Kimberly F Mcmanus

List of Publications by Year
in descending order

13 papers	1,892 citations	933447 10 h-index	1125743 13 g-index
17 all docs	17 docs citations	17 times ranked	3871 citing authors

#	Article	IF	Citations
1	Identification of novel risk loci, causal insights, and heritable risk for Parkinson's disease: a meta-analysis of genome-wide association studies. Lancet Neurology, The, 2019, 18, 1091-1102.	10.2	1,414
2	The Time Scale of Recombination Rate Evolution in Great Apes. Molecular Biology and Evolution, 2016, 33, 928-945.	8.9	92
3	Population genetic analysis of the DARC locus (Duffy) reveals adaptation from standing variation associated with malaria resistance in humans. PLoS Genetics, 2017, 13, e1006560.	3.5	87
4	Characterization of Prevalence and Health Consequences of Uniparental Disomy in Four Million Individuals from the General Population. American Journal of Human Genetics, 2019, 105, 921-932.	6.2	79
5	Inference of Gorilla Demographic and Selective History from Whole-Genome Sequence Data. Molecular Biology and Evolution, 2015, 32, 600-612.	8.9	55
6	Population History and Gene Divergence in Native Mexicans Inferred from 76 Human Exomes. Molecular Biology and Evolution, 2020, 37, 994-1006.	8.9	43
7	Genome-wide Study Identifies Association between HLA-Bâ—55:01 and Self-Reported Penicillin Allergy. American Journal of Human Genetics, 2020, 107, 612-621.	6.2	34
8	A Novel Test for Selection on cis-Regulatory Elements Reveals Positive and Negative Selection Acting on Mammalian Transcriptional Enhancers. Molecular Biology and Evolution, 2013, 30, 2509-2518.	8.9	25
9	Fast and Robust Identity-by-Descent Inference with the Templated Positional Burrowsâ€"Wheeler Transform. Molecular Biology and Evolution, 2021, 38, 2131-2151.	8.9	22
10	Mining Twitter Data to Improve Detection of Schizophrenia. AMIA Summits on Translational Science Proceedings, 2015, 2015, 122-6.	0.4	21
11	Sensor Data Integration: A New Cross-Industry Collaboration to Articulate Value, Define Needs, and Advance a Framework for Best Practices. Journal of Medical Internet Research, 2021, 23, e34493.	4.3	9
12	Bonsai: An efficient method for inferring large human pedigrees from genotype data. American Journal of Human Genetics, 2021, 108, 2052-2070.	6.2	7
13	popRange: a highly flexible spatially and temporally explicit Wright-Fisher simulator. Source Code for Biology and Medicine, 2015, 10, 6.	1.7	3