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logo

NTDB id 1014 ACIAD RS13855 WP 004924483.1     MSIRHFFTFFWATLLTLF.......SMQIVFAQGFDQQYDAWKTQQKAHDTRLNSSASHHSLSKPT....QQSGTGVKVN    69
NTDB id 1041 HON27 RS14740 WP 005124713.1     ...MQYSMRFWKNKYFLLIMLCLGVISPSVCAQSFDQNFQEWKAKQQMYDQKLNITKSSHSNGSKISQTKNFNDSTGQIH    77
NTDB id 91706 OCV19 RS11615 WP 048661473.1    .........MRTIYSTLLLS.FLMLLSPAV.FADSPTKAE..............................LYDGIEITVN    39
NTDB id 1159 ABDM36 RS04765 WP 001166105.1    .......MQIKTKIVTLFLSLCLPTLPLLANAEETAPAAQ..............................VEEGIVITVN    43
NTDB id 1150 A1552VC RS08580 WP 001166105.1   .......MQIKTKIVTLFLSLCLPTLPLLANAEETAPAAQ..............................VEEGIVITVN    43
NTDB id 1395 DSB67 RS04445 WP 010649155.1     ..............MKWMLTLCLLILAPMSWAE.TKTKAD..............................KYEGIEITVN    35
NTDB id 1294 VP RS04495 WP 021449594.1        ..............MKWILTLCLVVFAPLSLAA..DTKAD..............................KYEGIEITVN    34
NTDB id 1362 HI 1008 AAC22669.1               .........MKTL..FTSVVLCGALVVSSSFAEEKATXQTAQS.................VVTTQAEAQVAPAVVSDKLN    52
NTDB id 1382 A4U84 RS07375 WP 062924234.1     .........MKKLSRYTFSLLFG.LFSATTFAEEKVTTSTEHT.................PVVMQQ.VAATQKVNANVVN    52
consensus                                               **   * * ***** **  *  ***  * *                                  *** ****

logo

NTDB id 1014 ACIAD RS13855 WP 004924483.1     INSATAKELQQ.LHGVGEKKALAIVEYRQQYGKFASVDDLQKVKGIGPKFIEKNRDRLAI.   128
NTDB id 1041 HON27 RS14740 WP 005124713.1     LNQANVNEFQQ.LKGVGEKKAQAIVEYRQKNGSFKNIDEIKNVKGIGPAIFEKNKSRLAL.   136
NTDB id 91706 OCV19 RS11615 WP 048661473.1    INTATAEELSALLVGVGDKKAQEIVDYREQNGAFTTADGLASVKGIGEATVKKNRERIQL.    99
NTDB id 1159 ABDM36 RS04765 WP 001166105.1    INTASAEELATLLKGIGLKKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL.   103
NTDB id 1150 A1552VC RS08580 WP 001166105.1   INTASAEELATLLKGIGLKKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL.   103
NTDB id 1395 DSB67 RS04445 WP 010649155.1     VNSATAQEIATLLIGIGEKKAQDIVEYRNEHGPFKTAADLTKVKGIGEATVRKNEDRILL.    95
NTDB id 1294 VP RS04495 WP 021449594.1        INTASAEEIATMLKGIGEKKAQSIVDYRTEHGPFKTAADLTNVKGIGEATIKKNEDRILL.    94
NTDB id 1362 HI 1008 AAC22669.1               INTATASEIQKSLTGIGAKKAEAIVQYREKHGNFXNAEQLLEVQGIGKATLEKNRDRIIF.   112
NTDB id 1382 A4U84 RS07375 WP 062924234.1     INTATASEIQDKLVGIGAKKAQAIVEYREKNGKFLNVEQLTEVSGIGKATLEKNRDCIVIE   113
consensus                                     *!*!** !*  *! !*!*!!!**!!*!!* *! !  ***** !*!!!******!******
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X non conserved

X similar

X $\geq$ 50% conserved