

# Tiffany Amariuta

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7238840/publications.pdf>

Version: 2024-02-01

11  
papers

877  
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932766

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1281420

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#	ARTICLE	IF	CITATIONS
1	Single-cell eQTL models reveal dynamic T cell state dependence of disease loci. <i>Nature</i> , 2022, 606, 120-128.	13.7	75
2	Population-specific causal disease effect sizes in functionally important regions impacted by selection. <i>Nature Communications</i> , 2021, 12, 1098.	5.8	68
3	Annotations capturing cell type-specific TF binding explain a large fraction of disease heritability. <i>Human Molecular Genetics</i> , 2020, 29, 1057-1067.	1.4	16
4	Advances in genetics toward identifying pathogenic cell states of rheumatoid arthritis. <i>Immunological Reviews</i> , 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. <i>Nature Genetics</i> , 2020, 52, 1346-1354.	9.4	126
6	Large-scale genome-wide association study in a Japanese population identifies novel susceptibility loci across different diseases. <i>Nature Genetics</i> , 2020, 52, 669-679.	9.4	304
7	Allele-specific expression changes dynamically during T cell activation in HLA and other autoimmune loci. <i>Nature Genetics</i> , 2020, 52, 247-253.	9.4	85
8	Early progression to active tuberculosis is a highly heritable trait driven by 3q23 in Peruvians. <i>Nature Communications</i> , 2019, 10, 3765.	5.8	43
9	Benchmark: An Unbiased, Association-Data-Driven Strategy to Evaluate Gene Prioritization Algorithms. <i>American Journal of Human Genetics</i> , 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. <i>American Journal of Human Genetics</i> , 2019, 104, 879-895.	2.6	49
11	Discovering in vivo cytokine-eQTL interactions from a lupus clinical trial. <i>Genome Biology</i> , 2018, 19, 168.	3.8	36