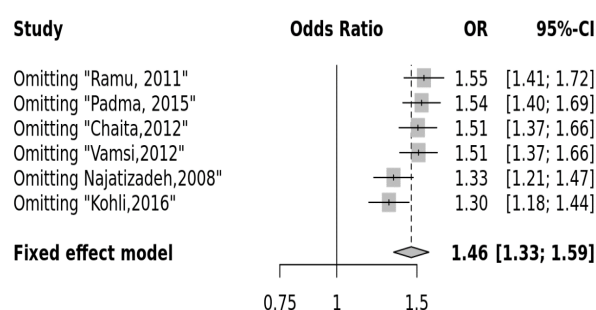
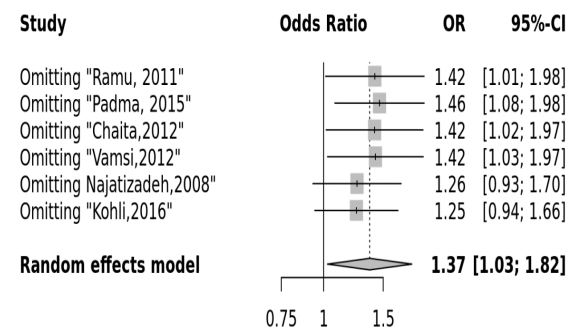


A

(i)

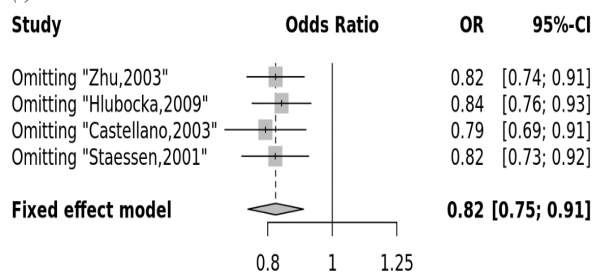


(ii)

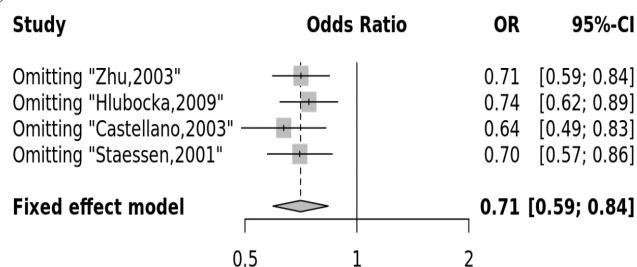


B

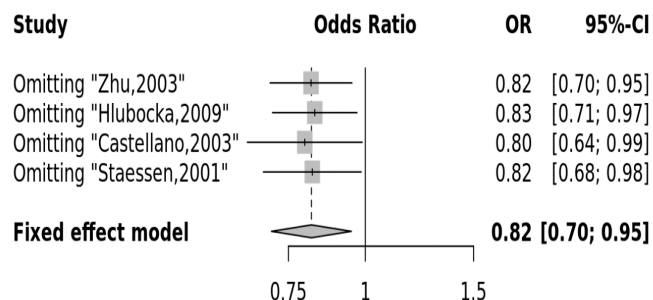
(i)



(ii)

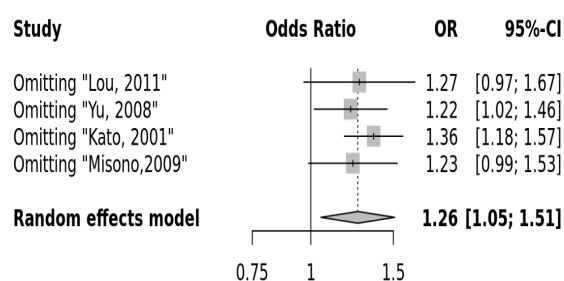


(iii)

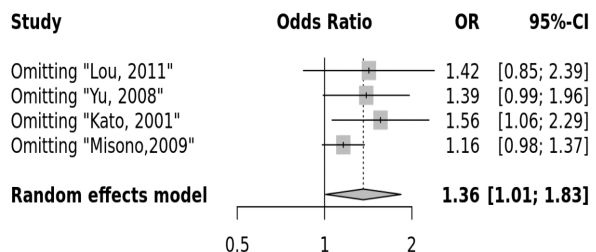


C

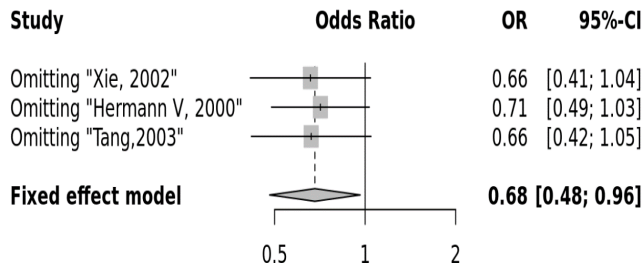
(i)



(ii)



(iii)



Supplementary Fig. 3. Sensitivity analysis of (A) *AGT*-rs699, India; north and south India (i) allele model and (ii) dominant model. (B) *CYP11B2*-rs1799998 (i) Allele model, (ii) recessive model, (iii) dominant model in Europe by excluding Bengra *et al*²⁶. (C) *ADRB2*-rs1042713, East Asia (i) allele model, (ii) recessive model and (iii) African America (dominant model) in Indian population ($P < 0.05$).