FINEMAP
ULTRAFAST SOFTWARE FOR HIGH-RESOLUTION FINE-MAPPING OF GENOMIC REGIONS

Implications: Identification of causal variant patterns in complex disease GWAS data
Unmet need: Current gold standard software is slow
IP Status: Licenses to software are available
Project status: Functional software is ready and in use

FINEMAP identifies causal SNP variants accurately

- Ultrafast identification of causal variants
- Superior performance in processing times
- Does not slow down when analyzing complex datasets