

Phylogenetic tree showing relationships between various GCA sequences. The tree is rooted at the bottom left. Bootstrap values are shown at the nodes. Some sequences are highlighted in blue.

Sequences (from top to bottom):

- GCA_020784275.1
- GCA_009649105.1
- GCA_004793875.1
- GCA_009649565.1
- GCA_020784525.1
- GCA_000179435.1
- GCA_910578625.1**
- GCA_010206285.1
- GCA_002254245.1
- GCA_002254195.1
- GCA_002254025.1
- GCA_002253765.1
- GCA_002253875.1
- GCA_002253945.1
- GCA_002253905.1
- GCA_002254185.1
- GCA_002253965.1
- GCA_002253785.1
- GCA_002253975.1
- GCA_002254255.1

Root

Scale bar: 250