Caroline M Nievergelt

List of Publications by Citations

Source: https://exaly.com/author-pdf/367011/caroline-m-nievergelt-publications-by-citations.pdf

Version: 2024-04-23

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

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

13,428 113 205 53 h-index g-index citations papers 6.32 17,108 242 7.9 L-index avg, IF ext. citations ext. papers

#	Paper	IF	Citations
205	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. <i>Nature Genetics</i> , 2013 , 45, 984-94	36.3	1628
204	Large-scale genome-wide association analysis of bipolar disorder identifies a new susceptibility locus near ODZ4. <i>Nature Genetics</i> , 2011 , 43, 977-83	36.3	1094
203	Dynamic changes in the copy number of pluripotency and cell proliferation genes in human ESCs and iPSCs during reprogramming and time in culture. <i>Cell Stem Cell</i> , 2011 , 8, 106-18	18	700
202	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , 2019 , 51, 793-803	36.3	662
201	Histopathology of pediatric nonalcoholic fatty liver disease. <i>Hepatology</i> , 2005 , 42, 641-9	11.2	549
200	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. <i>Cell</i> , 2019 , 179, 1469-1482.e11	56.2	402
199	Psychiatric Genomics: An Update and an Agenda. American Journal of Psychiatry, 2018, 175, 15-27	11.9	328
198	Post-traumatic stress disorder. <i>Nature Reviews Disease Primers</i> , 2015 , 1, 15057	51.1	322
197	Differential responses to lithium in hyperexcitable neurons from patients with bipolar disorder. <i>Nature</i> , 2015 , 527, 95-9	50.4	315
196	Genome-wide association study of bipolar disorder in European American and African American individuals. <i>Molecular Psychiatry</i> , 2009 , 14, 755-63	15.1	287
195	Assessment of plasma C-reactive protein as a biomarker of posttraumatic stress disorder risk. <i>JAMA Psychiatry</i> , 2014 , 71, 423-31	14.5	222
194	Suggestive evidence for association of the circadian genes PERIOD3 and ARNTL with bipolar disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2006 , 141B, 234-41	3.5	218
193	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
192	The resilience framework as a strategy to combat stress-related disorders. <i>Nature Human Behaviour</i> , 2017 , 1, 784-790	12.8	210
191	Circadian polymorphisms associated with affective disorders. <i>Journal of Circadian Rhythms</i> , 2009 , 7, 2	2.5	168
190	International meta-analysis of PTSD genome-wide association studies identifies sex- and ancestry-specific genetic risk loci. <i>Nature Communications</i> , 2019 , 10, 4558	17.4	151
189	A genomewide scan of male sexual orientation. <i>Human Genetics</i> , 2005 , 116, 272-8	6.3	146

188	Association between traumatic brain injury and risk of posttraumatic stress disorder in active-duty Marines. <i>JAMA Psychiatry</i> , 2014 , 71, 149-57	14.5	145
187	Biomarkers of PTSD: neuropeptides and immune signaling. <i>Neuropharmacology</i> , 2012 , 62, 663-73	5.5	135
186	Enrichment of cis-regulatory gene expression SNPs and methylation quantitative trait loci among bipolar disorder susceptibility variants. <i>Molecular Psychiatry</i> , 2013 , 18, 340-6	15.1	134
185	Gene networks specific for innate immunity define post-traumatic stress disorder. <i>Molecular Psychiatry</i> , 2015 , 20, 1538-45	15.1	129
184	Population structure, admixture, and aging-related phenotypes in African American adults: the Cardiovascular Health Study. <i>American Journal of Human Genetics</i> , 2005 , 76, 463-77	11	127
183	Genomic predictors of combat stress vulnerability and resilience in U.S. Marines: A genome-wide association study across multiple ancestries implicates PRTFDC1 as a potential PTSD gene. <i>Psychoneuroendocrinology</i> , 2015 , 51, 459-71	5	125
182	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
181	Singleton deletions throughout the genome increase risk of bipolar disorder. <i>Molecular Psychiatry</i> , 2009 , 14, 376-80	15.1	121
180	A survey of genomic studies supports association of circadian clock genes with bipolar disorder spectrum illnesses and lithium response. <i>PLoS ONE</i> , 2012 , 7, e32091	3.7	119
179	Genome-wide Association Studies of Posttraumatic Stress Disorder in 2 Cohorts of US Army Soldiers. <i>JAMA Psychiatry</i> , 2016 , 73, 695-704	14.5	114
178	Rare variants in neuronal excitability genes influence risk for bipolar disorder. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015 , 112, 3576-81	11.5	112
177	Traumatic stress and accelerated DNA methylation age: A meta-analysis. <i>Psychoneuroendocrinology</i> , 2018 , 92, 123-134	5	107
176	Paternally inherited cis-regulatory structural variants are associated with autism. <i>Science</i> , 2018 , 360, 327-331	33.3	106
175	Susceptibility and modifier genes in Portuguese transthyretin V30M amyloid polyneuropathy: complexity in a single-gene disease. <i>Human Molecular Genetics</i> , 2005 , 14, 543-53	5.6	99
174	The Psychiatric Genomics Consortium Posttraumatic Stress Disorder Workgroup: Posttraumatic Stress Disorder Enters the Age of Large-Scale Genomic Collaboration. <i>Neuropsychopharmacology</i> , 2015 , 40, 2287-97	8.7	94
173	Association of Predeployment Heart Rate Variability With Risk of Postdeployment Posttraumatic Stress Disorder in Active-Duty Marines. <i>JAMA Psychiatry</i> , 2015 , 72, 979-86	14.5	84
172	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
171	The cumulative effect of different childhood trauma types on self-reported symptoms of adult male depression and PTSD, substance abuse and health-related quality of life in a large active-duty military cohort. Journal of Psychiatric Research 2014, 58, 46-54	5.2	80

