

# Margarita Rivera

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

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	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> , <b>2022</b> , 12, 30	8.6	1
70	Mapping genomic loci implicates genes and synaptic biology in schizophrenia.. <i>Nature</i> , <b>2022</b> ,	50.4	35
69	Association of polygenic score for major depression with response to lithium in patients with bipolar disorder. <i>Molecular Psychiatry</i> , <b>2021</b> , 26, 2457-2470	15.1	17
68	Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. <i>Nature Genetics</i> , <b>2021</b> , 53, 817-829	36.3	83
67	Bipolar multiplex families have an increased burden of common risk variants for psychiatric disorders. <i>Molecular Psychiatry</i> , <b>2021</b> , 26, 1286-1298	15.1	17
66	The role of the FTO gene in the relationship between depression and obesity. A systematic review. <i>Neuroscience and Biobehavioral Reviews</i> , <b>2021</b> , 127, 630-637	9	2
65	Identifying the Common Genetic Basis of Antidepressant Response. <i>Biological Psychiatry Global Open Science</i> , <b>2021</b> ,		4
64	The Genetic Architecture of Depression in Individuals of East Asian Ancestry: A Genome-Wide Association Study. <i>JAMA Psychiatry</i> , <b>2021</b> , 78, 1258-1269	14.5	7
63	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. <i>Biological Psychiatry</i> , <b>2021</b> ,	7.9	11
62	Investigating rare pathogenic/likely pathogenic exonic variation in bipolar disorder. <i>Molecular Psychiatry</i> , <b>2021</b> , 26, 5239-5250	15.1	3
61	A phenome-wide association and Mendelian Randomisation study of polygenic risk for depression in UK Biobank. <i>Nature Communications</i> , <b>2020</b> , 11, 2301	17.4	31
60	Genome-wide gene-environment analyses of major depressive disorder and reported lifetime traumatic experiences in UK Biobank. <i>Molecular Psychiatry</i> , <b>2020</b> , 25, 1430-1446	15.1	47
59	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> , <b>2020</b> , 88, 169-184	7.9	57
58	Classical Human Leukocyte Antigen Alleles and C4 Haplotypes Are Not Significantly Associated With Depression. <i>Biological Psychiatry</i> , <b>2020</b> , 87, 419-430	7.9	9
57	Association of Polygenic Liabilities for Major Depression, Bipolar Disorder, and Schizophrenia With Risk for Depression in the Danish Population. <i>JAMA Psychiatry</i> , <b>2019</b> , 76, 516-525	14.5	51
56	Assessment of Bidirectional Relationships Between Physical Activity and Depression Among Adults: A 2-Sample Mendelian Randomization Study. <i>JAMA Psychiatry</i> , <b>2019</b> , 76, 399-408	14.5	165
55	Associations of major depressive disorder with chronic physical conditions, obesity and medication use: Results from the PISMA-ep study. <i>European Psychiatry</i> , <b>2019</b> , 60, 20-27	6	11

