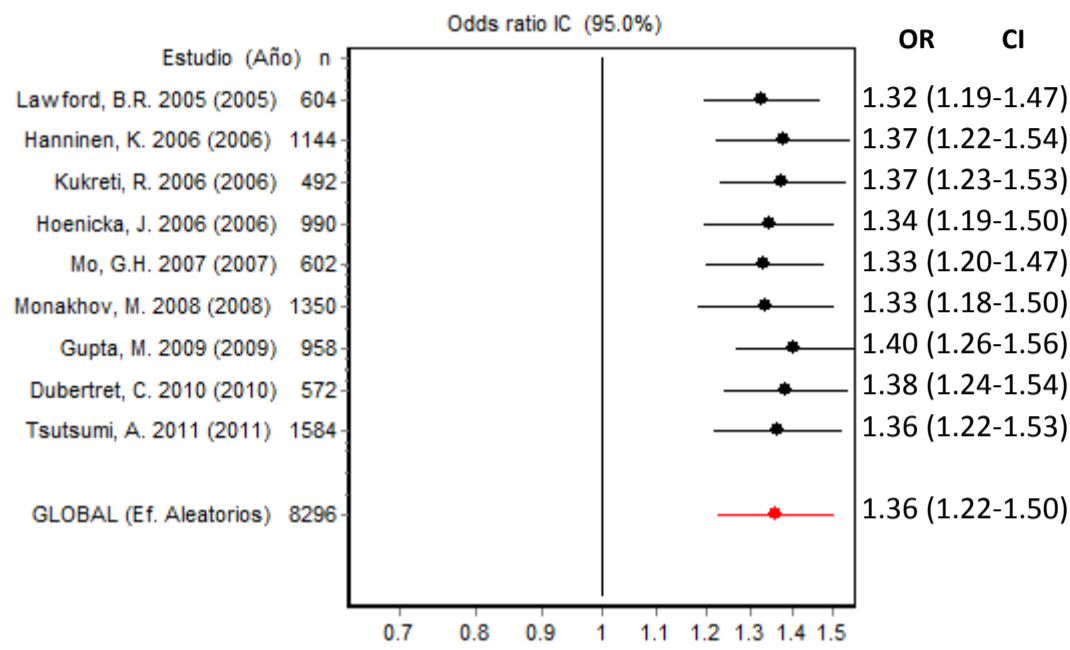
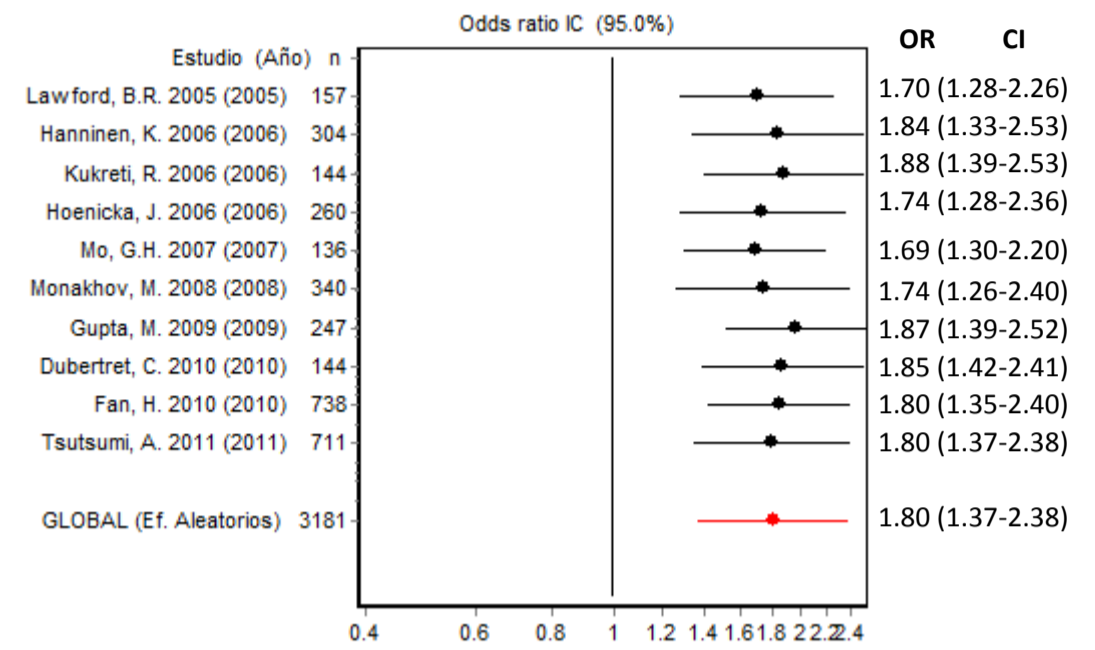


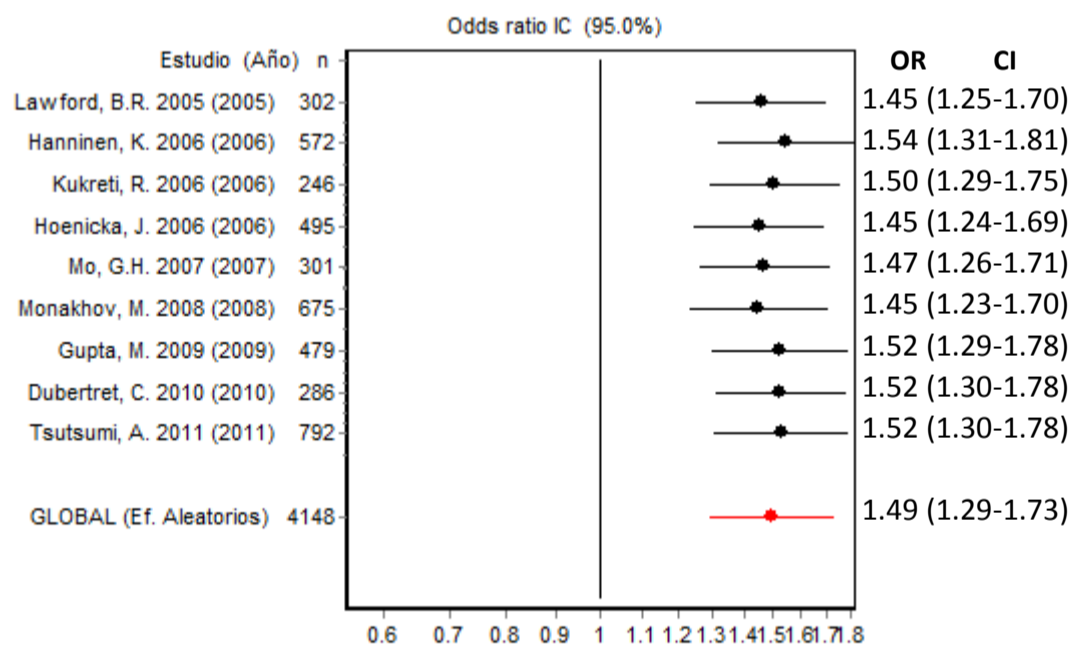
Figure S1. Sensitivity analysis of C957T polymorphism with SZ in whole population:
a) Allelic b) Additive c) Dominant and d) Recessive models.



