

Supplementary Materials

Absolute quantitative metagenomic analysis provides more accurate insights for the anti-colitis effect of berberine via modulation of gut microbiota

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Supplementary Figure S1

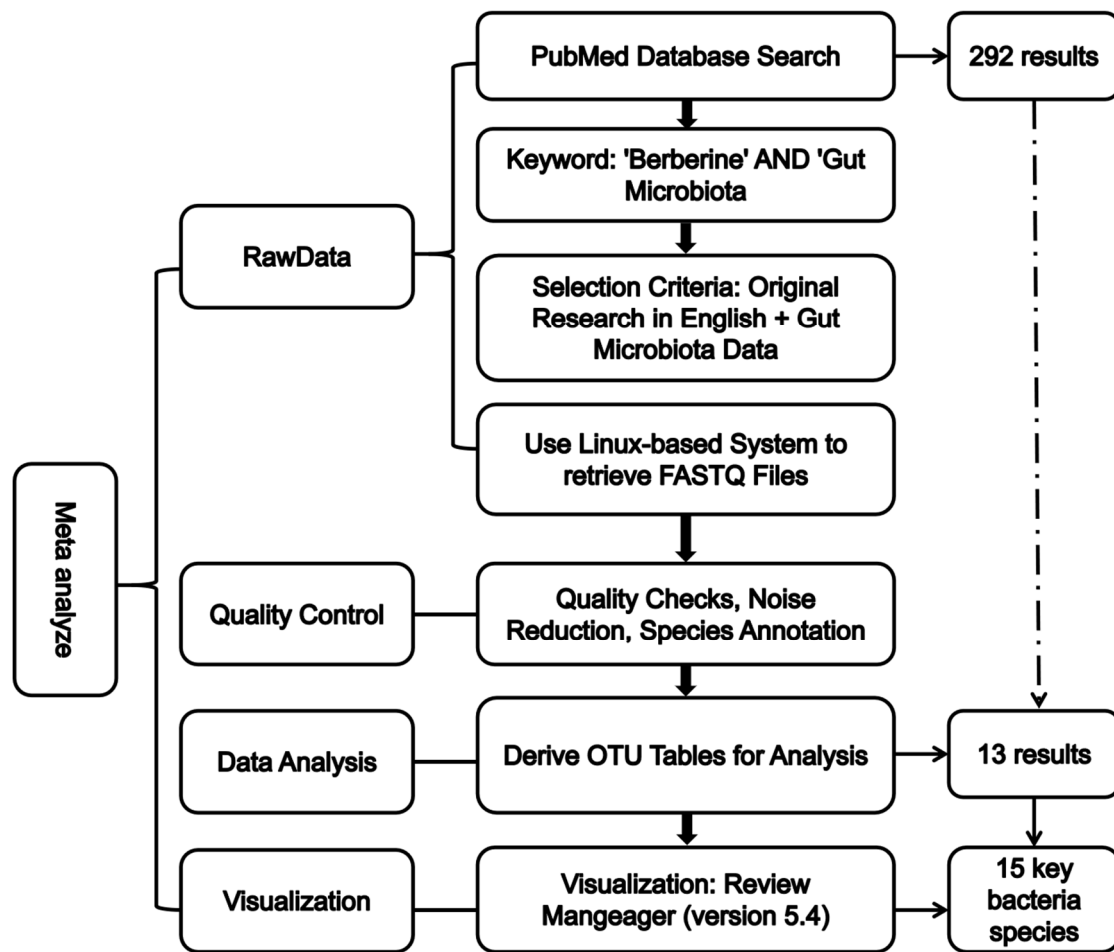


Figure S1. Flowchart of the meta-analysis.

