## Caitlin Uren

## List of Publications by Year in descending order

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1478505 1281871 12 214 6 11 citations h-index g-index papers 20 20 20 376 docs citations times ranked all docs citing authors

#	Article	IF	CITATIONS
1	Fine-Scale Human Population Structure in Southern Africa Reflects Ecogeographic Boundaries. Genetics, 2016, 204, 303-314.	2.9	93
2	Putting RFMix and ADMIXTURE to the test in a complex admixed population. BMC Genetics, 2020, 21, 40.	2.7	29
3	Population structure and infectious disease risk in southern Africa. Molecular Genetics and Genomics, 2017, 292, 499-509.	2.1	21
4	A post-GWAS analysis of predicted regulatory variants and tuberculosis susceptibility. PLoS ONE, 2017, 12, e0174738.	2.5	19
5	Primary immunodeficiency diseases in a tuberculosis endemic region: challenges and opportunities. Genes and Immunity, 2019, 20, 447-454.	4.1	14
6	Hyperphosphatasia with mental retardation syndrome type 4 in three unrelated South African patients. American Journal of Medical Genetics, Part A, 2020, 182, 2230-2235.	1.2	8
7	Prospective avenues for human population genomics and disease mapping in southern Africa. Molecular Genetics and Genomics, 2020, 295, 1079-1089.	2.1	7
8	Local Ancestry Adjusted Allelic Association Analysis Robustly Captures Tuberculosis Susceptibility Loci. Frontiers in Genetics, 2021, 12, 716558.	2.3	7
9	Inferring recombination patterns in African populations. Human Molecular Genetics, 2021, 30, R11-R16.	2.9	6
10	African Genetic Representation in the Context of SARS-CoV-2 Infection and COVID-19 Severity. Frontiers in Genetics, 2022, 13, .	2.3	5
11	<i>Mycobacterium tuberculosis</i> complex and human coadaptation: a two-way street complicating host susceptibility to TB. Human Molecular Genetics, 2021, 30, R146-R153.	2.9	3
12	Local ancestry inference in heterogeneous populations—Are recent recombination events more relevant?. Briefings in Bioinformatics, 2021, 22, .	6.5	2