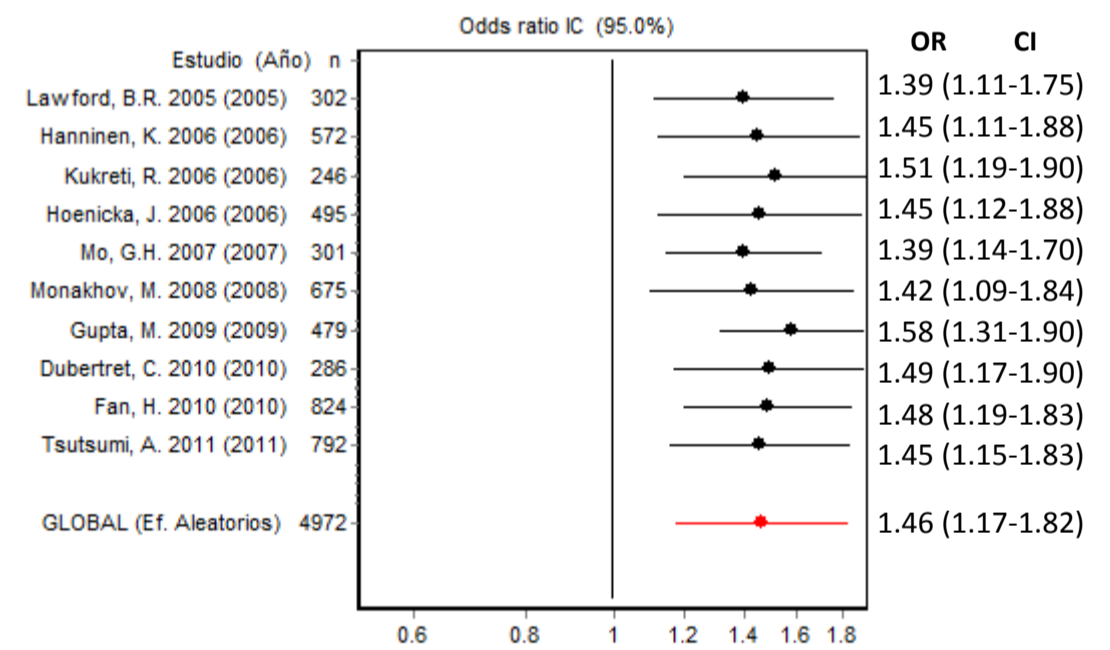
A)



B)

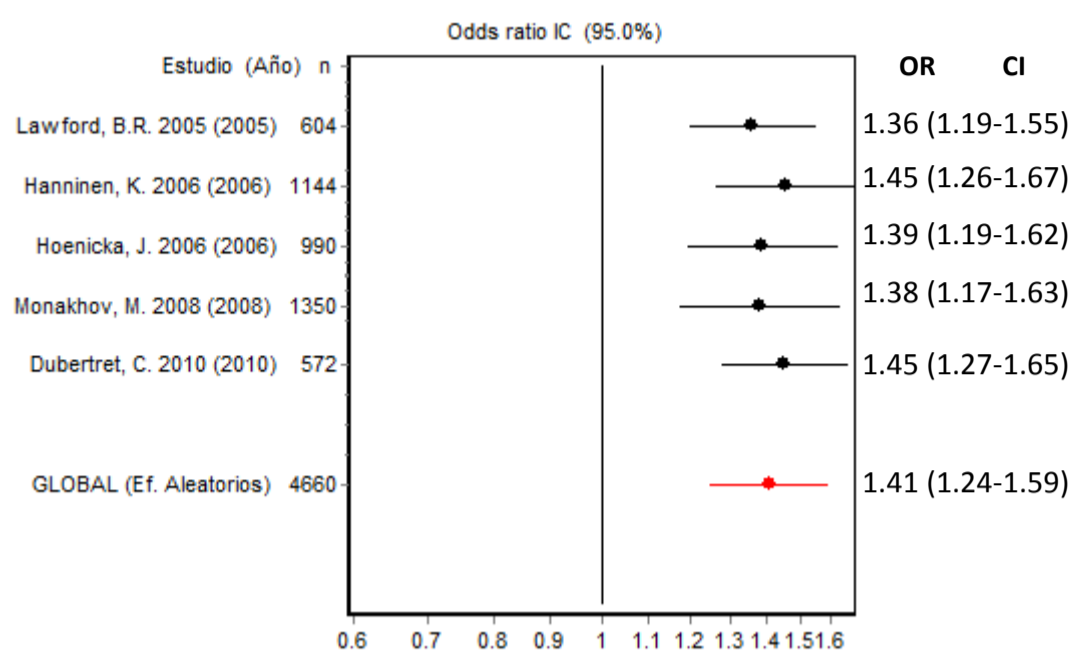


C)

