

Phylogenetic Tree (GlyA)

E. Griffiths and R.S. Gupta (2006) J Mol Evol. 63(2):283-296



Figure 3. A bootstrapped neighbor-joining distance tree based on SHMT (GlyA) homologs. The bootstrap scores that were >50 (out of 100) are indicated on various nodes. The chlamydiae and *Treponema* species were found to branch anomalously with the insert-containing SHMT homologs from *Streptomyces* and *Propionibacterium*, indicating lateral gene transfer between these groups.