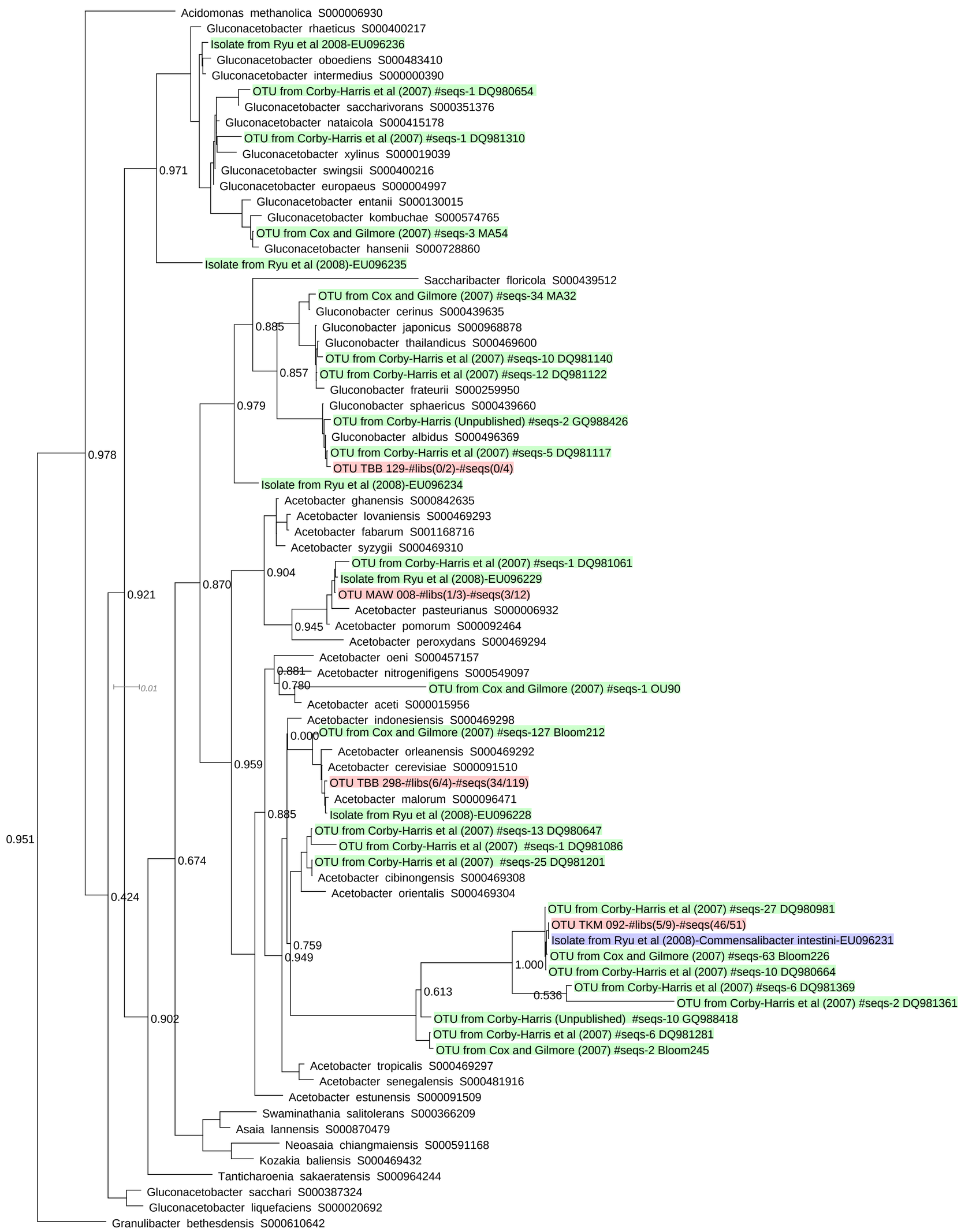
Acidomonas methanolica S000006930
Gluconacetobacter rhaeticus S000400217
Isolate from Ryu et al 2008-EU096236
Gluconacetobacter oboediens S000483410
Gluconacetobacter intermedius S000000390
OTU from Corby-Harris et al (2007) #seqs-1 DQ980654
Gluconacetobacter saccharivorans S000351376
Gluconacetobacter nataicola S000415178
OTU from Corby-Harris et al (2007) #seqs-1 DQ981310
Gluconacetobacter xylinus S000019039
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Gluconacetobacter swingsii S000400216
Gluconacetobacter europaeus S000004997
Gluconacetobacter entanii S000130015
Gluconacetobacter kombuchae S000574765
OTU from Cox and Gilmore (2007) #seqs-3 MA54
Gluconacetobacter hansenii S000728860
Isolate from Ryu et al (2008)-EU096235
Saccharibacter floricola S000439512
OTU from Cox and Gilmore (2007) #seqs-34 MA32
Gluconobacter cerinus S000439635
0.885
Gluconobacter japonicus S000968878
Gluconobacter thailandicus S000469600
OTU from Corby-Harris et al (2007) #seqs-10 DQ981140
0.857
OTU from Corby-Harris et al (2007) #seqs-12 DQ981122
Gluconobacter frateurii S000259950
0.979
Gluconobacter sphaericus S000439660
OTU from Corby-Harris (Unpublished) #seqs-2 GQ988426
Gluconobacter albidus S000496369
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OTU from Corby-Harris et al (2007) #seqs-5 DQ981117
OTU TBB 129-#libs(0/2)-#seqs(0/4)
Isolate from Ryu et al (2008)-EU096234
Acetobacter ghanensis S000842635
Acetobacter lovaniensis S000469293
Acetobacter fabarum S001168716
Acetobacter syzygii S000469310
OTU from Corby-Harris et al (2007) #seqs-1 DQ981061
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Isolate from Ryu et al (2008)-EU096229
0.870
OTU MAW 008-#libs(1/3)-#seqs(3/12)
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Acetobacter pasteurianus S000006932
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Acetobacter pomorum S000092464
Acetobacter peroxydans S000469294
