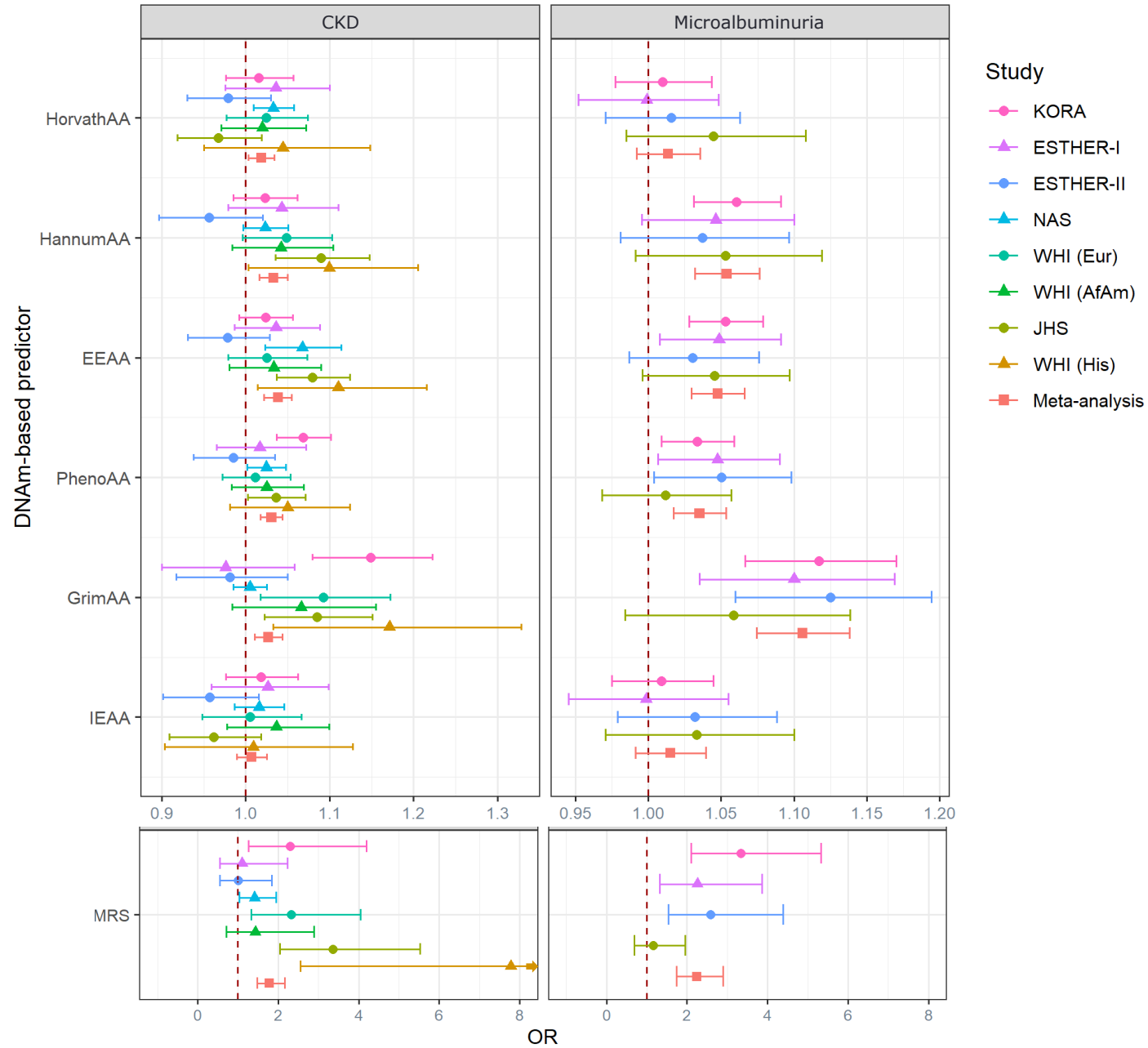


DNAmAge acceleration and renal function

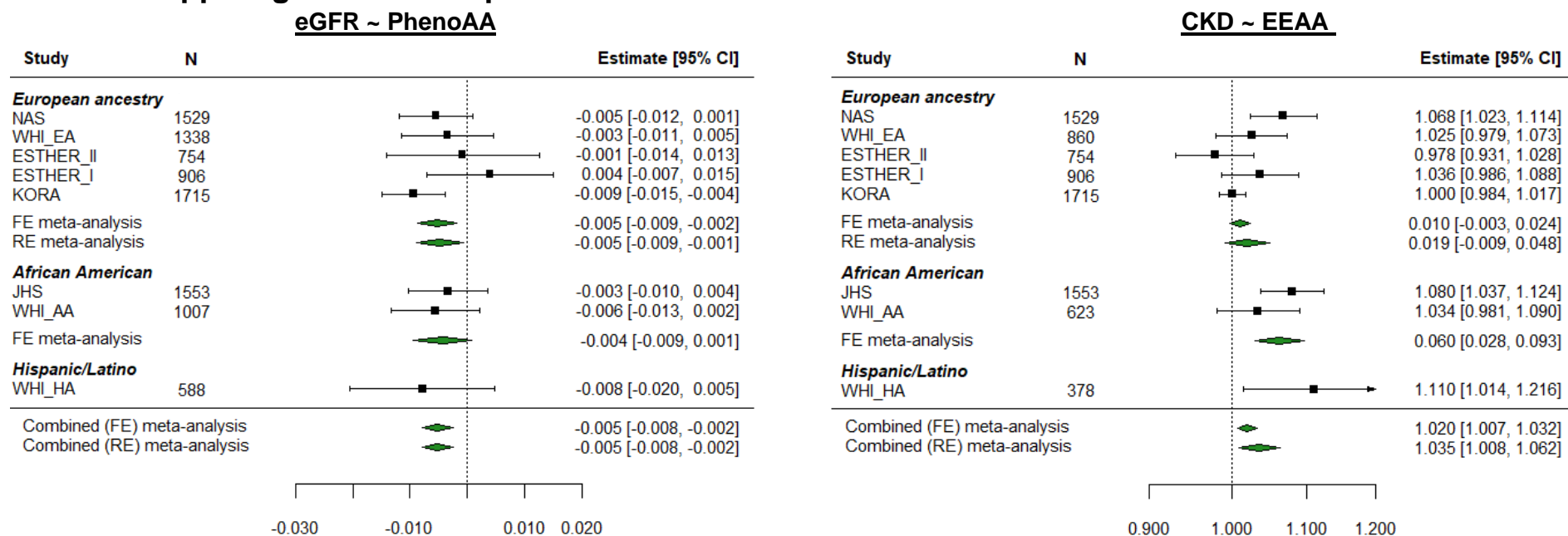
