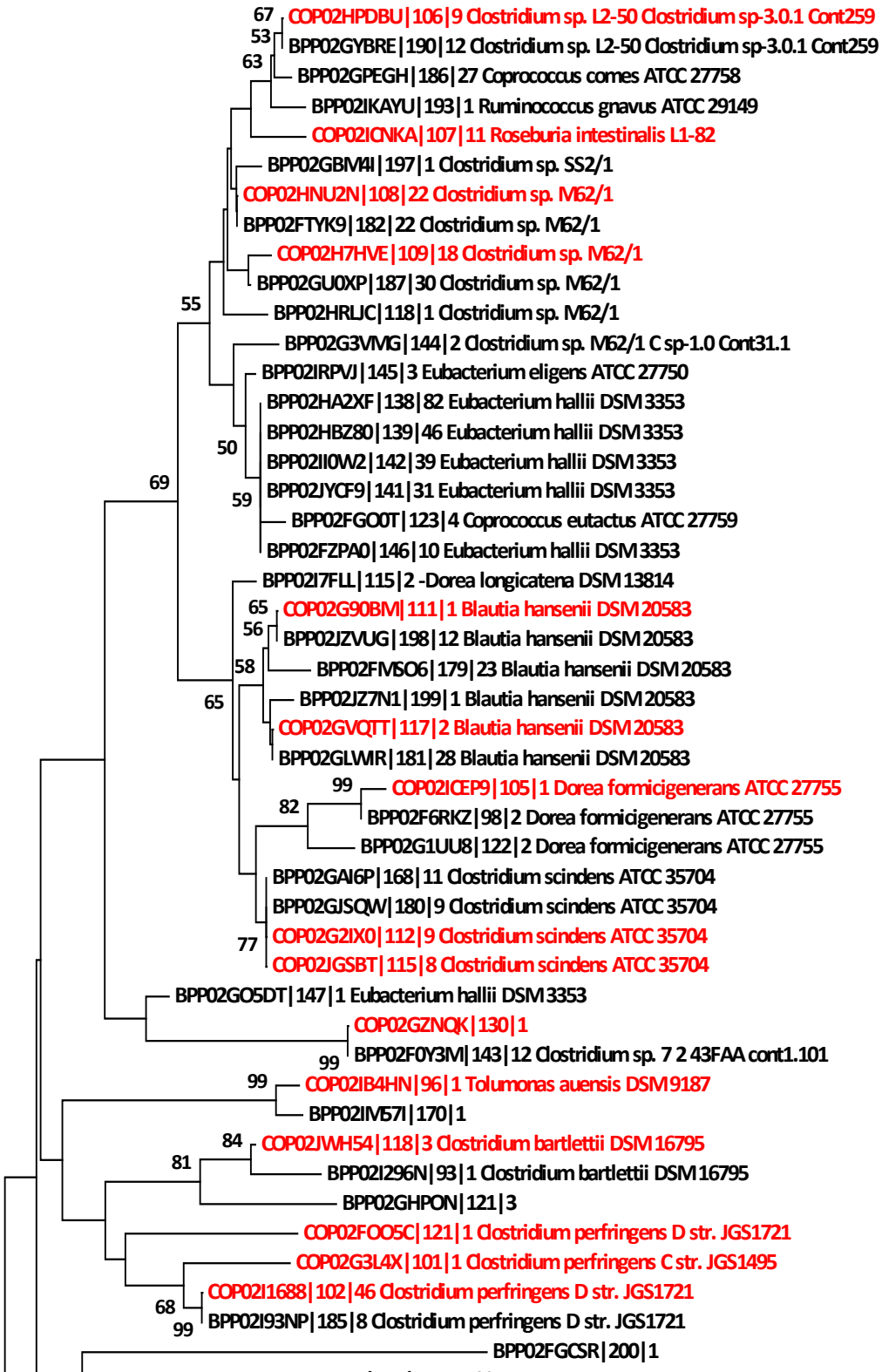
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BPP02IKAYU|193|1 Ruminococcus gnavus ATCC 29149
COP02ICNKA|107|11 Roseburia intestinalis L1-82
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COP02H7HVE|109|18 Clostridium sp. M62/1
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BPP02HRLJC|118|1 Clostridium sp. M62/1
BPP02G3VMG|144|2 Clostridium sp. M62/1 C sp-1.0 Cont31.1
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BPP02HA2XF|138|82 Eubacterium hallii DSM 3353
BPP02HBZ80|139|46 Eubacterium hallii DSM 3353
