

Supplementary materials for the article:

Gao Y. et al. Roles of Gut Microbiota and Associated Metabolites in *Clostridioides difficile* Infection.

Pol J Microbiol. 2025, Vol. 74, No 2, 206-217

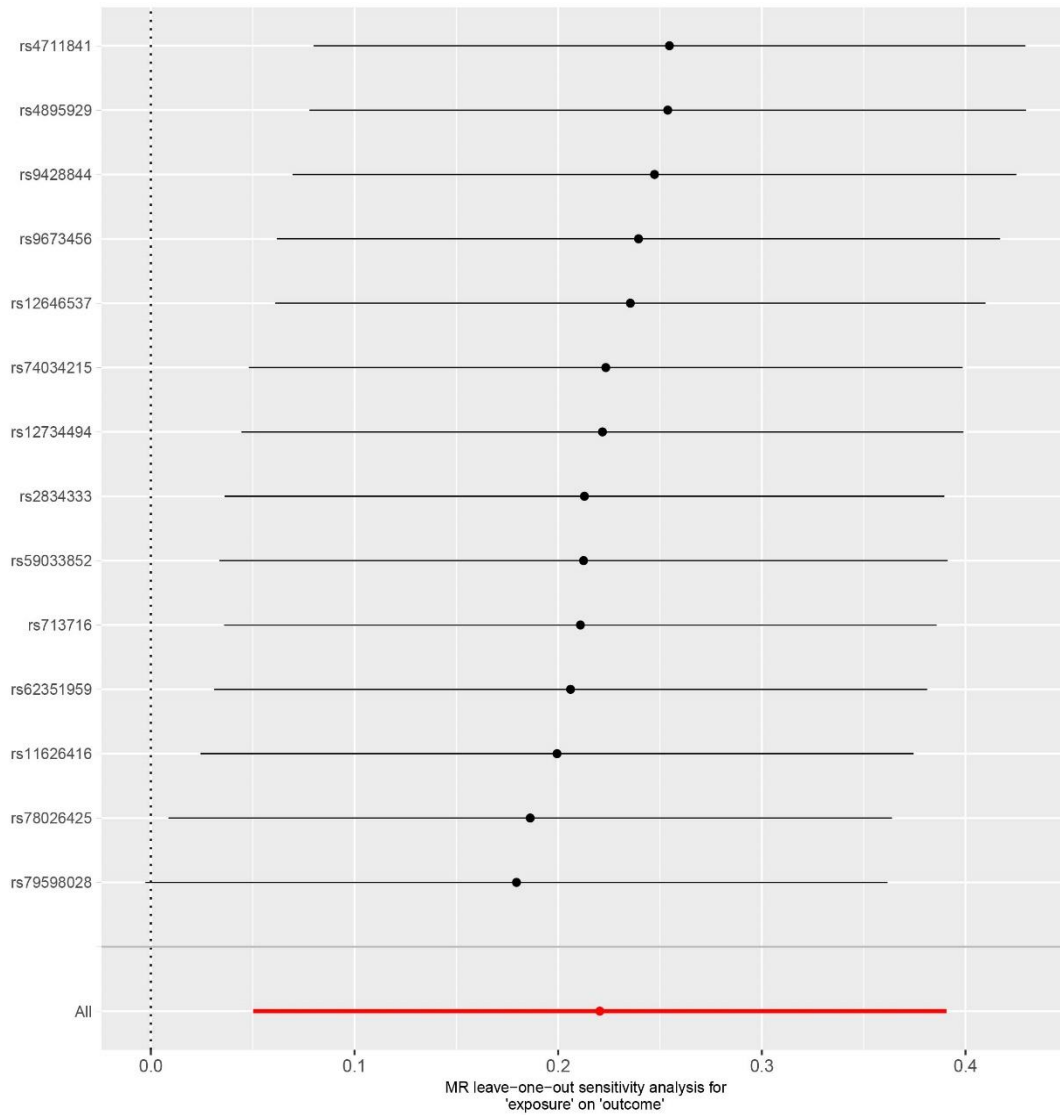


Fig. S1. Leave-one-out sensitivity of Burkholderiales.

