Kristin K Nicodemus

List of Publications by Year
in descending order

19 papers	2,094 citations	623734 14 h-index	794594 19 g-index
20 all docs	20 docs citations	20 times ranked	5043 citing authors

#	Article	IF	Citations
1	Genome-wide association study of antidepressant treatment resistance in a population-based cohort using health service prescription data and meta-analysis with GENDEP. Pharmacogenomics Journal, 2020, 20, 329-341.	2.0	45
2	A review of neuroeconomic gameplay in psychiatric disorders. Molecular Psychiatry, 2020, 25, 67-81.	7.9	32
3	A major role for common genetic variation in anxiety disorders. Molecular Psychiatry, 2020, 25, 3292-3303.	7.9	243
4	Using tree-based methods for detection of geneâ€“gene interactions in the presence of a polygenic signal: simulation study with application to educational attainment in the Generation Scotland Cohort Study. Bioinformatics, 2019, 35, 181-188.	4.1	10
5	The role of polygenic risk score gene-set analysis in the context of the omnigenic model of schizophrenia. Neuropsychopharmacology, 2019, 44, 1562-1569.	5.4	44
6	Pharmaco-epidemiology of antidepressant exposure in a UK cohort record-linkage study. Journal of Psychopharmacology, 2019, 33, 482-493.	4.0	11
7	Self-reported medication use validated through record linkage to national prescribing data. Journal of Clinical Epidemiology, 2018, 94, 132-142.	5.0	75
8	Phenotypic and genetic analysis of cognitive performance in Major Depressive Disorder in the Generation Scotland: Scottish Family Health Study. Translational Psychiatry, 2018, 8, 63.	4.8	11
9	Enhancing Psychosis-Spectrum Nosology Through an International Data Sharing Initiative. Schizophrenia Bulletin, 2018, 44, S460-S467.	4.3	15
10	Contribution of copy number variants to schizophrenia from a genome-wide study of 41,321 subjects. Nature Genetics, 2017, 49, 27-35.	21.4	838
11	Data science for mental health: a UK perspective on a global challenge. Lancet Psychiatry,the, 2016, 3, 993-998.	7.4	47
12	An examination of the language construct in NIMH's research domain criteria: Time for reconceptualization!. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2016, 171, 904-919.	1.7	25
13	Variability in Working Memory Performance Explained by Epistasis vs Polygenic Scores in theZNF804APathway. JAMA Psychiatry, 2014, 71, 778.	11.0	28
14	Category fluency, latent semantic analysis and schizophrenia: a candidate gene approach. Cortex, 2014, 55, 182-191.	2.4	67
15	Evidence of statistical epistasis between DISC1, CIT and NDEL1 impacting risk for schizophrenia: biological validation with functional neuroimaging. Human Genetics, 2010, 127, 441-452.	3.8	93
16	The behaviour of random forest permutation-based variable importance measures under predictor correlation. BMC Bioinformatics, 2010, 11, 110.	2.6	254
17	Biological Validation of Increased Schizophrenia Risk With NRG1, ERBB4, and AKT1 Epistasis via Functional Neuroimaging in Healthy Controls. Archives of General Psychiatry, 2010, 67, 991.	12.3	113
18	Predictor correlation impacts machine learning algorithms: implications for genomic studies. Bioinformatics, 2009, 25, 1884-1890.	4.1	123

#	Article	IF	Citations
19	Stability of variable importance scores and rankings using statistical learning tools on single-nucleotide polymorphisms and risk factors involved in gene Ã— gene and gene Ã— environment interactions. BMC Proceedings, 2007, 1, S58.	1.6	10