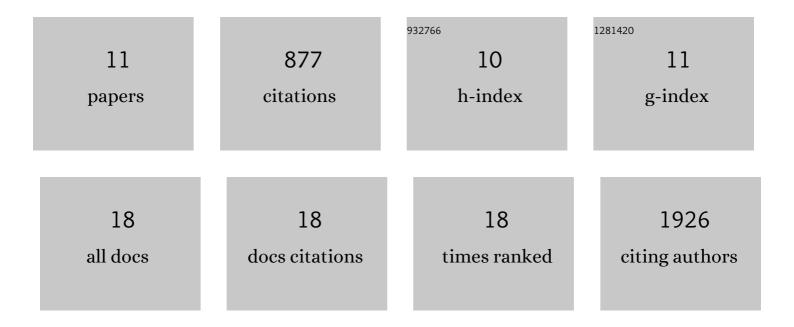
## **Tiffany Amariuta**

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7238840/publications.pdf Version: 2024-02-01



| #  | Article   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | Single-cell eQTL models reveal dynamic T cell state dependence of disease loci. Nature, 2022, 606, 120-128.   | 13.7 | 75        |
| 2  | Population-specific causal disease effect sizes in functionally important regions impacted by selection. Nature Communications, 2021, 12, 1098.   | 5.8  | 68        |
| 3  | Annotations capturing cell type-specific TF binding explain a large fraction of disease heritability.<br>Human Molecular Genetics, 2020, 29, 1057-1067.                                   | 1.4  | 16        |
| 4  | Advances in genetics toward identifying pathogenic cell states of rheumatoid arthritis.<br>Immunological Reviews, 2020, 294, 188-204.   | 2.8  | 23        |
| 5  | Improving the trans-ancestry portability of polygenic risk scores by prioritizing variants in predicted cell-type-specific regulatory elements. Nature Genetics, 2020, 52, 1346-1354.     | 9.4  | 126       |
| 6  | Large-scale genome-wide association study in a Japanese population identifies novel susceptibility loci<br>across different diseases. Nature Genetics, 2020, 52, 669-679.                 | 9.4  | 304       |
| 7  | Allele-specific expression changes dynamically during T cell activation in HLA and other autoimmune loci. Nature Genetics, 2020, 52, 247-253.   | 9.4  | 85        |
| 8  | Early progression to active tuberculosis is a highly heritable trait driven by 3q23 in Peruvians. Nature<br>Communications, 2019, 10, 3765.   | 5.8  | 43        |
| 9  | Benchmarker: An Unbiased, Association-Data-Driven Strategy to Evaluate Gene Prioritization<br>Algorithms. American Journal of Human Genetics, 2019, 104, 1025-1039.                       | 2.6  | 16        |
| 10 | IMPACT: Genomic Annotation of Cell-State-Specific Regulatory Elements Inferred from the Epigenome of Bound Transcription Factors. American Journal of Human Genetics, 2019, 104, 879-895. | 2.6  | 49        |
| 11 | Discovering in vivo cytokine-eQTL interactions from a lupus clinical trial. Genome Biology, 2018, 19,   | 3.8  | 36        |