BPP02II0W2|142|39 Eubacterium hallii DSM 3353
BPP02JYCF9|141|31 Eubacterium hallii DSM 3353
BPP02FGO0T|123|4 Coprococcus eutactus ATCC 27759
BPP02FZPA0|146|10 Eubacterium hallii DSM 3353
BPP02I7FLL|115|2 -Dorea longicatena DSM 13814
COP02G90BM|111|1 Blautia hansenii DSM 20583
BPP02JZVUG|198|12 Blautia hansenii DSM 20583
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BPP02JZ7N1|199|1 Blautia hansenii DSM 20583
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BPP02GLWIR|181|28 Blautia hansenii DSM 20583
COP02ICEP9|105|1 Dorea formicigenerans ATCC 27755
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COP02G3L4X|101|1 Clostridium perfringens C str. JGS1495
COP02I1688|102|46 Clostridium perfringens D str. JGS1721
BPP02I93NP|185|8 Clostridium perfringens D str. JGS1721
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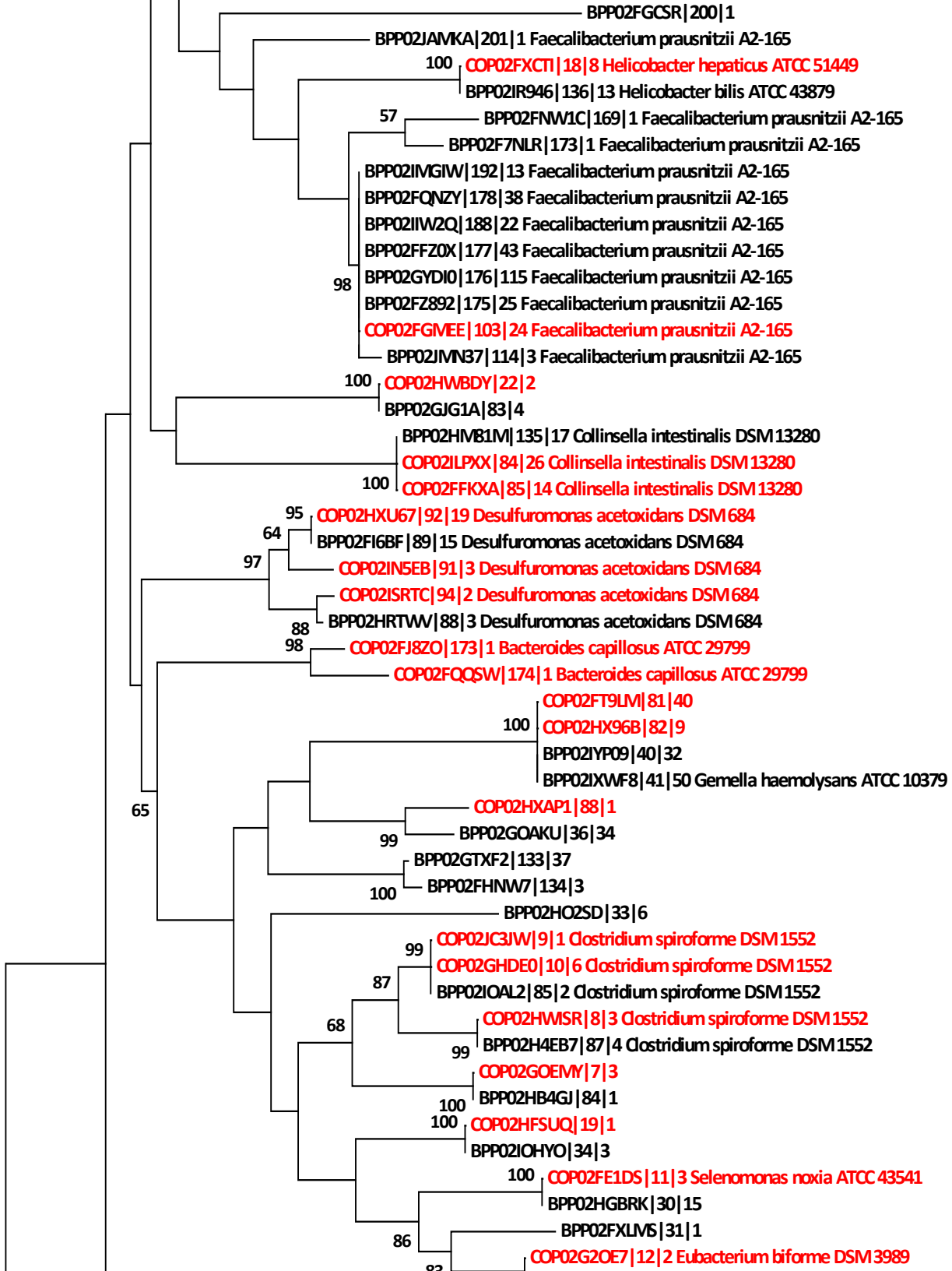
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COP02HWISR|8|3 Clostridium spiroforme DSM 1552
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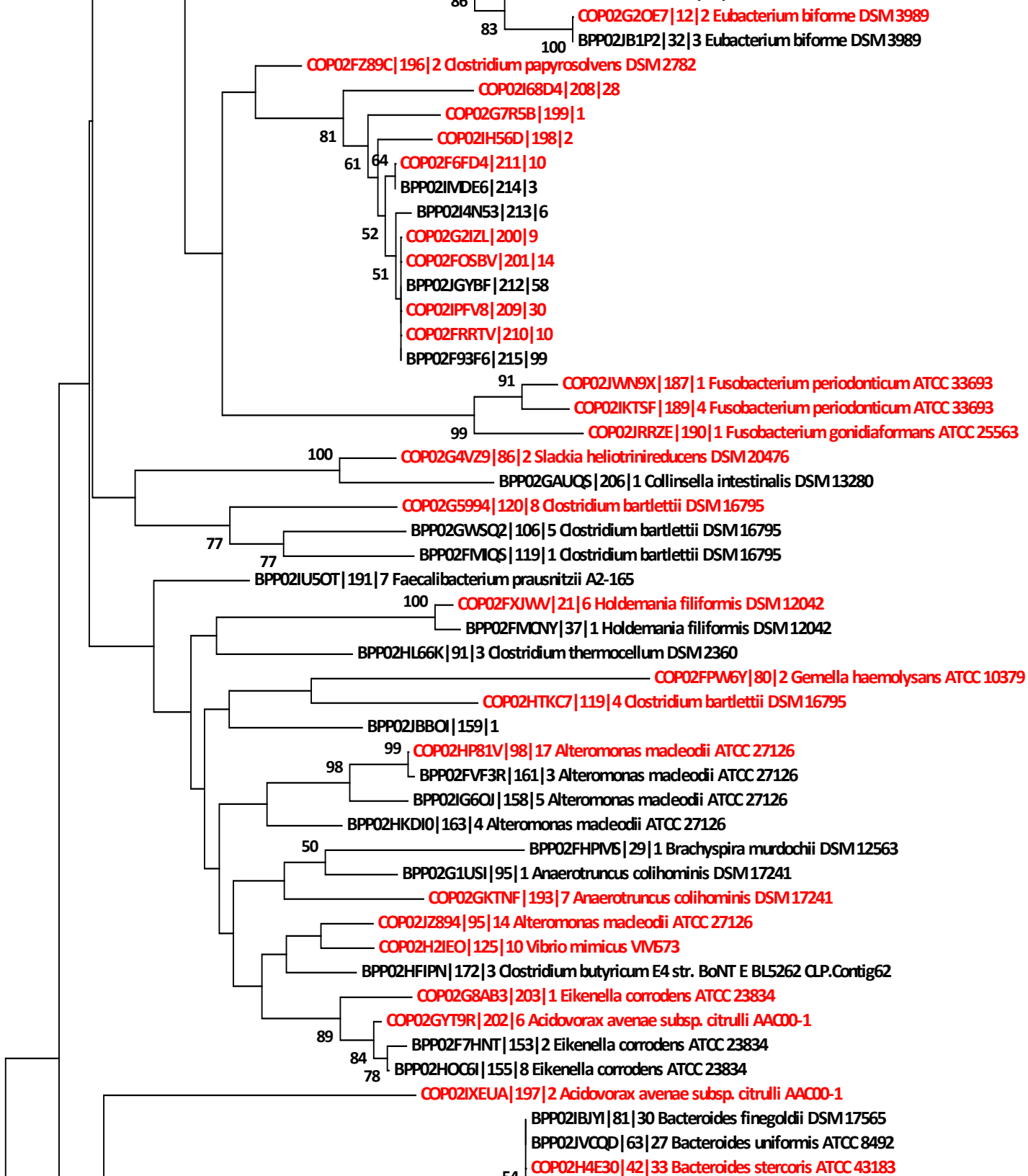
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COP02IPFV8|209|30
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91
