Roseann E Peterson

List of Publications by Year in descending order

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48 papers 5,051 citations

257450 24 h-index 206112 48 g-index

68 all docs 68
docs citations

68 times ranked 8886 citing authors

#	Article	IF	CITATIONS
1	Investigation of convergent and divergent genetic influences underlying schizophrenia and alcohol use disorder. Psychological Medicine, 2023, 53, 1196-1204.	4.5	7
2	Polygenic score for cigarette smoking is associated with ever electronicâ€cigarette use in a collegeâ€aged sample. Addiction, 2022, 117, 1071-1078.	3.3	4
3	History of Suicide Attempts and COVID-19 Infection in Veterans with Schizophrenia or Schizoaffective Disorder: Moderating Effects of Age and Body Mass Index. Complex Psychiatry, 2022, 8, 28-34.	0.9	2
4	Divergent changes: abstinence and higher-frequency substance use increase among racial/ethnic minority young adults during the COVID-19 global pandemic. American Journal of Drug and Alcohol Abuse, 2022, 48, 88-99.	2.1	10
5	Intermediate stable states in substance use. Addictive Behaviors, 2022, 129, 107252.	3.0	7
6	Indirect paths from genetics to education. Nature Genetics, 2022, 54, 372-373.	21.4	5
7	Genome-wide Association Meta-analysis of Childhood and Adolescent Internalizing Symptoms. Journal of the American Academy of Child and Adolescent Psychiatry, 2022, 61, 934-945.	0.5	26
8	Shared genetic risk between eating disorder†and substanceâ€useâ€related phenotypes: Evidence from genomeâ€wide association studies. Addiction Biology, 2021, 26, e12880.	2.6	28
9	The Genetics of Major Depression: Perspectives on the State of Research and Opportunities for Precision Medicine. Psychiatric Annals, 2021, 51, 165-169.	0.1	2
10	Genome-wide analyses of smoking behaviors in schizophrenia: Findings from the Psychiatric Genomics Consortium. Journal of Psychiatric Research, 2021, 137, 215-224.	3.1	10
11	Examining Sex-Differentiated Genetic Effects Across Neuropsychiatric and Behavioral Traits. Biological Psychiatry, 2021, 89, 1127-1137.	1.3	48
12	Genetic association study of childhood aggression across raters, instruments, and age. Translational Psychiatry, 2021, 11, 413.	4.8	31
13	The Genetic Architecture of Depression in Individuals of East Asian Ancestry. JAMA Psychiatry, 2021, 78, 1258.	11.0	88
14	Pathway-based polygene risk for severe depression implicates drug metabolism in CONVERGE. Psychological Medicine, 2020, 50, 793-798.	4.5	3
15	Contributions of common genetic variants to risk of schizophrenia among individuals of African and Latino ancestry. Molecular Psychiatry, 2020, 25, 2455-2467.	7.9	82
16	Classical Human Leukocyte Antigen Alleles and C4 Haplotypes Are Not Significantly Associated With Depression. Biological Psychiatry, 2020, 87, 419-430.	1.3	27
17	A large-scale genome-wide association study meta-analysis of cannabis use disorder. Lancet Psychiatry,the, 2020, 7, 1032-1045.	7.4	200
18	Leveraging genome-wide data to investigate differences between opioid use vs. opioid dependence in 41,176 individuals from the Psychiatric Genomics Consortium. Molecular Psychiatry, 2020, 25, 1673-1687.	7.9	82

