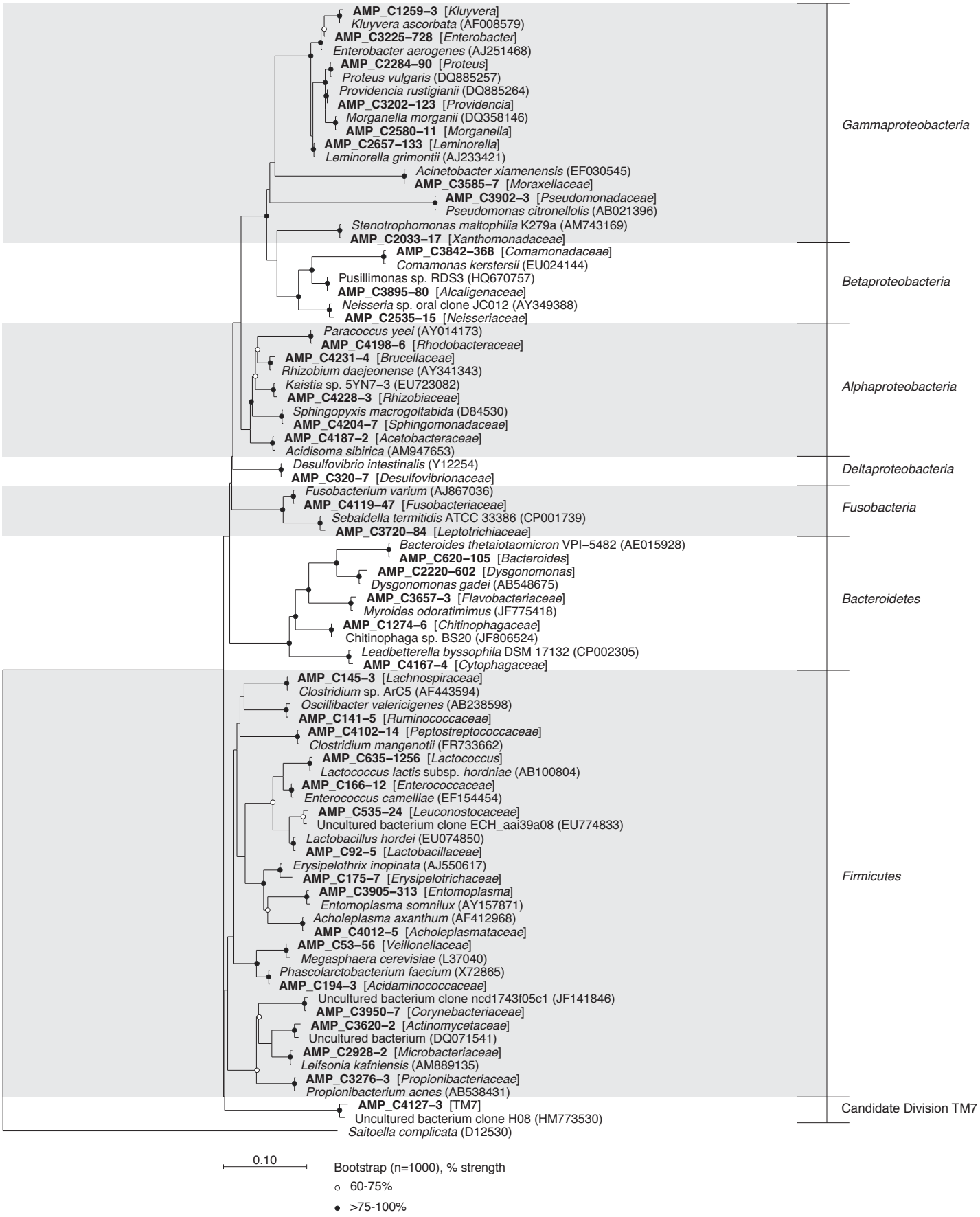
AMP_C1259–3 [Kluyvera]
Kluyvera ascorbata (AF008579)
AMP_C3225–728 [Enterobacter]
Enterobacter aerogenes (AJ251468)
AMP_C2284–90 [Proteus]
Proteus vulgaris (DQ885257)
Providencia rustigianii (DQ885264)
AMP_C3202–123 [Providencia]
Morganella morganii (DQ358146)
AMP_C2580–11 [Morganella]
AMP_C2657–133 [Leminorella]
Leminorella grimontii (AJ233421)
Acinetobacter xiamenensis (EF030545)
AMP_C3585–7 [Moraxellaceae]
AMP_C3902–3 [Pseudomonadaceae]
Pseudomonas citronellolis (AB021396)
Stenotrophomonas maltophilia K279a (AM743169)
AMP_C2033–17 [Xanthomonadaceae]
AMP_C3842–368 [Comamonadaceae]
Comamonas kerstersii (EU024144)
Pusillimonas sp. RDS3 (HQ670757)
AMP_C3895–80 [Alcaligenaceae]
Neisseria sp. oral clone JC012 (AY349388)
AMP_C2535–15 [Neisseriaceae]
Paracoccus yeei (AY014173)
AMP_C4198–6 [Rhodobacteraceae]
AMP_C4231–4 [Brucellaceae]
Rhizobium daejeonense (AY341343)
Kaistia sp. 5YN7–3 (EU723082)
AMP_C4228–3 [Rhizobiaceae]
Sphingopyxis macrogoltabida (D84530)
AMP_C4204–7 [Sphingomonadaceae]
AMP_C4187–2 [Acetobacteraceae]
Acidisoma sibirica (AM947653)
Desulfovibrio intestinalis (Y12254)
AMP_C320–7 [Desulfovibrionaceae]
Fusobacterium varium (AJ867036)
AMP_C4119–47 [Fusobacteriaceae]
Sebaldella termitidis ATCC 33386 (CP001739)
AMP_C3720–84 [Leptotrichiaceae]
Bacteroides thetaiotaomicron VPI–5482 (AE015928)
AMP_C620–105 [Bacteroides]
AMP_C2220–602 [Dysgonomonas]
Dysgonomonas gadei (AB548675)
AMP_C3657–3 [Flavobacteriaceae]
Myroides odoratimimus (JF775418)