Suppl. Figures

Suppl. Figure 1. Odds Ratios from DNAm predictors and binary kidney traits

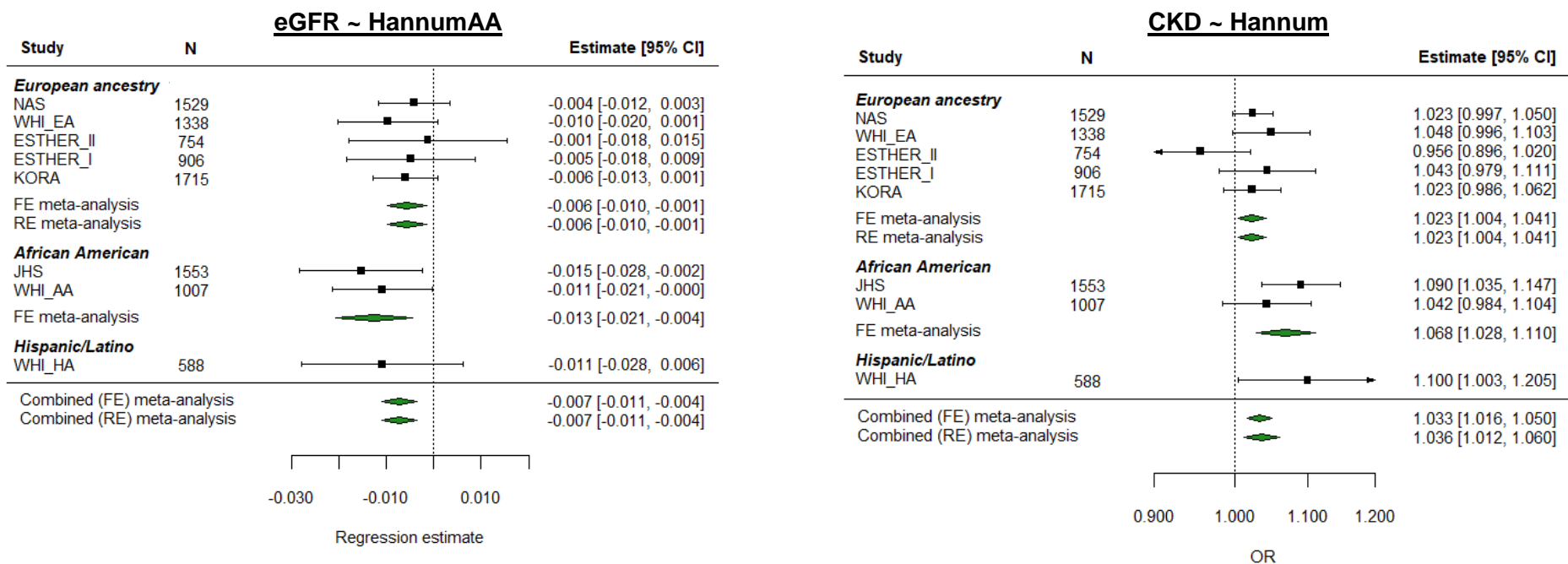


