```
logo
NTDB id 1398 DSB67 RS00645 WP 029388870.1     ..................................................MLSHQWQNIMHRVLSSQCGLCRFPIPHRQT    30
NTDB id 343825 EPB61 RS12815 WP 055465863.1  ..................................................MLTNRFQKTMFSWLAPACQLCGLALHA...    27
NTDB id 1154 A1552VC RS20525 WP 001884047.1  MPQPCTNTRSCPCPLTCQYNAMVTVSPSISIFIHSANWQDEAHYPDSWLAMLTHWLRKTTAPLLTPECHLCRLALDT...    77
consensus                                                                        !!** *****  *!** ! !!****

logo
NTDB id 1398 DSB67 RS00645 WP 029388870.1     PNLMRWCDSCFDHLTPIERCMRCGLKMTEEESQAATECGECLKEPPPWRRLYTLGDYDFPLSQQVQRFKDHGEVWHVGAL   110
NTDB id 343825 EPB61 RS12815 WP 055465863.1  PTKPRLCAACLAWYAPTPRCARCGLPCES....ATAQCGQCLSTPPPWHQLVCVGDYRFPLSHSVHQLKYQRQFWQARPL   103
NTDB id 1154 A1552VC RS20525 WP 001884047.1  NSPFGVCSACQAWLEHGYRCARCGLPTLT....PVDQCGQCLGQPPPWRKLMCVGDYRFPLSDAVHQLKYQRQFWQAPRL   153
consensus                                    **  **! *! *** *  !!*!!!!*  *    *  *!!*!!  !!!!**!***!!!*!!!!  !***!*****!**  !

logo
NTDB id 1398 DSB67 RS00645 WP 029388870.1     TQLLAERVEQPAPVITSVPVHWQRLLKRGFNQSHILAKHLAHHLDVRYENKVFKRVKNVQSQRGNKKASREQNLQGAFAL   190
NTDB id 343825 EPB61 RS12815 WP 055465863.1  AALLAPRITTPAPLLTCVPLHWRRQLRRGFNQSDVLARQLAQHLGCEYDPQLFSRPHATVPQHGLDKAQRRNLNRAFVL   183
NTDB id 1154 A1552VC RS20525 WP 001884047.1  AKLLATQINEPAPLLCSVPLHWRRRWQRGFNQSDLLARELANMLNVEYDHQLFARRRATPHQQGLSKAQRIHNLRDAFVL   233
consensus                                    * !!! **  !!!****!!*!!*! **!!!!!!*!!* !!**! **!* **! ! ***  ! !* !!*!  !!* !!*!

logo
NTDB id 1398 DSB67 RS00645 WP 029388870.1     RGEVDFSHVAIVDDVVTTGSTVRQLCHLLLEVGVESIDIYCICRTPAPGSL..   241
NTDB id 343825 EPB61 RS12815 WP 055465863.1  NHEPCQQHVAIVDDVVTTGSTVRQLCNLLLEVGVKSLDIYCICRTPEPSD...   233
NTDB id 1154 A1552VC RS20525 WP 001884047.1  NHPPNQPHVAIVDDVVTTGSTIRHLCDLLLDVGVQSIDIYCICRTPEPKDSHG   286
consensus                                    **** * !!!!!!!!!!!!!!*!*!! !!!*!!! !*!!!!!!!!!*! *
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X non conserved

X similar

X ≥ 50% conserved