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logo
NTDB id 404775 GNX71 RS27385 WP 206175326.1  MFSFFKKKPPAETPVAPAPTPPVAPPAAPVAEPA.P.....ARSVFSPSSWFGGK.PAAEEPAPAPVEAP.AVVVPAPSP   72
NTDB id 1118 NGFG RS11455 WP 003696286.1     MFSFFRRKKKQETPALEEAQVQE..TAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSE   78
consensus                                    !!!!!**!** !!!  * * ** ***!!*!  * *******   ! !*     !*** ! !* ***!! **  ! * !!*

logo

NTDB id 404775 GNX71 RS27385 WP 206175326.1  APVASPALIPAPVPAPAPVAAPAPVPVPPPAPASVPKPVAIEAAAAPERKGWFDKLKTGLRKTG....TGIQAVFVNAQI  148
NTDB id 1118 NGFG RS11455 WP 003696286.1     AGEAAE............RVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQI  146
consensus                                    !* ! *************   *!* *!*** *  !* *    !   *** !!  *!! !! !* ****  * *!!  *!!

logo

NTDB id 404775 GNX71 RS27385 WP 206175326.1  DDALYEELESALLMADTGVKATEYLLDDLKVRVKRQMATDASQVKVLLAEAIADLLRPLEKPLVIGQ.YTPTVIMVAGVN  227
NTDB id 1118 NGFG RS11455 WP 003696286.1     GEDLYEELETVLITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGIN  226
consensus                                     * !!!!!!* !* *! !* !!!!!* !** !!      !*  **  ! !!* !!**!!!!!!** *  ! !!!*!!*!

logo

NTDB id 404775 GNX71 RS27385 WP 206175326.1  GAGKTTSIGKLTKHLATEGASVLLAAADTFRAAAREQLLVWADRNTVEIVSNEGGDPSAVSFDAVNAGKARGKDVVLVDT  307
NTDB id 1118 NGFG RS11455 WP 003696286.1     GAGKTTSIGKLAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADT  306
consensus                                    !!!!!!!!!!! !*    ! !!!!!!*!!!!!!!!!!!  !* !! ! **!*  !!* !!*!!!!*!*!!!! !*!! !!

logo

NTDB id 404775 GNX71 RS27385 WP 206175326.1  AGRLPTQLHLMDELKKIKRVVTKADATAPHEVLLVIDGNTGQNALAQVKTFDETLGLTGLIVTKLDGTAKGGVLCAIARE  387
NTDB id 1118 NGFG RS11455 WP 003696286.1     AGRLPTQLHLMEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASD  386
consensus                                    !!!!!!!!!!!*!*!!*!!!* !! * !!!!***!*!*! !!!!* !!! !!* !!!!!!!!!!!!!!!!!!*!*!*! *

logo

NTDB id 404775 GNX71 RS27385 WP 206175326.1  RPIPVYFIGVGEKLEDLETFNAREFAQALLG   418
NTDB id 1118 NGFG RS11455 WP 003696286.1     RPVPVRYIGVGEGIDDLRPFDARAFVDALLD   417
consensus                                    !!*!! *!!!!! **!! *! !! !  !!!
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X non conserved

X similar

X ≥ 50% conserved