Marcella D C Rietschel

List of Publications by Citations

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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.

729 papers

45,223 citations

102 h-index 185 g-index

805 ext. papers

53,973 ext. citations

7.7 avg, IF

6.31 L-index

#	Paper	IF	Citations
729	Large recurrent microdeletions associated with schizophrenia. <i>Nature</i> , 2008 , 455, 232-6	50.4	1427
728	Common variants conferring risk of schizophrenia. <i>Nature</i> , 2009 , 460, 744-7	50.4	1350
727	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
726	City living and urban upbringing affect neural social stress processing in humans. <i>Nature</i> , 2011 , 474, 49	8-504	902
725	Identification of loci associated with schizophrenia by genome-wide association and follow-up. <i>Nature Genetics</i> , 2008 , 40, 1053-5	36.3	877
724	A mega-analysis of genome-wide association studies for major depressive disorder. <i>Molecular Psychiatry</i> , 2013 , 18, 497-511	15.1	853
723	Analysis of shared heritability in common disorders of the brain. <i>Science</i> , 2018 , 360,	33.3	666
722	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , 2019 , 51, 793-803	36.3	662
721	Common genetic variants influence human subcortical brain structures. <i>Nature</i> , 2015 , 520, 224-9	50.4	601
720	Microduplications of 16p11.2 are associated with schizophrenia. <i>Nature Genetics</i> , 2009 , 41, 1223-7	36.3	550
719	A genome-wide association study implicates diacylglycerol kinase eta (DGKH) and several other genes in the etiology of bipolar disorder. <i>Molecular Psychiatry</i> , 2008 , 13, 197-207	15.1	548
718	The ENIGMA Consortium: large-scale collaborative analyses of neuroimaging and genetic data. <i>Brain Imaging and Behavior</i> , 2014 , 8, 153-82	4.1	539
717	Contribution of copy number variants to schizophrenia from a genome-wide study of 41,321 subjects. <i>Nature Genetics</i> , 2017 , 49, 27-35	36.3	530
716	Identification of common variants associated with human hippocampal and intracranial volumes. <i>Nature Genetics</i> , 2012 , 44, 552-61	36.3	498
7 ¹ 5	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. <i>Cell</i> , 2019 , 179, 1469-1482.e11	56.2	402
714	The IMAGEN study: reinforcement-related behaviour in normal brain function and psychopathology. <i>Molecular Psychiatry</i> , 2010 , 15, 1128-39	15.1	401
713	Disruption of the neurexin 1 gene is associated with schizophrenia. <i>Human Molecular Genetics</i> , 2009 , 18, 988-96	5.6	376

(2011-2010)

712	A genome-wide association study of alcohol dependence. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010 , 107, 5082-7	11.5	367
711	BRAIN NETWORKS. Correlated gene expression supports synchronous activity in brain networks. <i>Science</i> , 2015 , 348, 1241-4	33.3	355
710	Genome scan meta-analysis of schizophrenia and bipolar disorder, part III: Bipolar disorder. <i>American Journal of Human Genetics</i> , 2003 , 73, 49-62	11	353
709	Neural mechanisms of a genome-wide supported psychosis variant. <i>Science</i> , 2009 , 324, 605	33.3	329
708	Genome-wide association analyses identify new risk variants and the genetic architecture of amyotrophic lateral sclerosis. <i>Nature Genetics</i> , 2016 , 48, 1043-8	36.3	328
707	Genome-wide association for major depressive disorder: a possible role for the presynaptic protein piccolo. <i>Molecular Psychiatry</i> , 2009 , 14, 359-75	15.1	322
706	Genome-wide association study of alcohol dependence. Archives of General Psychiatry, 2009, 66, 773-84		318
705	Support for association of schizophrenia with genetic variation in the 6p22.3 gene, dysbindin, in sib-pair families with linkage and in an additional sample of triad families. <i>American Journal of Human Genetics</i> , 2003 , 72, 185-90	11	318
704	Adolescent impulsivity phenotypes characterized by distinct brain networks. <i>Nature Neuroscience</i> , 2012 , 15, 920-5	25.5	304
703	A genome-wide association study confirms PNPLA3 and identifies TM6SF2 and MBOAT7 as risk loci for alcohol-related cirrhosis. <i>Nature Genetics</i> , 2015 , 47, 1443-8	36.3	303
702	Neuropsychosocial profiles of current and future adolescent alcohol misusers. <i>Nature</i> , 2014 , 512, 185-9	50.4	296
701	Multiple independent loci at chromosome 15q25.1 affect smoking quantity: a meta-analysis and comparison with lung cancer and COPD. <i>PLoS Genetics</i> , 2010 , 6, e1001053	6	291
700	Genome-wide pharmacogenetics of antidepressant response in the GENDEP project. <i>American Journal of Psychiatry</i> , 2010 , 167, 555-64	11.9	280
699	Examination of G72 and D-amino-acid oxidase as genetic risk factors for schizophrenia and bipolar affective disorder. <i>Molecular Psychiatry</i> , 2004 , 9, 203-7	15.1	270
698	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. <i>Nature Neuroscience</i> , 2018 , 21, 1656-1669	25.5	257
697	Evidence for a relationship between genetic variants at the brain-derived neurotrophic factor (BDNF) locus and major depression. <i>Biological Psychiatry</i> , 2005 , 58, 307-14	7.9	255
696	Genome-wide association study reveals two new risk loci for bipolar disorder. <i>Nature Communications</i> , 2014 , 5, 3339	17.4	248
695	High frequencies of de novo CNVs in bipolar disorder and schizophrenia. <i>Neuron</i> , 2011 , 72, 951-63	13.9	240

