

# Nicole M Ferraro

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9236057/publications.pdf>

Version: 2024-02-01

7  
papers

564  
citations

1684188

5  
h-index

1720034

7  
g-index

11  
all docs

11  
docs citations

11  
times ranked

1264  
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of rare-disease genes using blood transcriptome sequencing and large control cohorts. <i>Nature Medicine</i> , 2019, 25, 911-919.	30.7	221
2	Population-scale tissue transcriptomics maps long non-coding RNAs to complex disease. <i>Cell</i> , 2021, 184, 2633-2648.e19.	28.9	94
3	Transcriptomic signatures across human tissues identify functional rare genetic variation. <i>Science</i> , 2020, 369, .	12.6	89
4	A vast resource of allelic expression data spanning human tissues. <i>Genome Biology</i> , 2020, 21, 234.	8.8	68
5	Identification of rare and common regulatory variants in pluripotent cells using population-scale transcriptomics. <i>Nature Genetics</i> , 2021, 53, 313-321.	21.4	42
6	Integration of rare expression outlier-associated variants improves polygenic risk prediction. <i>American Journal of Human Genetics</i> , 2022, 109, 1055-1064.	6.2	8
7	Evaluating the Genomic Parameters Governing rAAV-Mediated Homologous Recombination. <i>Molecular Therapy</i> , 2021, 29, 1028-1046.	8.2	6