <i>Psychological Medicine</i> , 2016 , 46, 759-70	6.9	125
61	Genetic Association of Major Depression With Atypical Features and Obesity-Related Immunometabolic Dysregulations. <i>JAMA Psychiatry</i> , 2017 , 74, 1214-1225	14.5	109
60	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. <i>Molecular Psychiatry</i> , 2007 , 12, 748-55	15.1	102
59	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	36.3	83
58	Comorbid medical illness in bipolar disorder. <i>British Journal of Psychiatry</i> , 2014 , 205, 465-72	5.4	80
57	Pharmacogenetics of response to antipsychotics in patients with schizophrenia. <i>CNS Drugs</i> , 2011 , 25, 933-69	6.7	78
56	The protective effect of the obesity-associated rs9939609 A variant in fat mass- and obesity-associated gene on depression. <i>Molecular Psychiatry</i> , 2013 , 18, 1281-6	15.1	75
55	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. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2006 , 141B, 912-7	3.5	72

54	Genetic relationships between suicide attempts, suicidal ideation and major psychiatric disorders: a genome-wide association and polygenic scoring study. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2014 , 165B, 428-37	3.5	70
53	An Analysis of Two Genome-wide Association Meta-analyses Identifies a New Locus for Broad Depression Phenotype. <i>Biological Psychiatry</i> , 2017 , 82, 322-329	7.9	68
52	Molecular genetic gene-environment studies using candidate genes in schizophrenia: a systematic review. <i>Schizophrenia Research</i> , 2013 , 150, 356-65	3.6	68
51	Depressive disorder moderates the effect of the FTO gene on body mass index. <i>Molecular Psychiatry</i> , 2012 , 17, 604-11	15.1	59
50	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	7.9	57
49	Association of Polygenic Liabilities for Major Depression, Bipolar Disorder, and Schizophrenia With Risk for Depression in the Danish Population. <i>JAMA Psychiatry</i> , 2019 , 76, 516-525	14.5	51
48	Genetic effects influencing risk for major depressive disorder in China and Europe. <i>Translational Psychiatry</i> , 2017 , 7, e1074	8.6	48
47	Does Childhood Trauma Moderate Polygenic Risk for Depression? A Meta-analysis of 5765 Subjects From the Psychiatric Genomics Consortium. <i>Biological Psychiatry</i> , 2018 , 84, 138-147	7.9	48
46	Relationship between obesity and the risk of clinically significant depression: Mendelian randomisation study. <i>British Journal of Psychiatry</i> , 2014 , 205, 24-8	5.4	48
45	Genome-wide gene-environment analyses of major depressive disorder and reported lifetime traumatic experiences in UK Biobank. <i>Molecular Psychiatry</i> , 2020 , 25, 1430-1446	15.1	47
44	A genetic risk score combining 32 SNPs is associated with body mass index and improves obesity prediction in people with major depressive disorder. <i>BMC Medicine</i> , 2015 , 13, 86	11.4	45
43	Genome-wide association analysis of copy number variation in recurrent depressive disorder. <i>Molecular Psychiatry</i> , 2013 , 18, 183-9	15.1	40
42	Polymorphic variation at the serotonin 1-A receptor gene is associated with comorbid depression and generalized anxiety. <i>Psychiatric Genetics</i> , 2011 , 21, 195-201	2.9	39
41	High-activity variants of the uMAOA polymorphism increase the risk for depression in a large primary care sample. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2009 , 150B, 395-402	3.5	36
40	Mapping genomic loci implicates genes and synaptic biology in schizophrenia.. <i>Nature</i> , 2022 ,	50.4	35
39	The risk for major depression conferred by childhood maltreatment is multiplied by BDNF and SERT genetic vulnerability: a replication study. <i>Journal of Psychiatry and Neuroscience</i> , 2015 , 40, 187-96	4.5	34
38	Evidence of causal effect of major depression on alcohol dependence: findings from the psychiatric genomics consortium. <i>Psychological Medicine</i> , 2019 , 49, 1218-1226	6.9	33
37	Hair Cortisol in Twins: Heritability and Genetic Overlap with Psychological Variables and Stress-System Genes. <i>Scientific Reports</i> , 2017 , 7, 15351	4.9	33

