

Sandra Sanchez-roige

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.

46
papers

1,840
citations

19
h-index

42
g-index

63
ext. papers

2,930
ext. citations

9.7
avg, IF

4.69
L-index

#	Paper	IF	Citations
46	GWAS of lifetime cannabis use reveals new risk loci, genetic overlap with psychiatric traits, and a causal influence of schizophrenia. <i>Nature Neuroscience</i> , 2018 , 21, 1161-1170	25.5	270
45	Genome-wide association analyses of risk tolerance and risky behaviors in over 1 million individuals identify hundreds of loci and shared genetic influences. <i>Nature Genetics</i> , 2019 , 51, 245-257	36.3	259
44	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. <i>Nature Neuroscience</i> , 2018 , 21, 1656-1669	25.5	257
43	Genome-Wide Association Study Meta-Analysis of the Alcohol Use Disorders Identification Test (AUDIT) in Two Population-Based Cohorts. <i>American Journal of Psychiatry</i> , 2019 , 176, 107-118	11.9	156
42	The genetics of human personality. <i>Genes, Brain and Behavior</i> , 2018 , 17, e12439	3.6	88
41	Genome-Wide Association Studies of Impulsive Personality Traits (BIS-11 and UPPS-P) and Drug Experimentation in up to 22,861 Adult Research Participants Identify Loci in the and genes. <i>Journal of Neuroscience</i> , 2019 , 39, 2562-2572	6.6	77
40	Exaggerated waiting impulsivity associated with human binge drinking, and high alcohol consumption in mice. <i>Neuropsychopharmacology</i> , 2014 , 39, 2919-27	8.7	75
39	Genome-wide meta-analysis of problematic alcohol use in 435,563 individuals yields insights into biology and relationships with other traits. <i>Nature Neuroscience</i> , 2020 , 23, 809-818	25.5	69
38	Genome-wide association study of delay discounting in 23,217 adult research participants of European ancestry. <i>Nature Neuroscience</i> , 2018 , 21, 16-18	25.5	56
37	Genome-Wide Association Study of Loneliness Demonstrates a Role for Common Variation. <i>Neuropsychopharmacology</i> , 2017 , 42, 811-821	8.7	53
36	Genome-wide association study of alcohol use disorder identification test (AUDIT) scores in 20128 research participants of European ancestry. <i>Addiction Biology</i> , 2019 , 24, 121-131	4.6	49
35	Measuring impulsivity in mice: the five-choice serial reaction time task. <i>Psychopharmacology</i> , 2012 , 219, 253-70	4.7	46
34	A large-scale genome-wide association study meta-analysis of cannabis use disorder. <i>Lancet Psychiatry</i> , 2020 , 7, 1032-1045	23.3	43
33	Repeated ethanol exposure during early and late adolescence: double dissociation of effects on waiting and choice impulsivity. <i>Alcoholism: Clinical and Experimental Research</i> , 2014 , 38, 2579-89	3.7	30
32	Phenome-wide investigation of health outcomes associated with genetic predisposition to loneliness. <i>Human Molecular Genetics</i> , 2019 , 28, 3853-3865	5.6	29
31	Recent Efforts to Dissect the Genetic Basis of Alcohol Use and Abuse. <i>Biological Psychiatry</i> , 2020 , 87, 609-618	7.9	28
30	Heightened Impulsivity: Associated with Family History of Alcohol Misuse, and a Consequence of Alcohol Intake. <i>Alcoholism: Clinical and Experimental Research</i> , 2016 , 40, 2208-2217	3.7	25

