



LEGEND

Branch Length

- x1 branch length
- x10 branch length
- x100 branch length

Nodes

- gene node
- speciation node
- duplication node
- ambiguous node
- gene split event

Genes

- Gene ID gene of interest
- Gene ID within-sp. paralog

Collapsed nodes

- collapsed sub-tree
- collapsed (this gene)
- collapsed (paralog)
- (x10 branch length)
- (x100 branch length)

Collapsed Alignments

- 0 - 33% aligned seq
- 33 - 66% aligned seq
- 66 - 100% aligned seq

Expanded Alignments

- gap
- aligned seq