694	Depression symptom dimensions as predictors of antidepressant treatment outcome: replicable evidence for interest-activity symptoms. <i>Psychological Medicine</i> , 2012 , 42, 967-80	6.9	229
693	Psychiatric comorbidity and functional impairment in a clinically referred sample of adults with attention-deficit/hyperactivity disorder (ADHD). <i>European Archives of Psychiatry and Clinical Neuroscience</i> , 2007 , 257, 371-7	5.1	223
692	Genome-wide association study identifies genetic variation in neurocan as a susceptibility factor for bipolar disorder. <i>American Journal of Human Genetics</i> , 2011 , 88, 372-81	11	221
691	Genetic variants associated with response to lithium treatment in bipolar disorder: a genome-wide association study. <i>Lancet, The</i> , 2016 , 387, 1085-1093	40	216
690	Genetic variation in the PNPLA3 gene is associated with alcoholic liver injury in caucasians. <i>Hepatology</i> , 2011 , 53, 86-95	11.2	202
689	Common variants on 8p12 and 1q24.2 confer risk of schizophrenia. <i>Nature Genetics</i> , 2011 , 43, 1224-7	36.3	201
688	Copy number variations of chromosome 16p13.1 region associated with schizophrenia. <i>Molecular Psychiatry</i> , 2011 , 16, 17-25	15.1	194
687	Measuring depression: comparison and integration of three scales in the GENDEP study. <i>Psychological Medicine</i> , 2008 , 38, 289-300	6.9	193
686	Three circadian clock genes Per2, Arntl, and Npas2 contribute to winter depression. <i>Annals of Medicine</i> , 2007 , 39, 229-38	1.5	193
685	Pharmacogenetics of tardive dyskinesia: combined analysis of 780 patients supports association with dopamine D3 receptor gene Ser9Gly polymorphism. <i>Neuropsychopharmacology</i> , 2002 , 27, 105-19	8.7	193
684	Combined analysis from eleven linkage studies of bipolar disorder provides strong evidence of susceptibility loci on chromosomes 6q and 8q. <i>American Journal of Human Genetics</i> , 2005 , 77, 582-95	11	192
683	Age at onset in bipolar I affective disorder: further evidence for three subgroups. <i>American Journal of Psychiatry</i> , 2003 , 160, 999-1001	11.9	187
682	Systematic screening for mutations in the human serotonin-2A (5-HT2A) receptor gene: identification of two naturally occurring receptor variants and association analysis in schizophrenia. <i>Human Genetics</i> , 1996 , 97, 614-9	6.3	181
681	Lower ventral striatal activation during reward anticipation in adolescent smokers. <i>American Journal of Psychiatry</i> , 2011 , 168, 540-9	11.9	175
68o	Novel genetic loci associated with hippocampal volume. <i>Nature Communications</i> , 2017 , 8, 13624	17.4	173
679	Genetic predictors of response to antidepressants in the GENDEP project. <i>Pharmacogenomics Journal</i> , 2009 , 9, 225-33	3.5	172
678	The DTNBP1 (dysbindin) gene contributes to schizophrenia, depending on family history of the disease. <i>American Journal of Human Genetics</i> , 2003 , 73, 1438-43	11	167
677	Common variants at VRK2 and TCF4 conferring risk of schizophrenia. <i>Human Molecular Genetics</i> , 2011 , 20, 4076-81	5.6	162

