



NHLBI Trans-Omics for Precision Medicine

- [Get Started](#)
- [Centers](#)
 - [Administrative Coordinating Center \(ACC\)](#)
 - [Informatics Research Center \(IRC\)](#)
 - [NIH](#)
 - [NHLBI](#)
 - [Omics Centers](#)
 - [Baylor Human Genome Sequencing Center](#)
 - [Baylor-UTHealth Metabolomics Center](#)
 - [Broad Institute Genomics Platform](#)
 - [Broad Institute Metabolomics Platform](#)
 - [Broad Institute and Beth Israel Proteomics Platform](#)
 - [Illumina](#)
 - [Keck MGC](#)
 - [New York Genome Center Genomics](#)
 - [Northwest Genomics Center](#)
 - [New York Genome Center RNA-seq](#)
 - [Psomagen](#)
- [Projects/Studies](#)
 - [View All TOPMed Projects and their Parent Studies](#)
 - [Analysis Programs](#)
 - [NIH-sponsored investigator-initiated award centered on TOPMed data](#)
 - [U01 Integrative Omics Approaches for Analysis of TOPMed Data](#)
 - [R01 Integrative Computational Biology for Analysis of NHLBI TOPMed Data](#)
 - [Working Group Analysis Support Program](#)
 - [TOPMed Fellowship](#)
 - [Participant/Study Summary](#)
 - [Parent Study Descriptions & Ethics Statements](#)
 - [External Collaborations](#)
- [Working Groups](#)
 - [View All Working Groups](#)
- [Data](#)
 - [Data Access](#)
 - [Data Sets](#)
 - [DCC-harmonized phenotypes for the scientific community](#)
 - [Standards](#)
- [Publications](#)
 - [Publications](#)
 - [Abstracts](#)

- [Acknowledgements](#)
 - [TOPMed Acknowledgements](#)
 - [TOPMed Banner Authorship](#)
- [Recommendations on the use and reporting of race, ethnicity, and ancestry in TOPMed](#)
- [EEP](#)
- [ELSI](#)
- [Workshops](#)
 - [2020 ASHG Ancillary Session](#)
 - [2019 DCC Analysis Workshop in Seattle](#)
 - [2018 DCC Analysis Workshop in Seattle](#)
 - [2018 IRC Analysis Workshop at Michigan](#)
 - [2017 TOPMed DCC Analysis Workshop](#)

[Home](#) > [Abstracts](#)

Phenome-wide and molecular consequences of inbreeding