Acetobacter oeni S000457157
0.881
Acetobacter nitrogenifigens S000549097
0.01
0.780
OTU from Cox and Gilmore (2007) #seqs-1 OU90
Acetobacter aceti S000015956
Acetobacter indonesiensis S000469298
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OTU from Cox and Gilmore (2007) #seqs-127 Bloom212
Acetobacter orleanensis S000469292
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Acetobacter cerevisiae S000091510
OTU TBB 298-#libs(6/4)-#seqs(34/119)
Acetobacter malorum S000096471
0.885
Isolate from Ryu et al (2008)-EU096228
OTU from Corby-Harris et al (2007) #seqs-13 DQ980647
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OTU from Corby-Harris et al (2007) #seqs-1 DQ981086
0.674
OTU from Corby-Harris et al (2007) #seqs-25 DQ981201
Acetobacter cibinongensis S000469308
Acetobacter orientalis S000469304
0.424
OTU from Corby-Harris et al (2007) #seqs-27 DQ980981
OTU TKM 092-#libs(5/9)-#seqs(46/51)
Isolate from Ryu et al (2008)-Commensalibacter intestini-EU096231
0.759
0.949
OTU from Cox and Gilmore (2007) #seqs-63 Bloom226
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OTU from Corby-Harris et al (2007) #seqs-10 DQ980664
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OTU from Corby-Harris et al (2007) #seqs-6 DQ981369
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OTU from Corby-Harris et al (2007) #seqs-2 DQ981361
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OTU from Corby-Harris (Unpublished) #seqs-10 GQ988418
OTU from Corby-Harris et al (2007) #seqs-6 DQ981281
OTU from Cox and Gilmore (2007) #seqs-2 Bloom245
Acetobacter tropicalis S000469297
Acetobacter senegalensis S000481916
Acetobacter estunensis S000091509
Swaminathania salitolerans S000366209
Asaia lannensis S000870479
Neoasaia chiangmaiensis S000591168
Kozakia baliensis S000469432
Tanticharoenia sakaeratensis S000964244
Gluconacetobacter sacchari S000387324
Gluconacetobacter liquefaciens S000020692
Granulibacter bethesdensis S000610642