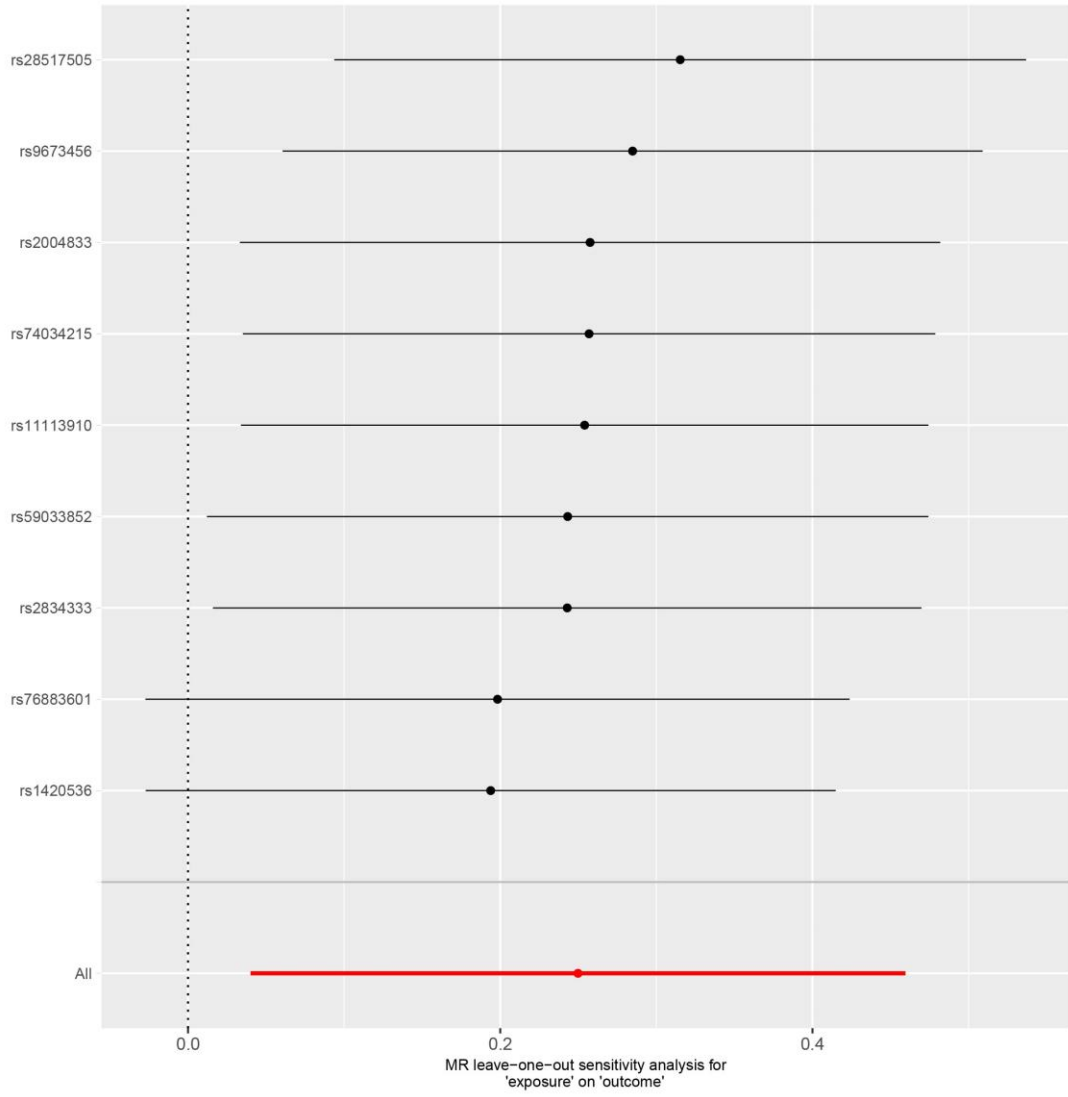


Fig. S2. Leave-one-out sensitivity of *Sutterellaceae*.

