```
logo

NTDB id 1014 ACIAD RS13855 WP 004924483.1      MSIRHFFTFFWATLLTLFSMQIVFAQGFDQQYDAWKTQQKAHDTRLNSSASHHSLSKPTQQSGTGVKVNINSATAKELQQ   80
NTDB id 1362 HI 1008 AAC22669.1                .....MKTLFTSVVLCGALV...VSS.......SFAEEKATXQT..AQSVVTTQAEAQVAPAVVSDKLNINTATASEIQK   63
NTDB id 1104970 ACN08P RS11445 WP 419237952.1  .MY.SYRLFKVIALASGLAL...VPL.......S.....HADE..............NT.HEGIVKTVNINEANVEQLDI   48
NTDB id 1159 ABDM36 RS04765 WP 001166105.1     MQI.KTKIVTLFLSLCLPTL...PLL.......ANAEETAPAA..............QV.EEGIVITVNINTASAEELAT   54
NTDB id 1150 A1552VC RS08580 WP 001166105.1    MQI.KTKIVTLFLSLCLPTL...PLL.......ANAEETAPAA..............QV.EEGIVITVNINTASAEELAT   54
NTDB id 1395 DSB67 RS04445 WP 010649155.1      ........MKWMLTLCLLIL...APM.......SWAE.TKTKA..............DK.YEGIEITVNVNSATAQEIAT   46
NTDB id 1294 VP RS04495 WP 021449594.1         ........MKWILTLCLVVF...APL.......SLAA..DTKA..............DK.YEGIEITVNINTASAEEIAT   45
consensus                                            * *    *****  *      *        * **     *               *    *** ***!*!*!*******

logo

NTDB id 1014 ACIAD RS13855 WP 004924483.1      .LHGVGEKKALAIVEYRQQYGKFASVDDLQKVKGIGPKFIEKNRDRLAI  128
NTDB id 1362 HI 1008 AAC22669.1                SLTGIGAKKAEAIVQYREKHGNFXNAEQLLEVQGIGKATLEKNRDRIIF  112
NTDB id 1104970 ACN08P RS11445 WP 419237952.1  LLDGIGAEKARAIVDYRNANGKFASADDLVKVKGIGKATVEKNRKIIEL   97
NTDB id 1159 ABDM36 RS04765 WP 001166105.1     LLKGIGLKKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL  103
NTDB id 1150 A1552VC RS08580 WP 001166105.1    LLKGIGLKKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL  103
NTDB id 1395 DSB67 RS04445 WP 010649155.1      LLIGIGEKKAQDIVEYRNEHGPFKTAADLTKVKGIGEATVRKNEDRILL   95
NTDB id 1294 VP RS04495 WP 021449594.1         MLKGIGEKKAQSIVDYRTEHGPFKTAADLTNVKGIGEATIKKNEDRILL   94
consensus                                      *! !*! *!!**!!*!!   !*! *****!* !*!!!******! *****
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X non conserved

X similar

X $\geq$ 50% conserved