Suppl. Figure 2. Forest plots of associations with eGFR and CKD

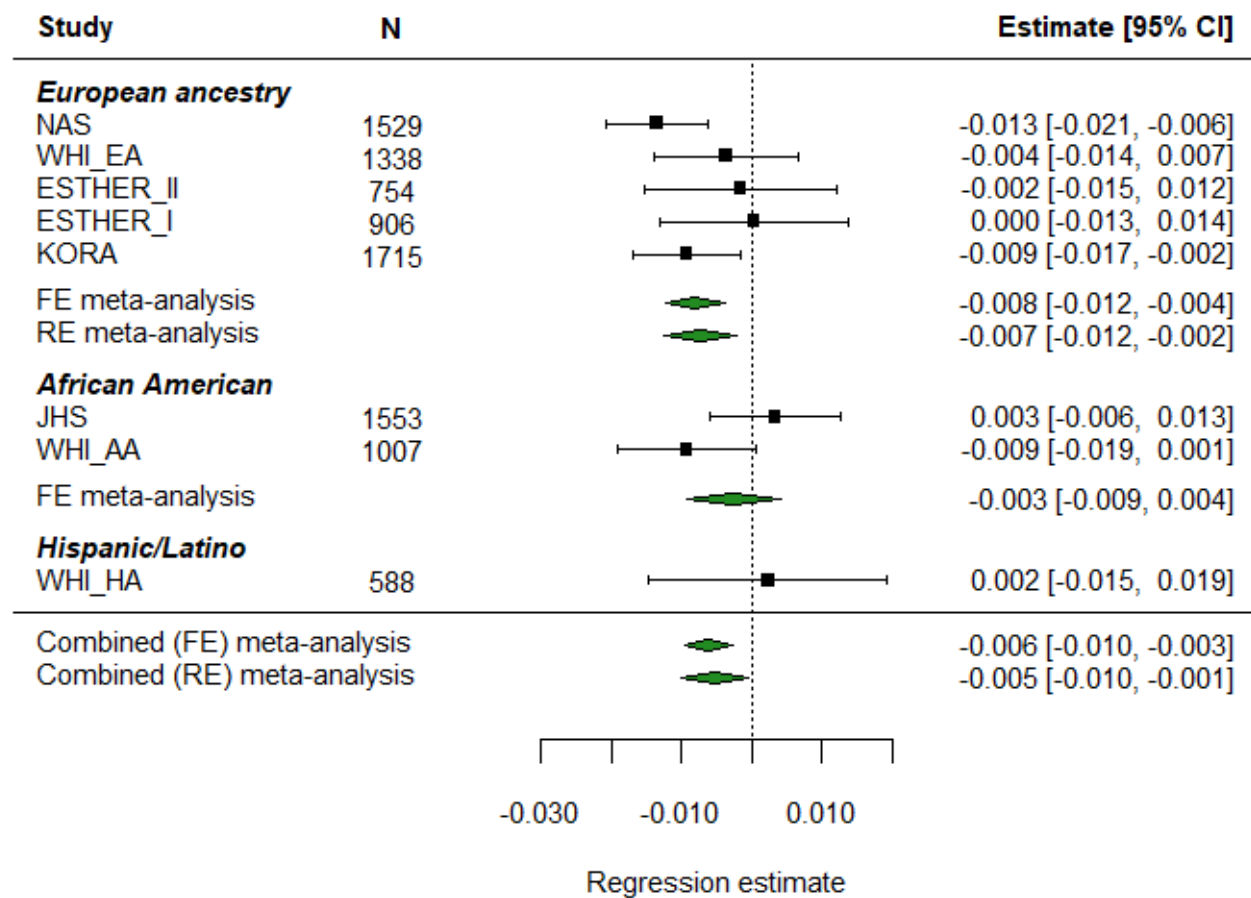
A.



B.

