

$\log_{10}(\text{AMR}/\text{bacterial genes})$

-2

-3

-4

Peptococcaceae
 Flavobacteriaceae
 Turicibacteraceae
 Lachnospiraceae
 Succinivibrionaceae
 Muribaculaceae
 Burkholderiaceae
 Ruminococcaceae
 CAG-382
 Acidaminococcaceae
 Erysipelotrichaceae
 Dialisteraceae
 Coriobacteriaceae
 Micrococcaceae
 Veillonellaceae
 CAG-508
 Eggerthellaceae
 Acetatifactoraceae
 Gemellaceae
 Anaerovoracaceae
 Atopobiaceae
 CAG-917
 CAG-74
 Neisseriaceae
 Peptostreptococcaceae
 CAG-138
 Erysipelatoclostridiaceae
 UBA1381
 UBA932
 Clostridiaceae
 Monoglobaceae
 Rikenellaceae
 Victivallaceae
 Streptococcaceae
 Marinifiliaceae
 CAG-274
 Tannerellaceae
 Bacteroidaceae
 CAG-239
 Lactobacillaceae
 Butyrivibrionaceae
 Oscillospiraceae
 Porphyromonadaceae
 CAG-611
 Actinomycetaceae
 Megaspheeraceae
 Selenomonadaceae
 Gastranaerophilaceae
 Akkermansiaceae
 CAG-302
 Desulfovibrionaceae
 Anaerotruncaceae
 Pasteurellaceae
 Bifidobacteriaceae
 Barnesiellaceae
 Enterococcaceae
 Enterobacteriaceae