COP02JWN9X|187|1 Fusobacterium periodonticum ATCC 33693
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COP02JZ894|95|14 Alteromonas macleodii ATCC 27126
COP02H2IEO|125|10 Vibrio mimicus VM573
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COP02GYT9R|202|6 Acidovorax avenae subsp. citrulli AAC00-1
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78
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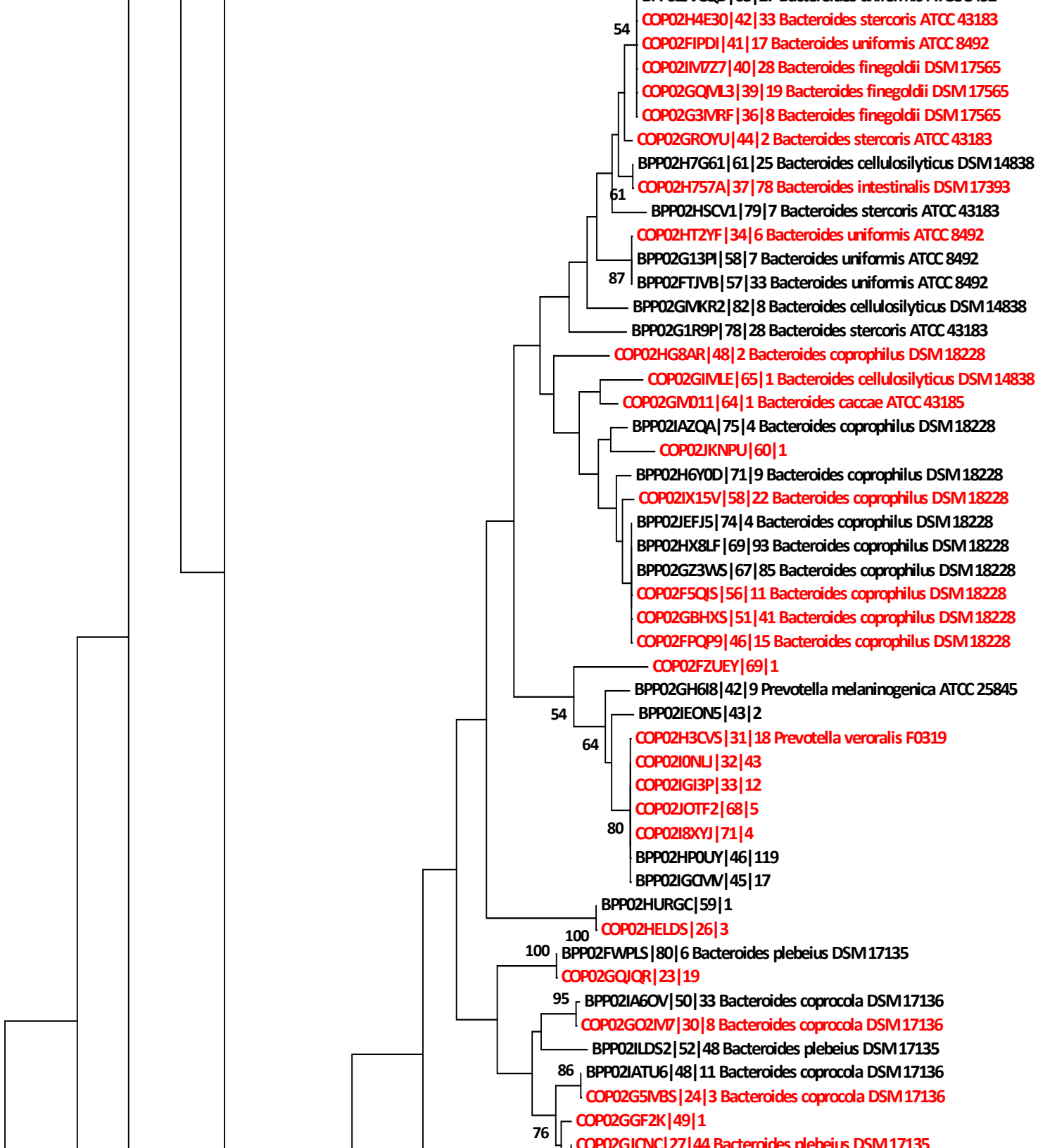
COP02IXEUA|197|2 Acidovorax avenae subsp. citrulli AAC00-1
BPP02IBJYI|81|30 Bacteroides finegoldii DSM 17565
BPP02JVCQD|63|27 Bacteroides uniformis ATCC 8492
