## Tiffany Amariuta

## List of Publications by Year

 in descending orderSource: https:/|exaly.com/author-pdf/7238840/publications.pdf
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## 1

Large-scale genome-wide association study in a Japanese population identifies novel susceptibility loci across different diseases. Nature Genetics, 2020, 52, 669-679.
9.4

304

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

Allele-specific expression changes dynamically during $T$ cell activation in HLA and other autoimmune
9.4

85
loci. Nature Genetics, 2020, 52, 247-253.

Single-cell eQTL models reveal dynamic T cell state dependence of disease loci. Nature, 2022, 606, 120-128.
13.7

75

5 Population-specific causal disease effect sizes in functionally important regions impacted by selection. Nature Communications, 2021, 12, 1098.
$5.8 \quad 68$

6 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

## Early progression to active tuberculosis is a highly heritable trait driven by $3 q 23$ in Peruvians. Nature

Communications, 2019, 10, 3765.
5.8

43

8
Discovering in vivo cytokine-eQTL interactions from a lupus clinical trial. Genome Biology, 2018, 19, 168.

Advances in genetics toward identifying pathogenic cell states of rheumatoid arthritis. Immunological Reviews, 2020, 294, 188-204.

Benchmarker: An Unbiased, Association-Data-Driven Strategy to Evaluate Gene Prioritization Algorithms. American Journal of Human Genetics, 2019, 104, 1025-1039.

