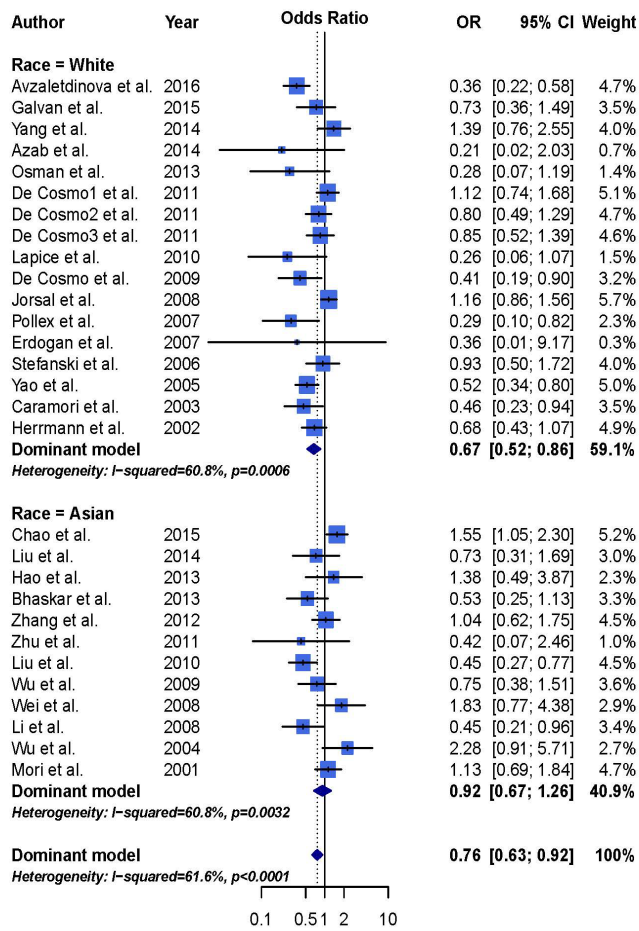


A

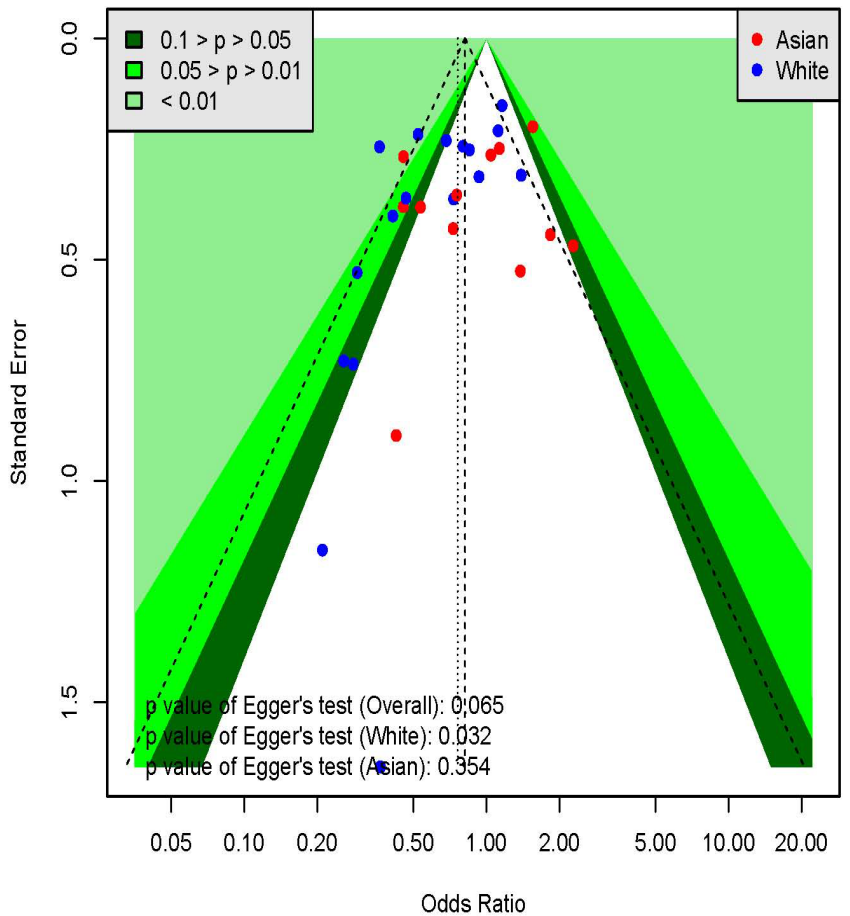
All



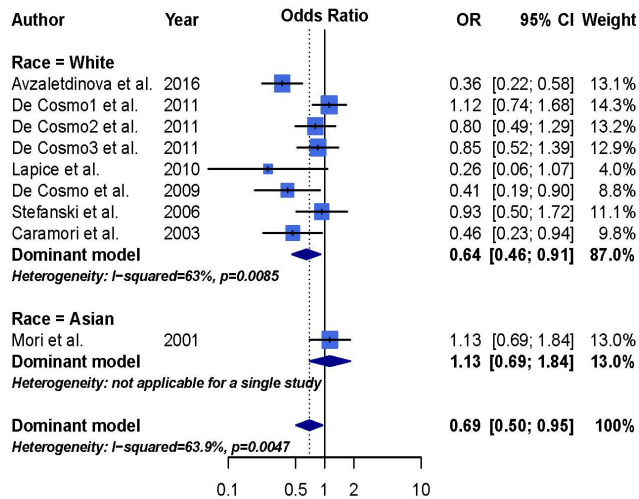
B

All

Dominant model

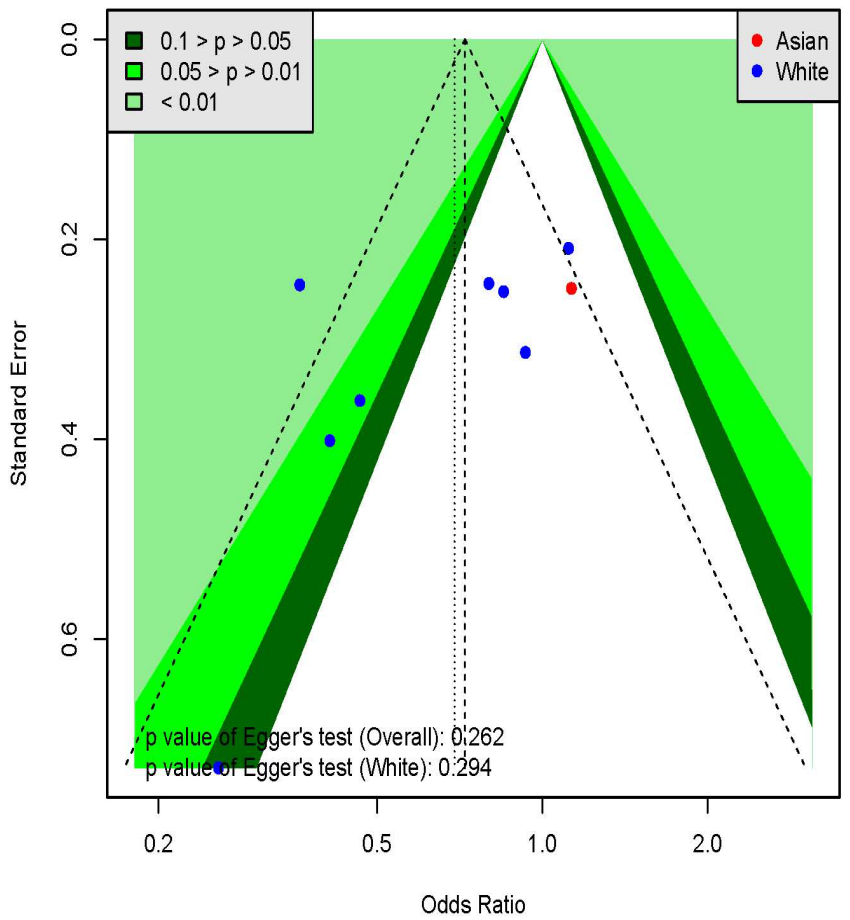