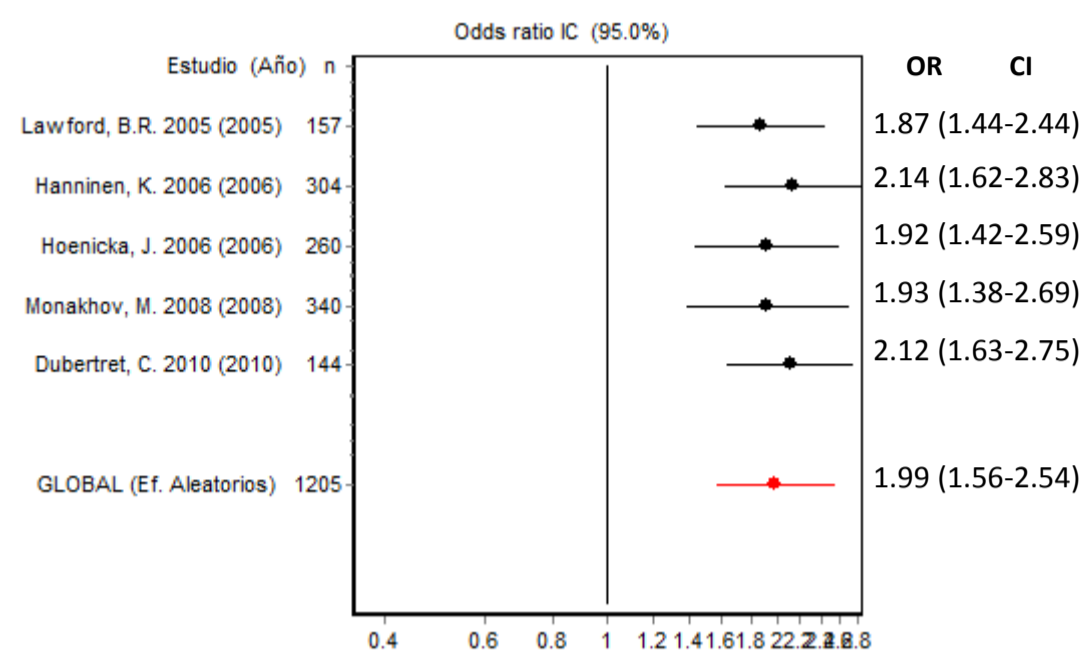


D)

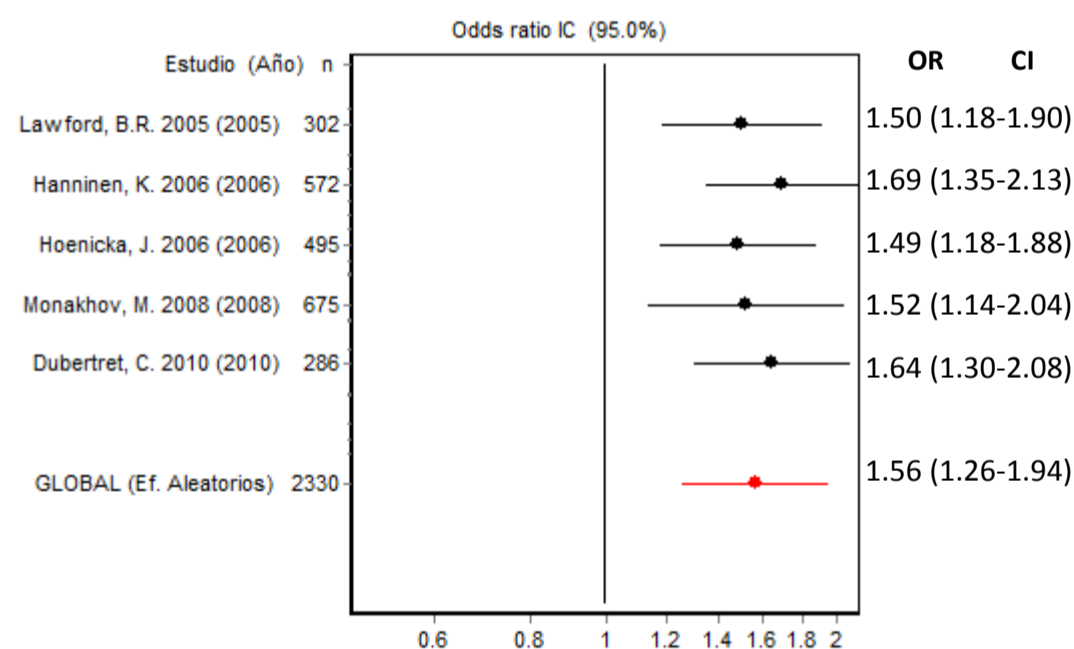
Figure S2. Sensitivity analysis of C957T polymorphism with SZ in Caucasians population:
a) Allelic b) Additive c) Dominant and d) Recessive models.



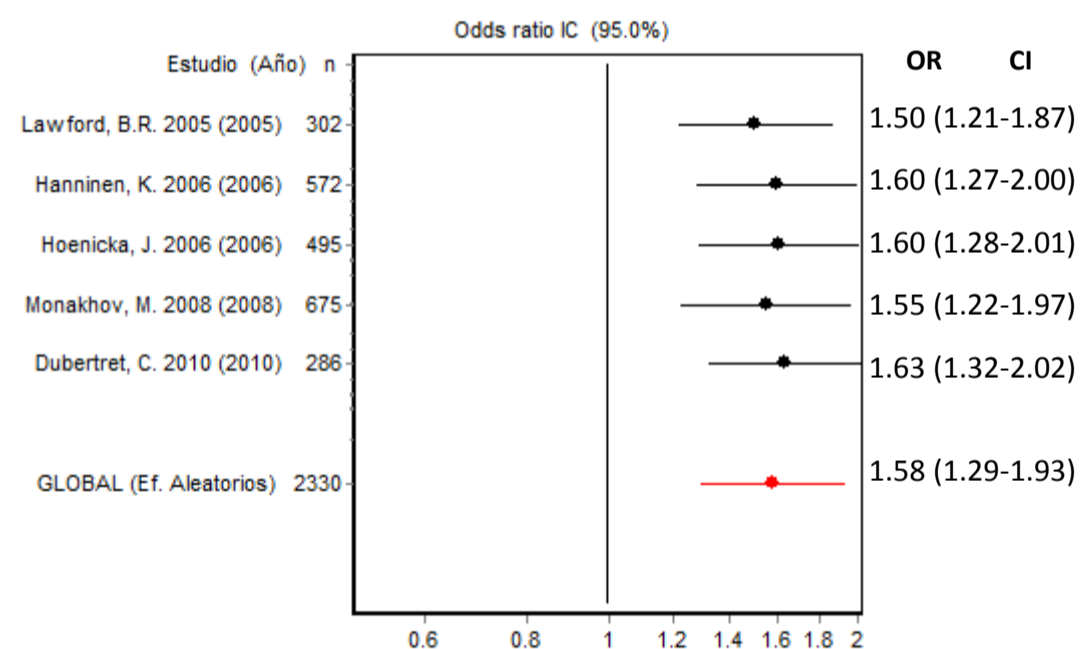
A)



