```
logo
NTDB id 381347 FYA87 RS13880 WP 005015354.1  MFSFFKKKTPPPAAQASPAPPPEPVVQPAAPVSPPA..........VPSP....PPAAAPAVPVPPPAVHPAAP.....P    61
NTDB id 1118 NGFG RS11455 WP 003696286.1     MFSFFRRKKKQETPALEEAQVQ....ETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMP    76
consensus                                    !!!!!**! *** *   *!******* *!!*! ** **********!*!******* ! *!!* ***!!**  ******!

logo

NTDB id 381347 FYA87 RS13880 WP 005015354.1  VQPEPAASPVPAVPEPVAAPVVPPAPSPSAIAPAAAAEPAAAPKKASWLNRLKQGLARTG....QSIGGIFVGVKVDENL   137
NTDB id 1118 NGFG RS11455 WP 003696286.1     SEAGEAAERVESAKEAVAET......VGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDL   150
consensus                                      * *!! *!*  *!*!! ******* * !***   ! *   **!  !  !!!!!!** **** !**!*! !  * ! !

logo

NTDB id 381347 FYA87 RS13880 WP 005015354.1  FEELESALIMADAGIEATEKLLTELRARVKKDRIEDAAQVKTALRQLLTQHLKPLERSFDLTR.TQPLVVMIAGVNGAGK   216
NTDB id 1118 NGFG RS11455 WP 003696286.1     YEELETVLITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGK   230
consensus                                    *!!!!* !! *! !*!!!! !* **!*!!    * !*  ** !!*  !  **!!!** !* *  ! !*!*!!*!!!!!

logo

NTDB id 381347 FYA87 RS13880 WP 005015354.1  TTSIGKLAHTFQRQGASVLLAAGDTFRAAAREQLVEWGTRNNVTVIAQEGGDPAAVAFDAVNAGRARGVGVVMIDTAGRL   296
NTDB id 1118 NGFG RS11455 WP 003696286.1     TTSIGKLAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRL   310
consensus                                    !!!!!!!!* !! !! !!!!!!!!!!!!!!!!!!  !! !!!!!!! !  !!*!!!*!!!!*!**!!!* *!* !!!!!!

logo

NTDB id 381347 FYA87 RS13880 WP 005015354.1  PTQLHLMEELKKIRRVISKADAAAPHEVLLVVDGNTGQNALAQIRAFDAAIQLTGLVVTKLDGTAKGGTLAAVAAGSQGV   376
NTDB id 1118 NGFG RS11455 WP 003696286.1     PTQLHLMEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALA....SD   386
consensus                                    !!!!!!!!!*!!**!!* !! **!!!!***!*!*! !!!!* !**!!! !* !!!!*!!!!!!!!!!! !!!*!****

logo

NTDB id 381347 FYA87 RS13880 WP 005015354.1  RPVPVYWIGVGEGLEDLQPFVAEEFASALLAD   408
NTDB id 1118 NGFG RS11455 WP 003696286.1     RPVPVRYIGVGEGIDDLRPFDARAFVDALLD.   417
consensus                                    !!!!! *!!!!!!**!! !! !  !  !!! *
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X non conserved

X similar

X ≥ 50% conserved