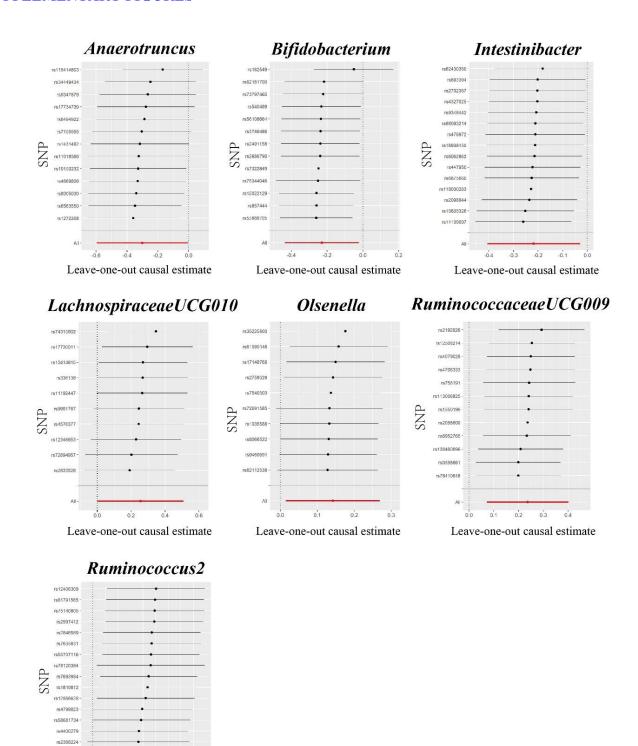
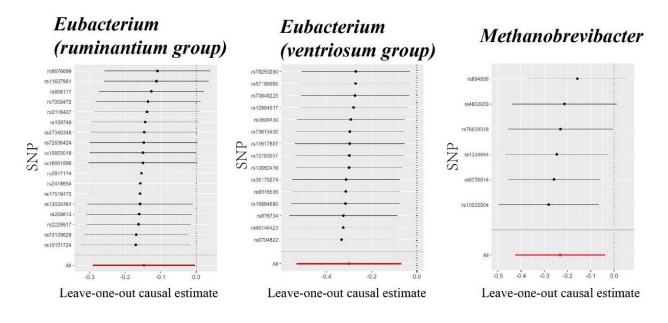
SUPPLEMENTARY FIGURES



Supplementary Figure 1. Leave-one-out plots for the causal association between gut microbiota and GH in forward MR analyses.

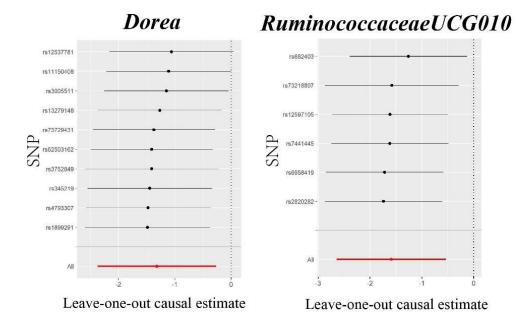
0.0 0.1 0.2 0.3 0.4

Leave-one-out causal estimate

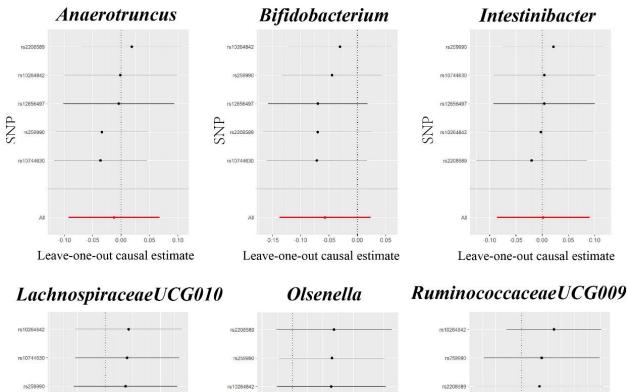


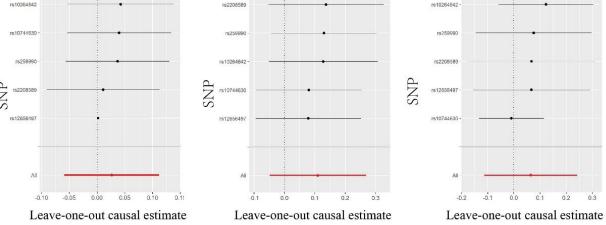
Tyzzerella3 RuminococcaceaeUCG002 rs113147300 rs1232220 rs7120052 rs77564310 rs55799124 rs12453378 rs6793778 rs7561370 rs10964441 rs882348 rs56030423 rs11607472 rs17706273 rs55793120 rs79016051 rs191093 rs11750293 rs67476743 rs7342369 rs15256 rs57079348 rs112102233 rs7333521 rs7155595 rs10927423 rs76847269 rs7249614 -rs116974815 -Leave-one-out causal estimate Leave-one-out causal estimate

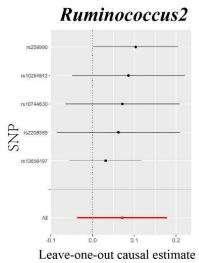
Supplementary Figure 2. Leave-one-out plots for the causal association between gut microbiota and PE in forward MR analyses.



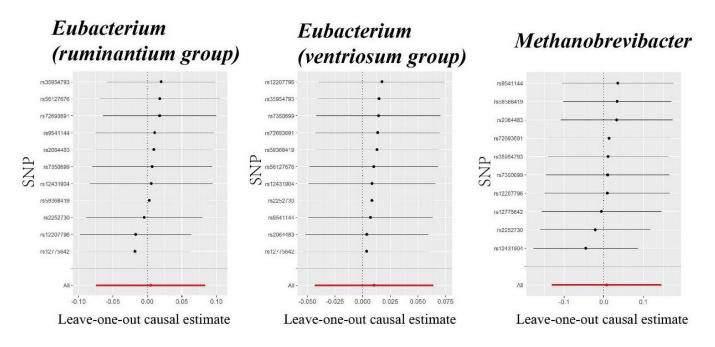
Supplementary Figure 3. Leave-one-out plots for the causal association between gut microbiota and eclampsia in forward MR analyses.

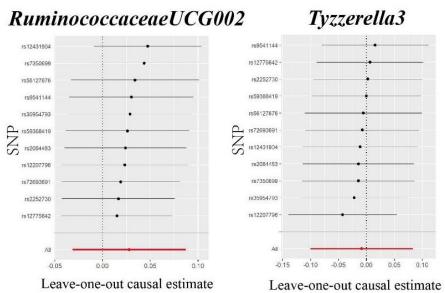




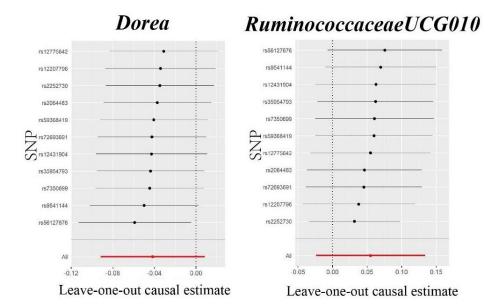


Supplementary Figure 4. Leave-one-out plots for the causal association between gut microbiota and GH in reverse MR analyses.





Supplementary Figure 5. Leave-one-out plots for the causal association between gut microbiota and PE in reverse MR analyses.



Supplementary Figure 6. Leave-one-out plots for the causal association between gut microbiota and eclampsia in reverse MR analyses.