



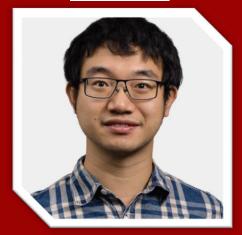
Join us for the NJ Acts Biostatistics and Epidemiology Workshop Series:

Thursday, April 13, 2023 12:30pm - 2:00pm

Integrating Multi-omics Summary Data Using Mendelian Randomization

This workshop is designed for anyone interested in learning about Mendelian randomization, a versatile tool for using genetic variation to identify how an exposure affects an outcome in observational studies. The presentation will focus on the use of multi-omics summary data to prioritize genes associated with a disease outcome.

Presenter:



Chong Jin, PhD
Assistant Professor, Mathematical
Sciences at NJIT

Target Audience:

Everyone is welcome, especially those interested in genetic and genomic data analysis.



Please click the <u>link</u> or scan the QR code to register Reach out to <u>natale.mazzaferro@rutgers.edu</u> with any questions