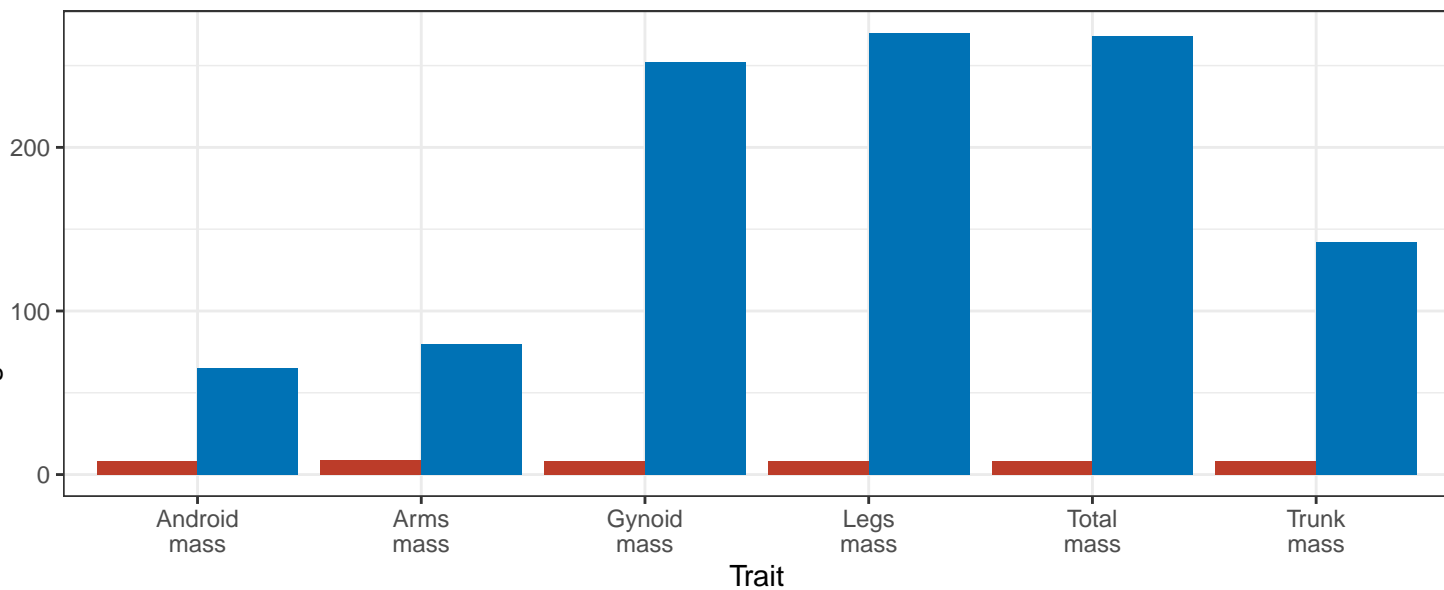


Method Standard SynSurr

A

Genome-wide
Significant Associations



B

Average χ^2 at
Genome-wide Significant Variants