B)



C)



D)

Figure S3. Sensitivity analysis of C957T polymorphism with SZ in Asians population:
a) Allelic b) Additive c) Dominant and d) Recessive models.

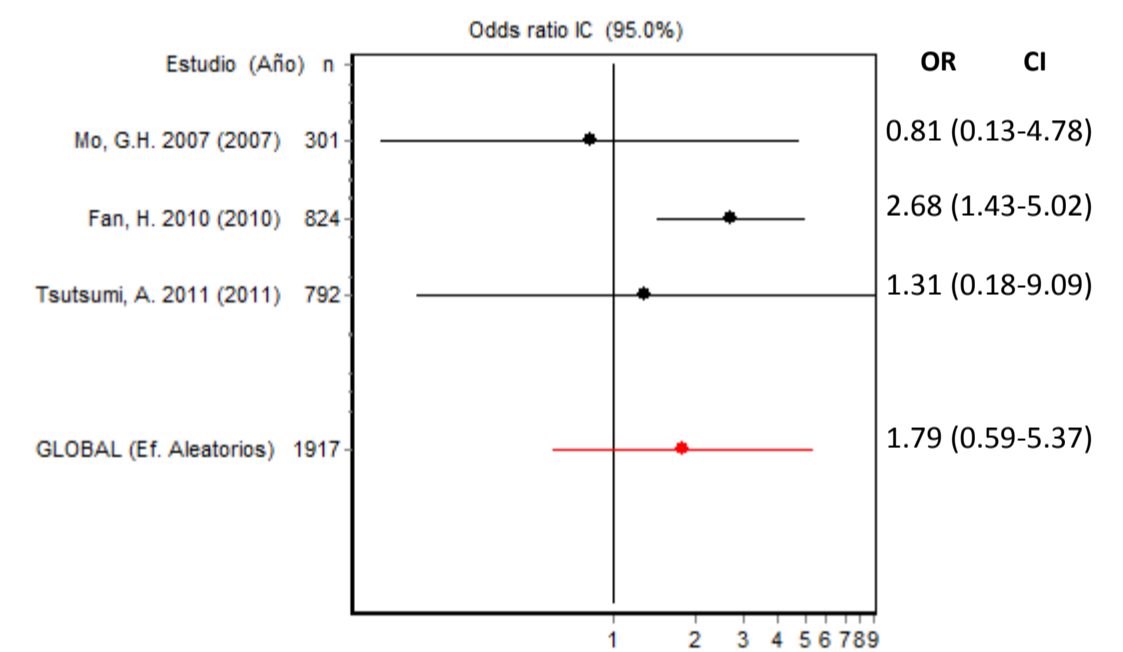
