Scott I Vrieze

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9029980/publications.pdf

Version: 2024-02-01

60 papers

12,195 citations

25 h-index 60 g-index

73 all docs

73 docs citations

73 times ranked

23743 citing authors

#	Article	IF	CITATIONS
1	Associations between polygenic risk of substance use and use disorder and alcohol, cannabis, and nicotine use in adolescence and young adulthood in a longitudinal twin study. Psychological Medicine, 2023, 53, 2296-2306.	2.7	7
2	Genetic correlation, pleiotropy, and causal associations between substance use and psychiatric disorder. Psychological Medicine, 2022, 52, 968-978.	2.7	41
3	Alcohol and nicotine polygenic scores are associated with the development of alcohol and nicotine use problems from adolescence to young adulthood. Addiction, 2022, 117, 1117-1127.	1.7	11
4	Bayesian Forecasting with a Regime-Switching Zero-Inflated Multilevel Poisson Regression Model: An Application to Adolescent Alcohol Use with Spatial Covariates. Psychometrika, 2022, , 1 .	1.2	3
5	Developmental and etiological patterns of substance use from adolescence to middle age: A longitudinal twin study. Drug and Alcohol Dependence, 2022, 233, 109378.	1.6	5
6	Sequencing of 53,831 diverse genomes from the NHLBI TOPMed Program. Nature, 2021, 590, 290-299.	13.7	1,069
7	Using multivariate endophenotypes to identify psychophysiological mechanisms associated with polygenic scores for substance use, schizophrenia, and education attainment. Psychological Medicine, 2021 , , $1-11$.	2.7	3
8	Genetic architecture of four smoking behaviors using partitioned SNP heritability. Addiction, 2021, 116, 2498-2508.	1.7	14
9	Model-based assessment of replicability for genome-wide association meta-analysis. Nature Communications, 2021, 12, 1964.	5.8	24
10	Associations between adolescent cannabis use and young-adult functioning in three longitudinal twin studies. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	28
11	Polygenic Score for Smoking Is Associated With Externalizing Psychopathology and Disinhibited Personality Traits but Not Internalizing Psychopathology in Adolescence. Clinical Psychological Science, 2021, 9, 1205-1213.	2.4	7
12	Mechanisms of parent–child transmission of tobacco and alcohol use with polygenic risk scores: Evidence for a genetic nurture effect Developmental Psychology, 2021, 57, 796-804.	1.2	6
13	Baseline brain function in the preadolescents of the ABCD Study. Nature Neuroscience, 2021, 24, 1176-1186.	7.1	48
14	Multivariate analysis of 1.5 million people identifies genetic associations with traits related to self-regulation and addiction. Nature Neuroscience, 2021, 24, 1367-1376.	7.1	137
15	Polygenic scores for smoking and educational attainment have independent influences on academic success and adjustment in adolescence and educational attainment in adulthood. PLoS ONE, 2021, 16, e0255348.	1.1	4
16	Adolescent cannabis use and adult psychoticism: A longitudinal co-twin control analysis using data from two cohorts Journal of Abnormal Psychology, 2021, 130, 691-701.	2.0	5
17	Substance use patterns in 9-10 year olds: Baseline findings from the adolescent brain cognitive development (ABCD) study. Drug and Alcohol Dependence, 2021, 227, 108946.	1.6	19
18	Validating Online Measures of Cognitive Ability in Genes for Good, a Genetic Study of Health and Behavior. Assessment, 2020, 27, 136-148.	1.9	7

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19	Meta-analysis of up to 622,409 individuals identifies 40 novel smoking behaviour associated genetic loci. Molecular Psychiatry, 2020, 25, 2392-2409.	4.1	83
20	A large-scale genome-wide association study meta-analysis of cannabis use disorder. Lancet Psychiatry,the, 2020, 7, 1032-1045.	3.7	200
21	Dissecting the genetic overlap of smoking behaviors, lung cancer, and chronic obstructive pulmonary disease: A focus on nicotinic receptors and nicotine metabolizing enzyme. Genetic Epidemiology, 2020, 44, 748-758.	0.6	7
22	Expanding the genetic architecture of nicotine dependence and its shared genetics with multiple traits. Nature Communications, 2020, 11, 5562.	5.8	80
23	Association Analysis and Meta-Analysis of Multi-Allelic Variants for Large-Scale Sequence Data. Genes, 2020, 11, 586.	1.0	3
24	Adolescent Externalizing Psychopathology and Its Prospective Relationship to Marijuana Use Development from Age 14 to 30: Replication Across Independent Longitudinal Twin Samples. Behavior Genetics, 2020, 50, 139-151.	1.4	6
25	Genes for Good: Engaging the Public in Genetics Research via Social Media. American Journal of Human Genetics, 2019, 105, 65-77.	2.6	16
26	Minnesota Center for Twin and Family Research. Twin Research and Human Genetics, 2019, 22, 746-752.	0.3	27
27	Association studies of up to 1.2 million individuals yield new insights into the genetic etiology of tobacco and alcohol use. Nature Genetics, 2019, 51, 237-244.	9.4	1,307
28	Exome Chip Meta-analysis Fine Maps Causal Variants and Elucidates the Genetic Architecture of Rare Coding Variants in Smoking and AlcoholÂUse. Biological Psychiatry, 2019, 85, 946-955.	0.7	69
29	Narrow-sense heritability estimation of complex traits using identity-by-descent information. Heredity, 2018, 121, 616-630.	1.2	20
30	Comparison of methods that use whole genome data to estimate the heritability and genetic architecture of complex traits. Nature Genetics, 2018, 50, 737-745.	9.4	205
31	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. Nature Neuroscience, 2018, 21, 1656-1669.	7.1	490
32	Genome-wide association meta-analysis in 269,867 individuals identifies new genetic and functional links to intelligence. Nature Genetics, 2018, 50, 912-919.	9.4	893
33	Proper conditional analysis in the presence of missing data: Application to large scale meta-analysis of tobacco use phenotypes. PLoS Genetics, 2018, 14, e1007452.	1.5	18
34	Behavioral impact of return of genetic test results for complex disease: Systematic review and meta-analysis Health Psychology, 2018, 37, 1134-1144.	1.3	45
35	Deep Sequencing of 71 Candidate Genes to Characterize Variation Associated with Alcohol Dependence. Alcoholism: Clinical and Experimental Research, 2017, 41, 711-718.	1.4	13
36	Endophenotype best practices. International Journal of Psychophysiology, 2017, 111, 115-144.	0.5	62