54	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , <b>2019</b> , 51, 793-803	36.3	662
53	Genome-wide Burden of Rare Short Deletions Is Enriched in Major Depressive Disorder in Four Cohorts. <i>Biological Psychiatry</i> , <b>2019</b> , 85, 1065-1073	7.9	14
52	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> , <b>2019</b> , 251, 263-269	6.6	7
51	Evidence of causal effect of major depression on alcohol dependence: findings from the psychiatric genomics consortium. <i>Psychological Medicine</i> , <b>2019</b> , 49, 1218-1226	6.9	33
50	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. <i>Cell</i> , <b>2019</b> , 179, 1469-1482.e11	56.2	402
49	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> , <b>2019</b> , 4, 91-100	3.4	12
48	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> , <b>2019</b> , 180, 35-45	3.5	10
47	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. <i>Nature Genetics</i> , <b>2018</b> , 50, 668-681	36.3	1301
46	Does Childhood Trauma Moderate Polygenic Risk for Depression? A Meta-analysis of 5765 Subjects From the Psychiatric Genomics Consortium. <i>Biological Psychiatry</i> , <b>2018</b> , 84, 138-147	7.9	48
45	Analysis of shared heritability in common disorders of the brain. <i>Science</i> , <b>2018</b> , 360,	33.3	666
44	Genome-wide interaction study of a proxy for stress-sensitivity and its prediction of major depressive disorder. <i>PLoS ONE</i> , <b>2018</b> , 13, e0209160	3.7	6
43	Reduction in the levels of CoQ biosynthetic proteins is related to an increase in lifespan without evidence of hepatic mitohormesis. <i>Scientific Reports</i> , <b>2018</b> , 8, 14013	4.9	6
42	A Cross-Sectional Study on the Prevalence and Risk Correlates of Mental Disorders: The GRANADP Study. <i>Journal of Nervous and Mental Disease</i> , <b>2018</b> , 206, 716-725	1.8	5
41	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> , <b>2017</b> , 81, 325-335	7.9	129
40	Genetic effects influencing risk for major depressive disorder in China and Europe. <i>Translational Psychiatry</i> , <b>2017</b> , 7, e1074	8.6	48
39	An Analysis of Two Genome-wide Association Meta-analyses Identifies a New Locus for Broad Depression Phenotype. <i>Biological Psychiatry</i> , <b>2017</b> , 82, 322-329	7.9	68
38	Genetic Association of Major Depression With Atypical Features and Obesity-Related Immunometabolic Dysregulations. <i>JAMA Psychiatry</i> , <b>2017</b> , 74, 1214-1225	14.5	109
37	Hair Cortisol in Twins: Heritability and Genetic Overlap with Psychological Variables and Stress-System Genes. <i>Scientific Reports</i> , <b>2017</b> , 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> , <b>2017</b> , 211, 70-76	5.4	33
35	Phenotypic Association Analyses With Copy Number Variation in Recurrent Depressive Disorder. <i>Biological Psychiatry</i> , <b>2016</b> , 79, 329-36	7.9	15
34	Immune signatures and disorder-specific patterns in a cross-disorder gene expression analysis. <i>British Journal of Psychiatry</i> , <b>2016</b> , 209, 202-8	5.4	23
33	Genome-wide assessment of Parkinson's disease in a Southern Spanish population. <i>Neurobiology of Aging</i> , <b>2016</b> , 45, 213.e3-213.e9	5.6	23
32	Polygenic interactions with environmental adversity in the aetiology of major depressive disorder. <i>Psychological Medicine</i> , <b>2016</b> , 46, 759-70	6.9	125
31	Association of CRTC1 polymorphisms with obesity markers in subjects from the general population with lifetime depression. <i>Journal of Affective Disorders</i> , <b>2016</b> , 198, 43-9	6.6	15
30	Protocol and methodology of Study epidemiological mental health in Andalusia: PISMA-ep. <i>Revista De Psiquiatría Y Salud Mental</i> , <b>2016</b> , 9, 185-194	2.7	12
29	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> , <b>2015</b> , 40, 187-96	4.5	34
28	Training and capacity building evaluation: Maximizing resources and results with Success Case Method. <i>Evaluation and Program Planning</i> , <b>2015</b> , 52, 126-32	1.7	9
27	Familiality and SNP heritability of age at onset and episodicity in major depressive disorder. <i>Psychological Medicine</i> , <b>2015</b> , 45, 2215-25	6.9	18
26	The interaction between stress and genetic factors in the etiopathogenesis of depression. <i>World Psychiatry</i> , <b>2015</b> , 14, 161-3	14.4	27
25	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> , <b>2015</b> , 11, 2743-54	3.1	12
24	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> , <b>2015</b> , 13, 86	11.4	45
23	Molecular signatures of major depression. <i>Current Biology</i> , <b>2015</b> , 25, 1146-56	6.3	162
22	The successful search for genetic loci associated with depression. <i>Genome Medicine</i> , <b>2015</b> , 7, 92	14.4	3
21	Comorbid medical illness in bipolar disorder. <i>British Journal of Psychiatry</i> , <b>2014</b> , 205, 465-72	5.4	80
20	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> , <b>2014</b> , 76, 510-2	7.9	125
19	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> , <b>2014</b> , 165B, 428-37	3.5	70

18	Investigating the genetic variation underlying episodicity in major depressive disorder: suggestive evidence for a bipolar contribution. <i>Journal of Affective Disorders</i> , <b>2014</b> , 155, 81-9	6.6	13
17	Relationship between obesity and the risk of clinically significant depression: Mendelian randomisation study. <i>British Journal of Psychiatry</i> , <b>2014</b> , 205, 24-8	5.4	48
16	Genome-wide association analysis of copy number variation in recurrent depressive disorder. <i>Molecular Psychiatry</i> , <b>2013</b> , 18, 183-9	15.1	40
15	Body mass index, but not FTO genotype or major depressive disorder, influences brain structure. <i>Neuroscience</i> , <b>2013</b> , 252, 109-17	3.9	32
14	Molecular genetic gene-environment studies using candidate genes in schizophrenia: a systematic review. <i>Schizophrenia Research</i> , <b>2013</b> , 150, 356-65	3.6	68
13	The protective effect of the obesity-associated rs9939609 A variant in fat mass- and obesity-associated gene on depression. <i>Molecular Psychiatry</i> , <b>2013</b> , 18, 1281-6	15.1	75
12	Depressive disorder moderates the effect of the FTO gene on body mass index. <i>Molecular Psychiatry</i> , <b>2012</b> , 17, 604-11	15.1	59
11	Pharmacogenetics of response to antipsychotics in patients with schizophrenia. <i>CNS Drugs</i> , <b>2011</b> , 25, 933-69	6.7	78
10	Polymorphic variation at the serotonin 1-A receptor gene is associated with comorbid depression and generalized anxiety. <i>Psychiatric Genetics</i> , <b>2011</b> , 21, 195-201	2.9	39
9	Genome-wide searches for bipolar disorder genes. <i>Current Psychiatry Reports</i> , <b>2011</b> , 13, 522-7	9.1	11
8	Genome-wide association study of major recurrent depression in the U.K. population. <i>American Journal of Psychiatry</i> , <b>2010</b> , 167, 949-57	11.9	194
7	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> , <b>2009</b> , 150B, 395-402	3.5	36
6	Variability in the COMT gene and modification of the risk of schizophrenia conferred by cannabis consumption. <i>Revista De Psiquiatria Y Salud Mental</i> , <b>2009</b> , 2, 89-94	2.7	8
5	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> , <b>2007</b> , 12, 748-55	15.1	102
4	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> , <b>2006</b> , 141B, 912-7	3.5	72
3	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depressive disorder		21
2	Genome-wide association study of over 40,000 bipolar disorder cases provides new insights into the underlying biology		11
1	Dissecting the shared genetic architecture of suicide attempt, psychiatric disorders and known risk factors		2

