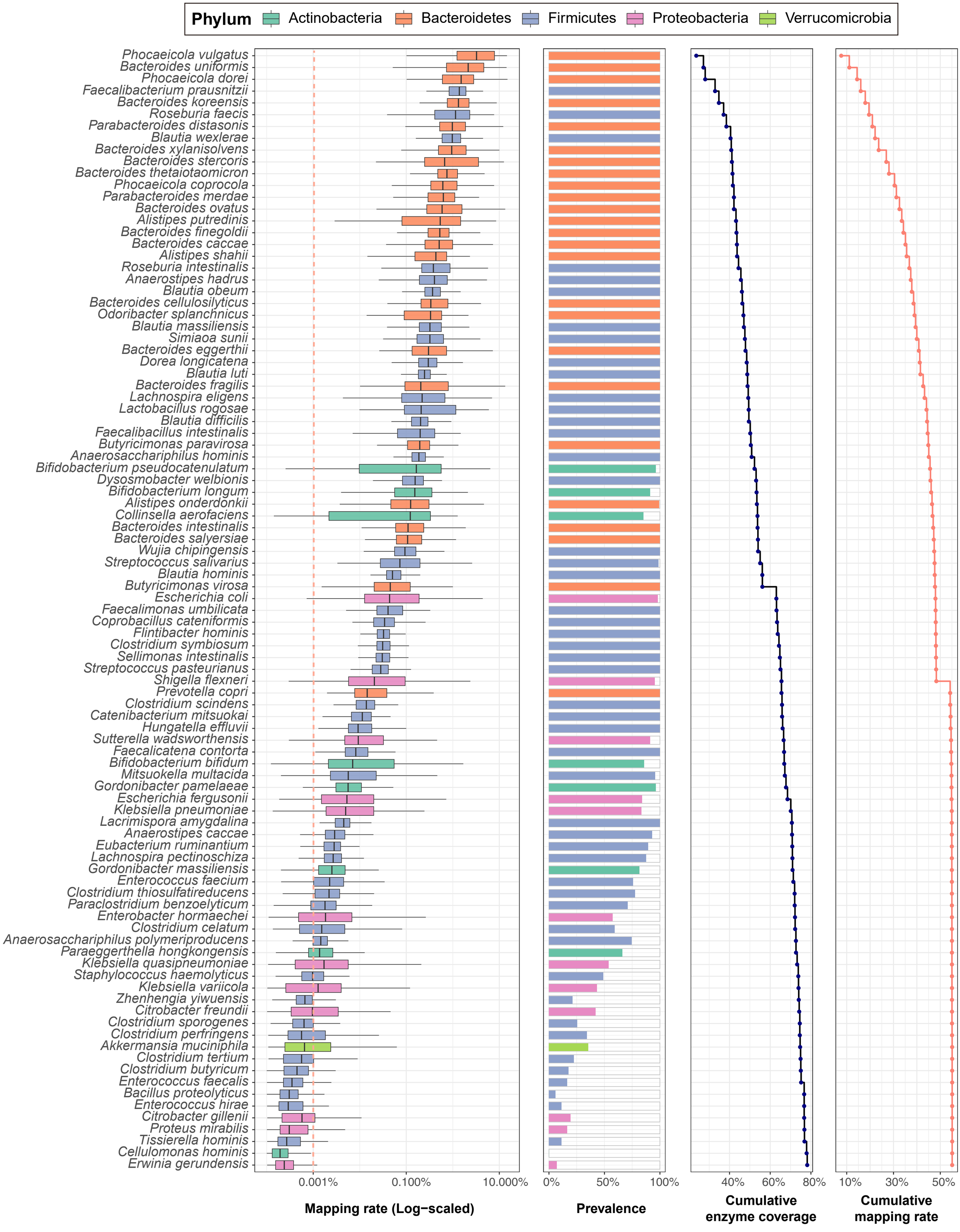
Phylum
Actinobacteria
Bacteroidetes
Firmicutes
Proteobacteria
Verrucomicrobia
Phocaeicola vulgatus
Bacteroides uniformis
Phocaeicola dorei
Faecalibacterium prausnitzii
Bacteroides koreensis
Roseburia faecis
Parabacteroides distasonis
Blautia wexlerae
Bacteroides xylanisolvens
Bacteroides stercoris
Bacteroides thetaiotaomicron
Phocaeicola coprocola
Parabacteroides merdae
Bacteroides ovatus
Alistipes putredinis
Bacteroides finegoldii
Bacteroides caccae
Alistipes shahii
Roseburia intestinalis
Anaerostipes hadrus
Blautia obeum
Bacteroides cellulosilyticus
Odoribacter splanchnicus
Blautia massiliensis
Simiaoa sunii
Bacteroides eggerthii
Dorea longicatena
Blautia luti
Bacteroides fragilis
Lachnospira eligens
Lactobacillus rogosae
Blautia difficilis
Faecalibacillus intestinalis
Butyricimonas paravirosa
Anaerosacchariphilus hominis
Bifidobacterium pseudocatenulatum
Dysosmobacter welbionis
Bifidobacterium longum
Alistipes onderdonkii
Collinsella aerofaciens
Bacteroides intestinalis
Bacteroides salyersiae
Wujia chipingensis
Streptococcus salivarius
Blautia hominis
Butyricimonas virosa
Escherichia coli
Faecalimonas umbilicata
Coprobacillus cateniformis
Flintibacter hominis
Clostridium symbiosum
Sellimonas intestinalis
Streptococcus pasteurianus
Shigella flexneri
Prevotella copri
Clostridium scindens
Catenibacterium mitsuokai
Hungatella effluvii
Sutterella wadsworthensis
Faecalicatena contorta
Bifidobacterium bifidum
Mitsuokella multacida
Gordonibacter pamelaeae
Escherichia fergusonii
Klebsiella pneumoniae
Lacrimispora amygdalina
Anaerostipes caccae
Eubacterium ruminantium
Lachnospira pectinoschiza
Gordonibacter massiliensis
Enterococcus faecium
Clostridium thiosulfatireducens
Paraclostridium benzoelyticum
Enterobacter hormaechei
Clostridium celatum
Anaerosacchariphilus polymeriproducens
Paraeggerthella hongkongensis
Klebsiella quasipneumoniae
Staphylococcus haemolyticus
Klebsiella variicola
Zhenhengia yiwuensis
Citrobacter freundii
Clostridium sporogenes
Clostridium perfringens
Akkermansia muciniphila
Clostridium tertium
Clostridium butyricum
Enterococcus faecalis
Bacillus proteolyticus
Enterococcus hirae
Citrobacter gillenii
Proteus mirabilis
Tissierella hominis
Cellulomonas hominis
Erwinia gerundensis
0.001%
0.100%
10.000%
Mapping rate (Log−scaled)
0%
50%
100%
Prevalence
40%
60%
80%
Cumulative enzyme coverage
10%
30%
50%
Cumulative mapping rate