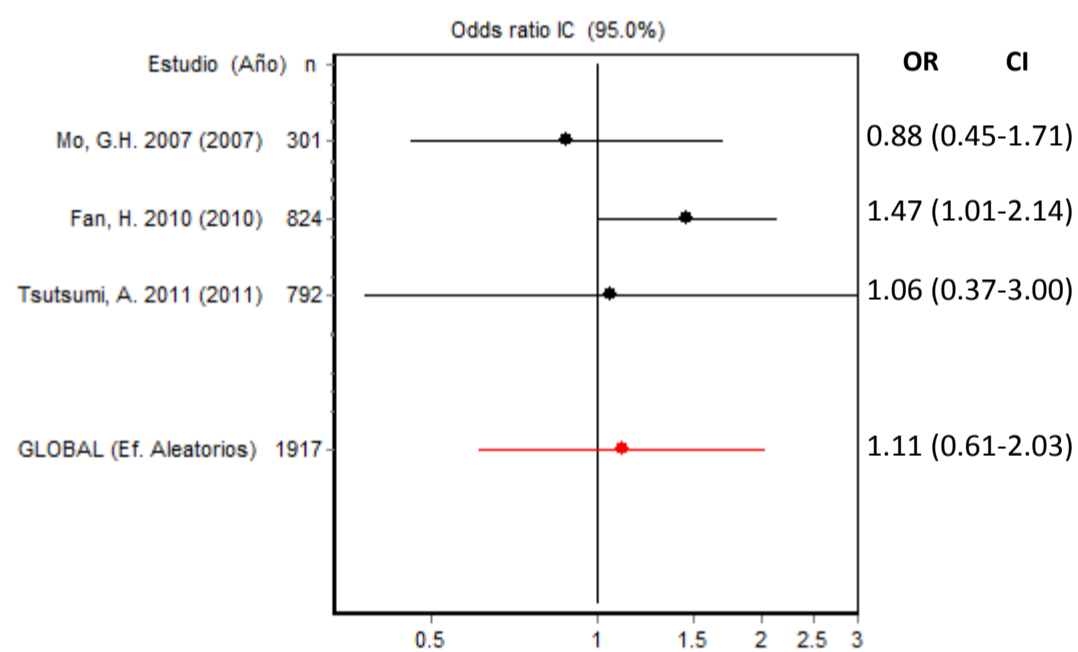
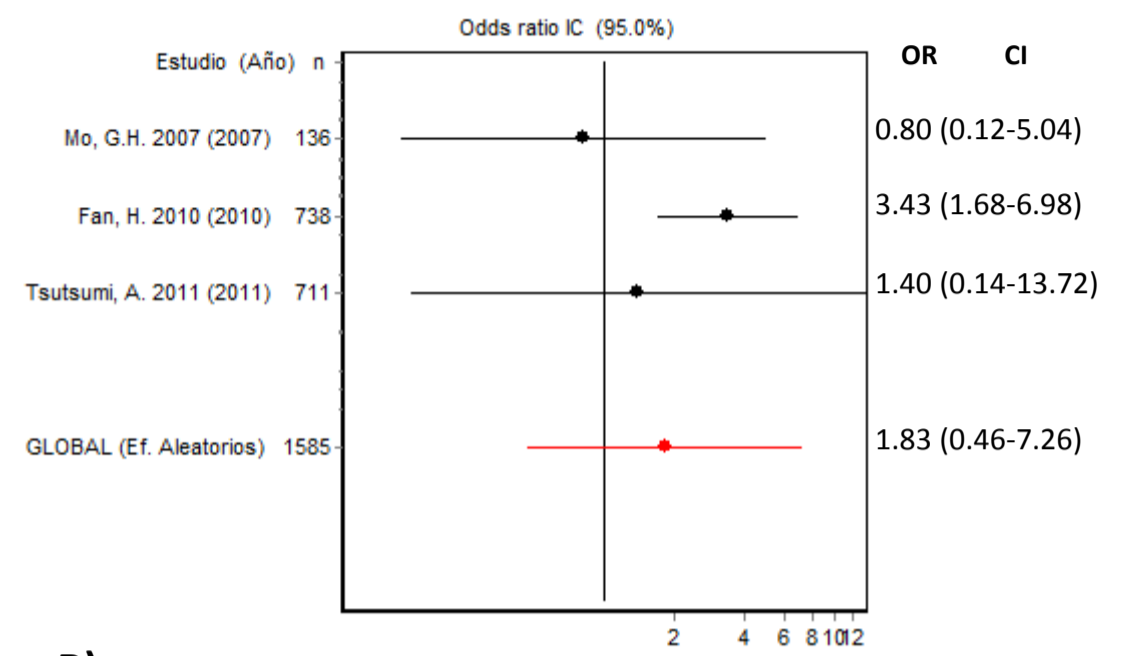
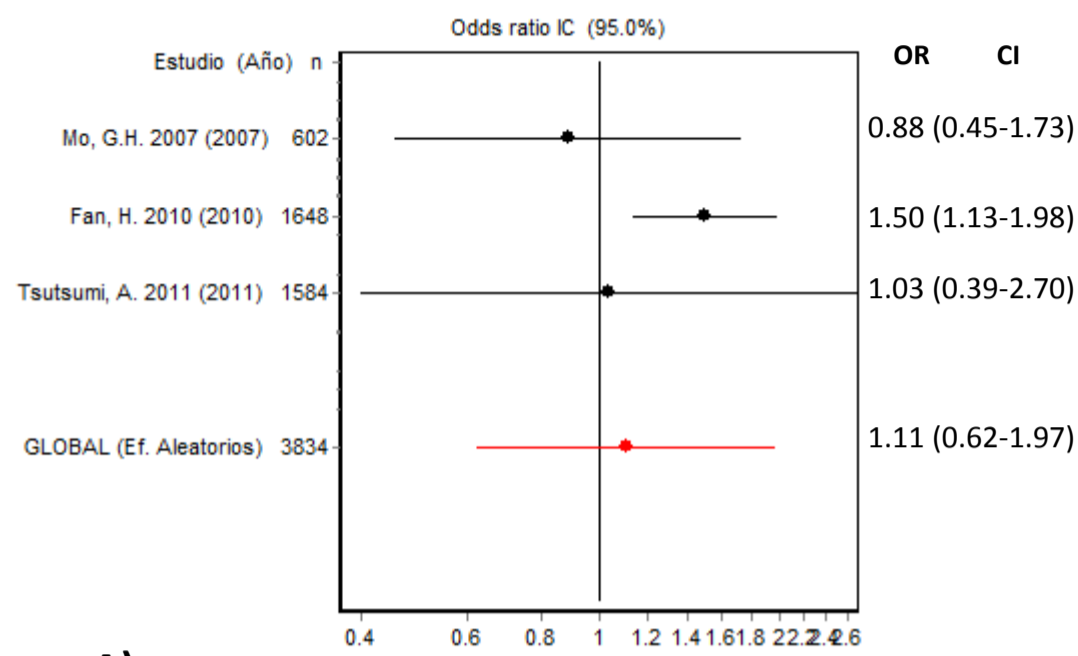
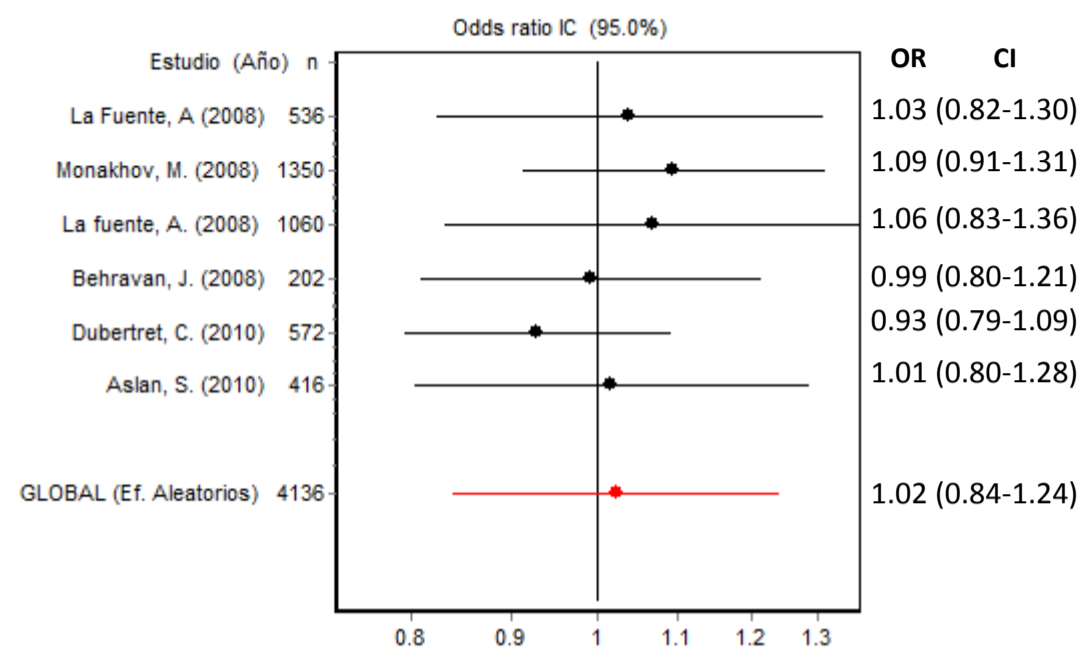
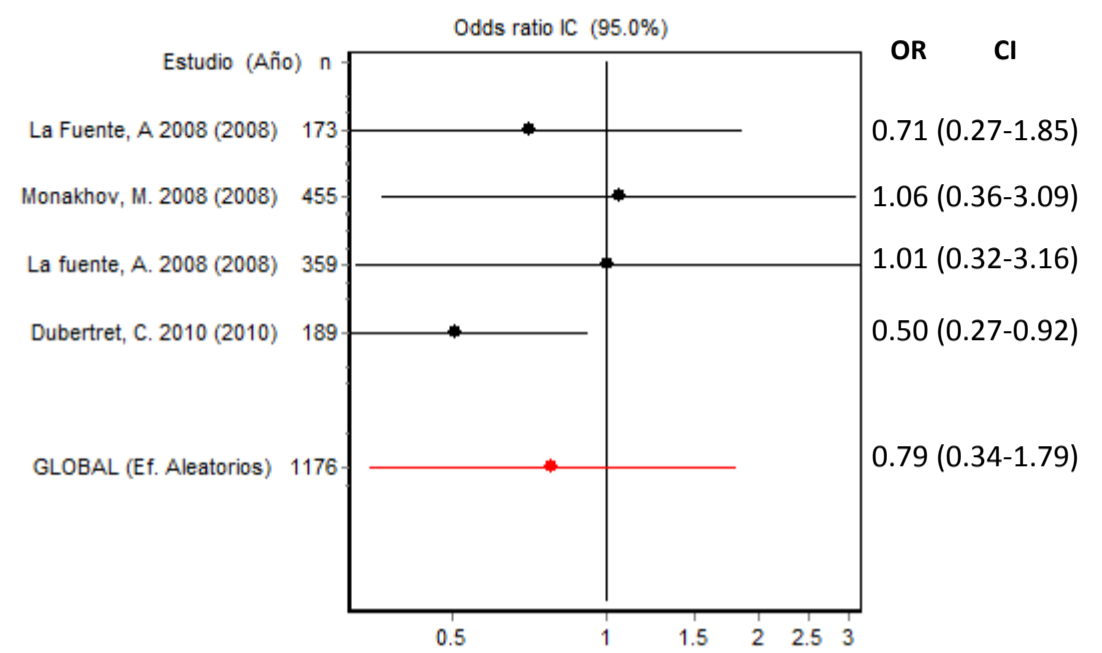


