

Phylogenetic tree showing the relationships between various GCA sequences. The tree is rooted at the bottom left. Bootstrap values are indicated by blue numbers above the branches, and posterior probabilities are indicated by red numbers below the branches. The scale bar at the bottom left represents 250 units.

Sequences and their associated values (Bootstrap/Posterior Probability):

- GCA_002735985.1 (393)
- GCA_015560055.1 (222)
- GCA_002250405.1 (229)
- GCA_018372595.1 (287)
- GCA_013487885.1 (321)
- GCA_002079465.1 (336)
- GCA_008016915.1 (388)
- GCA_019593595.1 (388)
- GCA_002079545.1 (0.591)
- GCA_002079565.1 (2.41)
- GCA_003052845.1 (0.569)
- GCA_003052875.1 (1.78)
- GCA_000159395.1 (0.981)
- GCA_003061585.1 (0.99)
- GCA_001435955.1 (0.0117)
- GCA_002736025.1 (0.00977)
- GCA_003061545.1 (0.17)
- GCA_013249205.1 (0.17)
- GCA_015159735.1 (0.17)
- GCA_015560415.1** (0.17)

Root: 0

Scale bar: 250