(2010-2009)

676	Adverse reactions to antidepressants. British Journal of Psychiatry, 2009, 195, 202-10	5.4	162
675	The neuronal transporter gene SLC6A15 confers risk to major depression. <i>Neuron</i> , 2011 , 70, 252-65	13.9	161
674	Gene variants associated with schizophrenia in a Norwegian genome-wide study are replicated in a large European cohort. <i>Journal of Psychiatric Research</i> , 2010 , 44, 748-53	5.2	161
673	Association of mouse Dlg4 (PSD-95) gene deletion and human DLG4 gene variation with phenotypes relevant to autism spectrum disorders and Williams' syndrome. <i>American Journal of Psychiatry</i> , 2010 , 167, 1508-17	11.9	159
672	The structure of psychopathology in adolescence and its common personality and cognitive correlates. <i>Journal of Abnormal Psychology</i> , 2016 , 125, 1039-1052	7	158
671	Genetic association of the human corticotropin releasing hormone receptor 1 (CRHR1) with binge drinking and alcohol intake patterns in two independent samples. <i>Molecular Psychiatry</i> , 2006 , 11, 594-60	0 2 5.1	158
670	The genetic architecture of the human cerebral cortex. <i>Science</i> , 2020 , 367,	33.3	156
669	Genome-wide association study meta-analysis of European and Asian-ancestry samples identifies three novel loci associated with bipolar disorder. <i>Molecular Psychiatry</i> , 2013 , 18, 195-205	15.1	155
668	A genome-wide autosomal screen for schizophrenia susceptibility loci in 71 families with affected siblings: support for loci on chromosome 10p and 6. <i>Molecular Psychiatry</i> , 2000 , 5, 638-49	15.1	152
667	Differential efficacy of escitalopram and nortriptyline on dimensional measures of depression. <i>British Journal of Psychiatry</i> , 2009 , 194, 252-9	5.4	150
666	Brain function in carriers of a genome-wide supported bipolar disorder variant. <i>Archives of General Psychiatry</i> , 2010 , 67, 803-11		147
665	Two variants in Ankyrin 3 (ANK3) are independent genetic risk factors for bipolar disorder. <i>Molecular Psychiatry</i> , 2009 , 14, 487-91	15.1	147
664	Association between a functional polymorphism in the monoamine oxidase A gene promoter and major depressive disorder. <i>American Journal of Medical Genetics Part A</i> , 2000 , 96, 801-803		147
663	Novel genetic loci underlying human intracranial volume identified through genome-wide association. <i>Nature Neuroscience</i> , 2016 , 19, 1569-1582	25.5	147
662	Genome-wide association-, replication-, and neuroimaging study implicates HOMER1 in the etiology of major depression. <i>Biological Psychiatry</i> , 2010 , 68, 578-85	7.9	139
661	Variability of 5-HT2C receptor cys23ser polymorphism among European populations and vulnerability to affective disorder. <i>Molecular Psychiatry</i> , 2001 , 6, 579-85	15.1	138
660	Cloning, genomic organization, alternative transcripts and mutational analysis of the gene responsible for autosomal recessive universal congenital alopecia. <i>Human Molecular Genetics</i> , 1998 , 7, 1671-9	5.6	136
659	Meta-analysis of genome-wide association data identifies a risk locus for major mood disorders on 3p21.1. <i>Nature Genetics</i> , 2010 , 42, 128-31	36.3	135