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37	Next-generation genotype imputation service and methods. Nature Genetics, 2016, 48, 1284-1287.	9.4	2,828
38	A reference panel of 64,976 haplotypes for genotype imputation. Nature Genetics, 2016, 48, 1279-1283.	9.4	2,421
39	A computational method for genotype calling in family-based sequencing data. BMC Bioinformatics, 2016, 17, 37.	1.2	8
40	The Art of Smart Science: Weaving Theory and Risky Study Design Into Psychopathology Research and RDOC. Psychological Inquiry, 2015, 26, 286-292.	0.4	2
41	The Power of Theory, Research Design, and Transdisciplinary Integration in Moving Psychopathology Forward. Psychological Inquiry, 2015, 26, 209-230.	0.4	25
42	Rare variant genotype imputation with thousands of study-specific whole-genome sequences: implications for cost-effective study designs. European Journal of Human Genetics, 2015, 23, 975-983.	1.4	92
43	Gammaâ€Aminobutyric Acid System Genesâ€"No Evidence for a Role in Alcohol Use and Abuse in a Communityâ€Based Sample. Alcoholism: Clinical and Experimental Research, 2014, 38, 938-947.	1.4	14
44	Genetic associations of nonsynonymous exonic variants with psychophysiological endophenotypes. Psychophysiology, 2014, 51, 1300-1308.	1.2	21
45	In search of rare variants: Preliminary results from whole genome sequencing of 1,325 individuals with psychophysiological endophenotypes. Psychophysiology, 2014, 51, 1309-1320.	1.2	25
46	The Role of Constraint in the Development of Nicotine, Marijuana, and Alcohol Dependence in Young Adulthood. Behavior Genetics, 2014, 44, 14-24.	1.4	9
47	Rare Nonsynonymous Exonic Variants in Addiction and Behavioral Disinhibition. Biological Psychiatry, 2014, 75, 783-789.	0.7	41
48	Three Mutually Informative Ways to Understand the Genetic Relationships Among Behavioral Disinhibition, Alcohol Use, Drug Use, Nicotine Use/Dependence, and Their Co-occurrence: Twin Biometry, GCTA, and Genome-Wide Scoring. Behavior Genetics, 2013, 43, 97-107.	1.4	91
49	A Genome-Wide Association Study of Behavioral Disinhibition. Behavior Genetics, 2013, 43, 363-373.	1.4	119
50	Best Practices: The Electronic Medical Record Is an Invaluable Clinical Tool: Let's Start Using It. Psychiatric Services, 2013, 64, 946-949.	1.1	7
51	Decline in Genetic Influence on the Co-Occurrence of Alcohol, Marijuana, and Nicotine Dependence Symptoms From Age 14 to 29. American Journal of Psychiatry, 2012, 169, 1073-1081.	4.0	65
52	Model selection and psychological theory: A discussion of the differences between the Akaike information criterion (AIC) and the Bayesian information criterion (BIC) Psychological Methods, 2012, 17, 228-243.	2.7	1,160
53	Confluence of genes, environment, development, and behavior in a post Genome-Wide Association Study world. Development and Psychopathology, 2012, 24, 1195-1214.	1.4	43
54	The interplay of genes and adolescent development in substance use disorders: leveraging findings from GWAS meta-analyses to test developmental hypotheses about nicotine consumption. Human Genetics, 2012, 131, 791-801.	1.8	44

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55	Is the Continuity of Externalizing Psychopathology the Same in Adolescents and Middle–Aged Adults? A Test of the Externalizing Spectrum's Developmental Coherence. Journal of Abnormal Child Psychology, 2012, 40, 459-470.	3.5	24
56	An Assessment of the Individual and Collective Effects of Variants on Height Using Twins and a Developmentally Informative Study Design. PLoS Genetics, 2011, 7, e1002413.	1.5	11
57	Multidimensional assessment of criminal recidivism: Problems, pitfalls, and proposed solutions Psychological Assessment, 2010, 22, 382-395.	1.2	26
58	An exploration of the base rate scores of the Millon Clinical Multiaxial Inventory–III Psychological Assessment, 2009, 21, 57-67.	1.2	15
59	Survey on the use of clinical and mechanical prediction methods in clinical psychology Professional Psychology: Research and Practice, 2009, 40, 525-531.	0.6	53
60	Predicting sex offender recidivism. I. Correcting for item overselection and accuracy overestimation in scale development. II. Sampling error-induced attenuation of predictive validity over base rate information Law and Human Behavior, 2008, 32, 266-278.	0.6	27