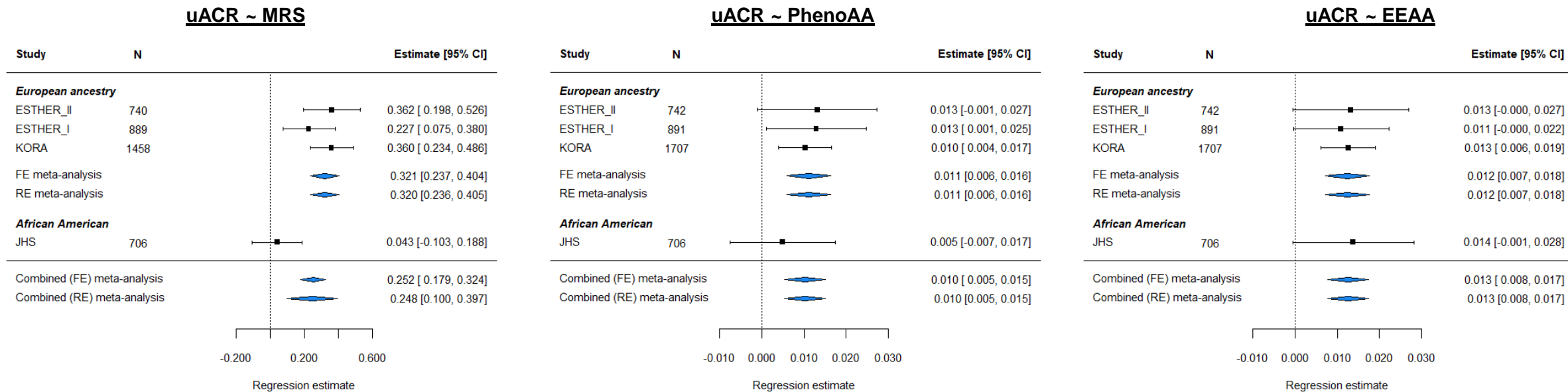


C.

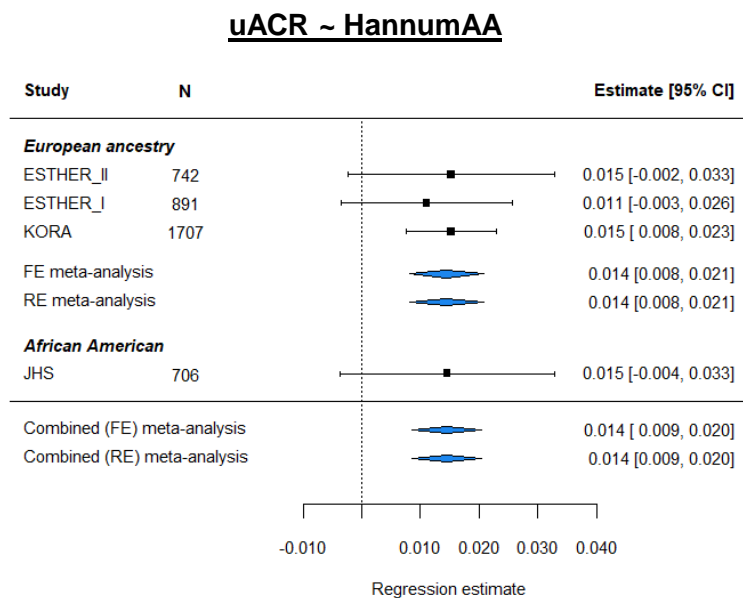
eGFR ~ Horvath

Suppl. Figure 3. Forest plots of associations with uACR

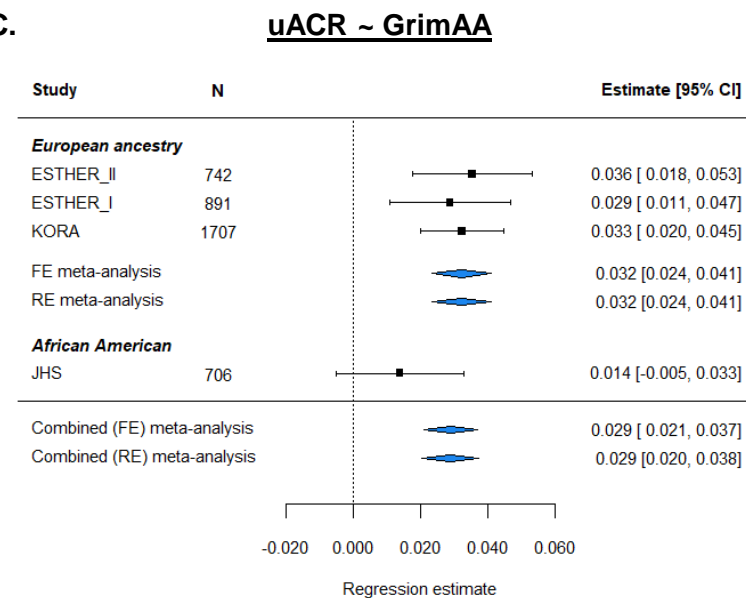
A.



