# PhenoPLIER: integrating transcriptome-wide association studies with machine learning

This manuscript (<u>permalink</u>) was automatically generated from <u>greenelab/phenoplier manuscript@11b238c</u> on July 3, 2020.

### **Authors**

#### Milton Pividori

Department of Systems Pharmacology and Translational Therapeutics, Perelman School of Medicine, University of Pennsylvania, Philadelphia, PA, USA 19104 · Funded by The Gordon and Betty Moore Foundation GBMF 4552; The National Human Genome Research Institute (R01 HG010067)

#### Casey S. Greene

Department of Systems Pharmacology and Translational Therapeutics, Perelman School of Medicine, University of Pennsylvania, Philadelphia, PA, 19104; Childhood Cancer Data Lab, Alex's Lemonade Stand Foundation, Philadelphia, PA, USA 19102 · Funded by The Gordon and Betty Moore Foundation GBMF 4552; The National Human Genome Research Institute (R01 HG010067); The National Cancer Institute (R01 CA237170)

## **Abstract**

## References