```
logo
NTDB id 1014 ACIAD RS13855 WP 004924483.1    MSIRHFFTFFWATLLTLFS...MQIVFAQGFDQQYDAWKTQQKAHDTRLNSSASHHSLSKPTQQSGTGVKVNINSATAKE   77
NTDB id 1440 BW25113 RS02290 WP 000680312.1  ..MKHGIKALLITLSLACAGMSHSALAAASVAK...PTAVETKAEAPAAQSK.AAV...PAKASDEEGTRVSINNASAEE   71
NTDB id 1294 VP RS04495 WP 021449594.1       .........MKWIL.TLCLVVFAPLSLAAD..T.........................KADKYEGIEITVNINTASAEE   42
NTDB id 1159 ABDM36 RS04765 WP 001166105.1   ..MQIKTKIVTLFL.SLCLPTLPLLANAEETAP.........................AAQVEEGIVITVNINTASAEE   51
NTDB id 1150 A1552VC RS08580 WP 001166105.1  ..MQIKTKIVTLFL.SLCLPTLPLLANAEETAP.........................AAQVEEGIVITVNINTASAEE   51
NTDB id 1382 A4U84 RS07375 WP 062924234.1    .....MKKLSRYTFSLLFG.LFSATTFAEEKVT...TST.EH...TPVVMQQ.......VAATQKVNANVVNINTATASE   60
NTDB id 1459 ASU2 RS10590 WP 015674357.1     .....MKTPSKLGLSLLIG.MLSTGVLAAEKTK...PSVTEP...TPVVQNAQATSPEHSHAEMTNNANLVNINIATAAE   68
NTDB id 26559 NT05HA RS10920 WP 005701687.1  ..MKAMKHLL..AAAFVFGSVFAGSAVAAEKTA.............EPVQLE.......QPAVQNSVGDKVNINTATAAE   56
NTDB id 1362 HI 1008 AAC22669.1              .....MKTLF..TSVVLCGALVVSSSFAEEKAT...XQTAQS...VVTTQAE.......AQVAPAVVSDKLNINTATASE   60
consensus                                      *     **   *  *** *      ! *                 *         *     **  **!!*!*! !
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logo
NTDB id 1014 ACIAD RS13855 WP 004924483.1    LQQ.LHGVGEKKALAIVEYRQQYGKFASVDDLQKVKGIGPKFIEKNRDRLAI.   128
NTDB id 1440 BW25113 RS02290 WP 000680312.1  LARAMNGVGLKKAQAIVSYREEYGPFKTVEDLKQVPGMGNSLVERNLAVLTL.   123
NTDB id 1294 VP RS04495 WP 021449594.1       IATMLKGIGEKKAQSIVDYRTEHGPFKTAADLTNVKGIGEATIKKNEDRILL.    94
NTDB id 1159 ABDM36 RS04765 WP 001166105.1   LATLLKGIGLKKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL.   103
NTDB id 1150 A1552VC RS08580 WP 001166105.1  LATLLKGIGLKKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL.   103
NTDB id 1382 A4U84 RS07375 WP 062924234.1    IQDKLVGIGAKKAQAIVEYREKNGKFLNVEQLTEVSGIGKATLEKNRDCIVIE   113
NTDB id 1459 ASU2 RS10590 WP 015674357.1     IQDKLVGIGAKKAQAIVEYRTKNGAFSSLEQLTDVSGIGKATLDKNRDRILLQ   121
NTDB id 26559 NT05HA RS10920 WP 005701687.1  IQKALVGIGAKKAEAVVQYREKHGNFSAVEQLLEVQGIGKATLEKNRDRIAL.   108
NTDB id 1362 HI 1008 AAC22669.1              IQKSLTGIGAKKAEAIVQYREKHGNFXNAEQLLEVQGIGKATLEKNRDRIIF.   112
consensus                                    **  * !*! !!!***!*!!*  ! !  ***!* ! !*! *****!******
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X non conserved
X similar
X ≥ 50% conserved