29	Alpha-synuclein deletion decreases motor impulsivity but does not affect risky decision making in a mouse Gambling Task. <i>Psychopharmacology</i> , 2014 , 231, 2493-506	4.7	22
28	Mouse and Human Genetic Analyses Associate Kalirin with Ventral Striatal Activation during Impulsivity and with Alcohol Misuse. <i>Frontiers in Genetics</i> , 2016 , 7, 52	4.5	16
27	Multivariate GWAS of psychiatric disorders and their cardinal symptoms reveal two dimensions of cross-cutting genetic liabilities		15
26	Emerging phenotyping strategies will advance our understanding of psychiatric genetics. <i>Nature Neuroscience</i> , 2020 , 23, 475-480	25.5	14
25	Alleviating waiting impulsivity and perseverative responding by Opioid receptor antagonism in two inbred mouse strains. <i>Psychopharmacology</i> , 2015 , 232, 1483-92	4.7	12
24	Shared genetic risk between eating disorder- and substance-use-related phenotypes: Evidence from genome-wide association studies. <i>Addiction Biology</i> , 2021 , 26, e12880	4.6	12
23	Multivariate analysis of 1.5 million people identifies genetic associations with traits related to self-regulation and addiction. <i>Nature Neuroscience</i> , 2021 , 24, 1367-1376	25.5	10
22	Genome-wide association study meta-analysis of the Alcohol Use Disorder Identification Test (AUDIT) in two population-based cohorts (N=141,932)		9
21	Genetic risk for major depressive disorder and loneliness in sex-specific associations with coronary artery disease. <i>Molecular Psychiatry</i> , 2021 , 26, 4254-4264	15.1	9
20	Steep Discounting of Future Rewards as an Impulsivity Phenotype: A Concise Review. <i>Current Topics in Behavioral Neurosciences</i> , 2020 , 47, 113-138	3.4	7
19	Long-term wheel running changes on sensorimotor activity and skeletal muscle in male and female mice of accelerated senescence. <i>Age</i> , 2014 , 36, 9697		7
18	Trans-ancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders		7
17	The Latent Genetic Structure of Impulsivity and Its Relation to Internalizing Psychopathology. <i>Psychological Science</i> , 2020 , 31, 1025-1035	7.9	7
16	Item-Level Genome-Wide Association Study of the Alcohol Use Disorders Identification Test in Three Population-Based Cohorts. <i>American Journal of Psychiatry</i> , 2021 , appiajp202020091390	11.9	7
15	Genomic basis of delayed reward discounting. <i>Behavioural Processes</i> , 2019 , 162, 157-161	1.6	6
14	Genetic influences on delayed reward discounting: A genome-wide prioritized subset approach. <i>Experimental and Clinical Psychopharmacology</i> , 2019 , 27, 29-37	3.2	6
13	Multivariate genomic analysis of 1.5 million people identifies genes related to addiction, antisocial behavior, and health		6
12	Meta-analysis of problematic alcohol use in 435,563 individuals identifies 29 risk variants and yields insights into biology, pleiotropy and causality		6

11	Electronic Health Records Are the Next Frontier for the Genetics of Substance Use Disorders. <i>Trends in Genetics</i> , 2019 , 35, 317-318	8.5	5
10	Polygenic contributions to alcohol use and alcohol use disorders across population-based and clinically ascertained samples. <i>Psychological Medicine</i> , 2021 , 51, 1147-1156	6.9	5
9	Genome-wide association study of Alcohol Use Disorder Identification Test (AUDIT) scores in 20,328 research participants of European ancestry		4
8	Phenome-wide Investigation of Health Outcomes Associated with Genetic Predisposition to Loneliness		4
7	Alcohol and cigarette smoking consumption as genetic proxies for alcohol misuse and nicotine dependence. <i>Drug and Alcohol Dependence</i> , 2021 , 221, 108612	4.9	4
6	Evaluation of delay discounting as a transdiagnostic research domain criteria indicator in 1388 general community adults.. <i>Psychological Medicine</i> , 2022 , 1-9	6.9	2
5	Two data-driven approaches to identifying the spectrum of problematic opioid use: A pilot study within a chronic pain cohort. <i>International Journal of Medical Informatics</i> , 2021 , 156, 104621	5.3	1
4	Multivariate GWAS elucidates the genetic architecture of alcohol consumption and misuse, corrects biases, and reveals novel associations with disease		1
3	Integration of evidence across human and model organism studies: A meeting report. <i>Genes, Brain and Behavior</i> , 2021 , 20, e12738	3.6	1
2	Mapping Pathways by Which Genetic Risk Influences Adolescent Externalizing Behavior: The Interplay Between Externalizing Polygenic Risk Scores, Parental Knowledge, and Peer Substance Use. <i>Behavior Genetics</i> , 2021 , 51, 543-558	3.2	1
1	A mutant allele of glycoprotein M6-B (Gpm6b) facilitates behavioral flexibility but increases delay discounting.. <i>Genes, Brain and Behavior</i> , 2022 , e12800	3.6	