```
logo
NTDB id 1154 A1552VC RS20525 WP 001884047.1  MPQPCTNTRSCPCPLTCQYNAMVTVSPSISIFIHSANWQDEAHYPDSWLAMLTHWLRKTTAPLLTPECHLCRLALDTN..   78
NTDB id 407708 GPY39 RS15155 WP 021448951.1 ..................................................MLSHHWQNIMHRVLSSQCGLCRFPIQATAQ   30
NTDB id 1398 DSB67 RS00645 WP 029388870.1    ..................................................MLSHQWQNIMHRVLSSQCGLCRFPIPHRQT   30
consensus                                                                                     !!*! ********!***!*!!!***

logo
NTDB id 1154 A1552VC RS20525 WP 001884047.1  .SPFGVCSACQAWLEHGYRCARCGLPTLT....PVDQCGQCLGQPPPWRKLMCVGDYRFPLSDAVHQLKYQRQFWQAPRL  153
NTDB id 407708 GPY39 RS15155 WP 021448951.1 PNALRWCDHCYQYLTPVKRCQRCGLSLKAEEANIESICGECLSEPPPWQRLFTLGDYDFPLSREVQRFKDHGQIWHVRAL  110
NTDB id 1398 DSB67 RS00645 WP 029388870.1   PNLMRWCDSCFDHLTPIERCMRCGLKMTEEESQAATECGECLKEPPPWRRLYTLGDYDFPLSQQVQRFKDHGEVWHVGAL  110
consensus                                   ** ***!* !* *!*** !! !!!! *  ** *  * !!*!! *!!!!**!***!!!*!!!!  !***!****!** *!

logo
NTDB id 1154 A1552VC RS20525 WP 001884047.1  AKLLATQINEPAPLLCSVPLHWRRRWQRGFNQSDLLARELANMLNVEYDHQLFARRRATPHQQGLSKAQRIHNLRDAFVL  233
NTDB id 407708 GPY39 RS15155 WP 021448951.1 TQLLAQRISTPAPLITTVPLHWQRYLYRGFNQSDILARHLAGHLNVRFDNHVFRRVKHAQSQRGYKKSSREQNLKGAFTL  190
NTDB id 1398 DSB67 RS00645 WP 029388870.1   TQLLAERVEQPAPVITSVPVHWQRLLKRGFNQSHILAKHLAHHLDVRYENKVFKRVKNVQSQRGNKKASREQNLQGAFAL  190
consensus                                   **!!! **  !!!****!!*!!*! * !!!!!!**!!**!! *!*!**** *!*!**  **!*! *!**!**!!**!! !

logo
NTDB id 1154 A1552VC RS20525 WP 001884047.1  NHPPNQPHVAIVDDVVTTGSTIRHLCDLLLDVGVQSIDIYCICRTPEPKDSHG   286
NTDB id 407708 GPY39 RS15155 WP 021448951.1 NQPPKYNHVAIVDDVVTTGSTVRQLCHLLLEVGVETVDIYCICRTPAPGAV..   241
NTDB id 1398 DSB67 RS00645 WP 029388870.1   RGEVDFSHVAIVDDVVTTGSTVRQLCHLLLEVGVESIDIYCICRTPAPGSL..   241
consensus                                   * ** * !!!!!!!!!!!!!!!*!*!!*!!!*!!!***!!!!!!!!!*!* *
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X non conserved

X similar

X ≥ 50% conserved