170	Social system of the Alaotran gentle lemur (Hapalemur griseus alaotrensis): genetic characterization of group composition and mating system. <i>American Journal of Primatology</i> , 2002 , 57, 157-76	2.5	78
169	Peptide YY (PYY) Gene Polymorphisms in the 3?-Untranslated Region and Proximal Promoter Regions Regulate Cellular Gene Expression and PYY Secretion and Metabolic Syndrome Traits in Vivo. <i>Molecular Endocrinology</i> , 2009 , 23, 2120-2120		78
168	Genetic Analysis of Group Composition and Breeding System in a Wild Common Marmoset (Callithrix jacchus) Population. <i>International Journal of Primatology</i> , 2000 , 21, 1-20	2	77
167	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
166	Group encounters and territoriality in wild Alaotran gentle lemurs (Hapalemur griseus alaotrensis). <i>American Journal of Primatology</i> , 1998 , 46, 251-8	2.5	74
165	Preliminary Field Data on Group Size, Diet and Activity in the Alaotran Gentle Lemur Hapalemur griseus alaotrensis. <i>Folia Primatologica</i> , 1998 , 69, 325-330	1.2	72
164	Functional genetic variation in the Rev-Erb[pathway and lithium response in the treatment of bipolar disorder. <i>Genes, Brain and Behavior</i> , 2011 , 10, 852-61	3.6	70
163	Inference of human continental origin and admixture proportions using a highly discriminative ancestry informative 41-SNP panel. <i>Investigative Genetics</i> , 2013 , 4, 13		69
162	Generalized analysis of molecular variance. <i>PLoS Genetics</i> , 2007 , 3, e51	6	66
161	Heart rate variability characteristics in a large group of active-duty marines and relationship to posttraumatic stress. <i>Psychosomatic Medicine</i> , 2014 , 76, 292-301	3.7	63
160	Delayed sleep phase syndrome is related to seasonal affective disorder. <i>Journal of Affective Disorders</i> , 2011 , 133, 573-9	6.6	63
159	Analysis of ABCG2 and other urate transporters in uric acid homeostasis in chronic kidney disease: potential role of remote sensing and signaling. <i>CKJ: Clinical Kidney Journal</i> , 2016 , 9, 444-53	4.5	62
158	Social organization of the Alaotran gentle lemur (Hapalemur griseus alaotrensis). <i>American Journal of Primatology</i> , 2000 , 50, 9-24	2.5	62
157	PTSD Blood Transcriptome Mega-Analysis: Shared Inflammatory Pathways across Biological Sex and Modes of Trauma. <i>Neuropsychopharmacology</i> , 2018 , 43, 469-481	8.7	61
156	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
155	New translational perspectives for blood-based biomarkers of PTSD: From glucocorticoid to immune mediators of stress susceptibility. <i>Experimental Neurology</i> , 2016 , 284, 133-140	5.7	56
154	Blood-based gene-expression predictors of PTSD risk and resilience among deployed marines: a pilot study. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2013 , 162B, 313-26	3.5	55
153	G-protein-coupled receptor kinase 4 polymorphisms and blood pressure response to metoprolol among African Americans: sex-specificity and interactions. <i>American Journal of Hypertension</i> , 2009 , 22, 332-8	2.3	54

(2015-2017)