B.



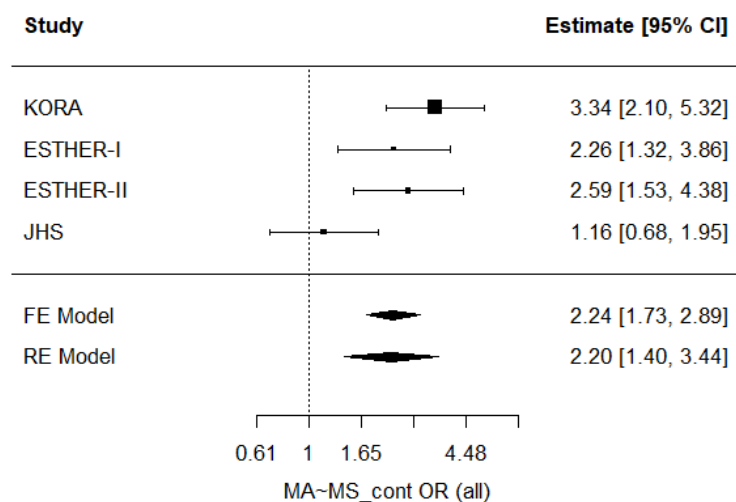
C.



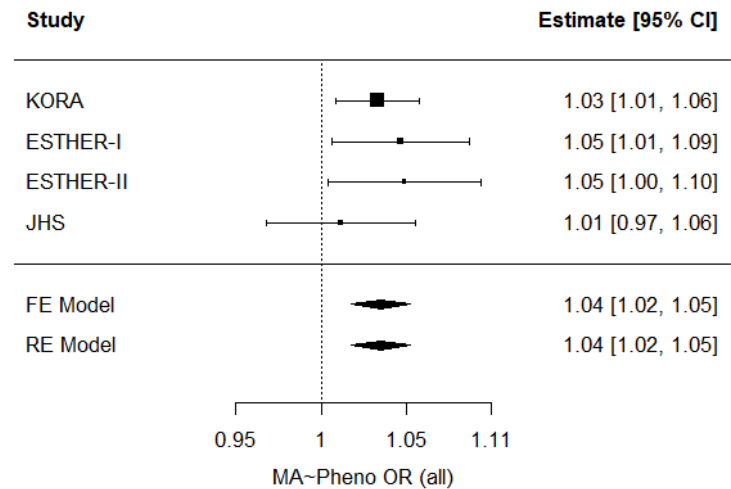
Suppl. Figure 4. Forest plots of associations with microalbuminuria

A.

