

Phylogenetic tree showing the relationships between various GCA sequences. The tree is rooted at the bottom left. The scale bar indicates a distance of 250. The sequences are labeled with their GCA IDs. Bootstrap values are shown at the nodes. The sequence GCA_001435955.1 is highlighted in blue.

Sequences (from top to bottom):

- GCA_002735985.1
- GCA_015560055.1
- GCA_002250405.1
- GCA_018372595.1
- GCA_013487885.1
- GCA_002079465.1
- GCA_008016915.1
- GCA_019593595.1
- GCA_002079545.1
- GCA_002079565.1
- GCA_003052845.1
- GCA_003052875.1
- GCA_000159395.1
- GCA_003061585.1
- GCA_001435955.1**
- GCA_002736025.1
- GCA_003061545.1
- GCA_013249205.1
- GCA_015159735.1
- GCA_015560415.1

Root

250