152	Epigenome-wide association of PTSD from heterogeneous cohorts with a common multi-site analysis pipeline. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017 , 174, 619	-6350	53	
151	Predictors of risk and resilience for posttraumatic stress disorder among ground combat Marines: methods of the Marine Resiliency Study. <i>Preventing Chronic Disease</i> , 2012 , 9, E97	3.7	53	
150	Genome-wide association of bipolar disorder suggests an enrichment of replicable associations in regions near genes. <i>PLoS Genetics</i> , 2011 , 7, e1002134	6	53	
149	Tracing sub-structure in the European American population with PCA-informative markers. <i>PLoS Genetics</i> , 2008 , 4, e1000114	6	53	
148	Genetic loci associated with heart rate variability and their effects on cardiac disease risk. <i>Nature Communications</i> , 2017 , 8, 15805	17.4	50	
147	Delayed sleep phase cases and controls. <i>Journal of Circadian Rhythms</i> , 2008 , 6, 6	2.5	50	
146	Restricted ethnic diversity in human embryonic stem cell lines. <i>Nature Methods</i> , 2010 , 7, 6-7	21.6	47	
145	Heritability and clinical features of multigenerational families with obsessive-compulsive disorder and hoarding. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2007 , 144B, 174-8	32 ^{3.5}	47	
144	Assessing Neuronal and Astrocyte Derived Exosomes From Individuals With Mild Traumatic Brain Injury for Markers of Neurodegeneration and Cytotoxic Activity. <i>Frontiers in Neuroscience</i> , 2019 , 13, 10	0 <i>5</i> ^{.1}	44	
143	Inflammation and stress-related candidate genes, plasma interleukin-6 levels, and longevity in older adults. <i>Experimental Gerontology</i> , 2009 , 44, 350-5	4.5	43	
142	Energy intake during reproduction in captive common marmosets (Callithrix jacchus). <i>Physiology and Behavior</i> , 1999 , 65, 849-54	3.5	43	
141	Chronotype and cellular circadian rhythms predict the clinical response to lithium maintenance treatment in patients with bipolar disorder. <i>Neuropsychopharmacology</i> , 2019 , 44, 620-628	8.7	43	
140	PROC, PROCR and PROS1 polymorphisms, plasma anticoagulant phenotypes, and risk of cardiovascular disease and mortality in older adults: the Cardiovascular Health Study. <i>Journal of Thrombosis and Haemostasis</i> , 2008 , 6, 1625-32	15.4	42	
139	The Pharmacogenomics of Bipolar Disorder study (PGBD): identification of genes for lithium response in a prospective sample. <i>BMC Psychiatry</i> , 2016 , 16, 129	4.2	42	
138	Blood-based gene-expression biomarkers of post-traumatic stress disorder among deployed marines: A pilot study. <i>Psychoneuroendocrinology</i> , 2015 , 51, 472-94	5	41	
137	Loss of GSTM1, a NRF2 target, is associated with accelerated progression of hypertensive kidney disease in the African American Study of Kidney Disease (AASK). <i>American Journal of Physiology - Renal Physiology</i> , 2013 , 304, F348-55	4.3	41	
136	Diagnostic Utility of the Posttraumatic Stress Disorder (PTSD) Checklist for Identifying Full and Partial PTSD in Active-Duty Military. <i>Assessment</i> , 2015 , 22, 289-97	3.7	40	
135	Conditioned fear and extinction learning performance and its association with psychiatric symptoms in active duty Marines. <i>Psychoneuroendocrinology</i> , 2015 , 51, 495-505	5	40	

