A)
B)
Streptomyces LCA
Tree scale: 0.1
Assembly accession
Num complete genomes
AVG Chrom. protein
Complete genomes
AVG Num. Accessory proteins
BGC(s) completed genomes
Fragmented genomes
Fragmented BGCs
AVG chrom. protein per species (213 sp.)
Core proteins per species (61 sp.)
AVG Num of Accessory prot. per species (61 sp.)
Relaxed fingerprints per species (61 sp.)
Max smBGCs per species (213 sp.)
Intra-species smBGC AVG Jaccard dist (12 sp.)
Max CAZYmes per species (213 sp.)
Intra-species CAZYmes AVG Jaccard dist (12 sp.)
TTA-bearing proteins per species (213 sp.)
Intra-species AVG Jaccard distance (12 sp.) based on presence of TTA-bearing proteins in Orthogroups