

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

-2

-3

-4

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