134	The role of emotional numbing in sexual functioning among veterans of the Iraq and Afghanistan wars. <i>Military Medicine</i> , 2010 , 175, 424-8	1.3	39
133	Common genetic variants in ARNTL and NPAS2 and at chromosome 12p13 are associated with objectively measured sleep traits in the elderly. <i>Sleep</i> , 2013 , 36, 431-46	1.1	38
132	A candidate gene study of CYP19 (aromatase) and male sexual orientation. <i>Behavior Genetics</i> , 2004 , 34, 243-50	3.2	38
131	Angiotensin-converting enzyme gene polymorphism predicts the time-course of blood pressure response to angiotensin converting enzyme inhibition in the AASK trial. <i>Journal of Hypertension</i> , 2007 , 25, 2082-92	1.9	37
130	Matrix metalloproteinase haplotypes associated with coronary artery aneurysm formation in patients with Kawasaki disease. <i>Journal of Human Genetics</i> , 2010 , 55, 779-84	4.3	35
129	Large-scale integration of human genetic and physical maps. <i>Genome Research</i> , 2004 , 14, 1199-205	9.7	35
128	Autonomic and hemodynamic origins of pre-hypertension: central role of heredity. <i>Journal of the American College of Cardiology</i> , 2012 , 59, 2206-16	15.1	34
127	Genetic analysis of hybridization and introgression between wild mongoose and brown lemurs. <i>BMC Evolutionary Biology</i> , 2009 , 9, 32	3	34
126	Epigenome-wide meta-analysis of PTSD across 10 military and civilian cohorts identifies methylation changes in AHRR. <i>Nature Communications</i> , 2020 , 11, 5965	17.4	34
125	Genomic Approaches to Posttraumatic Stress Disorder: The Psychiatric Genomic Consortium Initiative. <i>Biological Psychiatry</i> , 2018 , 83, 831-839	7.9	33
124	Genome-wide Association of Endophenotypes for Schizophrenia From the Consortium on the Genetics of Schizophrenia (COGS) Study. <i>JAMA Psychiatry</i> , 2019 , 76, 1274-1284	14.5	32
123	An epigenome-wide association study of posttraumatic stress disorder in US veterans implicates several new DNA methylation loci. <i>Clinical Epigenetics</i> , 2020 , 12, 46	7.7	31
122	Post-traumatic stress disorder: emerging concepts of pharmacotherapy. <i>Expert Opinion on Emerging Drugs</i> , 2009 , 14, 251-72	3.7	31
121	Genetic variants associated with sleep disorders. Sleep Medicine, 2015 , 16, 217-24	4.6	29
120	Heritability of Biomarkers of Oxidized Lipoproteins: Twin Pair Study. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2015 , 35, 1704-11	9.4	29
119	Examination of the clock gene Cryptochrome 1 in bipolar disorder: mutational analysis and absence of evidence for linkage or association. <i>Psychiatric Genetics</i> , 2005 , 15, 45-52	2.9	28
118	Genome-wide association study identifies 30 Loci Associated with Bipolar Disorder		28
117	Genetic implication of a novel thiamine transporter in human hypertension. <i>Journal of the American College of Cardiology</i> , 2014 , 63, 1542-55	15.1	27

(2020-2013)

116	Characterization of cerebrospinal fluid (CSF) and plasma NPY levels in normal volunteers over a 24-h timeframe. <i>Psychoneuroendocrinology</i> , 2013 , 38, 2378-82	5	27	
115	Female veterans of the OEF/OIF conflict: concordance of PTSD symptoms and substance misuse. <i>Addictive Behaviors</i> , 2010 , 35, 655-9	4.2	27	
114	Successful treatment of post-traumatic stress disorder reverses DNA methylation marks. <i>Molecular Psychiatry</i> , 2021 , 26, 1264-1271	15.1	27	
113	Epigenetic meta-analysis across three civilian cohorts identifies NRG1 and HGS as blood-based biomarkers for post-traumatic stress disorder. <i>Epigenomics</i> , 2018 , 10, 1585-1601	4.4	27	
112	Common genetic variants of the human uromodulin gene regulate transcription and predict plasma uric acid levels. <i>Kidney International</i> , 2013 , 83, 733-40	9.9	26	
111	Tractor uses local ancestry to enable the inclusion of admixed individuals in GWAS and to boost power. <i>Nature Genetics</i> , 2021 , 53, 195-204	36.3	26	
110	Longitudinal epigenome-wide association studies of three male military cohorts reveal multiple CpG sites associated with post-traumatic stress disorder. <i>Clinical Epigenetics</i> , 2020 , 12, 11	7.7	24	
109	Chromogranin A regulates renal function by triggering Weibel-Palade body exocytosis. <i>Journal of the American Society of Nephrology: JASN</i> , 2009 , 20, 1623-32	12.7	24	
108	Suggestive linkage of a chromosomal locus on 18p11 to cyclothymic temperament in bipolar disorder families. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008 , 147, 326-	-325	24	
107	A putative causal relationship between genetically determined female body shape and posttraumatic stress disorder. <i>Genome Medicine</i> , 2017 , 9, 99	14.4	23	
106	Fear learning alterations after traumatic brain injury and their role in development of posttraumatic stress symptoms. <i>Depression and Anxiety</i> , 2017 , 34, 723-733	8.4	23	
105	Genome-wide analyses of psychological resilience in U.S. Army soldiers. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2019 , 180, 310-319	3.5	22	
104	Prevalence of suicidal ideation and other suicide warning signs in veterans attending an urgent care psychiatric clinic. <i>Comprehensive Psychiatry</i> , 2015 , 60, 149-55	7.3	22	
103	The catecholamine biosynthetic enzyme dopamine Ehydroxylase (DBH): first genome-wide search positions trait-determining variants acting additively in the proximal promoter. <i>Human Molecular Genetics</i> , 2014 , 23, 6375-84	5.6	22	
102	Common variants in the CRP gene in relation to longevity and cause-specific mortality in older adults: the Cardiovascular Health Study. <i>Atherosclerosis</i> , 2008 , 197, 922-30	3.1	21	
101	Analysis of Genetically Regulated Gene Expression Identifies a Prefrontal PTSD Gene, SNRNP35, Specific to Military Cohorts. <i>Cell Reports</i> , 2020 , 31, 107716	10.6	21	
100	Measuring novel antecedents of mental illness: the Questionnaire of Unpredictability in Childhood. <i>Neuropsychopharmacology</i> , 2019 , 44, 876-882	8.7	21	
99	Genomic influences on self-reported childhood maltreatment. <i>Translational Psychiatry</i> , 2020 , 10, 38	8.6	20	