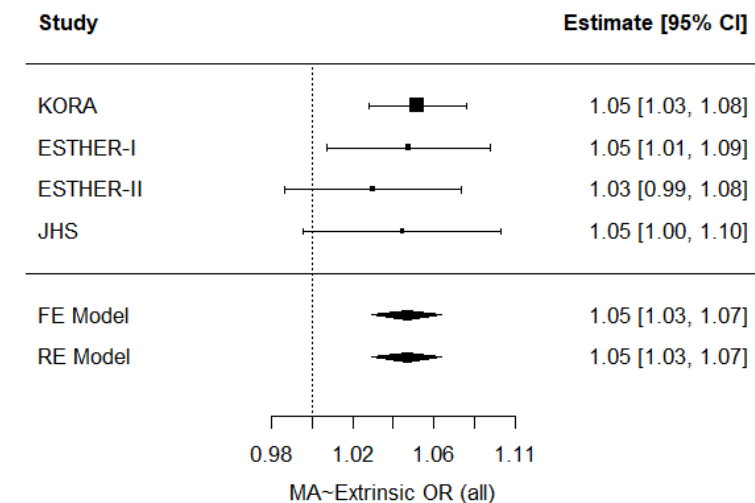
microalbuminuria ~ MRS



microalbuminuria ~ PhenoAA

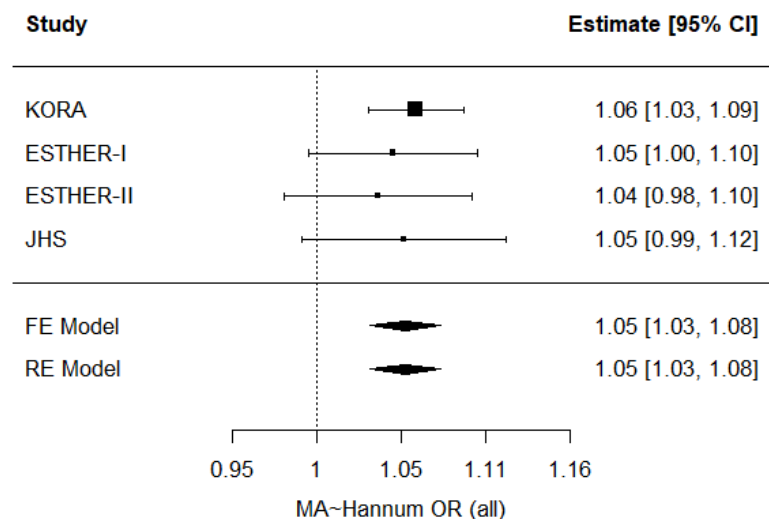


microalbuminuria ~ EEAA



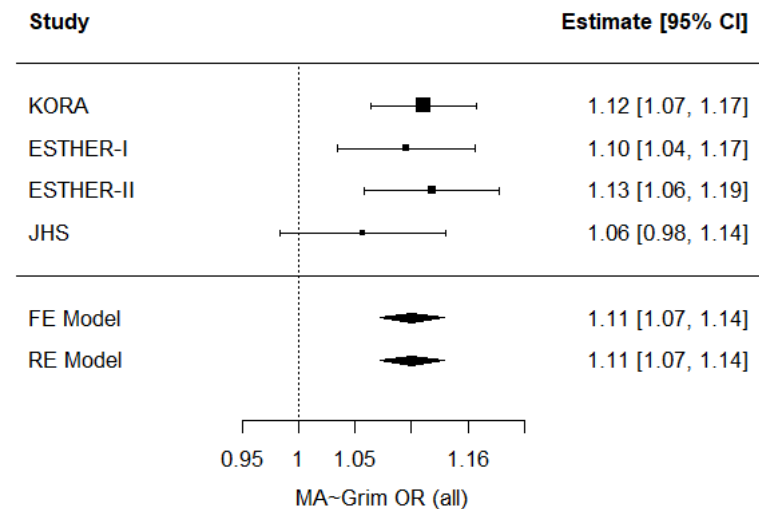
B.

microalbuminuria ~ HannumAA



C.

