```
logo
NTDB id 1154 A1552VC RS20525 WP 001884047.1  MPQPCTNTRSCPCPLTCQYNAMVTVSPSISIFIHSANWQDEAHYPDSWLAMLTHWLRKTTAPLLTPECHLCRLALDT...   77
NTDB id 407766 GPY29 RS02585 WP 031781654.1  ..................................................MLSHHWQNIMHRVLSSQCGLCRFPILAAAQ   30
NTDB id 1398 DSB67 RS00645 WP 029388870.1    ..................................................MLSHQWQNIMHRVLSSQCGLCRFPIPHRQT   30
consensus                                                                                     !!*! ********!***!*!!!***

logo
NTDB id 1154 A1552VC RS20525 WP 001884047.1  NSPFGVCSACQAWLEHGYRCARCGLPTLT....PVDQCGQCLGQPPPWRKLMCVGDYRFPLSDAVHQLKYQRQFWQAPRL  153
NTDB id 407766 GPY29 RS02585 WP 031781654.1  PNALRWCDHCYQYLTPVKRCQRCGLSLKAEEANIESICGECLSEPPPWQRLFTLGDYDFPLSREVQRFKDHGQTWHVRAL  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 407766 GPY29 RS02585 WP 031781654.1  TQLLAQRISTPAPLITTVPLHWQRYLYRGFNQSDILARHLAGHLNVRFDNHVFRRVKHAQSQRGNKKSSREQNLKGAFTL  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 407766 GPY29 RS02585 WP 031781654.1  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