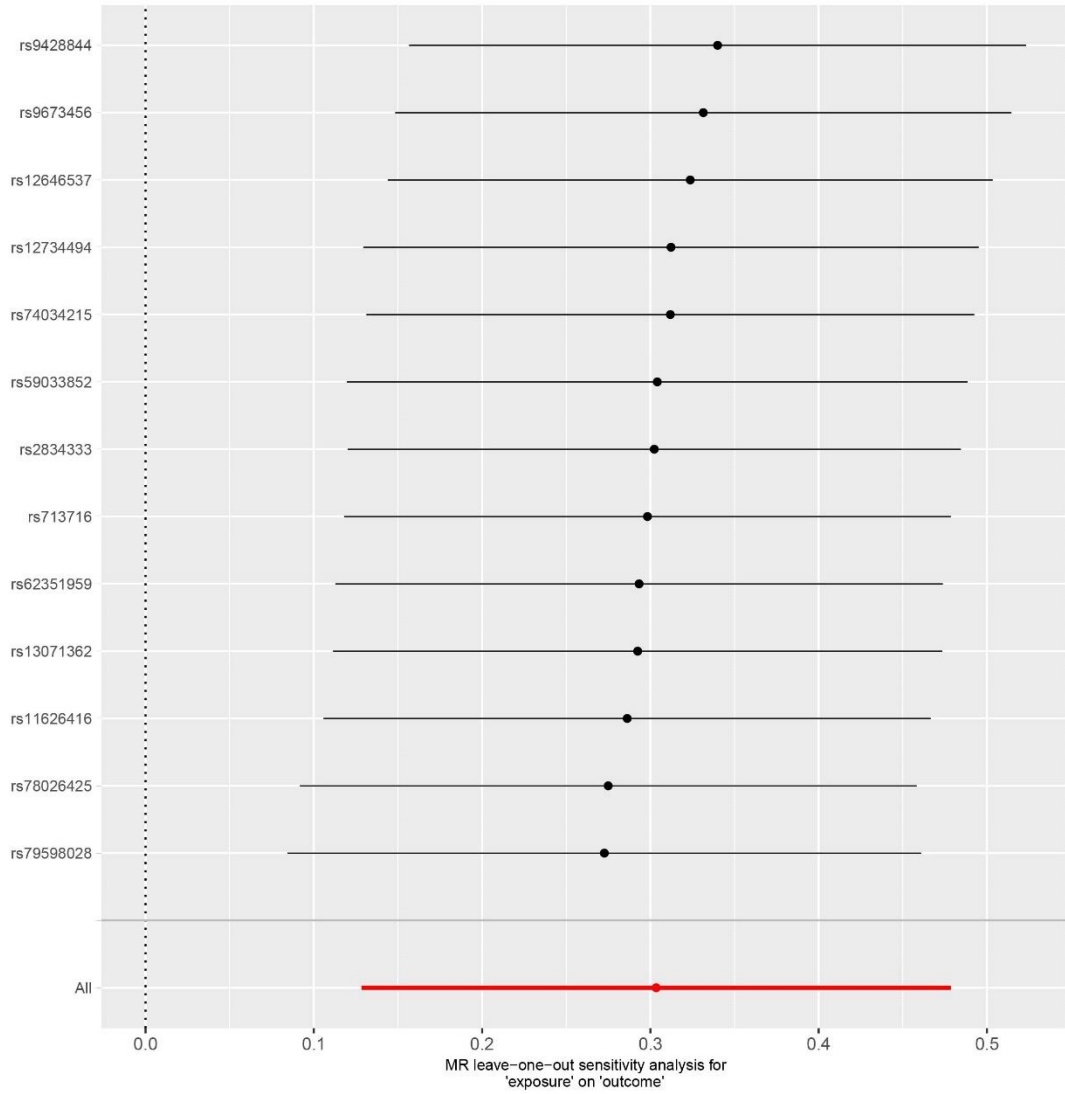


Fig. S3. Leave-one-out sensitivity of Betaproteobacteria.

