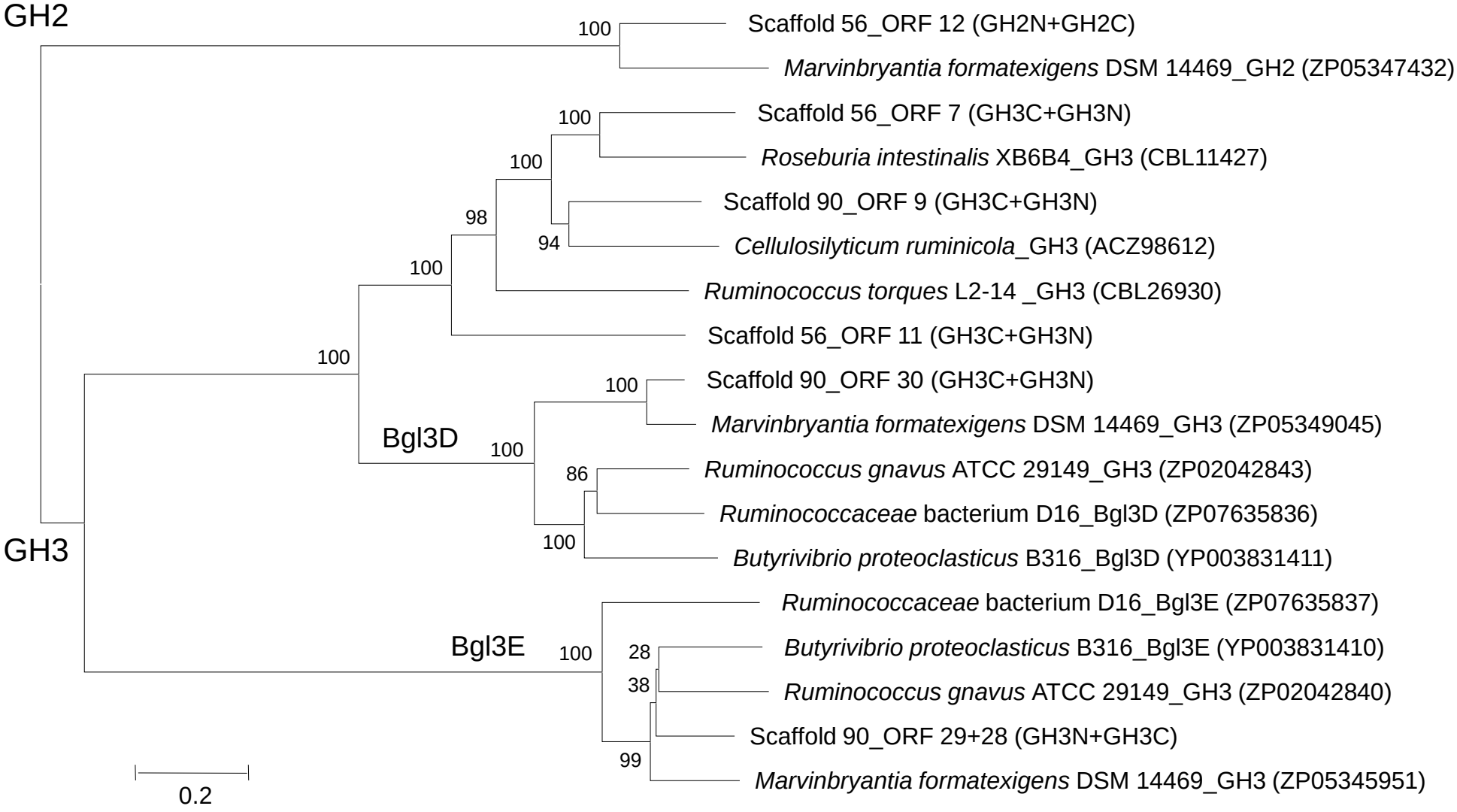
GH2
100
Scaffold 56_ORF 12 (GH2N+GH2C)
Marvinbryantia formatexigens DSM 14469_GH2 (ZP05347432)
100
Scaffold 56_ORF 7 (GH3C+GH3N)
100
Roseburia intestinalis XB6B4_GH3 (CBL11427)
Scaffold 90_ORF 9 (GH3C+GH3N)
98
94
Cellulosilyticum ruminicola_GH3 (ACZ98612)
100
Ruminococcus torques L2-14 _GH3 (CBL26930)
Scaffold 56_ORF 11 (GH3C+GH3N)
100
Scaffold 90_ORF 30 (GH3C+GH3N)
100
Marvinbryantia formatexigens DSM 14469_GH3 (ZP05349045)
Bgl3D
100
86
Ruminococcus gnavus ATCC 29149_GH3 (ZP02042843)
Ruminococcaceae bacterium D16_Bgl3D (ZP07635836)
100
Butyrivibrio proteoclasticus B316_Bgl3D (YP003831411)
GH3
Ruminococcaceae bacterium D16_Bgl3E (ZP07635837)
Bgl3E
100
28
Butyrivibrio proteoclasticus B316_Bgl3E (YP003831410)
38
Ruminococcus gnavus ATCC 29149_GH3 (ZP02042840)
Scaffold 90_ORF 29+28 (GH3N+GH3C)
99
Marvinbryantia formatexigens DSM 14469_GH3 (ZP05345951)
0.2