Records identified through database searching and manual searching (n=294 abstracts)

Excluded duplicates (n=59)

Titles/abstracts screened (n=236)

Records excluded (no genotyping, not original research) (n=176)

Full-text articles assessed for inclusion/exclusion (n=60)

Full-text articles excluded: (n=44)
  No risk endpoint = 27
  Partial genotyping = 8
  Not screening population = 3
  Not original research = 2
  No genotyping = 2
  Modeling study = 2

Studies included in synthesis (n=16)