

# Richard Border

## List of Publications by Year in descending order

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Version: 2024-02-01

13  
papers

814  
citations

1040056

9  
h-index

1125743

13  
g-index

21  
all docs

21  
docs citations

21  
times ranked

1751  
citing authors

#	ARTICLE	IF	CITATIONS
1	No Support for Historical Candidate Gene or Candidate Gene-by-Interaction Hypotheses for Major Depression Across Multiple Large Samples. <i>American Journal of Psychiatry</i> , 2019, 176, 376-387.	7.2	436
2	No Evidence That Schizophrenia Candidate Genes Are More Associated With Schizophrenia Than Noncandidate Genes. <i>Biological Psychiatry</i> , 2017, 82, 702-708.	1.3	170
3	Assortative mating biases marker-based heritability estimators. <i>Nature Communications</i> , 2022, 13, 660.	12.8	35
4	Genetic association study of childhood aggression across raters, instruments, and age. <i>Translational Psychiatry</i> , 2021, 11, 413.	4.8	31
5	Commentary: Fundamental problems with candidate gene-by-environment interaction studies – reflections on Moore and Thoenes (2016). <i>Journal of Child Psychology and Psychiatry and Allied Disciplines</i> , 2017, 58, 328-330.	5.2	26
6	Genome-wide Association Meta-analysis of Childhood and Adolescent Internalizing Symptoms. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2022, 61, 934-945.	0.5	26
7	Independent predictors of mortality in adolescents ascertained for conduct disorder and substance use problems, their siblings and community controls. <i>Addiction</i> , 2018, 113, 2107-2115.	3.3	15
8	Imputation of behavioral candidate gene repeat variants in 486,551 publicly-available UK Biobank individuals. <i>European Journal of Human Genetics</i> , 2019, 27, 963-969.	2.8	15
9	Continuity of Genetic Risk for Aggressive Behavior Across the Life-Course. <i>Behavior Genetics</i> , 2021, 51, 592-606.	2.1	13
10	Predictors of adult outcomes in clinically- and legally-ascertained youth with externalizing problems. <i>PLoS ONE</i> , 2018, 13, e0206442.	2.5	9
11	Stochastic Lanczos estimation of genomic variance components for linear mixed-effects models. <i>BMC Bioinformatics</i> , 2019, 20, 411.	2.6	5
12	Measurement Error Cannot Account for Failed Replications of Historic Candidate Gene-by-Environment Hypotheses: Response to Vrshek-Schallhorn et al.. <i>American Journal of Psychiatry</i> , 2019, 176, 668-669.	7.2	4
13	Little Evidence of Modified Genetic Effect of rs16969968 on Heavy Smoking Based on Age of Onset of Smoking. <i>Nicotine and Tobacco Research</i> , 2021, 23, 1055-1063.	2.6	4