

# Kimberly F Mcmanus

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4296923/publications.pdf>

Version: 2024-02-01

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