658	Genome-wide study of association and interaction with maternal cytomegalovirus infection suggests new schizophrenia loci. <i>Molecular Psychiatry</i> , 2014 , 19, 325-33	15.1	133
657	Genome-wide significant association between alcohol dependence and a variant in the ADH gene cluster. <i>Addiction Biology</i> , 2012 , 17, 171-80	4.6	133
656	Whole genome linkage scan of recurrent depressive disorder from the depression network study. <i>Human Molecular Genetics</i> , 2005 , 14, 3337-45	5.6	133
655	Familial occurrence of primary premature ejaculation. <i>Psychiatric Genetics</i> , 1998 , 8, 37-40	2.9	133
654	Expanding the range of ZNF804A variants conferring risk of psychosis. <i>Molecular Psychiatry</i> , 2011 , 16, 59-66	15.1	129
653	Association of the functional V158M catechol-O-methyl-transferase polymorphism with panic disorder in women. <i>International Journal of Neuropsychopharmacology</i> , 2004 , 7, 183-8	5.8	129
652	Association between C-reactive protein (CRP) with depression symptom severity and specific depressive symptoms in major depression. <i>Brain, Behavior, and Immunity,</i> 2017 , 62, 344-350	16.6	128
651	Genome-wide association study of 40,000 individuals identifies two novel loci associated with bipolar disorder. <i>Human Molecular Genetics</i> , 2016 , 25, 3383-3394	5.6	125
650	The complement control-related genes CSMD1 and CSMD2 associate to schizophrenia. <i>Biological Psychiatry</i> , 2011 , 70, 35-42	7.9	124
649	Moderation of antidepressant response by the serotonin transporter gene. <i>British Journal of Psychiatry</i> , 2009 , 195, 30-8	5.4	124
648	Impact of polymorphisms of cytochrome-P450 isoenzymes 2C9, 2C19 and 2D6 on plasma concentrations and clinical effects of antidepressants in a naturalistic clinical setting. <i>European Journal of Clinical Pharmacology</i> , 2004 , 60, 329-36	2.8	124
647	Risk taking and the adolescent reward system: a potential common link to substance abuse. <i>American Journal of Psychiatry</i> , 2012 , 169, 39-46	11.9	123
646	Effects of a genome-wide supported psychosis risk variant on neural activation during a theory-of-mind task. <i>Molecular Psychiatry</i> , 2011 , 16, 462-70	15.1	122
645	The power of sample size and homogenous sampling: association between the 5-HTTLPR serotonin transporter polymorphism and major depressive disorder. <i>Biological Psychiatry</i> , 2005 , 57, 247-51	7.9	121
644	Evaluation of linkage of bipolar affective disorder to chromosome 18 in a sample of 57 German families. <i>Molecular Psychiatry</i> , 1999 , 4, 76-84	15.1	121
643	TMEM132D, a new candidate for anxiety phenotypes: evidence from human and mouse studies. <i>Molecular Psychiatry</i> , 2011 , 16, 647-63	15.1	120
642	Association between COMT (Val158Met) functional polymorphism and early onset in patients with major depressive disorder in a European multicenter genetic association study. <i>Molecular Psychiatry</i> , 2005 , 10, 598-605	15.1	119
641	Impact of age at first drink on vulnerability to alcohol-related problems: testing the marker hypothesis in a prospective study of young adults. <i>Journal of Psychiatric Research</i> , 2009 , 43, 1205-12	5.2	118