AMP_C1274–6 [Chitinophagaceae]
Chitinophaga sp. BS20 (JF806524)
Leadbetterella byssophila DSM 17132 (CP002305)
AMP_C4167–4 [Cytophagaceae]
AMP_C145–3 [Lachnospiraceae]
Clostridium sp. ArC5 (AF443594)
Oscillibacter valericigenes (AB238598)
AMP_C141–5 [Ruminococcaceae]
AMP_C4102–14 [Peptostreptococcaceae]
Clostridium mangenotii (FR733662)
AMP_C635–1256 [Lactococcus]
Lactococcus lactis subsp. hordniae (AB100804)
AMP_C166–12 [Enterococcaceae]
Enterococcus camelliae (EF154454)
AMP_C535–24 [Leuconostocaceae]
Uncultured bacterium clone ECH_aai39a08 (EU774833)
Lactobacillus hordei (EU074850)
AMP_C92–5 [Lactobacillaceae]
Erysipelothrix inopinata (AJ550617)
AMP_C175–7 [Erysipelotrichaceae]
AMP_C3905–313 [Entomoplasma]
Entomoplasma somnilux (AY157871)
Acholeplasma axanthum (AF412968)
AMP_C4012–5 [Acholeplasmataceae]
AMP_C53–56 [Veillonellaceae]
Megasphaera cerevisiae (L37040)
Phascolarctobacterium faecium (X72865)
AMP_C194–3 [Acidaminococcaceae]
Uncultured bacterium clone ncd1743f05c1 (JF141846)
AMP_C3950–7 [Corynebacteriaceae]
AMP_C3620–2 [Actinomycetaceae]
Uncultured bacterium (DQ071541)
AMP_C2928–2 [Microbacteriaceae]
Leifsonia kafniensis (AM889135)
AMP_C3276–3 [Propionibacteriaceae]
Propionibacterium acnes (AB538431)
AMP_C4127–3 [TM7]
Uncultured bacterium clone H08 (HM773530)
Saitoella complicata (D12530)
Gammaproteobacteria
Betaproteobacteria
Alphaproteobacteria
Deltaproteobacteria
Fusobacteria
Bacteroidetes
Firmicutes
Candidate Division TM7
0.10
Bootstrap (n=1000), % strength
60-75%
>75-100%