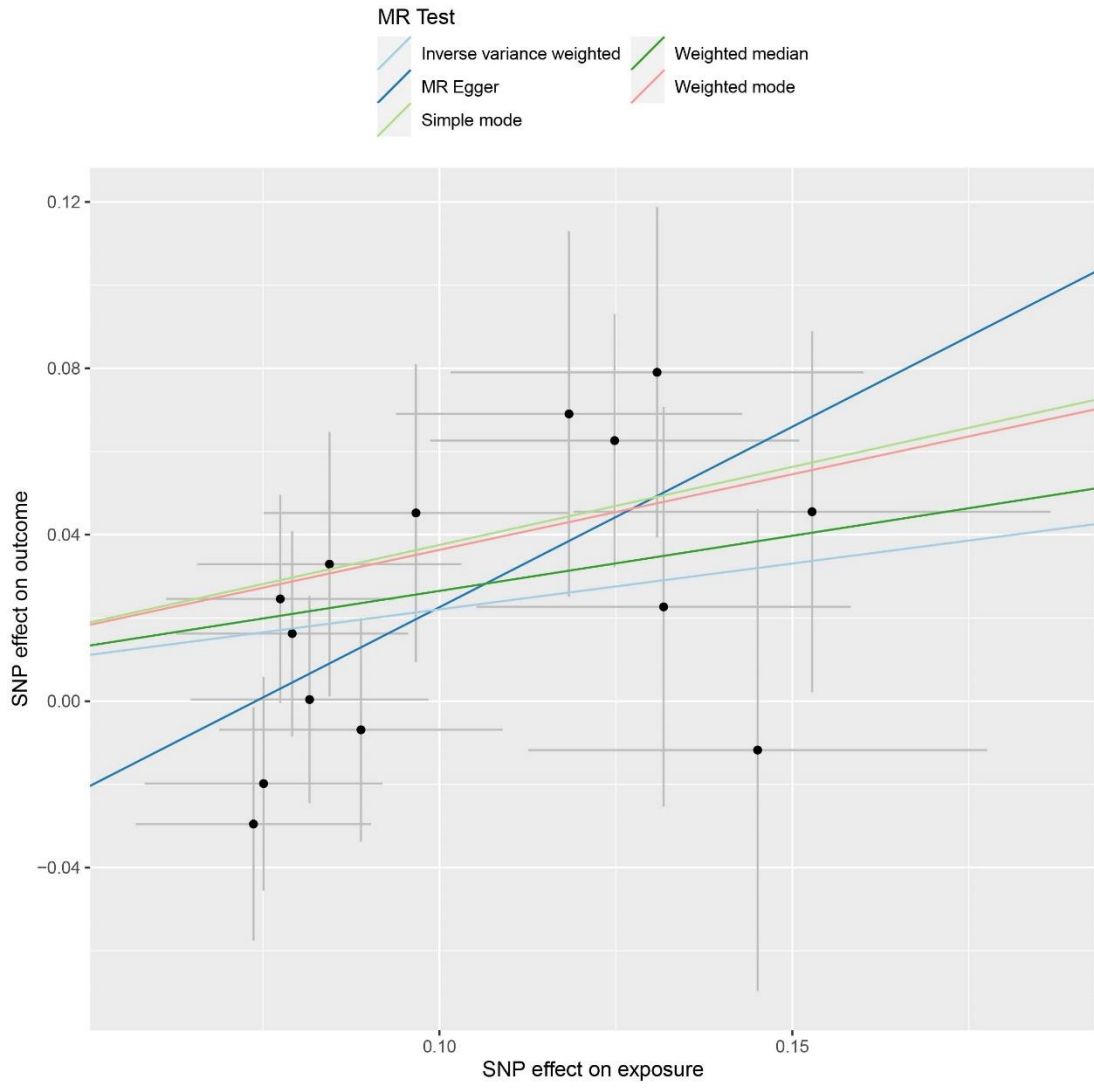


Fig. S4. The scatter plot of Burkholderiales.