COP02H4E30|42|33 Bacteroides stercoris ATCC 43183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BPP02H7G61|61|25 Bacteroides cellulosilyticus DSM 14838
COP02H757A|37|78 Bacteroides intestinalis DSM 17393
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BPP02G13PI|58|7 Bacteroides uniformis ATCC 8492
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BPP02FTJVB|57|33 Bacteroides uniformis ATCC 8492
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COP02GIMLE|65|1 Bacteroides cellulosilyticus DSM 14838
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86
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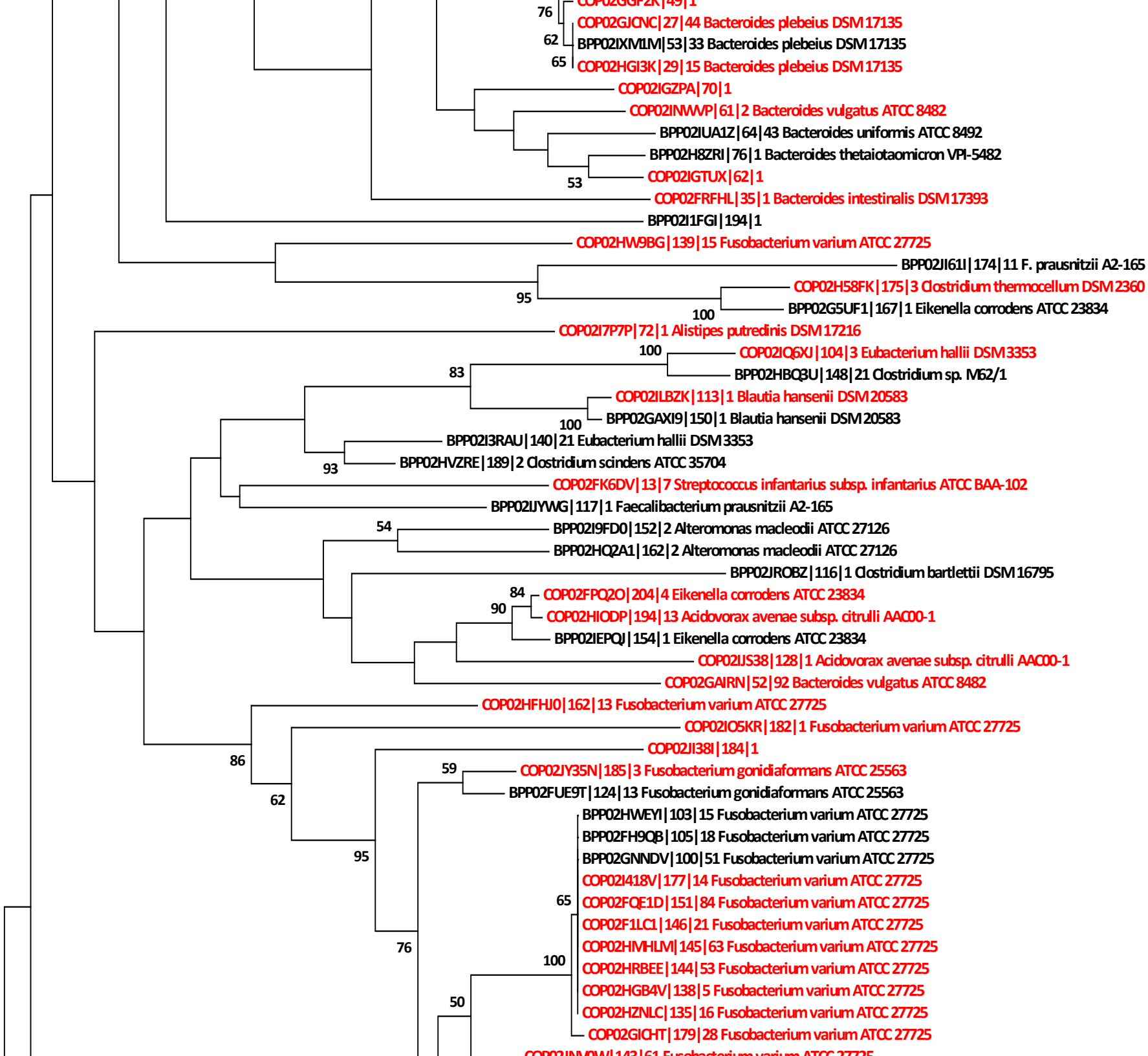
COP02GICNC|27|44 Bacteroides plebeius DSM 17135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COP02IO5KR|182|1 Fusobacterium varium ATCC 27725
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COP02JY35N|185|3 Fusobacterium gonidiaformans ATCC 25563
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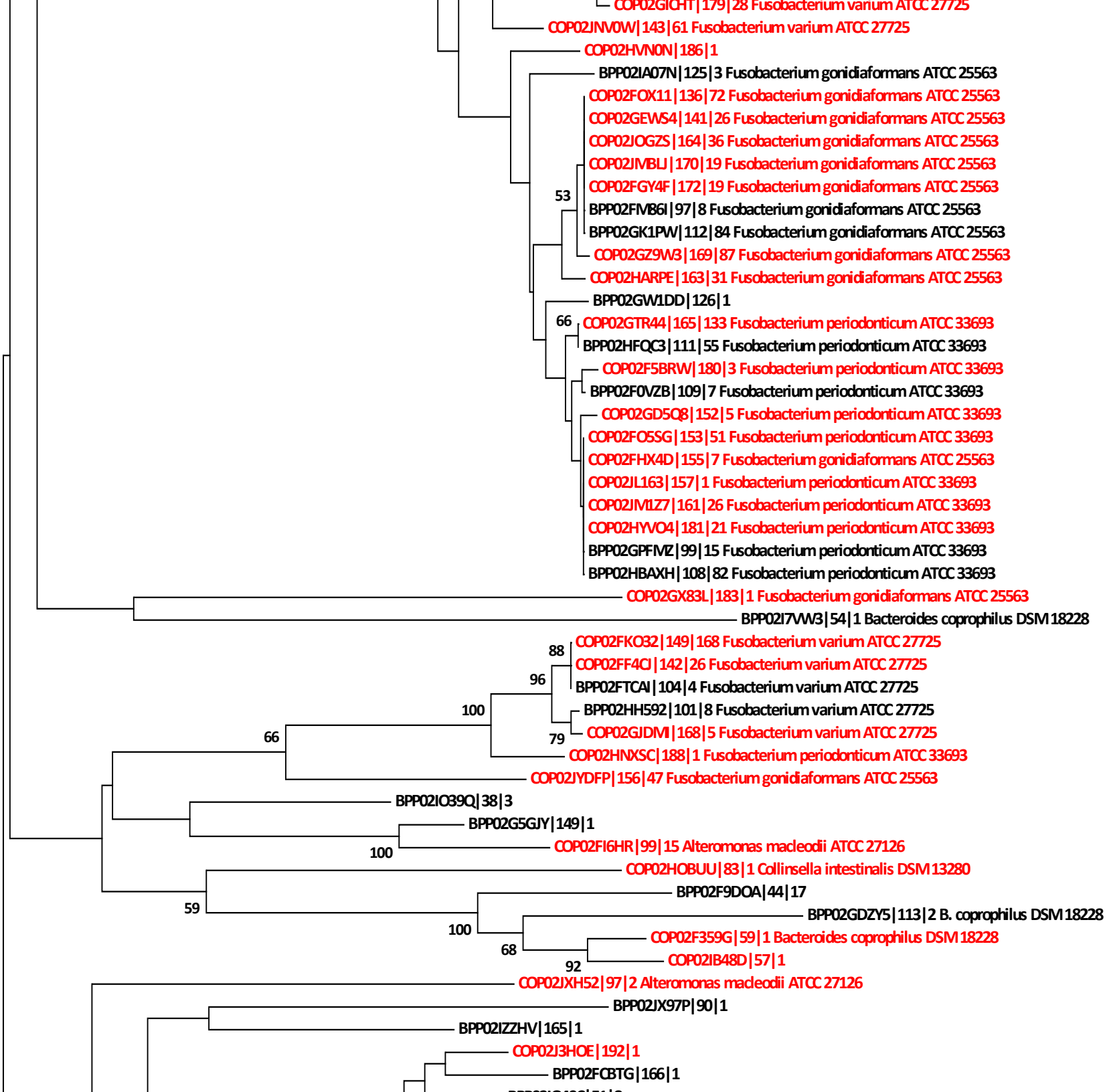
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COP02FHX4D|155|7 Fusobacterium gonidiaformans ATCC 25563