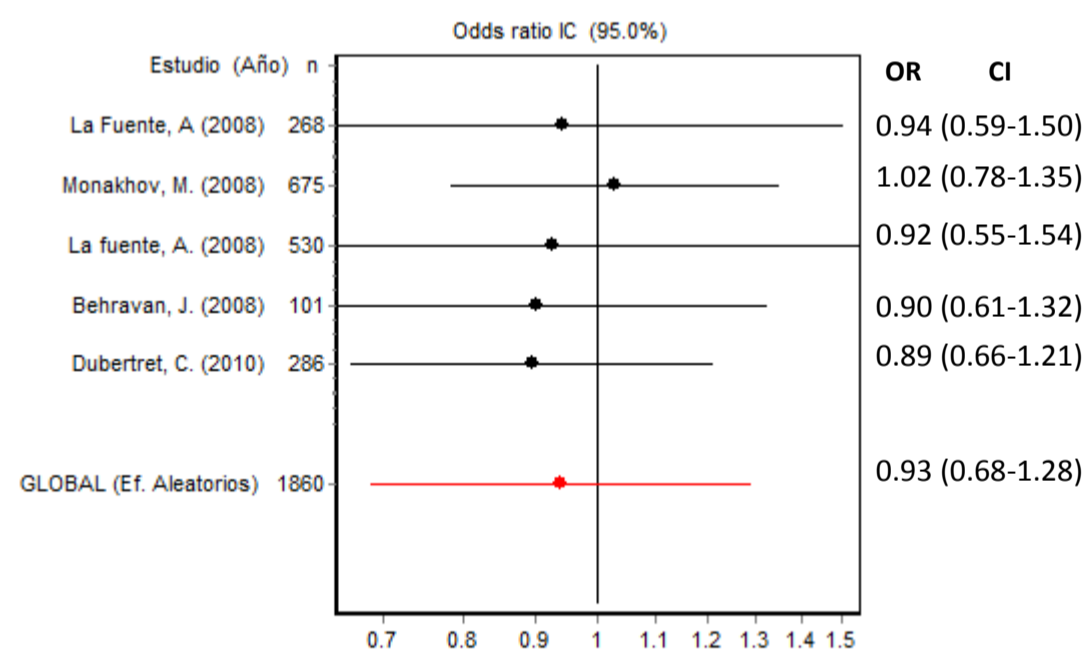
Figure S5. Sensitivity analysis of TaqI polymorphism with SZ in whole population:
a) Allelic b) Additive c) Dominant and d) Recessive models.