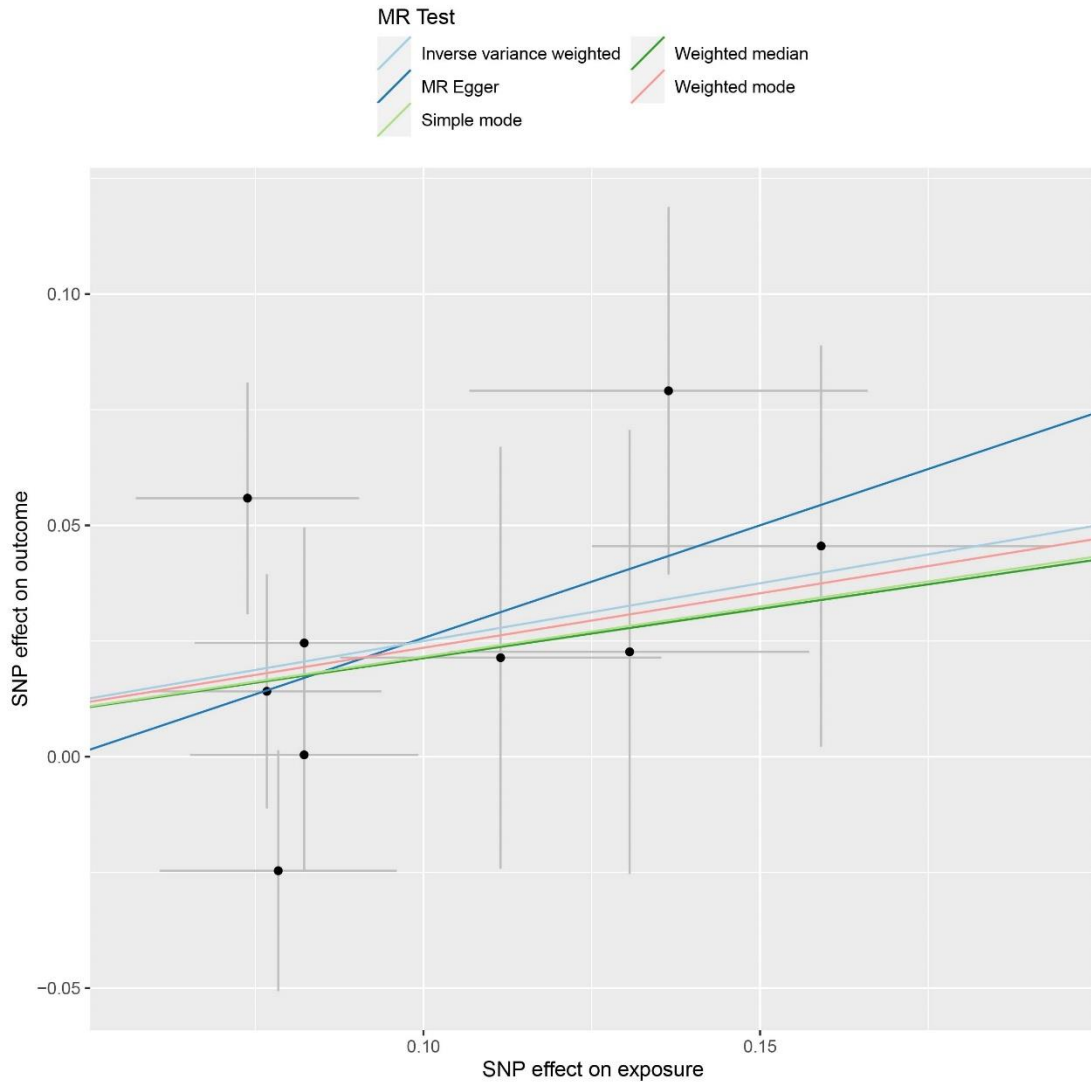


Fig. S5. The scatter plot of *Sutterellaceae*.