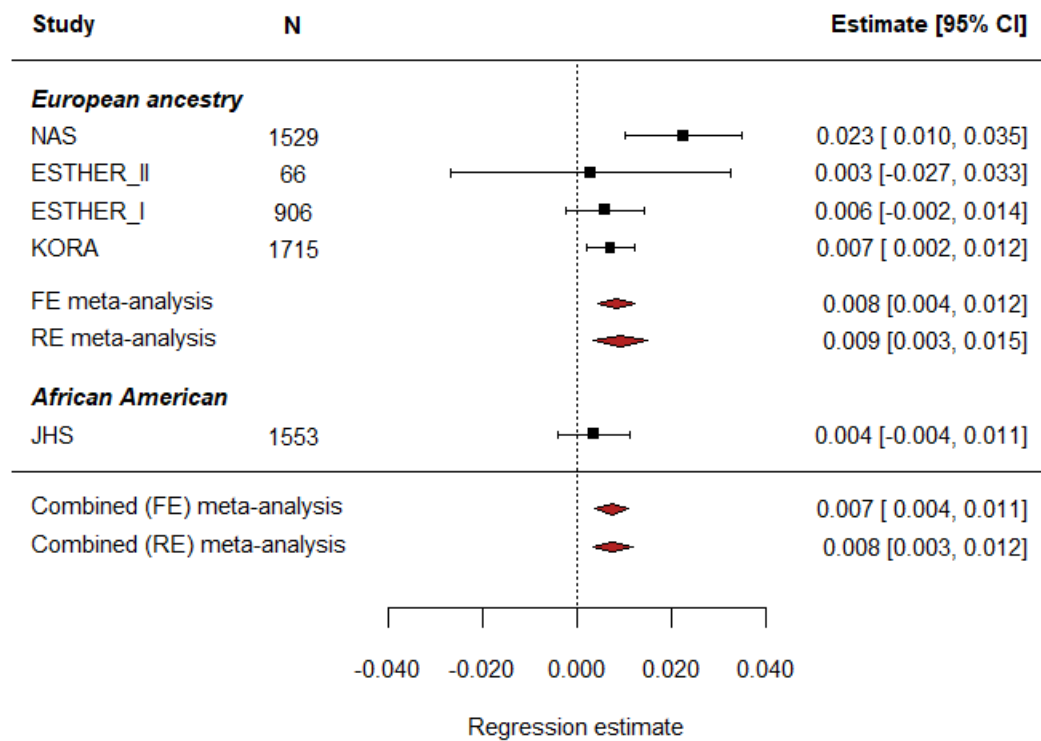
microalbuminuria ~ GrimAA



Suppl. Figure 5. Forest plots of associations with urate

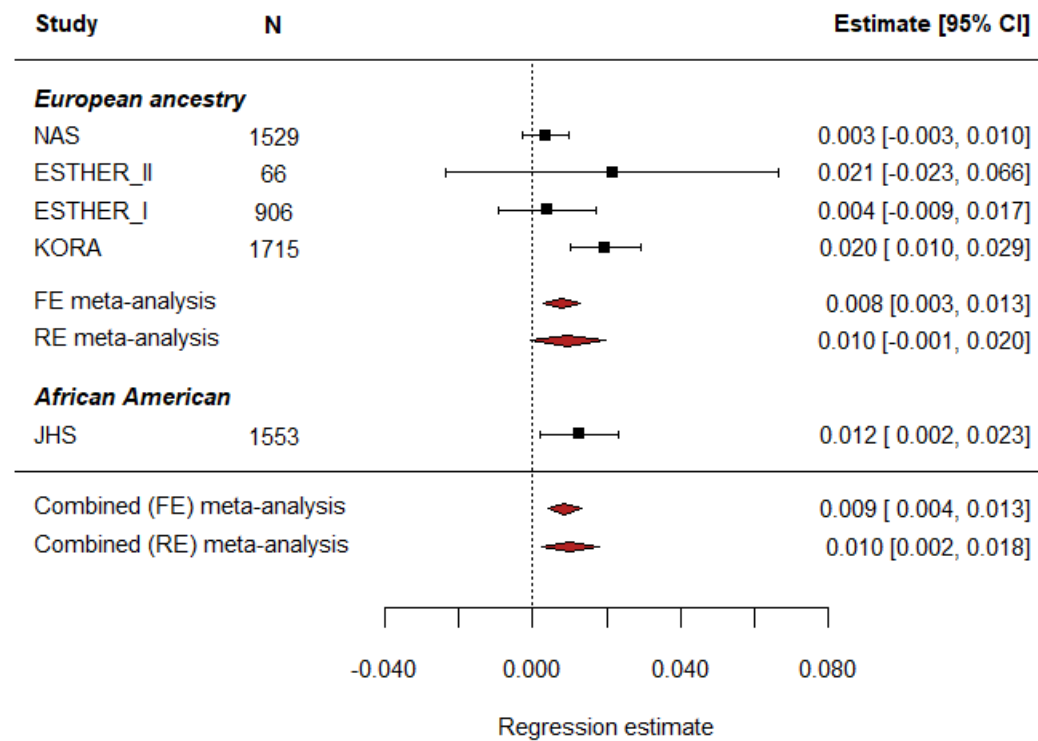
A.

urate ~ EEAA



B.