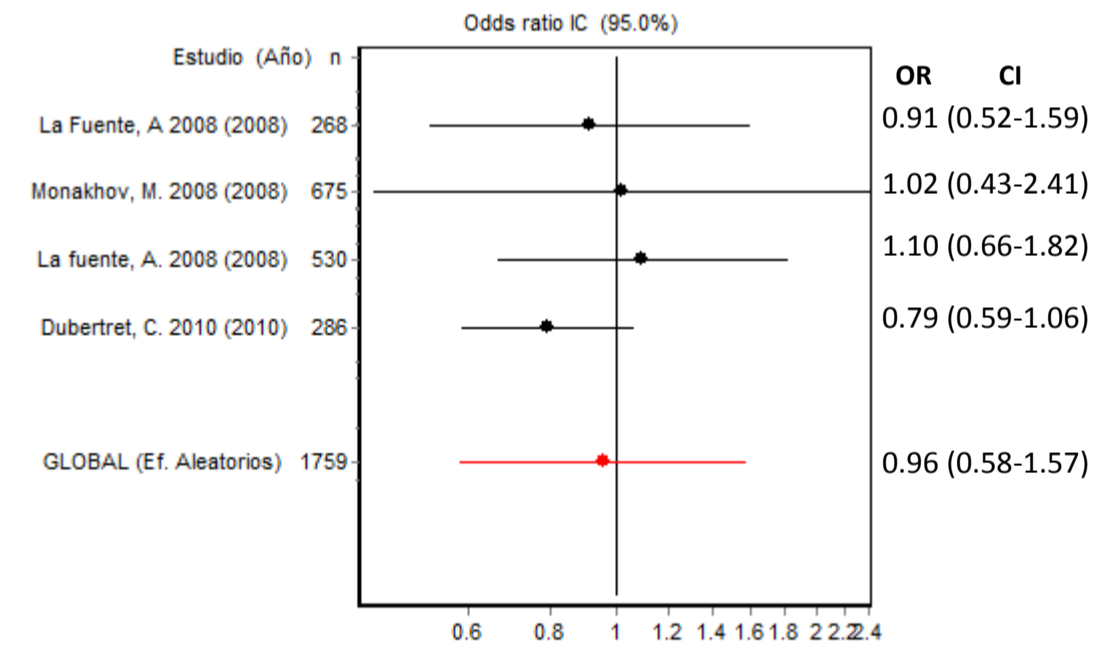
A)



B)

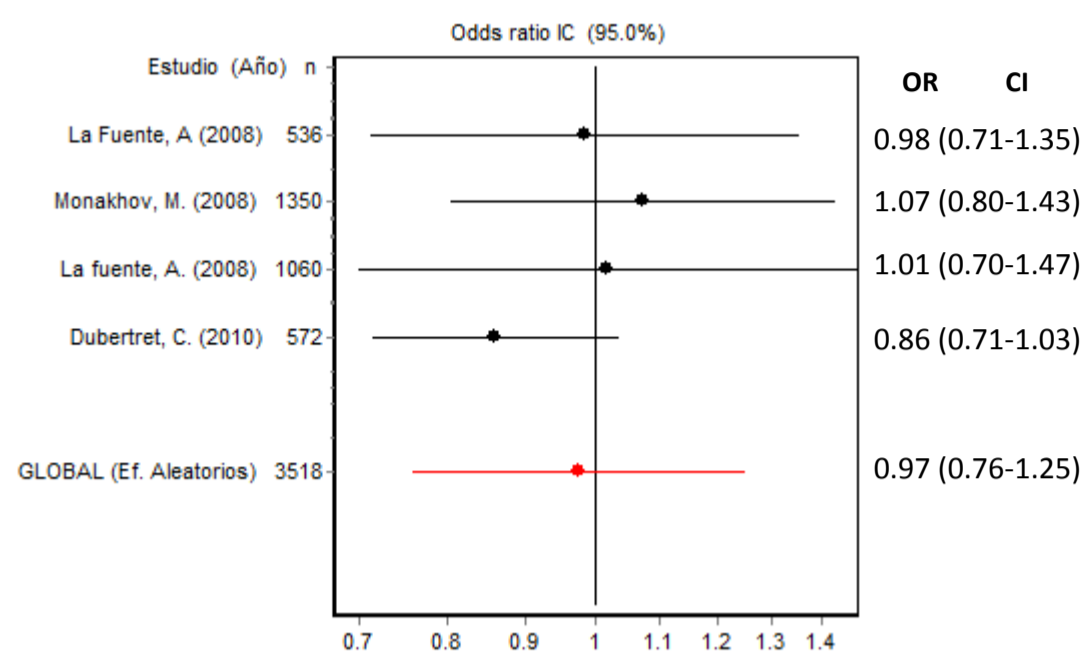


C)