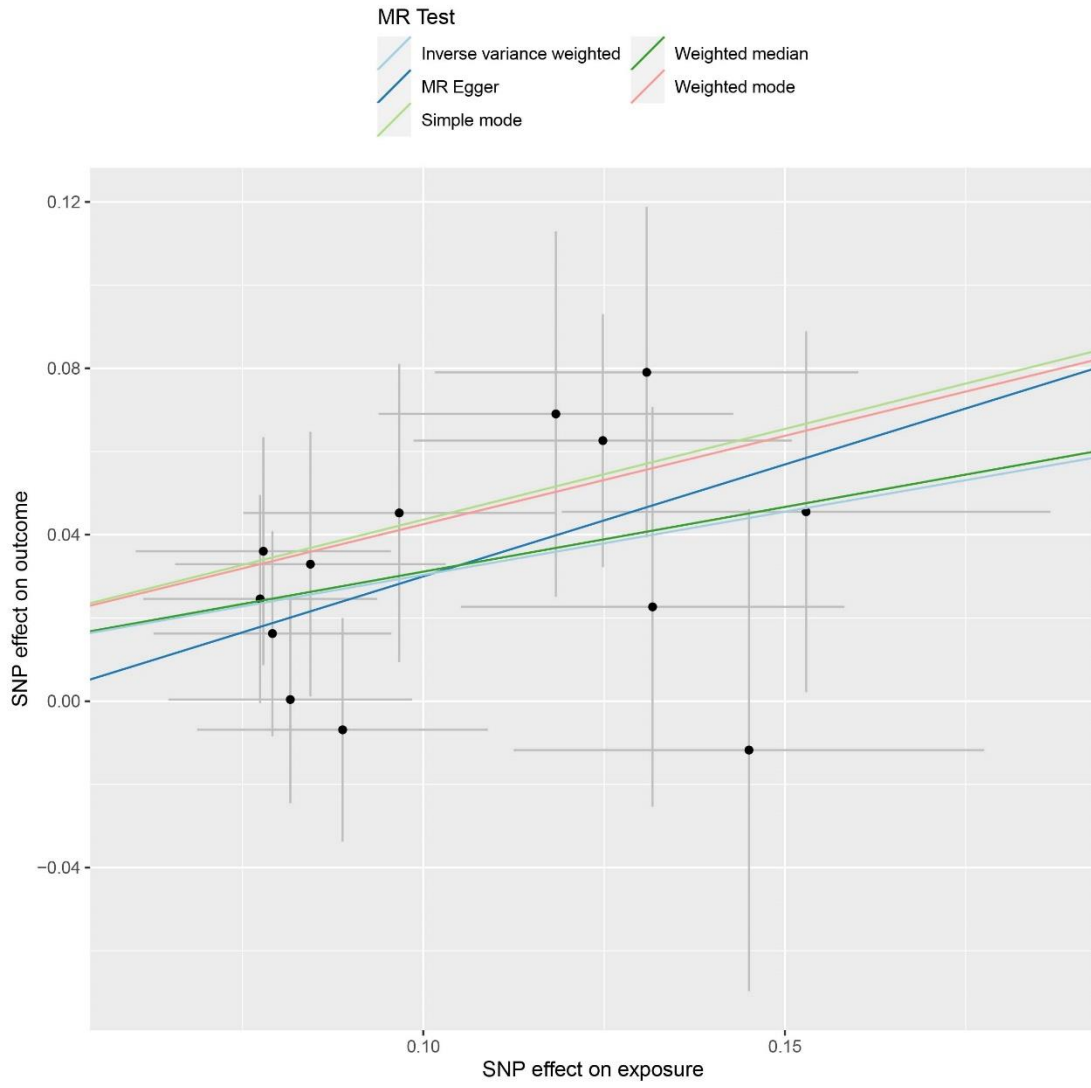


Fig. S6. The scatter plot of Betaproteobacteria.