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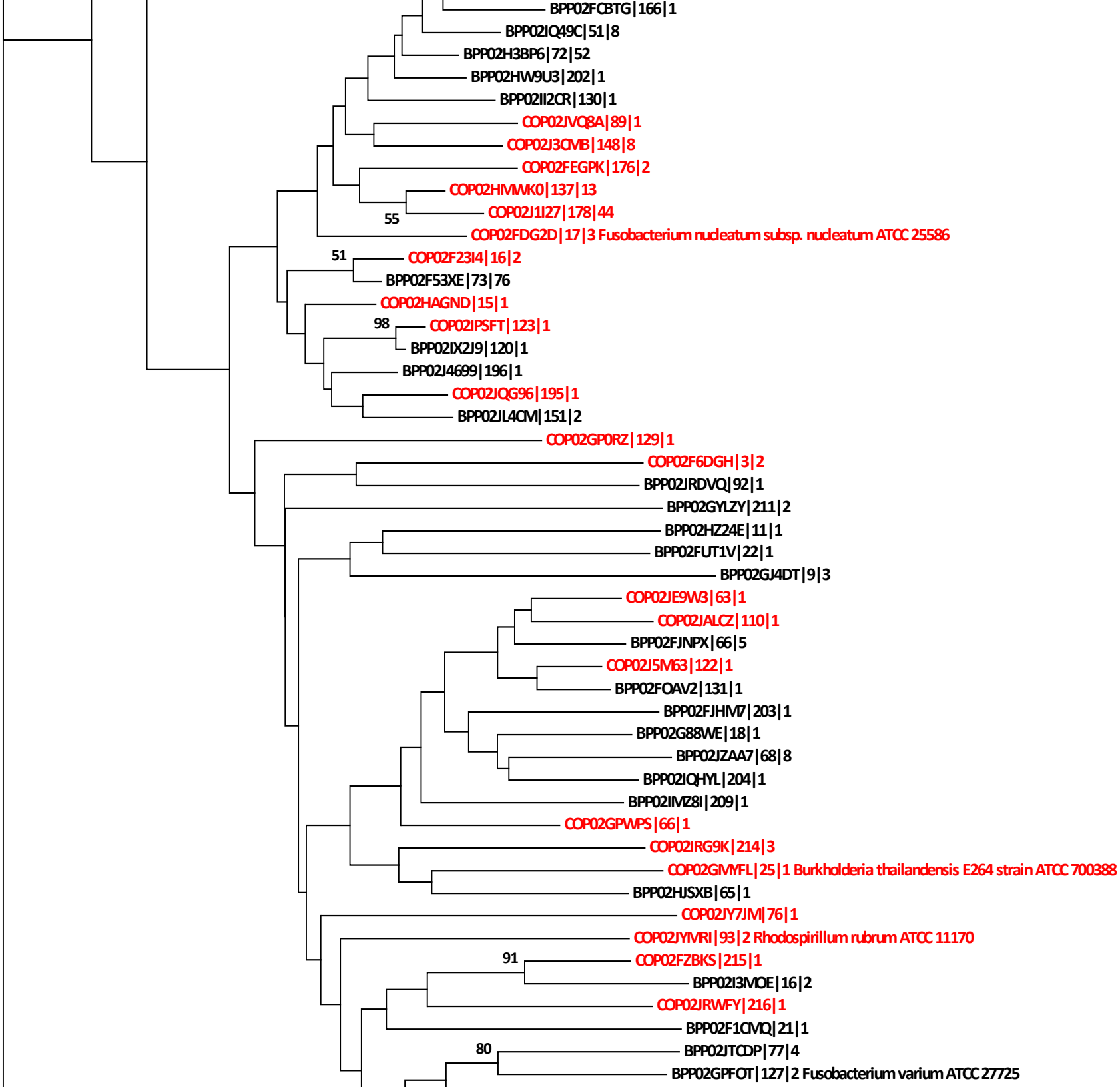
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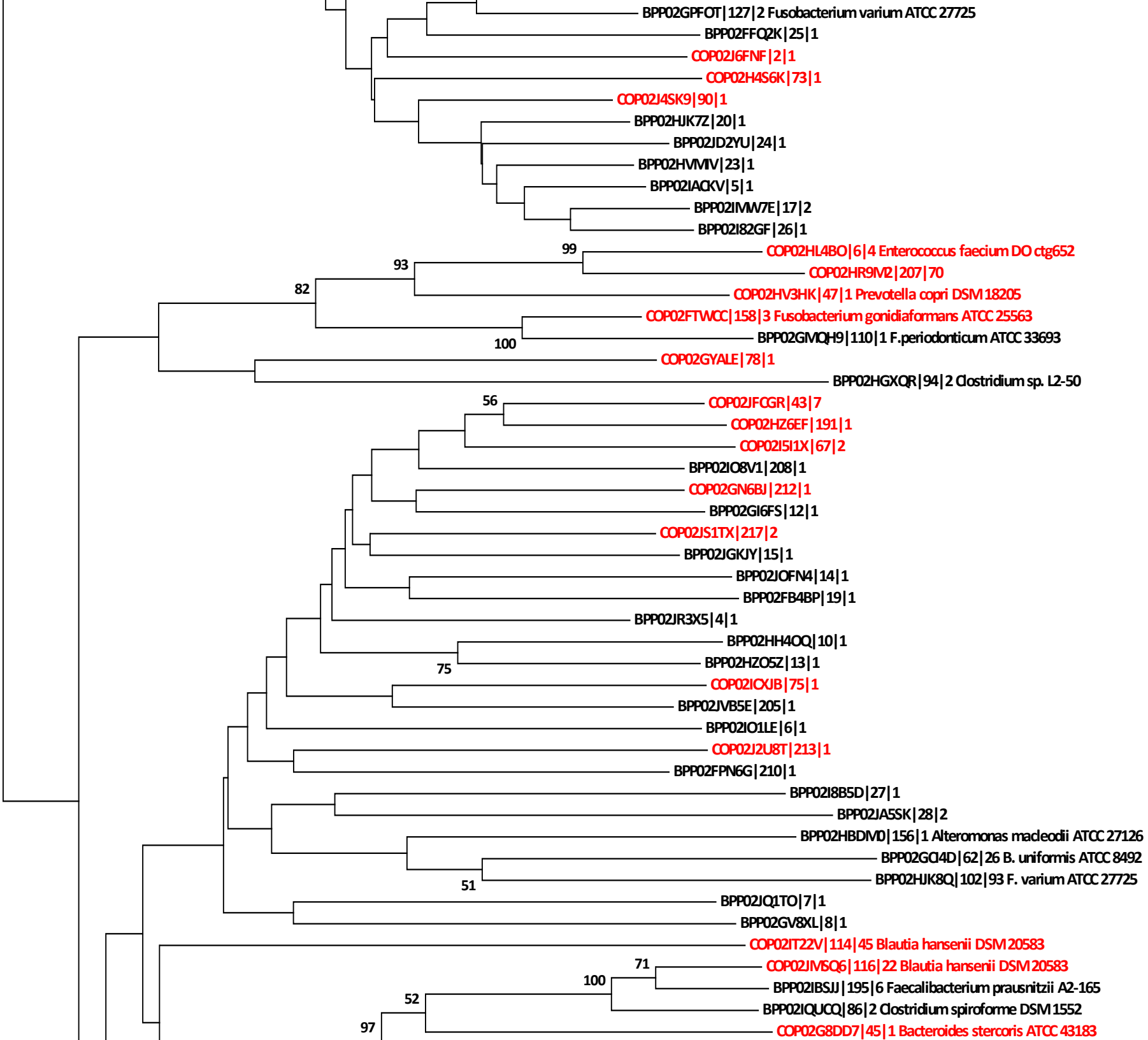
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COP02HL4BO|6|4 Enterococcus faecium DO ctg652
93
COP02HR9IV2|207|70
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COP02HV3HK|47|1 Prevotella copri DSM 18205
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BPP02I8B5D|27|1
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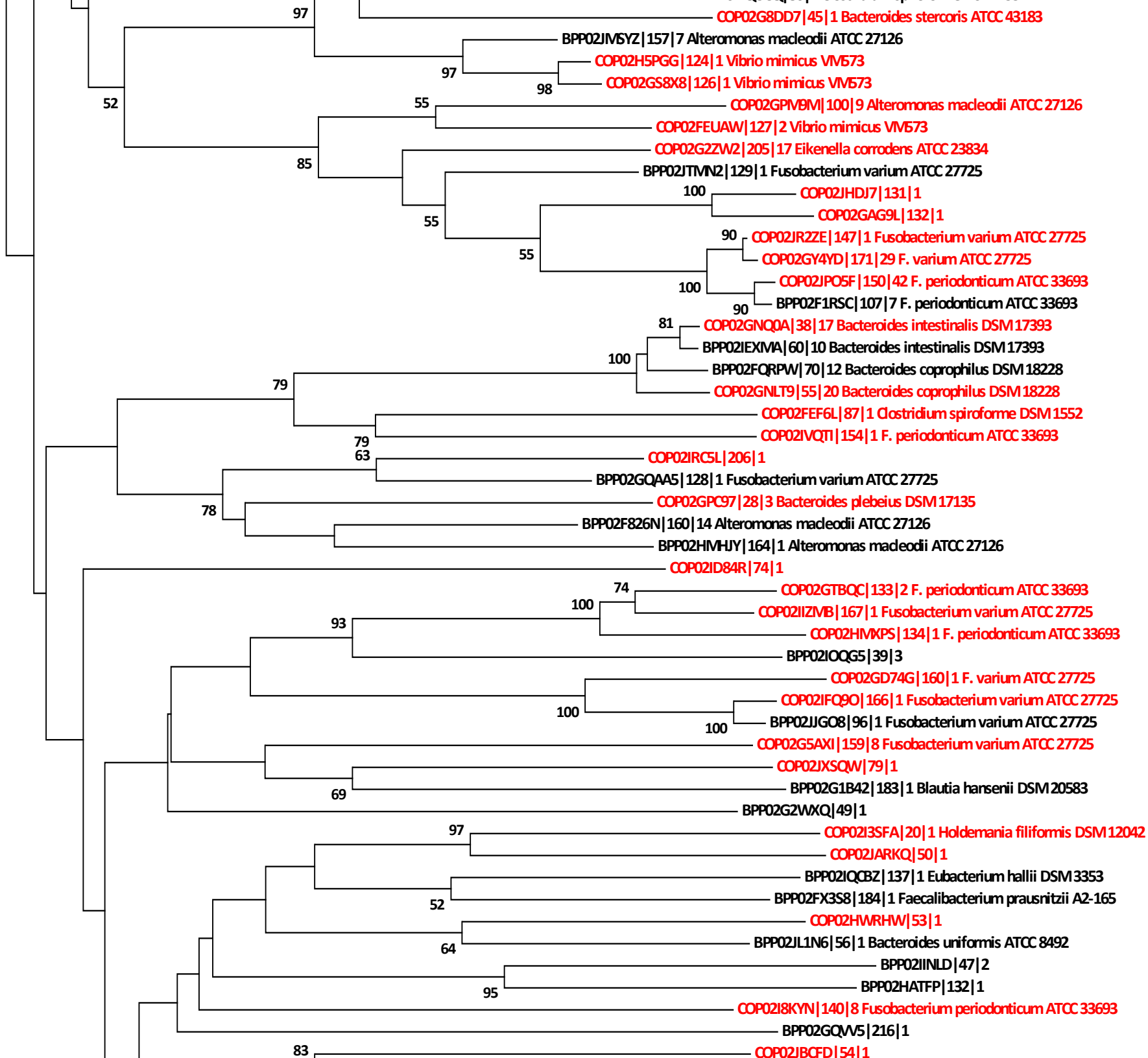
BPP02HBDMD|156|1 Alteromonas macleodii ATCC 27126