640	A genome-wide association study of attempted suicide. <i>Molecular Psychiatry</i> , 2012 , 17, 433-44	15.1	117	
639	Genotype-phenotype studies in bipolar disorder showing association between the DAOA/G30 locus and persecutory delusions: a first step toward a molecular genetic classification of psychiatric phenotypes. <i>American Journal of Psychiatry</i> , 2005 , 162, 2101-8	11.9	116	
638	Early Cannabis Use, Polygenic Risk Score for Schizophrenia and Brain Maturation in Adolescence. <i>JAMA Psychiatry</i> , 2015 , 72, 1002-11	14.5	115	
637	The neural basis of video gaming. <i>Translational Psychiatry</i> , 2011 , 1, e53	8.6	114	
636	Genome-wide association study of suicide attempts in mood disorder patients. <i>American Journal of Psychiatry</i> , 2010 , 167, 1499-507	11.9	113	
635	A genome screen for genes predisposing to bipolar affective disorder detects a new susceptibility locus on 8q. <i>Human Molecular Genetics</i> , 2001 , 10, 2933-44	5.6	113	
634	Assessment of Response to Lithium Maintenance Treatment in Bipolar Disorder: A Consortium on Lithium Genetics (ConLiGen) Report. <i>PLoS ONE</i> , 2013 , 8, e65636	3.7	113	
633	Genetic Association of Major Depression With Atypical Features and Obesity-Related Immunometabolic Dysregulations. <i>JAMA Psychiatry</i> , 2017 , 74, 1214-1225	14.5	109	
632	Combining clinical variables to optimize prediction of antidepressant treatment outcomes. <i>Journal of Psychiatric Research</i> , 2016 , 78, 94-102	5.2	106	
631	Further evidence for a susceptibility locus on chromosome 10p14\bar{\bar{\bar{\bar{\bar{\bar{\bar{		104	
630	GWAS of Suicide Attempt in Psychiatric Disorders and Association With Major Depression Polygenic Risk Scores. <i>American Journal of Psychiatry</i> , 2019 , 176, 651-660	11.9	103	
629	Determinants of early alcohol use in healthy adolescents: the differential contribution of neuroimaging and psychological factors. <i>Neuropsychopharmacology</i> , 2012 , 37, 986-95	8.7	103	
628	Systematic analysis of glutamatergic neurotransmission genes in alcohol dependence and adolescent risky drinking behavior. <i>Archives of General Psychiatry</i> , 2008 , 65, 826-38		103	
627	Cognitive state and connectivity effects of the genome-wide significant psychosis variant in ZNF804A. <i>NeuroImage</i> , 2011 , 54, 2514-23	7.9	102	
626	At-risk variant in TCF7L2 for type II diabetes increases risk of schizophrenia. <i>Biological Psychiatry</i> , 2011 , 70, 59-63	7.9	101	
625	Efficacy and side-effects of clozapine not associated with variation in the 5-HT2C receptor. <i>NeuroReport</i> , 1997 , 8, 1999-2003	1.7	101	
624	Genome-wide association study of borderline personality disorder reveals genetic overlap with bipolar disorder, major depression and schizophrenia. <i>Translational Psychiatry</i> , 2017 , 7, e1155	8.6	100	
623	Effects of the circadian rhythm gene period 1 (per1) on psychosocial stress-induced alcohol drinking. <i>American Journal of Psychiatry</i> , 2011 , 168, 1090-8	11.9	100	

622	Subtype differences in adults with attention-deficit/hyperactivity disorder (ADHD) with regard to ADHD-symptoms, psychiatric comorbidity and psychosocial adjustment. <i>European Psychiatry</i> , 2008 , 23, 142-9	6	99
621	Genetic variation of the 5-HT2A receptor and response to clozapine. <i>Lancet, The</i> , 1995 , 346, 908-9	40	97
620	Association of a functional 1019C>G 5-HT1A receptor gene polymorphism with panic disorder with agoraphobia. <i>International Journal of Neuropsychopharmacology</i> , 2004 , 7, 189-92	5.8	95
619	Left prefrontal high-frequency repetitive transcranial magnetic stimulation for the treatment of schizophrenia with predominant negative symptoms: a sham-controlled, randomized multicenter trial. <i>Biological Psychiatry</i> , 2015 , 77, 979-88	7.9	94
618	Volition diminishes genetically mediated amygdala hyperreactivity. <i>NeuroImage</i> , 2010 , 53, 943-51	7.9	94
617	Genetic predictors of response to serotonergic and noradrenergic antidepressants in major depressive disorder: a genome-wide analysis of individual-level data and a meta-analysis. <i>PLoS Medicine</i> , 2012 , 9, e1001326	11.6	94
616	Systematic screening for DNA sequence variation in the coding region of the human dopamine transporter gene (DAT1). <i>Molecular Psychiatry</i> , 2000 , 5, 275-82	15.1	93
615	A Genetic Investigation of Sex Bias in the Prevalence of Attention-Deficit/Hyperactivity Disorder. <i>Biological Psychiatry</i> , 2018 , 83, 1044-1053	7.9	93
614	Neuroimaging evidence for a role of neural social stress processing in ethnic minority-associated environmental risk. <i>JAMA Psychiatry</i> , 2014 , 71, 672-80	14.5	92
613	Brain-specific tryptophan hydroxylase 2 (TPH2): a functional Pro206Ser substitution and variation in the 5'-region are associated with bipolar affective disorder. <i>Human Molecular Genetics</i> , 2008 , 17, 87-	97 ^{5.6}	92
612	Association between genetic variation in a region on chromosome 11 and schizophrenia in large samples from Europe. <i>Molecular Psychiatry</i> , 2012 , 17, 906-17	15.1	91
611	Meta-analysis of two genome-wide association studies of bipolar disorder reveals important points of agreement. <i>Molecular Psychiatry</i> , 2008 , 13, 466-7	15.1	91
610	Trajectories of change in depression severity during treatment with antidepressants. <i>Psychological Medicine</i> , 2010 , 40, 1367-77	6.9	90
609	Mechanisms of disturbed emotion processing and social interaction in borderline personality disorder: state of knowledge and research agenda of the German Clinical Research Unit. <i>Borderline Personality Disorder and Emotion Dysregulation</i> , 2014 , 1, 12	4.9	88
608	Increased medial orbitofrontal and amygdala activation: evidence for a systems-level endophenotype of bipolar I disorder. <i>American Journal of Psychiatry</i> , 2012 , 169, 316-25	11.9	88
607	What is familial about familial bipolar disorder? Resemblance among relatives across a broad spectrum of phenotypic characteristics. <i>Archives of General Psychiatry</i> , 2006 , 63, 1368-76		88
606	Impact of psychosocial adversity on alcohol intake in young adults: moderation by the LL genotype of the serotonin transporter polymorphism. <i>Biological Psychiatry</i> , 2009 , 66, 102-9	7.9	87
605	Genetic predictors of increase in suicidal ideation during antidepressant treatment in the GENDEP project. <i>Neuropsychopharmacology</i> , 2009 , 34, 2517-28	8.7	87