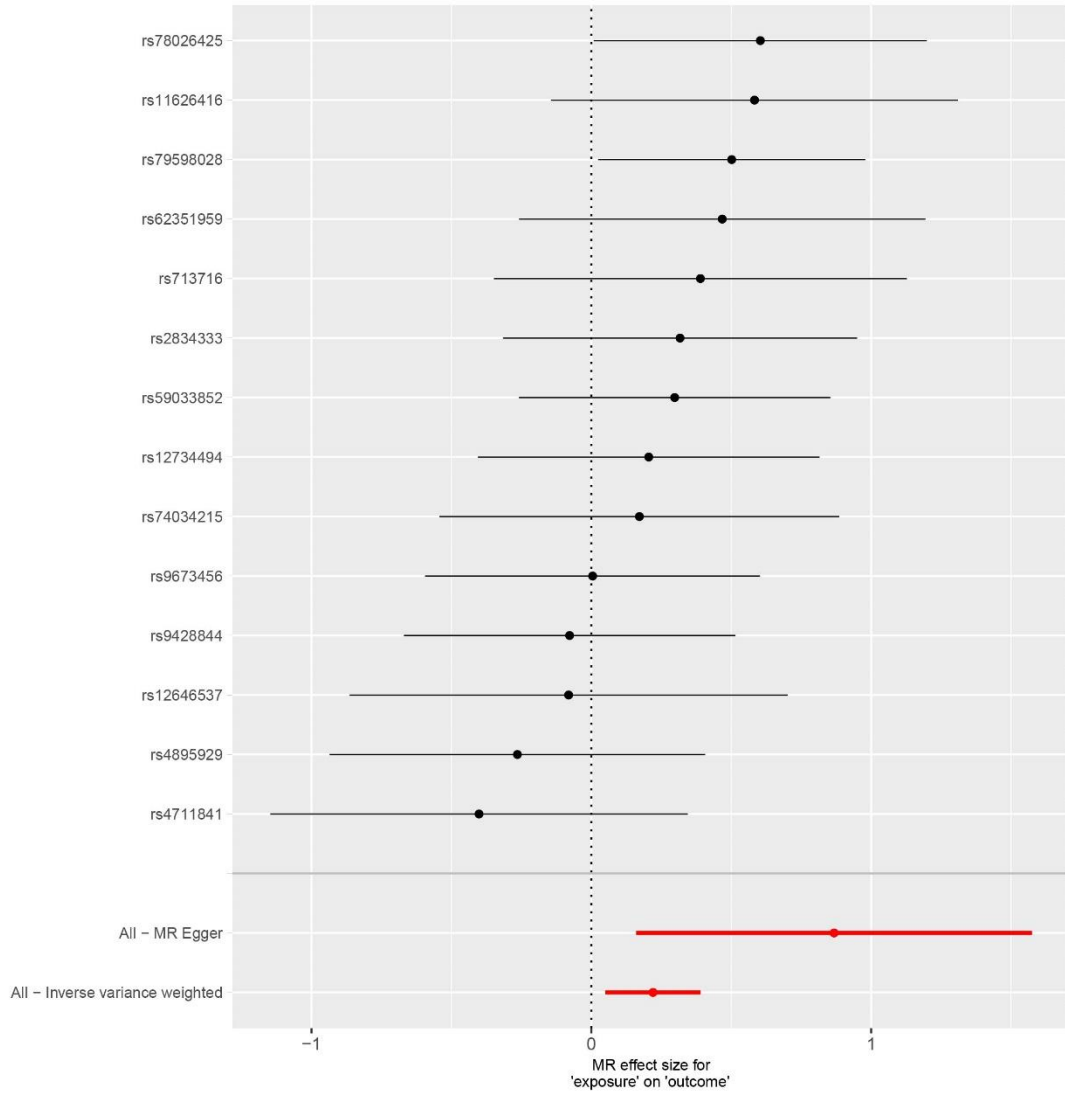


Fig. S7. Forest plot of Burkholderiales.

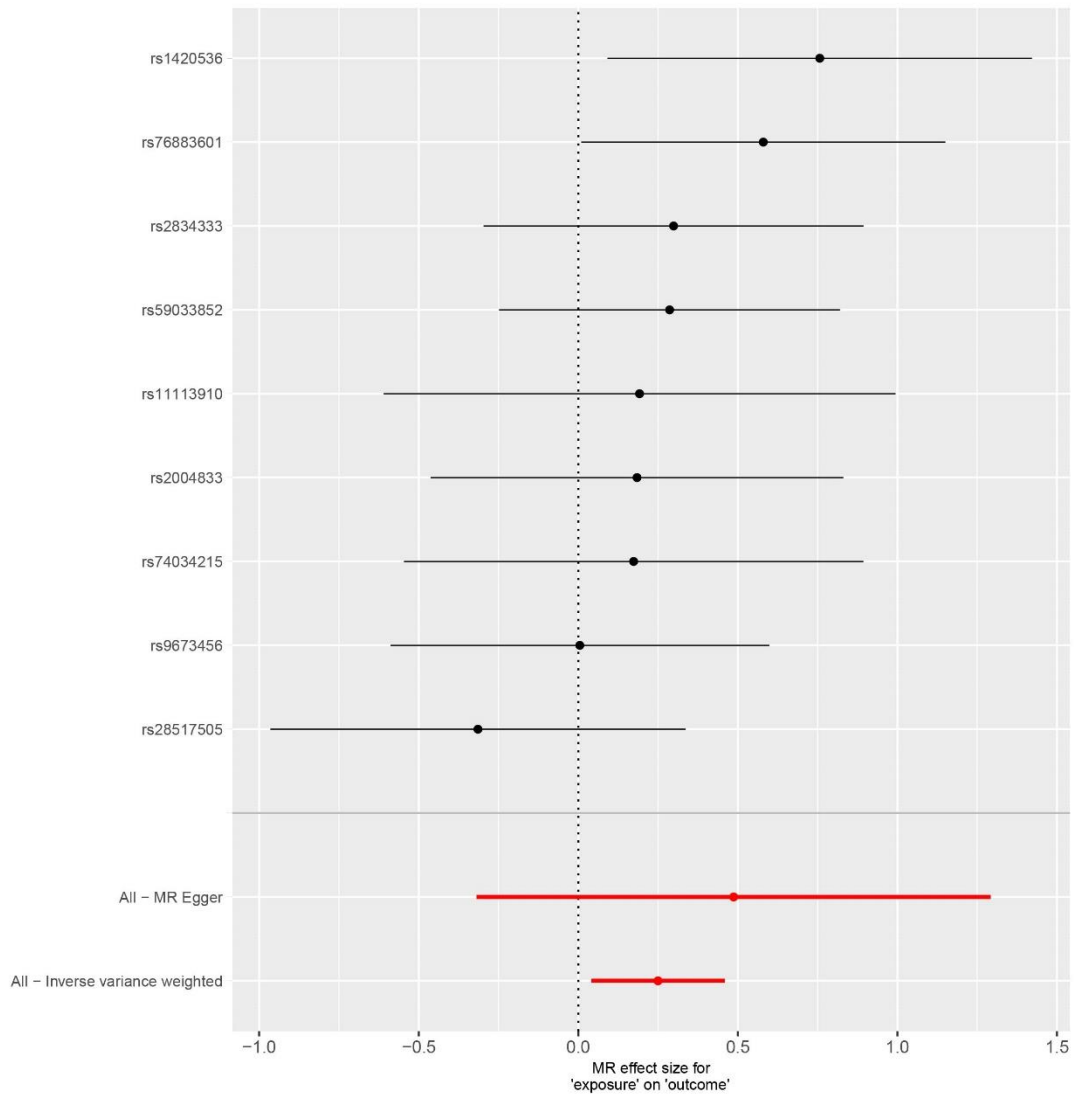


Fig. S8. Forest plot of *Sutterellaceae*.



