

logo

NTDB id 249653 CQJ38 RS00190 WP 198404711.1
NTDB id 80 BSU 00370 NP 387918.1
consensus

MLMKSTGIVRKVDELGRVVIPIELRRTLGLIAEKDALEIYVDDEKIILKKYKPNMTCQVTGEVSDDNLKLAGGKLVLSKEG
MF

80
80

logo

NTDB id 249653 CQJ38 RS00190 WP 198404711.1
NTDB id 80 BSU 00370 NP 387918.1
consensus

Sequence logo for the 12th position. The top row shows amino acids: A, E, Q, I, S, E, I, Q, N, H, L, Q, N, Q, K. The bottom row shows their frequencies: A, E, Q, I, S, E, I, Q, N, Q, L, Q, N, L, K. The 10th position (H) is highlighted with a red asterisk below it.

96
96

96

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