

logo

NTDB id 289399 DDG67 RS05165 WP 000287265.1
NTDB id 39 MW RS04875 WP 000287265.1
NTDB id 23 SA RS04995 WP 000287265.1
consensus

80
80
80

logo

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NTDB id 39 MW RS04875 WP 000287265.1
NTDB id 23 SA RS04995 WP 000287265.1
consensus

LTPLFPTYFFPTHSDRQNENIWLNMHYIESIKELKNRKCKVTFINNESILHVSYHSLWHQYNNSIFYYYMVDKQSRMIS

LTPLFPTYFFPTHSDRQNENIWLNMHYIESIKELKNRKCKVTFINNESILHVSYHSLWHQYNNSIFYYYMVDKQSRMIS

LTPLFPTYFFPTHSDRQNENIWLNMHYIESIKELKNRKCKVTFINNESILHVSYHSLWHQYNNSIFYYYMVDKQSRMIS

LTPLFPTYFFPTHSDRQNENIWLNMHYIESIKELKNRKCKVTFINNESILHVSYHSLWHQYNNSIFYYYMVDKQSRMIS

160
160
160

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Sequence logo for the 16-mer motif KNPDQPIDYNKATLNVFEALTRYSLFEDK. The logo shows the conservation of each amino acid at each position. The sequence is repeated four times below the logo.

189
189
189

- X non conserved
- X similar
- X > 50% conserved