Supplementary Figure S2

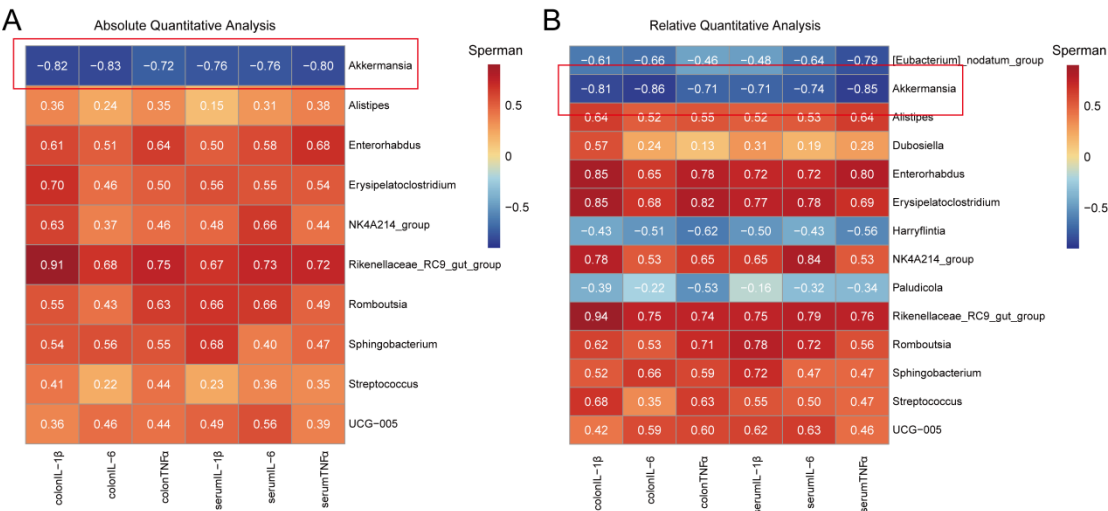


Figure S2. Correlation analyses comparing relative and absolute quantitative analyses with inflammatory factors.

Supplementary Figure S3

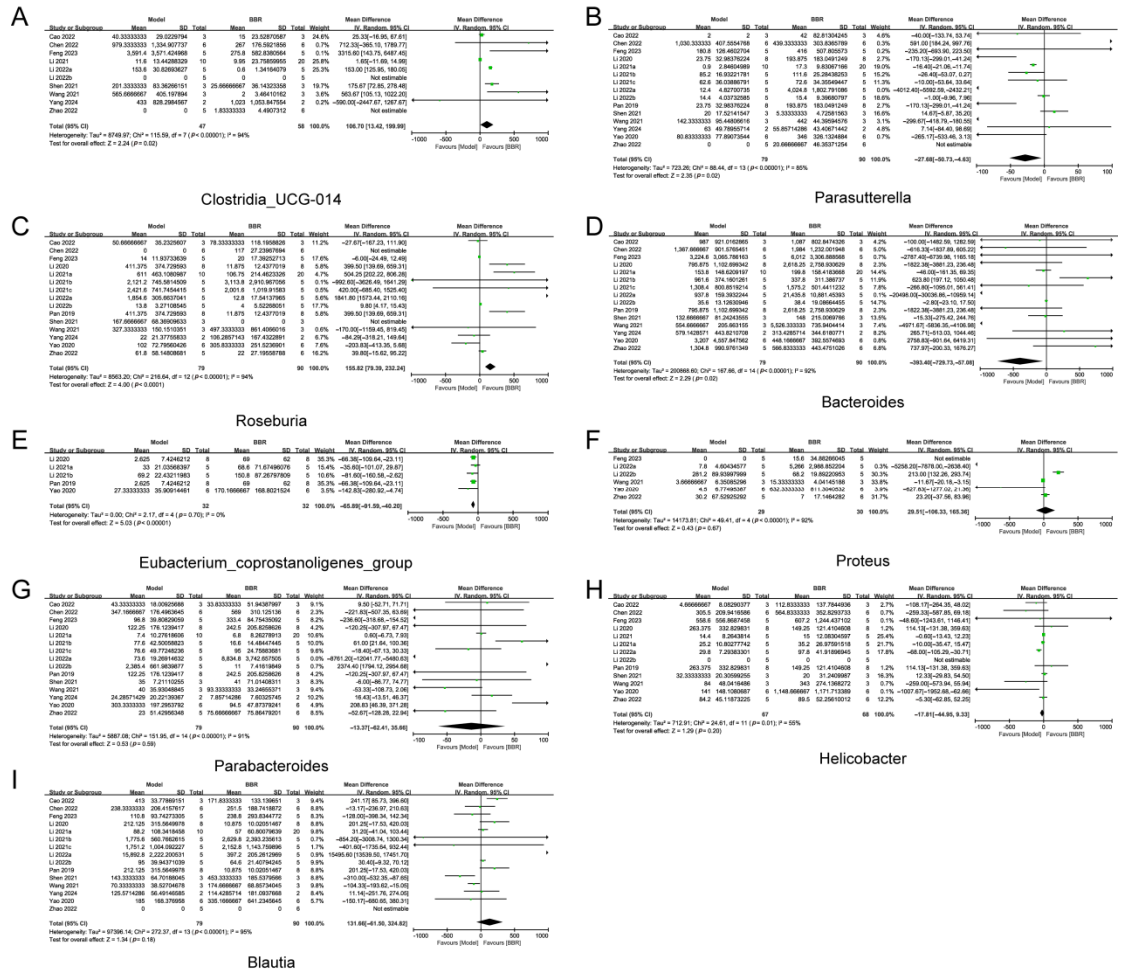


Figure S3. BBR regulated Gut bacteria in meta-analysis. (A-I) Forest plot illustrating a meta-analysis of bacteria regulated by BBR. (A) *Clostridia_UCG-014*. (B) *Parasuterella*. (C) *Roseburia*. (D) *Bacteroides*. (E) *Eubacterium_coprostanoligenes_group*. (F) *Proteus*. (G) *Parabacteroides*. (H) *Helicobacter* and (I) *Blautia*.