Tree scale: 0.1

bootstrap

- 873
- 3154.75
- **5436.5**
- 7718.25
- 10000





StaphylococcusAureus YP 500978.1

LactobacillusRhamnosus WP 005686703.1

ListeriaMonocytogenes NP 466154.1

BacillusSubtilis NP 387998.1

NitrospiraDefluvii WP 213041942.1

GloeobacterViolaceus WP 011140090.1

CyanobiumGracile WP 015109401.1

TheobromaCacao XP 017984011.1

HelianthusAnnuus XP 021973724.1

MicromonasPusilla XP 003061099.1

ChlamydomonasReinhardtii XP 001697380.1

StreptomycesCoelicolor NP 628862.1

FusariumOxysporum XP 031036278.1

YarrowiaLipolytica XP 504189.1

SaccharomycesCerevisiae NP 013687.1

NitrosomonasEutropha WP 011633674.1

PseudomonasAeruginosa NP 252952.1

KlebsiellaPneumoniae YP 005229167.1

EscherichiaColi NP 417778.1

EnterobacterCloacae WP 000424395.1

ChlamydomonasReinhardtii XP 042918467.1

HydraVulgaris XP 047133073.1

DrosophilaMelanogaster NP 524939.1

CaenorhabditisElegans NP 505181.1

RhodobacterCapsulatus WP 013066045.1

HelianthusAnnuus XP 035830377.1

TheobromaCacao XP 007026182.2

HelianthusAnnuus XP 021975671.1

