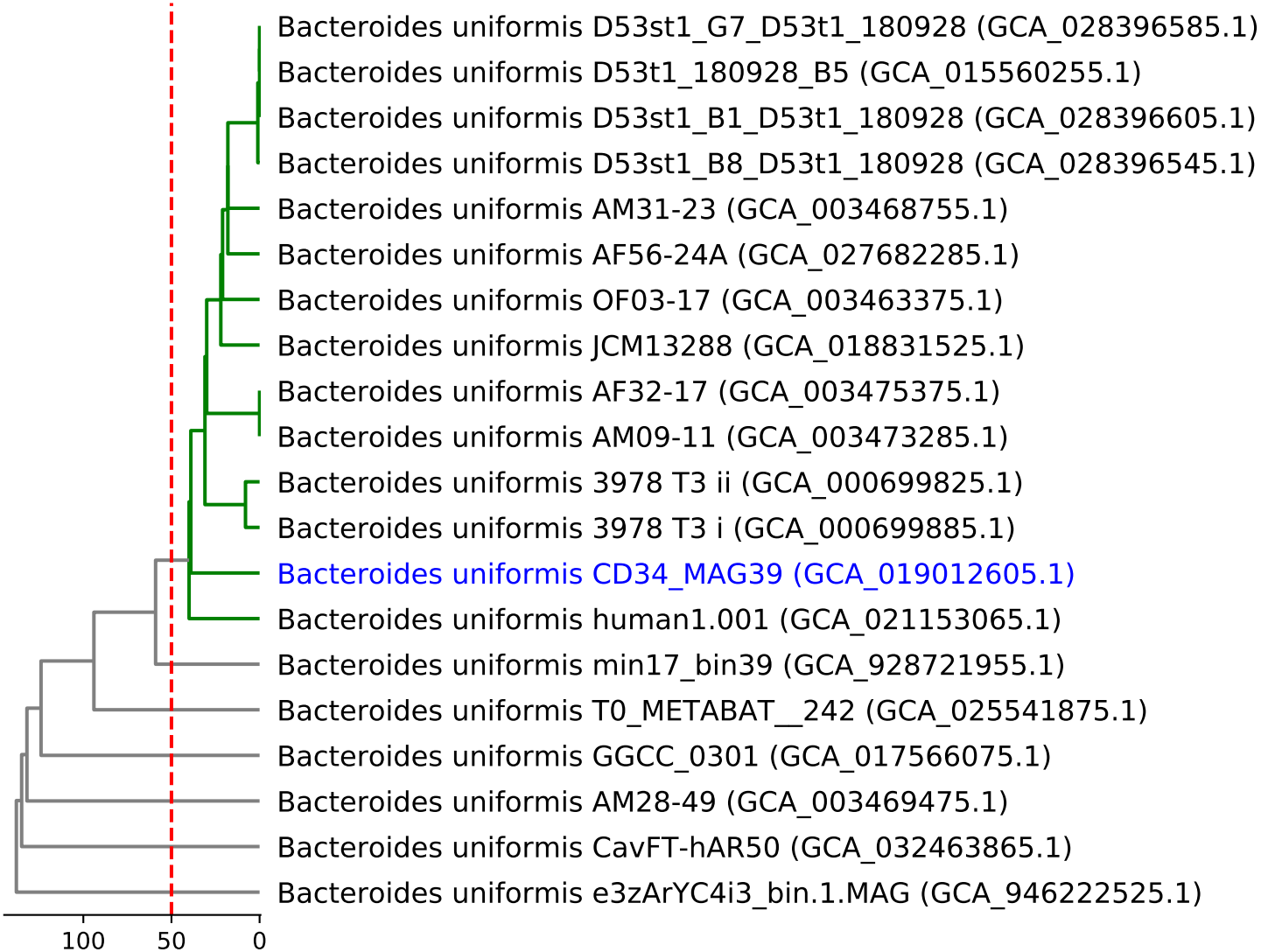
Bacteroides uniformis D53st1_G7_D53t1_180928 (GCA_028396585.1)
Bacteroides uniformis D53t1_180928_B5 (GCA_015560255.1)
Bacteroides uniformis D53st1_B1_D53t1_180928 (GCA_028396605.1)
Bacteroides uniformis D53st1_B8_D53t1_180928 (GCA_028396545.1)
Bacteroides uniformis AM31-23 (GCA_003468755.1)
Bacteroides uniformis AF56-24A (GCA_027682285.1)
Bacteroides uniformis OF03-17 (GCA_003463375.1)
Bacteroides uniformis JCM13288 (GCA_018831525.1)
Bacteroides uniformis AF32-17 (GCA_003475375.1)
Bacteroides uniformis AM09-11 (GCA_003473285.1)
Bacteroides uniformis 3978 T3 ii (GCA_000699825.1)
Bacteroides uniformis 3978 T3 i (GCA_000699885.1)
Bacteroides uniformis CD34_MAG39 (GCA_019012605.1)
Bacteroides uniformis human1.001 (GCA_021153065.1)
Bacteroides uniformis min17_bin39 (GCA_928721955.1)
Bacteroides uniformis T0_METABAT__242 (GCA_025541875.1)
Bacteroides uniformis GGCC_0301 (GCA_017566075.1)
Bacteroides uniformis AM28-49 (GCA_003469475.1)
Bacteroides uniformis CavFT-hAR50 (GCA_032463865.1)
Bacteroides uniformis e3zArYC4i3_bin.1.MAG (GCA_946222525.1)
100
50
0