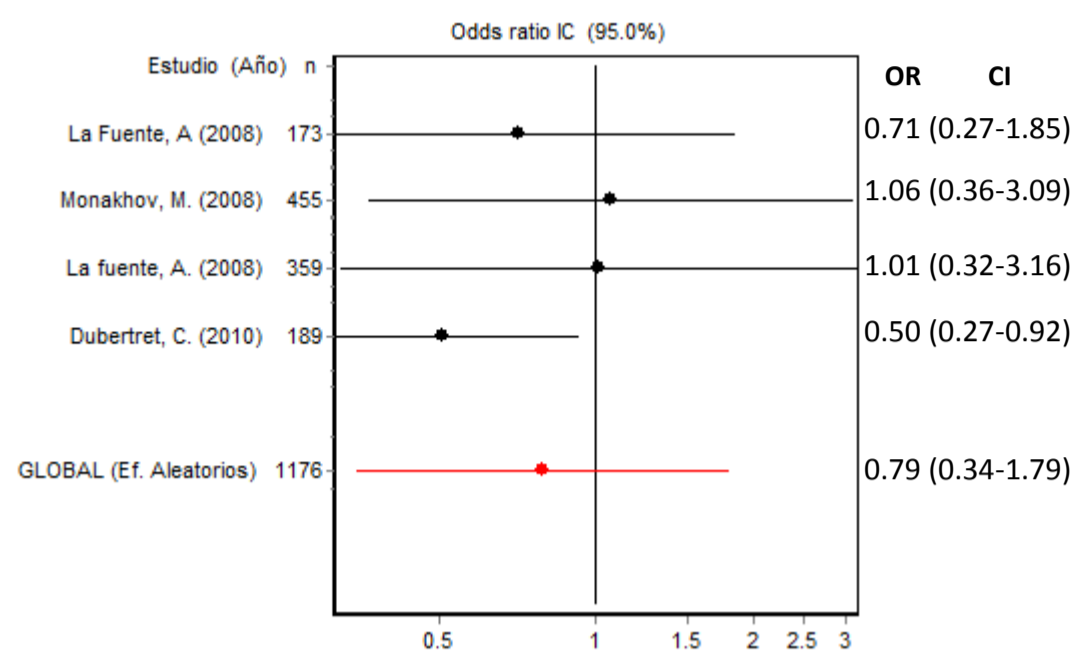


D)

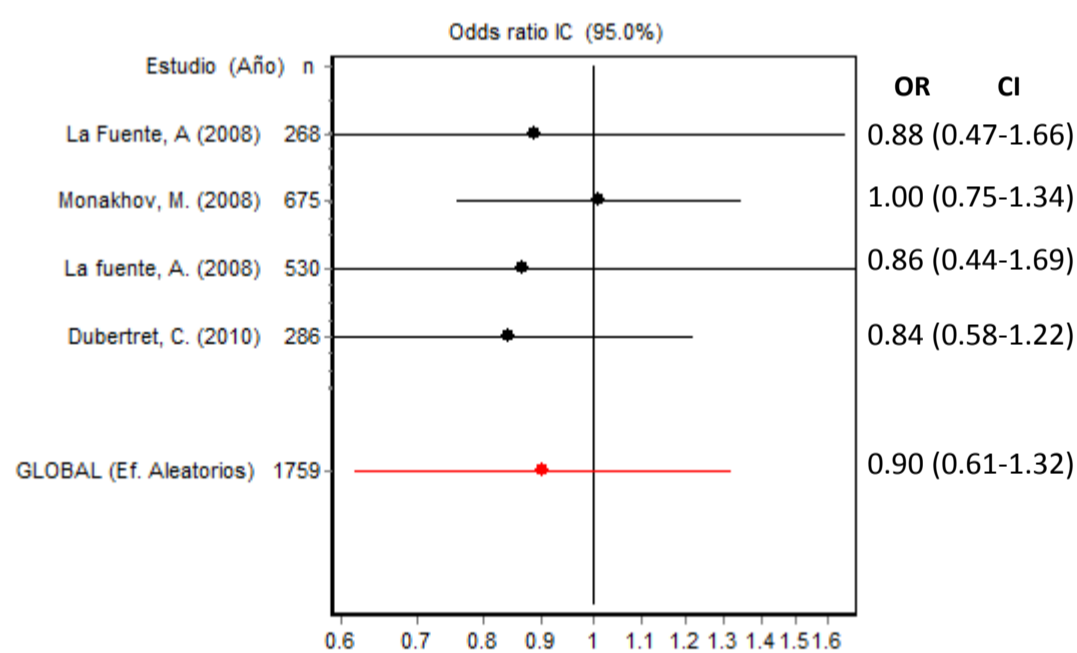
Figure S6. Sensitivity analysis of TaqI polymorphism with SZ in Caucasians population:
a) Allelic b) Additive c) Dominant and d) Recessive models.



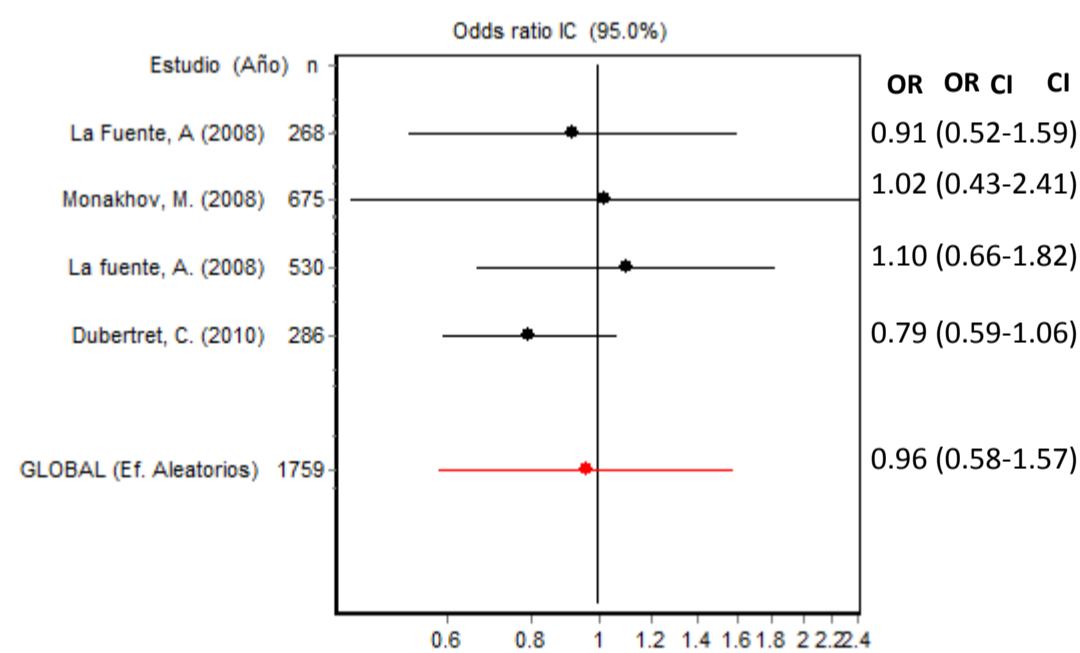
A)



B)



C)



D)