logo

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NTDB id 1041 H0N27 RS14740 WP 005124713.1    MQYSMRFWKNKYFLLIMLCLGVISPSVCAQSFDQNFQEWKAKQQMYDQKLNITKSSHSNGSKISQTKNFNDSTGQIHLNQ   80
NTDB id 1159 ABDM36 RS04765 WP 001166105.1   MQIK......TKIVTL......FLSL..................CLPT.LPLLANAEETA.PAAQ..VEEGIVITVNINT   46
NTDB id 1150 A1552VC RS08580 WP 001166105.1  MQIK......TKIVTL......FLSL..................CLPT.LPLLANAEETA.PAAQ..VEEGIVITVNINT   46
NTDB id 1362 HI 1008 AAC22669.1              MKTL........FTSVVLCGALVVSSSFA........EEKATXQTAQS.V.VTT.......QAEAQVAPAVVSDKLNINT   55
NTDB id 78841 ASU1 RS10645 WP 015674357.1    MKTP......SKLGLSLLIG.MLSTGVLA........AEKTKPSVTEP.TPVVQNAQATS.PEHSHAEMTNNANLVNINI   63
NTDB id 1459 ASU2 RS10590 WP 015674357.1     MKTP......SKLGLSLLIG.MLSTGVLA........AEKTKPSVTEP.TPVVQNAQATS.PEHSHAEMTNNANLVNINI   63
NTDB id 1382 A4U84 RS07375 WP 062924234.1    MKKL......SRYTFSLLFG.LFSATTFA........EEKVTTST.EH.TPVVMQQ........VAATQKVNANVVNINT   55
consensus                                    !*          ***  ***  *  ****    *          **       *   ****  **   *   *              ***!*
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logo

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NTDB id 1041 H0N27 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 1362 HI 1008 AAC22669.1              ATASEIQKSLTGIGAKKAEAIVQYREKHGNFXNAEQLLEVQGIGKATLEKNRDRIIF.   112
NTDB id 78841 ASU1 RS10645 WP 015674357.1    ATAAEIQDKLVGIGAKKAQAIVEYRTKNGAFSSLEQLTDVSGIGKATLDKNRDRILLQ   121
NTDB id 1459 ASU2 RS10590 WP 015674357.1     ATAAEIQDKLVGIGAKKAQAIVEYRTKNGAFSSLEQLTDVSGIGKATLDKNRDRILLQ   121
NTDB id 1382 A4U84 RS07375 WP 062924234.1    ATASEIQDKLVGIGAKKAQAIVEYREKNGKFLNVEQLTEVSGIGKATLEKNRDCIVIE   113
consensus                                    !**  !**   !  !*!*!!!*!!!*!!***!  !* *******!  !!!*!*!****!******
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X non conserved

X similar

X ≥ 50% conserved