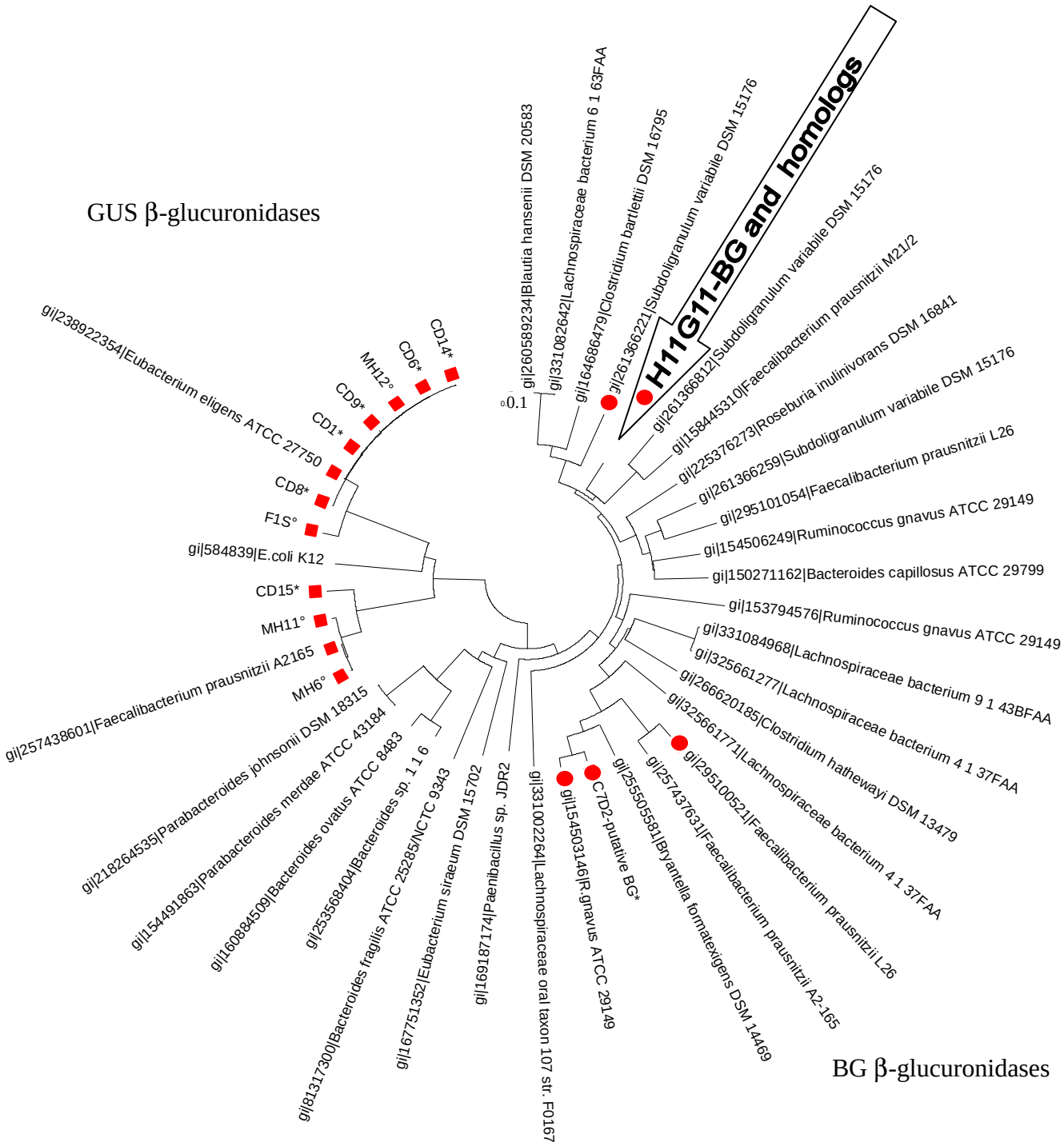
GUS β-glucuronidases
BG β-glucuronidases
H11G11-BG and homologs
0.1
gi|260589234|Blautia hansenii DSM 20583
gi|331082642|Lachnospiraceae bacterium 6 1 63FAA
gi|164686479|Clostridium bartlettii DSM 16795
gi|261366221|Subdoligranulum variabile DSM 15176
gi|261366812|Subdoligranulum variabile DSM 15176
gi|158445310|Faecalibacterium prausnitzii M21/2
gi|225376273|Roseburia inulinivorans DSM 16841
gi|261366259|Subdoligranulum variabile DSM 15176
gi|295101054|Faecalibacterium prausnitzii L26
gi|154506249|Ruminococcus gnavus ATCC 29149
gi|150271162|Bacteroides capillosus ATCC 29799
gi|153794576|Ruminococcus gnavus ATCC 29149
gi|331084968|Lachnospiraceae bacterium 9 1 43BFAA
gi|325661277|Lachnospiraceae bacterium 4 1 37FAA
gi|266620185|Clostridium hathewayi DSM 13479
gi|325661771|Lachnospiraceae bacterium 4 1 37FAA
gi|295100521|Faecalibacterium prausnitzii L26
gi|257437631|Faecalibacterium prausnitzii A2-165
gi|255505581|Bryantella formatexigens DSM 14469
C7D2-putative BG*
gi|154503146|R.gnavus ATCC 29149
gi|331002264|Lachnospiraceae oral taxon 107 str. F0167
gi|169187174|Paenibacillus sp. JDR2
gi|167751352|Eubacterium siraeum DSM 15702
gi|253568404|Bacteroides sp. 1 1 6
gi|83117300|Bacteroides fragilis ATCC 25285/NCTC 9343
gi|160884509|Bacteroides ovatus ATCC 8483
gi|154491863|Parabacteroides merdae ATCC 43184
gi|218264535|Parabacteroides johnsonii DSM 18315
gi|257438601|Faecalibacterium prausnitzii A2165
MH6°
MH11°
CD15*
gi|584839|E.coli K12
F1S°
CD8*
CD1*
CD9*
MH12°
CD6*
CD14*
gi|238922354|Eubacterium eligens ATCC 27750

S4 Figure