```
logo
NTDB id 1154 A1552VC RS20525 WP 001884047.1  MPQPCTNTRSCPCPLTCQYNAMVTVSPSISIFIHSANWQDEAHYPDSWLAMLTHWLRKTTAPLLTPECHLCRLALDTNS.   79
NTDB id 407452 GPY59 RS09680 WP 158173560.1 ..................................................MLSHHWQNIMHRVLSSQCGLCRFPIQTPSH   30
NTDB id 1398 DSB67 RS00645 WP 029388870.1    ..................................................MLSHQWQNIMHRVLSSQCGLCRFPIPHRQT   30
consensus                                                                                     !!*! ********!***!*!!!*** * *

logo
NTDB id 1154 A1552VC RS20525 WP 001884047.1  ..PFGVCSACQAWLEHGYRCARCGLPTL....TPVDQCGQCLGQPPPWRKLMCVGDYRFPLSDAVHQLKYQRQFWQAPRL  153
NTDB id 407452 GPY59 RS09680 WP 158173560.1 TNTLRWCEHCLKYLTPVKRCQRCGLSLQKEEMSTDSICGECLSKPPPWQRLYTLGDYDFPLSREVQRFKDHGESWHVKAL  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 407452 GPY59 RS09680 WP 158173560.1 TEQLAQRITTPAPIITSVPLHWQRYLRRGFNQSDVLARQLAKQLQTNFDAKVFRRVRLAQSQRGNTKTSREQNLKGAFIL  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 407452 GPY59 RS09680 WP 158173560.1 NKRPHSSHVAIVDDVVTTGSTVRQLCHLLLEVGVESIDIYCICRTPAPRSL..   241
NTDB id 1398 DSB67 RS00645 WP 029388870.1   RGEVDFSHVAIVDDVVTTGSTVRQLCHLLLEVGVESIDIYCICRTPAPGSL..   241
consensus                                    *  *  *!!!!!!!!!!!!!!*!*!!*!!!*!!!*!!!!!!!!!!!*!***
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X non conserved

X similar

X ≥ 50% conserved