(2012-2011)

604	Early and delayed onset of response to antidepressants in individual trajectories of change during treatment of major depression: a secondary analysis of data from the Genome-Based Therapeutic Drugs for Depression (GENDEP) study. <i>Journal of Clinical Psychiatry</i> , 2011 , 72, 1478-84	4.6	87
603	Interacting effects of CRHR1 gene and stressful life events on drinking initiation and progression among 19-year-olds. <i>International Journal of Neuropsychopharmacology</i> , 2010 , 13, 703-14	5.8	85
602	Sex-specific association between the 5-HTT gene-linked polymorphic region and basal cortisol secretion. <i>Psychoneuroendocrinology</i> , 2009 , 34, 972-82	5	84
601	Maternally derived microduplications at 15q11-q13: implication of imprinted genes in psychotic illness. <i>American Journal of Psychiatry</i> , 2011 , 168, 408-17	11.9	84
600	Genomewide association scan of suicidal thoughts and behaviour in major depression. <i>PLoS ONE</i> , 2011 , 6, e20690	3.7	83
599	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
598	Molecular genetic overlap in bipolar disorder, schizophrenia, and major depressive disorder. <i>World Journal of Biological Psychiatry</i> , 2014 , 15, 200-8	3.8	82
597	Effects of a CACNA1C genotype on attention networks in healthy individuals. <i>Psychological Medicine</i> , 2011 , 41, 1551-61	6.9	82
596	The genetics of alcohol dependence. Annals of the New York Academy of Sciences, 2013, 1282, 39-70	6.5	81
595	Melancholic, atypical and anxious depression subtypes and outcome of treatment with escitalopram and nortriptyline. <i>Journal of Affective Disorders</i> , 2011 , 132, 112-20	6.6	81
594	Genetic architecture of subcortical brain structures in 38,851 individuals. <i>Nature Genetics</i> , 2019 , 51, 16	24 3 663	6 81
593	Genetic Differences in the Immediate Transcriptome Response to Stress Predict Risk-Related Brain Function and Psychiatric Disorders. <i>Neuron</i> , 2015 , 86, 1189-202	13.9	79
592	Neural and cognitive correlates of the common and specific variance across externalizing problems in young adolescence. <i>American Journal of Psychiatry</i> , 2014 , 171, 1310-9	11.9	79
591	S13. IMPACT OF POLYGENIC AND POLY-ENVIRONMENTAL RISK FACTORS ON A PSYCHOSIS RISK PHENOTYPE EXPLAINED THROUGH BRAIN STRUCTURE. <i>Schizophrenia Bulletin</i> , 2020 , 46, S35-S36	1.3	78
590	Involvement of the atrial natriuretic peptide transcription factor GATA4 in alcohol dependence, relapse risk and treatment response to acamprosate. <i>Pharmacogenomics Journal</i> , 2011 , 11, 368-74	3.5	78
589	From nature versus nurture, via nature and nurture, to gene x environment interaction in mental disorders. <i>European Child and Adolescent Psychiatry</i> , 2010 , 19, 199-210	5.5	78
588	Association of the OPRM1 Variant rs1799971 (A118G) with Non-Specific Liability to Substance Dependence in a Collaborative de novo Meta-Analysis of European-Ancestry Cohorts. <i>Behavior Genetics</i> , 2016 , 46, 151-69	3.2	77
587	Genome-wide association study of increasing suicidal ideation during antidepressant treatment in the GENDEP project. <i>Pharmacogenomics Journal</i> , 2012 , 12, 68-77	3.5	76

