```
logo
NTDB id 1112102 BN940 RS16575 WP 043684826.1  MFSFFKRKKAPAPQEAAPGRDEALGAAASPDAPEVAPVAPTELPDTWPADAGGLPET..GADAPAHEAPAAAA....EPV   74
NTDB id 1118 NGFG RS11455 WP 003696286.1      MFSFFRRKKKQETPALEEAQVQETAAK...VESEVAQIV.....GNIKEDVESLAESVKGRAESAVETVSGAVEQVKETV   72
consensus                                     !!!!!*!!! * **   **     *! ***  *!!!** *****   * !   !*!***!   *!*! * *! ****!*!

logo
NTDB id 1112102 BN940 RS16575 WP 043684826.1  SEFVSGAGPAPAAAPVQPPDAWSDRPDETADPLVRLDRAASAEAAVPASSGSWLGRLRRGLSRT....GQSLGGLFTGVK   150
NTDB id 1118 NGFG RS11455 WP 003696286.1      AEMPSEAGEAAER..VES...AKEAVAETVGE..AVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQ   145
consensus                                      ! *! !!*!*  **! ****  * * !!  *** *     ! !  * *   ! *!!* !! ******* !!*!*! !

logo
NTDB id 1112102 BN940 RS16575 WP 043684826.1  VDEALFEELETALIMADAGVEATEKLLEQLRARVRRQRIERPEAVRAALRDILAEHLKVLEKPFVVGE.AAPMVVMIAGV   229
NTDB id 1118 NGFG RS11455 WP 003696286.1      IGEDLYEELETVLITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGI   225
consensus                                     * ! !*!!!!! !! *! !*!!!! !*  *!*!!    *  *  *!*!!** ! ***!*!!!! !**!*  ! !*!*!!*

logo
NTDB id 1112102 BN940 RS16575 WP 043684826.1  NGAGKTTSIGKLAHALRERQASVLLAAGDTFRAAAREQLVAWGARNGVTVIAQEGGDPAAVAFDAVNAGRARSADVVLVD   309
NTDB id 1118 NGFG RS11455 WP 003696286.1      NGAGKTTSIGKLAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLAD   305
consensus                                     !!!!!!!!!!!!!*       !!!!!!!!!!!!!!!!!! !!!*!! !!!! !   !!*!!!*!!!!*!**!!   !*!! !

logo
NTDB id 1112102 BN940 RS16575 WP 043684826.1  TAGRLPTQLHLMEELKKIKRVIGKADPAAPHEILLVVDGNTGQNALAQIRAFDAALGLTGLVVTKLDGTAKGGTLAAVAA   389
NTDB id 1118 NGFG RS11455 WP 003696286.1      TAGRLPTQLHLMEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALA.   384
consensus                                     !!!!!!!!!!!!!!*!!*!!!* !! !*!!!!!**!*!*! !!!!* !**!!! !!!!!!!*!!!!!!!!!!! !!!*!*

logo
NTDB id 1112102 BN940 RS16575 WP 043684826.1  GSQGVRPVPVYWIGVGEGMQDLQPFVAAEFADALLGGPETDRA   432
NTDB id 1118 NGFG RS11455 WP 003696286.1      ...SDRPVPVRYIGVGEGIDDLRPFDARAFVDALLD.......   417
consensus                                     ***  !!!!! *!!!!!!* !! !! !  ! !!!! *******
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X non conserved

X similar

X ≥50% conserved