

- 1 - piece 1, NC 000913, qor dnaB+, config: linear, direction: +, begin: 4262225, end: 4262356

Sequence alignment diagram showing the amino acid sequence of a protein across 12 different species. The sequence is color-coded by residue type: hydrophobic (black), polar (green), acidic (red), basic (blue), and cysteine (orange). Amino acids are aligned horizontally, with gaps indicated by dashes. Red dots above the sequence indicate positions where the amino acid is conserved across all species. A red box highlights a specific region in the C-terminal half of the sequence.

----- 1 NC 000913.com

p35 6.1 bits

{-----} sd-(11)-ir 4262291 Gap 3.0 bits

|-----| sd-ir 4262291 qor\_dnaB+ total 5.1 bits

புது புது p10 3.8 bits

62271 Gap 2.3 bits  
total 7.6 bits

5' \*4262310 \* \*4262320 \* \*4262330 \* \*4262340 \* \*4262350 \* 3'

ccccattctttatctcggttaactccattcactatggcaggaaataaacccctt

- pro - ile - leu - tyr - leu - gly - asn - ser - ile - his - tyr - gly - arg - lys - - - fMet - ala - gly - asn - lys - pro -

[##> orf 20 codons

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