586	Cortical thickness of superior frontal cortex predicts impulsiveness and perceptual reasoning in adolescence. <i>Molecular Psychiatry</i> , 2013 , 18, 624-30	15.1	75
585	Association of Polygenic Score for Schizophrenia and HLA Antigen and Inflammation Genes With Response to Lithium in Bipolar Affective Disorder: A Genome-Wide Association Study. <i>JAMA Psychiatry</i> , 2018 , 75, 65-74	14.5	75
584	Human NPY promoter variation rs16147:T>C as a moderator of prefrontal NPY gene expression and negative affect. <i>Human Mutation</i> , 2010 , 31, E1594-608	4.7	75
583	Striatal response to reward anticipation: evidence for a systems-level intermediate phenotype for schizophrenia. <i>JAMA Psychiatry</i> , 2014 , 71, 531-9	14.5	74
582	Association between a serotonin transporter length polymorphism and primary insomnia. <i>Sleep</i> , 2010 , 33, 343-7	1.1	73
581	Serotonin transporter 5HTTLPR polymorphism and affective disorders: no evidence of association in a large European multicenter study. <i>European Journal of Human Genetics</i> , 2004 , 12, 377-82	5.3	73
580	A systems medicine research approach for studying alcohol addiction. <i>Addiction Biology</i> , 2013 , 18, 883-9	6 4.6	72
579	Body weight as a predictor of antidepressant efficacy in the GENDEP project. <i>Journal of Affective Disorders</i> , 2009 , 118, 147-54	6.6	72
578	How the serotonin transporter 5-HTTLPR polymorphism influences amygdala function: the roles of in vivo serotonin transporter expression and amygdala structure. <i>Translational Psychiatry</i> , 2011 , 1, e37	8.6	72
577	RASGRF2 regulates alcohol-induced reinforcement by influencing mesolimbic dopamine neuron activity and dopamine release. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, 21128-33	11.5	72
576	The DISC locus and schizophrenia: evidence from an association study in a central European sample and from a meta-analysis across different European populations. <i>Human Molecular Genetics</i> , 2009 , 18, 2719-27	5.6	71
575	Genetic models of schizophrenia and bipolar disorder: overlapping inheritance or discrete genotypes?. <i>European Archives of Psychiatry and Clinical Neuroscience</i> , 2005 , 255, 159-66	5.1	71
574	Efficacy and side-effects of clozapine: testing for association with allelic variation in the dopamine D4 receptor gene. <i>Neuropsychopharmacology</i> , 1996 , 15, 491-6	8.7	70
573	Addiction Research Consortium: Losing and regaining control over drug intake (ReCoDe)-From trajectories to mechanisms and interventions. <i>Addiction Biology</i> , 2020 , 25, e12866	4.6	70
572	The CACNA1C risk variant for bipolar disorder influences limbic activity. <i>Molecular Psychiatry</i> , 2010 , 15, 1126-7	15.1	69
571	Familial occurrence of tardive dyskinesia. <i>Acta Psychiatrica Scandinavica</i> , 2001 , 104, 375-9	6.5	69
570	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
569	Genome-wide association study of pathological gambling. <i>European Psychiatry</i> , 2016 , 36, 38-46	6	68

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(2010-2014)

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(2016-1991)

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(1998-2019)

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(2021-2019)

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(2000-2019)

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60	Assessing 42 inflammatory markers in 321 control subjects and 887 major depressive disorder cases: BMI and other confounders and overall predictive ability for current depression		2
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58	Generative network models of altered structural brain connectivity in schizophrenia. <i>NeuroImage</i> , 2021 , 225, 117510	7.9	2
57	Rhythm of Fetoplacental 11 Hydroxysteroid Dehydrogenase Type 2 - Fetal Protection From Morning Maternal Glucocorticoids. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2021 , 106, 1630-16	35 6	2
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54	Examining sex differences in neurodevelopmental and psychiatric genetic risk in anxiety and depression. <i>PLoS ONE</i> , 2021 , 16, e0248254	3.7	2
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Geneties of Schizophrenia and Bipolar Affective Disorder **2010**, 759-776

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