

REVIEW ARTICLE

TNFA gene polymorphisms and the risk of endometriosis: An updated meta-analysis of genetic association studies

Supplementary File

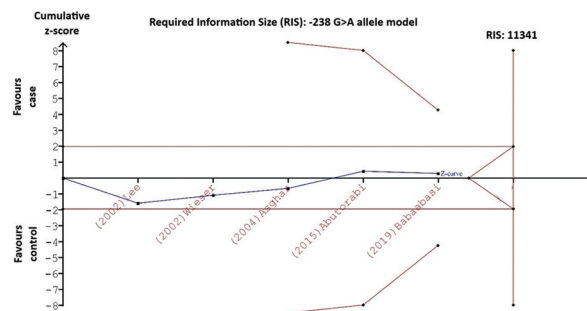


Figure S1. -238 G>A: TSA graph shows the z-curve under the allele model did not reach the RIS threshold, indicating a small sample size. Abbreviations: TSA: Trial sequential analysis; RIS: Required information size.

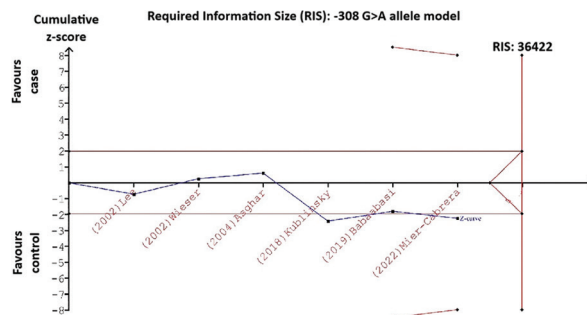


Figure S2. -308 G>A: TSA graph shows the z-curve under the allele model did not reach the RIS threshold, indicating a small sample size. Abbreviations: TSA: Trial sequential analysis; RIS: Required information size.



Figure S3. -857 C>T: TSA graph shows the z-curve under the allele model did not reach the RIS threshold, indicating a small sample size. Abbreviations: TSA: Trial sequential analysis; RIS: Required information size.

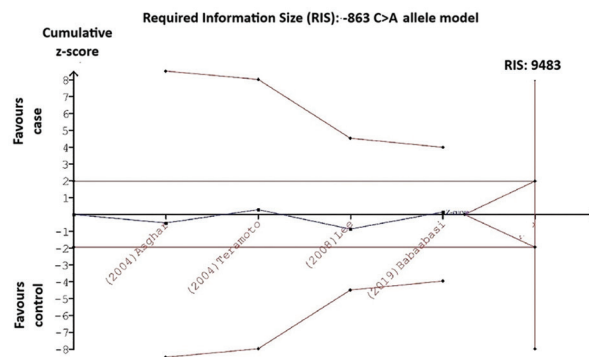


Figure S4. -863 C>A: TSA graph shows the z-curve under the allele model did not reach the RIS threshold, indicating a small sample size. Abbreviations: TSA: Trial sequential analysis; RIS: Required information size.

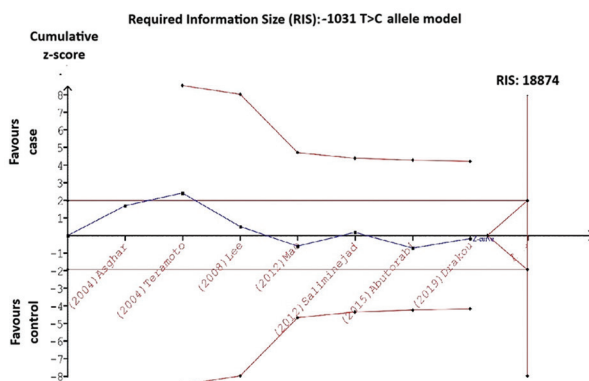


Figure S5. -1031 T>C: TSA graph shows the z-curve under the allele model did not reach the RIS threshold, indicating a small sample size. Abbreviations: TSA: Trial sequential analysis; RIS: Required information size.