

logo

```
NTDB id 32 MW RS07995 WP 000472256.1
NTDB id 16 SA RS07765 WP 000472256.1
NTDB id 1063381 ACGLYC RS11990 WP 003153087.1
NTDB id 98 BSU 24710 NP 390351.1
consensus
```

[illegible]

logo

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Sequence logo for the 103 bp region. The logo shows the conservation of nucleotides at each position. The top part shows the sequence KKAQKT CPSKTIIT SNGEVWNLAKAEQ. The bottom part shows the sequence KEAQKTCXSGETITISNGEAVAN. The conservation is highest at positions 1-10 and 18-20. The sequence is aligned with the reference sequence KEAQKTCXSGETITISNGEAVAN. The conservation is highest at positions 1-10 and 18-20.

-  non conserved
-  similar
-  $\geq 50\%$ conserved