Robert C Culverhouse

List of Publications by Year
in descending order

19 papers	856 citations	933447 10 h-index	794594 19 g-index
21 all docs	21 docs citations	21 times ranked	2365 citing authors

#	Article	IF	Citations
1	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. Nature Neuroscience, 2018, 21, 1656-1669.	14.8	490
2	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. Behavior Genetics, 2016, 46, 151-169.	2.1	98
3	E-cigarette Usage Is Associated With Increased Past-12-Month Quit Attempts and Successful Smoking Cessation in Two US Population–Based Surveys. Nicotine and Tobacco Research, 2019, 21, 1331-1338.	2.6	43
4	Protocol for a collaborative meta-analysis of 5-HTTLPR, stress, and depression. BMC Psychiatry, 2013, 13, 304.	2.6	35
5	Multiple distinct CHRNB3-CHRNA6 variants are genetic risk factors for nicotine dependence in African Americans and European Americans. Addiction, 2014, 109, 814-822.	3.3	34
6	Daily Drinking Is Associated with Increased Mortality. Alcoholism: Clinical and Experimental Research, 2018, 42, 2246-2255.	2.4	31
7	Genome-Wide Association Study of Heavy Smoking and Daily/Nondaily Smoking in the Hispanic Community Health Study/Study of Latinos (HCHS/SOL). Nicotine and Tobacco Research, 2018, 20, 448-457.	2.6	21
8	Genetic Variant in CHRNA5 and Response to Varenicline and Combination Nicotine Replacement in a Randomized Placebo–Controlled Trial. Clinical Pharmacology and Therapeutics, 2020, 108, 1315-1325.	4.7	17
9	The state of knowledge about the relationship between 5-HTTLPR, stress, and depression. Journal of Affective Disorders, 2018, 228, 205-206.	4.1	13
10	Genetic Analysis Workshop 18: Methods and strategies for analyzing human sequence and phenotype data in members of extended pedigrees. BMC Proceedings, 2014, 8, S1.	1.6	12
11	A Comparison of Methods Sensitive to Interactions With Small Main Effects. Genetic Epidemiology, 2012, 36, 303-311.	1.3	10
12	Uncovering hidden variance: pair-wise SNP analysis accounts for additional variance in nicotine dependence. Human Genetics, 2011, 129, 177-188.	3.8	8
13	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.	1.3	7
14	Studying the Utility of Using Genetics to Predict Smoking-Related Outcomes in a Population-Based Study and a Selected Cohort. Nicotine and Tobacco Research, 2021, 23, 2110-2116.	2.6	6
15	Stratify or adjust? Dealing with multiple populations when evaluating rare variants. BMC Proceedings, 2011, 5, S101.	1.6	5
16	Genotypic discrepancies arising from imputation. BMC Proceedings, 2014, 8, S17.	1.6	5
17	Gene by environment interactions. Genetic Epidemiology, 2007, 31, S68-S74.	1.3	4
18	Variants in the CHRNA5–CHRNA3–CHRNB4 Region of Chromosome 15 Predict Gastrointestinal Adverse Events in the Transdisciplinary Tobacco Use Research Center Smoking Cessation Trial. Nicotine and Tobacco Research, 2020, 22, 248-255.	2.6	4

#	Article	IF	Citations
19	Identifying cryptic population structure in multigenerational pedigrees in a Mexican American sample. BMC Proceedings, 2014, 8, S4.	1.6	1