

# Nicole M Ferraro

## List of Publications by Year in descending order

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

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1264  
citing authors

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