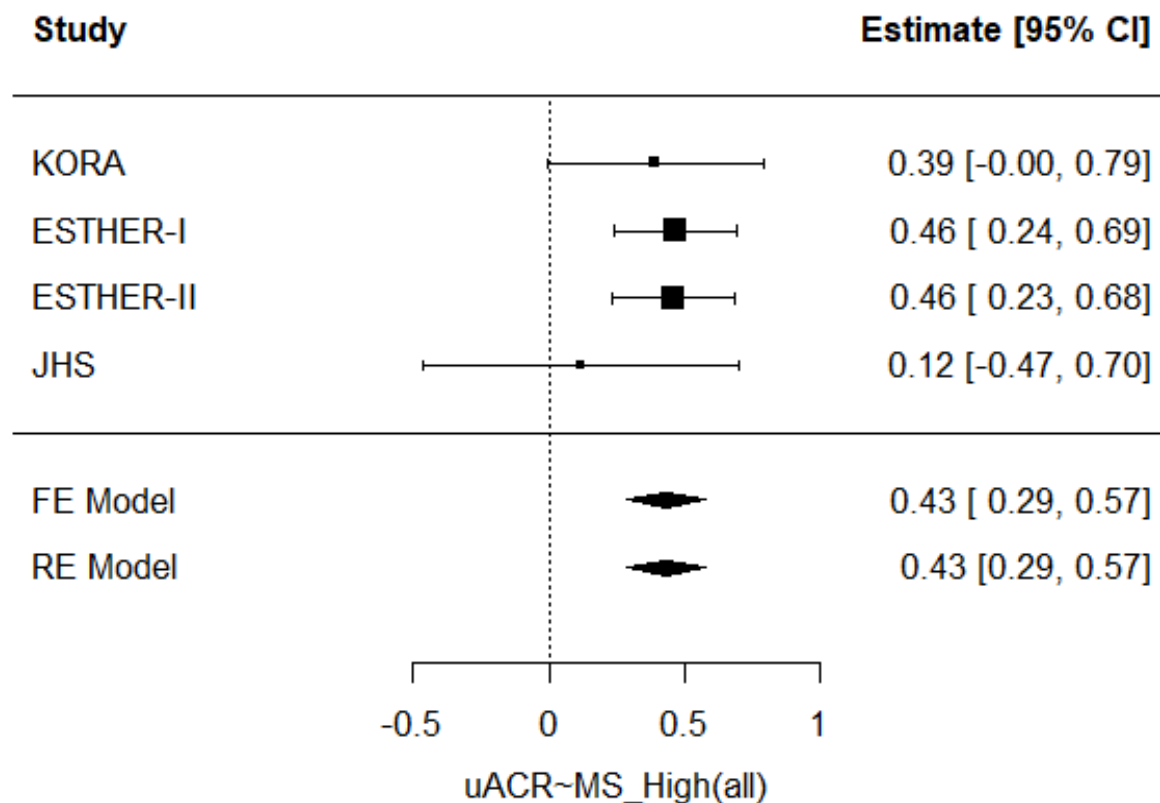
urate ~ GrimAA



Suppl. Figure 6. Forest plots for secondary associations with uACR and serum urate

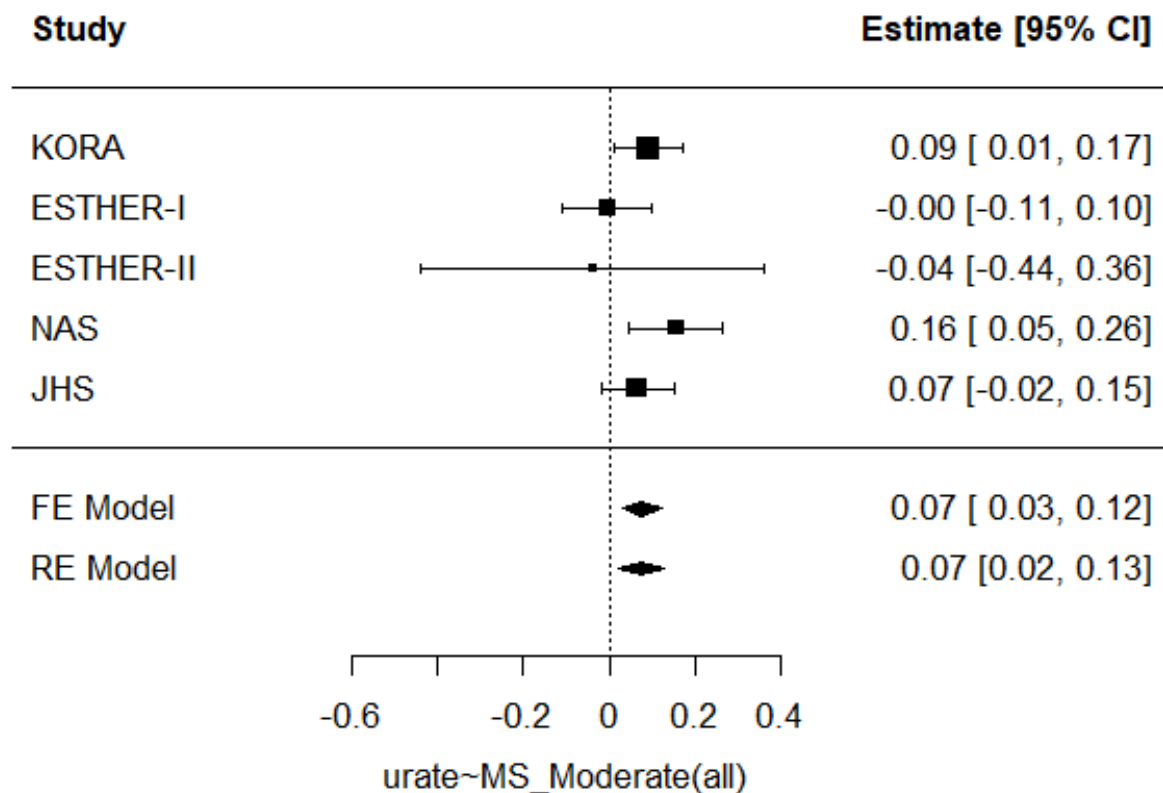
A.

uACR ~ MRS high risk



B.

Serum urate ~ MRS moderate risk



Suppl. Figure 7. Effect estimates from MRS and renal traits

