

logo

NTDB id 422875 G3T50 RS07280 WP 002880445.1
NTDB id 1231 CJJ81176 RS07070 WP 002855438.1
consensus

MCFSTLFAQDNFEEYFKLIDKENR**ID**FNTLQNPFVNPTFEKLRHIKITAIMLDKVKIYDRWYKKGDMIDDALH**E**INTKE
MCFSTLFAQDNFEEYFKLIDKENR**IN**FNTLQNPFVNPTFEKLRHIKITAIMLDKVKIYDRWYKKGDMIDDAL**I*******YEINTKE

80
80

logo

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Sequence logo for the motif IKFQYDNL EIAIQINKNDKININ. The logo shows the conservation of each amino acid position. The sequence is displayed in two rows: IKFQYDNL EIAIQINKNDKININ and IKFQYDNL EIAIQINKNDKININ. Below the sequence, a red bar indicates the conservation score for each position.

103
103

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