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logo

NTDB id 1041 HON27 RS14740 WP 005124713.1    MQYSMRFWKNKYFLLIMLCLGVISPSVCAQSFDQNFQEWKAKQQMYDQKLNITKSSHSNGSKISQTKNFNDSTGQIHLNQ   80
NTDB id 1159 ABDM36 RS04765 WP 001166105.1   ....MQIKTKIVTLFLSLCLPTLPLLANAE...ETA....PAA.......................QVEEGIVITVNINT   46
NTDB id 1150 A1552VC RS08580 WP 001166105.1  ....MQIKTKIVTLFLSLCLPTLPLLANAE...ETA....PAA.......................QVEEGIVITVNINT   46
NTDB id 321748 D0812 RS08735 WP 020196004.1  ...........MKWMLTLCLLLLAPMSWAEEGKSVT....TKA.......................AKYEGIEITVNVNT   42
NTDB id 1395 DSB67 RS04445 WP 010649155.1    ...........MKWMLTLCLLILAPMSWAETK........TKA.......................DKYEGIEITVNVNS   38
NTDB id 1294 VP RS04495 WP 021449594.1       ...........MKWILTLCLVVFAPLSLAAD.........TKA.......................DKYEGIEITVNINT   37
NTDB id 1362 HI 1008 AAC22669.1              ......MKTL..FTSVVLCGALVVSSSFAEEKATXQ....TAQS....VVTTQAEAQ.........VAPAVVSDKLNINT   55
NTDB id 1382 A4U84 RS07375 WP 062924234.1    ......MKKLSRYTFSLLFG.LFSATTFAEEKVTTS....TEHT....PVVMQQ.VA.........ATQKVNANVVNINT   55
consensus                                              *   **!** **  **  !*          * *                      *** *****!*
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logo

NTDB id 1041 HON27 RS14740 WP 005124713.1    ANVNEF.QQLKGVGEKKAQAIVEYRQKNGSFKNIDEIKNVKGIGPAIFEKNKSRLAL.   136
NTDB id 1159 ABDM36 RS04765 WP 001166105.1   ASAEELATLLKGIGLKKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL.   103
NTDB id 1150 A1552VC RS08580 WP 001166105.1  ASAEELATLLKGIGLKKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL.   103
NTDB id 321748 D0812 RS08735 WP 020196004.1  ASAEEIATLLNGIGTKKAQDIVEYRNEHGPFKTAADLSKVKGIGEATVKKNQDRILL.    99
NTDB id 1395 DSB67 RS04445 WP 010649155.1    ATAQEIATLLIGIGEKKAQDIVEYRNEHGPFKTAADLTKVKGIGEATVRKNEDRILL.    95
NTDB id 1294 VP RS04495 WP 021449594.1       ASAEEIATMLKGIGEKKAQSIVDYRTEHGPFKTAADLTNVKGIGEATIKKNEDRILL.    94
NTDB id 1362 HI 1008 AAC22669.1              ATASEIQKSLTGIGAKKAEAIVQYREKHGNFXNAEQLLEVQGIGKATLEKNRDRIIF.   112
NTDB id 1382 A4U84 RS07375 WP 062924234.1    ATASEIQDKLVGIGAKKAQAIVEYREKNGKFLNVEQLTEVSGIGKATLEKNRDCIVIE   113
consensus                                    !** !****! !*! !!!**!!*!!   !*!   **** !*!!!*!****! *****
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X non conserved

X similar

X ≥ 50% conserved