98	Comparison of genetic distance measures using human SNP genotype data. <i>Human Biology</i> , 2009 , 81, 389-406	1.2	20
97	Associations of PER3 and RORA Circadian Gene Polymorphisms and Depressive Symptoms in Older Adults. <i>American Journal of Geriatric Psychiatry</i> , 2015 , 23, 1075-87	6.5	19
96	Circadian polymorphisms in night owls, in bipolars, and in non-24-hour sleep cycles. <i>Psychiatry Investigation</i> , 2014 , 11, 345-62	3.1	19
95	Association of Economic Status and Educational Attainment With Posttraumatic Stress Disorder: A Mendelian Randomization Study. <i>JAMA Network Open</i> , 2019 , 2, e193447	10.4	18
94	Toward a Global Roadmap for Precision Medicine in Psychiatry: Challenges and Opportunities. <i>OMICS A Journal of Integrative Biology</i> , 2016 , 20, 557-564	3.8	18
93	Polymorphisms in melatonin synthesis pathways: possible influences on depression. <i>Journal of Circadian Rhythms</i> , 2011 , 9, 8	2.5	18
92	Further evidence for association of GRK3 to bipolar disorder suggests a second disease mutation. <i>Psychiatric Genetics</i> , 2007 , 17, 315-22	2.9	18
91	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
90	Neuropeptide Y (NPY): genetic variation in the human promoter alters glucocorticoid signaling, yielding increased NPY secretion and stress responses. <i>Journal of the American College of Cardiology</i> , 2012 , 60, 1678-89	15.1	17
89	Gene-environment correlations and causal effects of childhood maltreatment on physical and mental health: a genetically informed approach. <i>Lancet Psychiatry,the</i> , 2021 , 8, 373-386	23.3	17
88	Single nucleotide polymorphisms and haplotypes in Native American populations. <i>American Journal of Physical Anthropology</i> , 2011 , 146, 495-502	2.5	16
87	Analysis of the Influence of microRNAs in Lithium Response in Bipolar Disorder. <i>Frontiers in Psychiatry</i> , 2018 , 9, 207	5	15
86	Investigating polygenic burden in age at disease onset in bipolar disorder: Findings from an international multicentric study. <i>Bipolar Disorders</i> , 2019 , 21, 68-75	3.8	15
85	Effects of military service and deployment on clinical symptomatology: The role of trauma exposure and social support. <i>Journal of Psychiatric Research</i> , 2017 , 95, 121-128	5.2	15
84	Candidate SNP associations of optimism and resilience in older adults: exploratory study of 935 community-dwelling adults. <i>American Journal of Geriatric Psychiatry</i> , 2014 , 22, 997-1006.e5	6.5	15
83	Biogeographic Ancestry in the African Descent and Glaucoma Evaluation Study (ADAGES): Association With Corneal and Optic Nerve Structure 2015 , 56, 2043-9		14
82	Peptide YY (PYY) gene polymorphisms in the 3Suntranslated and proximal promoter regions regulate cellular gene expression and PYY secretion and metabolic syndrome traits in vivo. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2009 , 94, 4557-66	5.6	14
81	Novel Risk Loci in Tinnitus and Causal Inference With Neuropsychiatric Disorders Among Adults of European Ancestry. <i>JAMA Otolaryngology - Head and Neck Surgery</i> , 2020 , 146, 1015-1025	3.9	14

(2012-2019)

