

[illegible][illegible]

COG3186

Phylogenetic tree showing relationships between various bacterial species. The tree is rooted at 'root 36' and branches out into numerous clades. The species names are listed on the right side of the tree, with green circles indicating 'OG presence' and grey circles indicating 'OG absence'. The tree is labeled with 'Dscore = 0.925', 'Pvalue = NA', and 'AdjPvalue = NA'.

Legend:

- OG presence (Green circle)
- OG absence (Grey circle)
- Inferred root (Red diamond)

Dscore = 0.925  
Pvalue = NA  
AdjPvalue = NA