BPP02GCI4D|62|26 B. uniformis ATCC 8492
51
BPP02HJK8Q|102|93 F. varium ATCC 27725
BPP02JQ1TO|7|1
BPP02GV8XL|8|1
COP02IT22V|114|45 Blautia hansenii DSM 20583
71
COP02JMSQ6|116|22 Blautia hansenii DSM 20583
100
BPP02IBSJJ|195|6 Faecalibacterium prausnitzii A2-165
52
BPP02IQUCQ|86|2 Clostridium spiroforme DSM 1552
97
COP02G8DD7|45|1 Bacteroides stercoris ATCC 43183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COP02G2ZW2|205|17 Eikenella corrodens ATCC 23834
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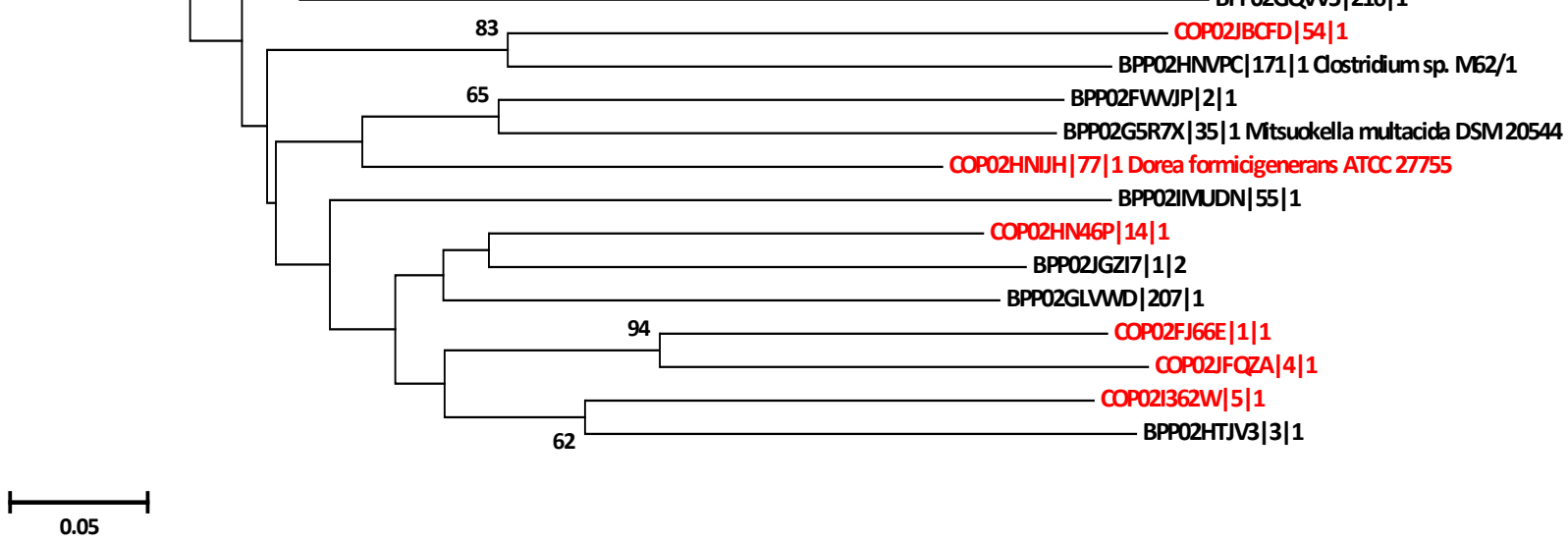
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