80	Concordance of genetic variation that increases risk for anxiety disorders and posttraumatic stress disorders and that influences their underlying neurocircuitry. <i>Journal of Affective Disorders</i> , 2019 , 245, 885-896	6.6	14
79	Genome wide association study identifies variants in NBEA associated with migraine in bipolar disorder. <i>Journal of Affective Disorders</i> , 2015 , 172, 453-61	6.6	12
78	Progression of chronic kidney disease: Adrenergic genetic influence on glomerular filtration rate decline in hypertensive nephrosclerosis. <i>American Journal of Nephrology</i> , 2010 , 32, 23-30	4.6	12
77	Admixture mapping as a gene discovery approach for complex human traits and diseases. <i>Current Hypertension Reports</i> , 2005 , 7, 31-7	4.7	12
76	Biological profiling of plasma neuropeptide Y in relation to posttraumatic stress symptoms in two combat cohorts. <i>Biological Psychology</i> , 2018 , 134, 72-79	3.2	11
75	COMT val158met polymorphism links to altered fear conditioning and extinction are modulated by PTSD and childhood trauma. <i>Depression and Anxiety</i> , 2018 , 35, 32-42	8.4	11
74	FMR1, circadian genes and depression: suggestive associations or false discovery?. <i>Journal of Circadian Rhythms</i> , 2013 , 11, 3	2.5	11
73	Identification of novel loci affecting circulating chromogranins and related peptides. <i>Human Molecular Genetics</i> , 2017 , 26, 233-242	5.6	11
72	Sleep disturbance at pre-deployment is a significant predictor of post-deployment re-experiencing symptoms. <i>H</i> gre <i>Utbildning</i> , 2019 , 10, 1679964	5	11
71	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. <i>Biological Psychiatry</i> , 2021 ,	7.9	11
70	MicroRNAs in Post-traumatic Stress Disorder. Current Topics in Behavioral Neurosciences, 2018, 38, 23-46	53.4	10
69	Polymorphisms at the F12 and KLKB1 loci have significant trait association with activation of the renin-angiotensin system. <i>BMC Medical Genetics</i> , 2016 , 17, 21	2.1	10
68	Who are the Okinawans? Ancestry, genome diversity, and implications for the genetic study of human longevity from a geographically isolated population. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2014 , 69, 1474-84	6.4	10
67	Chip-based direct genotyping of coding variants in genome wide association studies: utility, issues and prospects. <i>Gene</i> , 2014 , 540, 104-9	3.8	10
66	Evidence for association of bipolar disorder to haplotypes in the 22q12.3 region near the genes stargazin, IFT27 and parvalbumin. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2012 , 159B, 941-50	3.5	10
65	Dopamine D1 receptor (DRD1) genetic polymorphism: pleiotropic effects on heritable renal traits. <i>Kidney International</i> , 2009 , 76, 1070-80	9.9	10
64	Adrenergic beta-1 receptor genetic variation predicts longitudinal rate of GFR decline in hypertensive nephrosclerosis. <i>Nephrology Dialysis Transplantation</i> , 2009 , 24, 3677-86	4.3	10
63	Genes and environment: novel, functional polymorphism in the human cathepsin L (CTSL1) promoter disrupts a xenobiotic response element (XRE) to alter transcription and blood pressure. Journal of Hypertension, 2012, 30, 1961-9	1.9	10

62	Dissecting the genetic association of C-reactive protein with PTSD, traumatic events, and social support. <i>Neuropsychopharmacology</i> , 2021 , 46, 1071-1077	8.7	10
61	Neuroepigenetics of Post-Traumatic Stress Disorder. <i>Progress in Molecular Biology and Translational Science</i> , 2018 , 158, 227-253	4	9
60	APOEI and slow wave sleep in older adults. <i>PLoS ONE</i> , 2018 , 13, e0191281	3.7	9
59	Molecular Mechanism for Hypertensive Renal Disease: Differential Regulation of Chromogranin A Expression at 3SUntranslated Region Polymorphism C+87T by MicroRNA-107. <i>Journal of the American Society of Nephrology: JASN</i> , 2015 , 26, 1816-25	12.7	9
58	Human heart rate: heritability of resting and stress values in twin pairs, and influence of genetic variation in the adrenergic pathway at a microribonucleic acid (microrna) motif in the 3SUTR of cytochrome b561 [corrected]. <i>Journal of the American College of Cardiology</i> , 2014 , 63, 358-68	15.1	9
57	Heritable influence of DBH on adrenergic and renal function: twin and disease studies. <i>PLoS ONE</i> , 2013 , 8, e82956	3.7	9
56	Molecular genetic overlap between posttraumatic stress disorder and sleep phenotypes. <i>Sleep</i> , 2020 , 43,	1.1	9
55	Post-traumatic stress following military deployment: Genetic associations and cross-disorder genetic correlations. <i>Journal of Affective Disorders</i> , 2019 , 252, 350-357	6.6	8
54	HIGH AND LOW THRESHOLD FOR STARTLE REACTIVITY ASSOCIATED WITH PTSD SYMPTOMS BUT NOT PTSD RISK: EVIDENCE FROM A PROSPECTIVE STUDY OF ACTIVE DUTY MARINES. <i>Depression and Anxiety</i> , 2016 , 33, 192-202	8.4	8
53	Common charge-shift mutation Glu65Lys in K+ channel Ebubunit KCNMB1: pleiotropic consequences for glomerular filtration rate and progressive renal disease. <i>American Journal of Nephrology</i> , 2010 , 32, 414-24	4.6	8
52	Clinical predictors of non-response to lithium treatment in the Pharmacogenomics of Bipolar Disorder (PGBD) study. <i>Bipolar Disorders</i> , 2021 ,	3.8	8
51	Comparison of the genetic variation of captive ring-tailed lemurs with a wild population in Madagascar. <i>Zoo Biology</i> , 2015 , 34, 463-72	1.6	7
50	Naturally occurring genetic variants in human chromogranin A (CHGA) associated with hypertension as well as hypertensive renal disease. <i>Cellular and Molecular Neurobiology</i> , 2010 , 30, 1395-	400	7
49	Polygenic risk for anxiety influences anxiety comorbidity and suicidal behavior in bipolar disorder. Translational Psychiatry, 2020 , 10, 298	8.6	7
48	Circadian rhythms in bipolar disorder patient-derived neurons predict lithium response: preliminary studies. <i>Molecular Psychiatry</i> , 2021 , 26, 3383-3394	15.1	7
47	The association between lithium use and neurocognitive performance in patients with bipolar disorder. <i>Neuropsychopharmacology</i> , 2020 , 45, 1743-1749	8.7	6
46	CYBA Gene Polymorphisms and Adverse Outcomes in Acute Kidney Injury: A Prospective Cohort Study. <i>Nephron Extra</i> , 2011 , 1, 112-23		6
45	Largest genome-wide association study for PTSD identifies genetic risk loci in European and African ancestries and implicates novel biological pathways		6