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19	Genome-wide gene-environment analyses of major depressive disorder and reported lifetime traumatic experiences in UK Biobank. Molecular Psychiatry, 2020, 25, 1430-1446.	7.9	116
20	Genome-wide Association Studies in Ancestrally Diverse Populations: Opportunities, Methods, Pitfalls, and Recommendations. Cell, 2019, 179, 589-603.	28.9	428
21	International Society of Psychiatric Genetics Ethics Committee: Issues facing us. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2019, 180, 543-554.	1.7	16
22	Evidence of causal effect of major depression on alcohol dependence: findings from the psychiatric genomics consortium. Psychological Medicine, 2019, 49, 1218-1226.	4.5	74
23	Molecular Genetic Analysis Subdivided by Adversity Exposure Suggests Etiologic Heterogeneity in Major Depression. American Journal of Psychiatry, 2018, 175, 545-554.	7.2	69
24	F217. Genome-Wide Analyses of Smoking Behaviors in Schizophrenia: Findings From the Psychiatric Genomics Consortium. Biological Psychiatry, 2018, 83, S323.	1.3	0
25	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. Nature Genetics, 2018, 50, 668-681.	21.4	2,224
26	Robust symptom networks in recurrent major depression across different levels of genetic and environmental risk. Journal of Affective Disorders, 2018, 227, 313-322.	4.1	34
27	Polygenic risk for severe psychopathology among Europeans is associated with major depressive disorder in Han Chinese women. Psychological Medicine, 2018, 48, 777-789.	4.5	8
28	Polygenic prediction of the phenome, across ancestry, in emerging adulthood. Psychological Medicine, 2018, 48, 1814-1823.	4.5	29
29	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. Nature Neuroscience, 2018, 21, 1656-1669.	14.8	490
30	Replication of the Interaction of PRKG1 and Trauma Exposure on Alcohol Misuse in an Independent African American Sample. Journal of Traumatic Stress, 2018, 31, 927-932.	1.8	9
31	Age of onset and family history as indicators of polygenic risk for major depression. Depression and Anxiety, 2017, 34, 446-452.	4.1	19
32	Deep Sequencing of 71 Candidate Genes to Characterize Variation Associated with Alcohol Dependence. Alcoholism: Clinical and Experimental Research, 2017, 41, 711-718.	2.4	13
33	Genetic effects influencing risk for major depressive disorder in China and Europe. Translational Psychiatry, 2017, 7, e1074-e1074.	4.8	64
34	The Genetic Architecture of Major Depressive Disorder in Han Chinese Women. JAMA Psychiatry, 2017, 74, 162.	11.0	82
35	Genetic Association of Major Depression With Atypical Features and Obesity-Related Immunometabolic Dysregulations. JAMA Psychiatry, 2017, 74, 1214.	11.0	174
36	Leveraging Molecular Genetic Approaches to Yield Insights Into Major Depression Etiology and Clinical Presentation. JAMA Psychiatry, 2017, 74, 1189.	11.0	0

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37	The utility of empirically assigning ancestry groups in crossâ€population genetic studies of addiction. American Journal on Addictions, 2017, 26, 494-501.	1.4	46
38	Hair Cortisol in Twins: Heritability and Genetic Overlap with Psychological Variables and Stress-System Genes. Scientific Reports, 2017, 7, 15351.	3.3	50
39	CHRONICITY OF DEPRESSION AND MOLECULAR MARKERS IN A LARGE SAMPLE OF HAN CHINESE WOMEN. Depression and Anxiety, 2016, 33, 1048-1054.	4.1	18
40	SNP-based heritability estimates of the personality dimensions and polygenic prediction of both neuroticism and major depression: findings from CONVERGE. Translational Psychiatry, 2016, 6, e926-e926.	4.8	33
41	Genome-Wide Meta-Analysis of Cotinine Levels in Cigarette Smokers Identifies Locus at 4q13.2. Scientific Reports, 2016, 6, 20092.	3.3	42
42	Deep Sequencing of Three Loci Implicated in Large-Scale Genome-Wide Association Study Smoking Meta-Analyses. Nicotine and Tobacco Research, 2016, 18, 626-631.	2.6	10
43	On the association of common and rare genetic variation influencing body mass index: a combined SNP and CNV analysis. BMC Genomics, 2014, 15, 368.	2.8	18
44	A comparison of the accuracy of self-reported intake with measured intake of a laboratory overeating episode in overweight and obese women with and without binge eating disorder. European Journal of Nutrition, 2013, 52, 193-202.	3.9	29
45	Comparisons of Energy Intake and Energy Expenditure in Overweight and Obese Women With and Without Binge Eating Disorder. Obesity, 2012, 20, 765-772.	3.0	26
46	Binge Eating Disorder Mediates Links between Symptoms of Depression, Anxiety, and Caloric Intake in Overweight and Obese Women. Journal of Obesity, 2012, 2012, 1-8.	2.7	41
47	Genetic risk sum score comprised of common polygenic variation is associated with body mass index. Human Genetics, 2011, 129, 221-230.	3.8	62
48	A comparison of energy intake and food selection during laboratory binge eating episodes in obese women with and without a binge eating disorder diagnosis. International Journal of Eating Disorders, 2007, 40, 67-71.	4.0	57