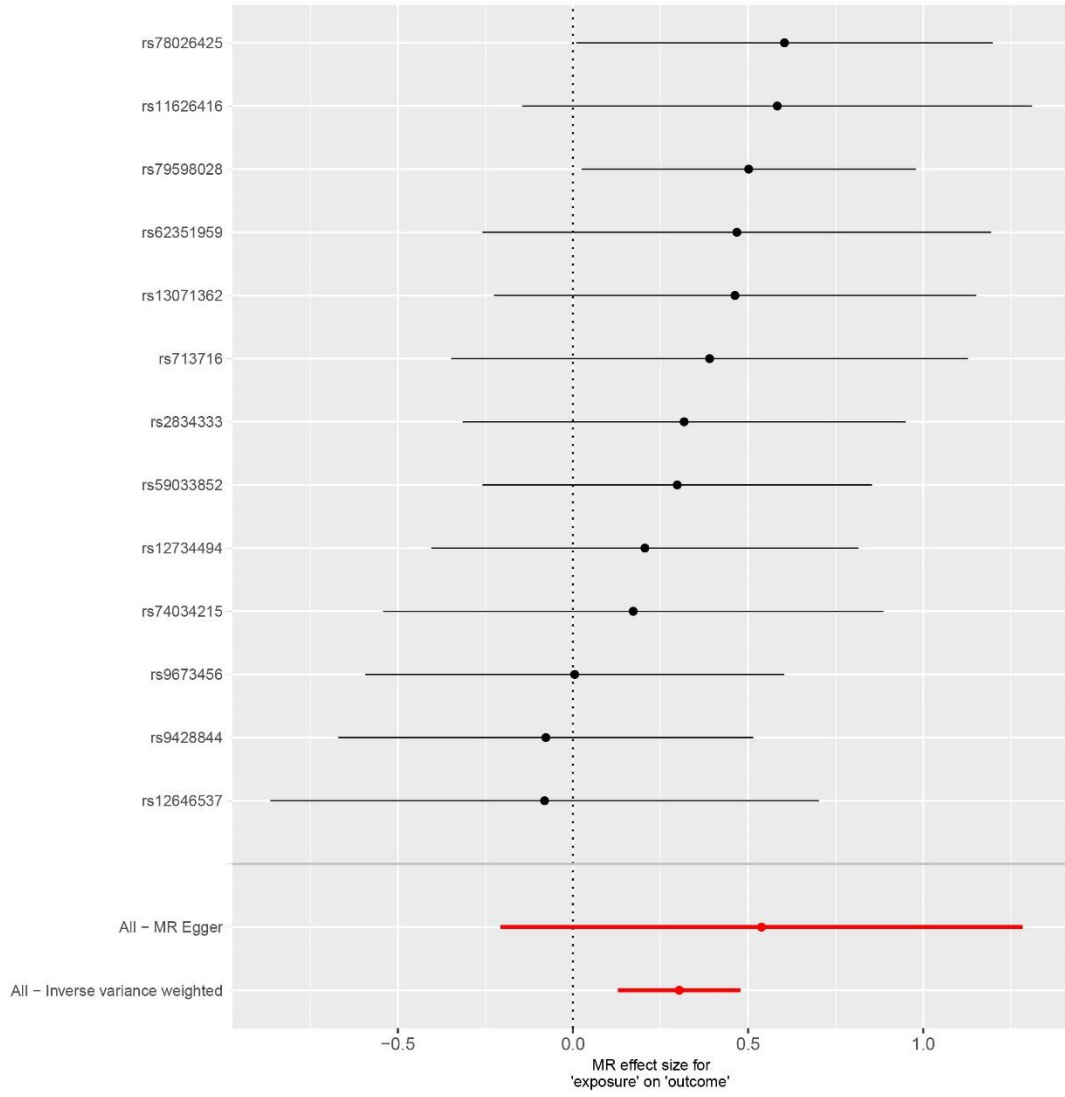


Fig. S9. Forest plot of Betaproteobacteria.