44	Problematic alcohol use associates with sodium channel and clathrin linker 1 (SCLT1) in trauma-exposed populations. <i>Addiction Biology</i> , 2018 , 23, 1145-1159	4.6	5	
43	Paraoxonase 1 (PON1) C/T-108 association with longitudinal mean arterial blood pressure. American Journal of Hypertension, 2012 , 25, 1188-94	2.3	5	
42	Integrated computational and experimental analysis of the neuroendocrine transcriptome in genetic hypertension identifies novel control points for the cardiometabolic syndrome. <i>Circulation: Cardiovascular Genetics</i> , 2012 , 5, 430-40		5	
41	Rationale and Design of the Genetic Contribution to Drug Induced Renal Injury[(DIRECT) Study. <i>Kidney International Reports</i> , 2016 , 1, 288-298	4.1	5	
40	Efficient region-based test strategy uncovers genetic risk factors for functional outcome in bipolar disorder. <i>European Neuropsychopharmacology</i> , 2019 , 29, 156-170	1.2	5	
39	Entrainment of Circadian Rhythms to Temperature Reveals Amplitude Deficits in Fibroblasts from Patients with Bipolar Disorder and Possible Links to Calcium Channels. <i>Molecular Neuropsychiatry</i> , 2019 , 5, 115-124	4.9	4	
38	Further evidence for linkage of bipolar disorder to chromosomes 6 and 17 in a new independent pedigree series. <i>Bipolar Disorders</i> , 2012 , 14, 71-9	3.8	4	
37	Associations of Genetically Determined Continental Ancestry With CD4+ Count and Plasma HIV-1 RNA Beyond Self-Reported Race and Ethnicity. <i>Journal of Acquired Immune Deficiency Syndromes</i> (1999), 2016 , 71, 544-50	3.1	4	
36	Detecting significant genotype-phenotype association rules in bipolar disorder: market research meets complex genetics. <i>International Journal of Bipolar Disorders</i> , 2018 , 6, 24	5.4	4	
35	A new common functional coding variant at the DDC gene change renal enzyme activity and modify renal dopamine function. <i>Scientific Reports</i> , 2019 , 9, 5055	4.9	3	
34	Association of Race and Major Adverse Cardiac Events (MACE): The Atherosclerosis Risk in Communities (ARIC) Cohort. <i>Journal of Aging Research</i> , 2020 , 2020, 7417242	2.3	3	
33	Epigenome-wide meta-analysis of PTSD symptom severity in three military cohorts implicates DNA methylation changes in genes involved in immune system and oxidative stress <i>Molecular Psychiatry</i> , 2022 ,	15.1	3	
32	Psychiatric Genomics: An Update and an Agenda		3	
31	Epigenome-wide meta-analysis of PTSD across 10 military and civilian cohorts identifies novel methylation loci		3	
30	Enhancing Discovery of Genetic Variants for Posttraumatic Stress Disorder Through Integration of Quantitative Phenotypes and Trauma Exposure Information. <i>Biological Psychiatry</i> , 2021 ,	7.9	3	
29	Epigenetic Biomarkers Of PTSD: Updates From The EWAS Working Group of The PTSD PGC. European Neuropsychopharmacology, 2019 , 29, S750	1.2	2	
28	Genetically determined ancestry is more informative than self-reported race in HIV-infected and -exposed children. <i>Medicine (United States)</i> , 2016 , 95, e4733	1.8	2	
27	Agreement between self and psychiatrist reporting of suicidal ideation at a Veterans Administration psychiatric emergency clinic. <i>Depression and Anxiety</i> , 2018 , 35, 1114-1121	8.4	2	