36	Interaction between the gene, body mass index and depression: meta-analysis of 13701 individuals. <i>British Journal of Psychiatry</i> , 2017 , 211, 70-76	5.4	33
35	Body mass index, but not FTO genotype or major depressive disorder, influences brain structure. <i>Neuroscience</i> , 2013 , 252, 109-17	3.9	32
34	A phenome-wide association and Mendelian Randomisation study of polygenic risk for depression in UK Biobank. <i>Nature Communications</i> , 2020 , 11, 2301	17.4	31
33	The interaction between stress and genetic factors in the etiopathogenesis of depression. <i>World Psychiatry</i> , 2015 , 14, 161-3	14.4	27
32	Immune signatures and disorder-specific patterns in a cross-disorder gene expression analysis. <i>British Journal of Psychiatry</i> , 2016 , 209, 202-8	5.4	23
31	Genome-wide assessment of Parkinson's disease in a Southern Spanish population. <i>Neurobiology of Aging</i> , 2016 , 45, 213.e3-213.e9	5.6	23
30	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depressive disorder		21
29	Familiality and SNP heritability of age at onset and episodicity in major depressive disorder. <i>Psychological Medicine</i> , 2015 , 45, 2215-25	6.9	18
28	Association of polygenic score for major depression with response to lithium in patients with bipolar disorder. <i>Molecular Psychiatry</i> , 2021 , 26, 2457-2470	15.1	17
27	Bipolar multiplex families have an increased burden of common risk variants for psychiatric disorders. <i>Molecular Psychiatry</i> , 2021 , 26, 1286-1298	15.1	17
26	Phenotypic Association Analyses With Copy Number Variation in Recurrent Depressive Disorder. <i>Biological Psychiatry</i> , 2016 , 79, 329-36	7.9	15
25	Association of CRT1 polymorphisms with obesity markers in subjects from the general population with lifetime depression. <i>Journal of Affective Disorders</i> , 2016 , 198, 43-9	6.6	15
24	Genome-wide Burden of Rare Short Deletions Is Enriched in Major Depressive Disorder in Four Cohorts. <i>Biological Psychiatry</i> , 2019 , 85, 1065-1073	7.9	14
23	Investigating the genetic variation underlying episodicity in major depressive disorder: suggestive evidence for a bipolar contribution. <i>Journal of Affective Disorders</i> , 2014 , 155, 81-9	6.6	13
22	Epidemiological support for genetic variability at hypothalamic-pituitary-adrenal axis and serotonergic system as risk factors for major depression. <i>Neuropsychiatric Disease and Treatment</i> , 2015 , 11, 2743-54	3.1	12
21	Protocol and methodology of Study epidemiological mental health in Andalusia: PISMA-ep. <i>Revista De Psiquiatría Y Salud Mental</i> , 2016 , 9, 185-194	2.7	12
20	Association of Whole-Genome and NETRIN1 Signaling Pathway-Derived Polygenic Risk Scores for Major Depressive Disorder and White Matter Microstructure in the UK Biobank. <i>Biological Psychiatry: Cognitive Neuroscience and Neuroimaging</i> , 2019 , 4, 91-100	3.4	12
19	Associations of major depressive disorder with chronic physical conditions, obesity and medication use: Results from the PISMA-ep study. <i>European Psychiatry</i> , 2019 , 60, 20-27	6	11

18	Genome-wide searches for bipolar disorder genes. <i>Current Psychiatry Reports</i> , 2011 , 13, 522-7	9.1	11
17	Genome-wide association study of over 40,000 bipolar disorder cases provides new insights into the underlying biology		11
16	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. <i>Biological Psychiatry</i> , 2021 ,	7.9	11
15	Evidence for increased genetic risk load for major depression in patients assigned to electroconvulsive therapy. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2019 , 180, 35-45	3.5	10
14	Training and capacity building evaluation: Maximizing resources and results with Success Case Method. <i>Evaluation and Program Planning</i> , 2015 , 52, 126-32	1.7	9
13	Classical Human Leukocyte Antigen Alleles and C4 Haplotypes Are Not Significantly Associated With Depression. <i>Biological Psychiatry</i> , 2020 , 87, 419-430	7.9	9
12	Variability in the COMT gene and modification of the risk of schizophrenia conferred by cannabis consumption. <i>Revista De Psiquiatria Y Salud Mental</i> , 2009 , 2, 89-94	2.7	8
11	Physical exercise and body mass index as correlates of major depressive disorder in community-dwelling adults: Results from the PISMA-ep study. <i>Journal of Affective Disorders</i> , 2019 , 251, 263-269	6.6	7
10	The Genetic Architecture of Depression in Individuals of East Asian Ancestry: A Genome-Wide Association Study. <i>JAMA Psychiatry</i> , 2021 , 78, 1258-1269	14.5	7
9	Genome-wide interaction study of a proxy for stress-sensitivity and its prediction of major depressive disorder. <i>PLoS ONE</i> , 2018 , 13, e0209160	3.7	6
8	Reduction in the levels of CoQ biosynthetic proteins is related to an increase in lifespan without evidence of hepatic mitohormesis. <i>Scientific Reports</i> , 2018 , 8, 14013	4.9	6
7	A Cross-Sectional Study on the Prevalence and Risk Correlates of Mental Disorders: The GRANADP Study. <i>Journal of Nervous and Mental Disease</i> , 2018 , 206, 716-725	1.8	5
6	Identifying the Common Genetic Basis of Antidepressant Response. <i>Biological Psychiatry Global Open Science</i> , 2021 ,		4
5	The successful search for genetic loci associated with depression. <i>Genome Medicine</i> , 2015 , 7, 92	14.4	3
4	Investigating rare pathogenic/likely pathogenic exonic variation in bipolar disorder. <i>Molecular Psychiatry</i> , 2021 , 26, 5239-5250	15.1	3
3	Dissecting the shared genetic architecture of suicide attempt, psychiatric disorders and known risk factors		2
2	The role of the FTO gene in the relationship between depression and obesity. A systematic review. <i>Neuroscience and Biobehavioral Reviews</i> , 2021 , 127, 630-637	9	2
1	Body mass index interacts with a genetic-risk score for depression increasing the risk of the disease in high-susceptibility individuals.. <i>Translational Psychiatry</i> , 2022 , 12, 30	8.6	1