C Incomplete genotype

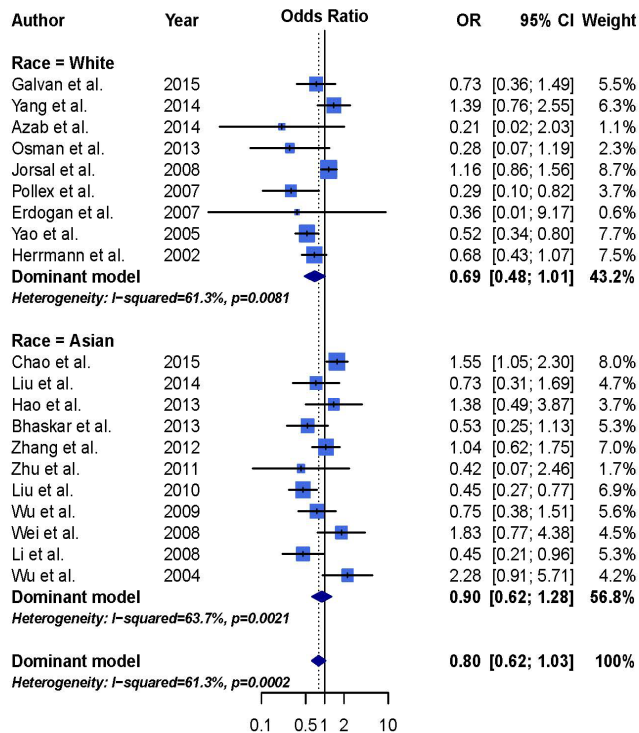


D Incomplete genotype

Dominant model



E Complete genotype



F

Complete genotype

Dominant model

