

# Robert C Culverhouse

## List of Publications by Year in descending order

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Version: 2024-02-01

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	Studying the Utility of Using Genetics to Predict Smoking-Related Outcomes in a Population-Based Study and a Selected Cohort. <i>Nicotine and Tobacco Research</i> , 2021, 23, 2110-2116.	2.6	6
2	Variants in the CHRNA5-CHRNA3-CHRNA4 Region of Chromosome 15 Predict Gastrointestinal Adverse Events in the Transdisciplinary Tobacco Use Research Center Smoking Cessation Trial. <i>Nicotine and Tobacco Research</i> , 2020, 22, 248-255.	2.6	4
3	Dissecting the genetic overlap of smoking behaviors, lung cancer, and chronic obstructive pulmonary disease: A focus on nicotinic receptors and nicotine metabolizing enzyme. <i>Genetic Epidemiology</i> , 2020, 44, 748-758.	1.3	7
4	Genetic Variant in CHRNA5 and Response to Varenicline and Combination Nicotine Replacement in a Randomized Placebo-Controlled Trial. <i>Clinical Pharmacology and Therapeutics</i> , 2020, 108, 1315-1325.	4.7	17
5	E-cigarette Usage Is Associated With Increased Past-12-Month Quit Attempts and Successful Smoking Cessation in Two US Population-Based Surveys. <i>Nicotine and Tobacco Research</i> , 2019, 21, 1331-1338.	2.6	43
6	The state of knowledge about the relationship between 5-HTTLPR, stress, and depression. <i>Journal of Affective Disorders</i> , 2018, 228, 205-206.	4.1	13
7	Genome-Wide Association Study of Heavy Smoking and Daily/Nondaily Smoking in the Hispanic Community Health Study/Study of Latinos (HCHS/SOL). <i>Nicotine and Tobacco Research</i> , 2018, 20, 448-457.	2.6	21
8	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. <i>Nature Neuroscience</i> , 2018, 21, 1656-1669.	14.8	490
9	Daily Drinking Is Associated with Increased Mortality. <i>Alcoholism: Clinical and Experimental Research</i> , 2018, 42, 2246-2255.	2.4	31
10	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. <i>Behavior Genetics</i> , 2016, 46, 151-169.	2.1	98
11	Multiple distinct CHRNA3-CHRNA6 variants are genetic risk factors for nicotine dependence in African Americans and European Americans. <i>Addiction</i> , 2014, 109, 814-822.	3.3	34
12	Genetic Analysis Workshop 18: Methods and strategies for analyzing human sequence and phenotype data in members of extended pedigrees. <i>BMC Proceedings</i> , 2014, 8, S1.	1.6	12
13	Identifying cryptic population structure in multigenerational pedigrees in a Mexican American sample. <i>BMC Proceedings</i> , 2014, 8, S4.	1.6	1
14	Genotypic discrepancies arising from imputation. <i>BMC Proceedings</i> , 2014, 8, S17.	1.6	5
15	Protocol for a collaborative meta-analysis of 5-HTTLPR, stress, and depression. <i>BMC Psychiatry</i> , 2013, 13, 304.	2.6	35
16	A Comparison of Methods Sensitive to Interactions With Small Main Effects. <i>Genetic Epidemiology</i> , 2012, 36, 303-311.	1.3	10
17	Uncovering hidden variance: pair-wise SNP analysis accounts for additional variance in nicotine dependence. <i>Human Genetics</i> , 2011, 129, 177-188.	3.8	8
18	Stratify or adjust? Dealing with multiple populations when evaluating rare variants. <i>BMC Proceedings</i> , 2011, 5, S101.	1.6	5

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19	Gene by environment interactions. <i>Genetic Epidemiology</i> , 2007, 31, S68-S74.	1.3	4