Complete coding Allignment 2022.fasta
Sequencher(tm) "Untitled Project"

4213-1, 1 to 1,413

4213-32, 1 to 1,454

OX20-8, 1 to 1,413

PI105B, 1 to 1,454

WMS-82, 1 to 1,413

1   1,454

Diagram Key

☐ Hole in contig
▨ Single fragment
▦ Multiple fragments same direction
▬ Both strands
▬ Both strands plus
⊢—P————P←Start codon frame 1
⊢——P↲—Stop codon frame 2

Bumps on
fragments
show motifs,
hollow
rectangles
show features