ACC2 (PF00364): gene tree
Rooted at midpoint
Tree is ultrametric
Scale: 0.5 substitutions per site
Tree includes 1000 sequences, 1000 sequences in the alignment, 1000 sequences in the tree including PDs
Assembly: job 30 PSJ2KP (assembly=LongQuick)
Alignment: 1000 x 2000
Generated: 2 of 100
Analysed by HomeBuilt-Bacteria v2
Date: Tue Jul 11 10:00:47 2006

gi|167.0.1.0.0 [Bifidobacterium adolescentis Synthetase protein Bacto_0167.0.0]