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logo

NTDB id 1014 ACIAD RS13855 WP 004924483.1     MSI.RHFFTFFWATLL.TLFSMQIVFAQGFDQQYDAWKTQQKAHDTRLNSSASHHSLSKPTQQSGTGVKVNINSATAKEL   78
NTDB id 1294 VP RS04495 WP 021449594.1        .......MK.WILTLCLVVFAPL.....SLAAD...........TK.............ADKYEGIEITVNINTASAEEI   43
NTDB id 1159 ABDM36 RS04765 WP 001166105.1    MQIKTKIVT.LFLSLCLPTLPLL.....ANAEET.........APA.............AQVEEGIVITVNINTASAEEL   52
NTDB id 1150 A1552VC RS08580 WP 001166105.1   MQIKTKIVT.LFLSLCLPTLPLL.....ANAEET.........APA.............AQVEEGIVITVNINTASAEEL   52
NTDB id 1382 A4U84 RS07375 WP 062924234.1     ...MKKLSRYTFSLLF.GLFSAT.....TFAEEK.........VTTSTEHTPVVM.QQVAATQKVNANVVNINTATASEI   61
NTDB id 1121600 EL144 RS05035 WP 005705031.1 MKAMKHLL..ATAFVFGSVFAGS.....AVAAEK.........TA.......EPVQLEQPAVQNSVGDKVNINTATAAEI   57
NTDB id 1362 HI 1008 AAC22669.1               ...MKTLF..TSVVLCGALVVSS.....SFAEEK.........ATXQTAQSVVTTQAEAQVAPAVVSDKLNINTATASEI   61
consensus                                     *   * **   *****  **        * ***                        *    ** * *!!!*!*! !*

logo

NTDB id 1014 ACIAD RS13855 WP 004924483.1     QQ.LHGVGEKKALAIVEYRGQYGKFASVDDLQKVKGIGPKFIEKNRDRLAI.   128
NTDB id 1294 VP RS04495 WP 021449594.1        ATMLKGIGEKKAQSIVDYRTEHGPFKTAADLTNVKGIGEATIKKNEDRILL.    94
NTDB id 1159 ABDM36 RS04765 WP 001166105.1    ATLLKGIGLKKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL.   103
NTDB id 1150 A1552VC RS08580 WP 001166105.1   ATLLKGIGLKKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL.   103
NTDB id 1382 A4U84 RS07375 WP 062924234.1     QDKLVGIGAKKAQAIVEYREKNGKFLNVEQLTEVSGIGKATLEKNRDCIVIE   113
NTDB id 1121600 EL144 RS05035 WP 005705031.1 QKALVGIGAKKAEAVVQYREKHGNFSAVEQLLEVQGIGKATLEKNRDRIAL.   108
NTDB id 1362 HI 1008 AAC22669.1               QKSLTGIGAKKAEAIVQYREKHGNFXNAEQLLEVQGIGKATLEKNRDRIIF.   112
consensus                                     *  ! !*! !!!***!*!!*  ! !  ***!* !*!!! *****!*****
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X non conserved

X similar

X ≥50% conserved