26	Genetic variation at the delta-sarcoglycan (SGCD) locus elevates heritable sympathetic nerve activity in human twin pairs. <i>Journal of Neurochemistry</i> , 2013 , 127, 750-61	6	2
25	223. Genome-Wide Association Study of Posttraumatic Stress Disorder Symptom Domains in Two Cohorts of United States Army Soldiers. <i>Biological Psychiatry</i> , 2017 , 81, S91-S92	7.9	2
24	Genetic variability and phylogeography in the wild Alaotran gentle lemur population. <i>Evolutionary Anthropology</i> , 2003 , 11, 175-179	4.7	2
23	Tractor: A framework allowing for improved inclusion of admixed individuals in large-scale association studies		2
22	Critical evaluation of copy number variant calling methods using DNA methylation. <i>Genetic Epidemiology</i> , 2020 , 44, 148-158	2.6	2
21	A 7 Tesla Amygdalar-Hippocampal Shape Analysis of Lithium Response in Bipolar Disorder. <i>Frontiers in Psychiatry</i> , 2021 , 12, 614010	5	2
20	Trauma and posttraumatic stress disorder modulate polygenic predictors of hippocampal and amygdala volume <i>Translational Psychiatry</i> , 2021 , 11, 637	8.6	1
19	Combining schizophrenia and depression polygenic risk scores improves the genetic prediction of lithium response in bipolar disorder patients. <i>Translational Psychiatry</i> , 2021 , 11, 606	8.6	1
18	Correction of depression-associated circadian rhythm abnormalities is associated with lithium response in bipolar disorder. <i>Bipolar Disorders</i> , 2021 ,	3.8	1
17	Associations between the development of PTSD symptoms and longitudinal changes in the DNA methylome of deployed military servicemen: A comparison with polygenic risk scores. <i>Comprehensive Psychoneuroendocrinology</i> , 2020 , 4, 100018	1.1	1
16	Dissecting the association of C-reactive protein levels with PTSD, traumatic events, and social support		1
15	Polygenic scores for major depressive disorder and depressive symptoms predict response to lithium in patients with bipolar disorder		1
14	Analysis of Genetically Regulated Gene Expression identifies a trauma type specific PTSD gene, SNRNP	35	1
13	Examining Individual and Synergistic Contributions of PTSD and Genetics to Blood Pressure: A Trans-Ethnic Meta-Analysis. <i>Frontiers in Neuroscience</i> , 2021 , 15, 678503	5.1	1
12	Consequences of Lakeside Living for the Diet and Social Ecology of the Lake Alaotran Gentle Lemur 2019 , 276-278		1
11	Hapalemur alaotrensis: A Conservation Case Study from the Swamps of Alaotra, Madagascar 2019 , 293-	-296	1
10	HLA-DRB1 and HLA-DQB1 genetic diversity modulates response to lithium in bipolar affective disorders. <i>Scientific Reports</i> , 2021 , 11, 17823	4.9	1
9	Using polygenic scores and clinical data for bipolar disorder patient stratification and lithium response prediction: machine learning approach <i>British Journal of Psychiatry</i> , 2022 , 1-10	5.4	1

LIST OF PUBLICATIONS

8	Association of polygenic risk scores, traumatic life events and coping strategies with war-related PTSD diagnosis and symptom severity in the South Eastern Europe (SEE)-PTSD cohort. <i>Journal of Neural Transmission</i> , 2021 , 1	4.3	О
7	Reply to: On powerful GWAS in admixed populations. <i>Nature Genetics</i> , 2021 , 53, 1634-1635	36.3	0
6	Dissociable impact of childhood trauma and deployment trauma on affective modulation of startle. <i>Neurobiology of Stress</i> , 2021 , 15, 100362	7.6	O
5	Prospective examination of pre-trauma anhedonia as a risk factor for post-traumatic stress symptoms <i>H</i> gre Utbildning, 2022 , 13, 2015949	5	
4	NPY: Low Cerebrospinal Fluid (CSF) Levels in Posttraumatic Stress Disorder in Comparison to Combat and Civilian, non-combat Control Subjects 2014 , 224		
3	Current progress and future direction in the genetics of PTSD: Focus on the development and contributions of the PGC-PTSD working group 2020 , 285-296		
2	Disentangling sex differences in the shared genetic architecture of posttraumatic stress disorder, traumatic experiences, and social support with body size and composition. <i>Neurobiology of Stress</i> , 2021 , 15, 100400	7.6	
1	Alcohol use and alcohol use disorder differ in their genetic relationships with PTSD: A genomic structural equation modelling approach <i>Drug and Alcohol Dependence</i> , 2022 , 234, 109430	4.9	