```
logo
NTDB id 205805 BO996 RS08570 WP 013803937.1  MFSFFKKKPTPSAAEISPTDSS.....AAPEKAGIPATPPQDSPPALPPAA..PAAPAADTGSAWWRKPFGGGKGDA..A   71
NTDB id 1118 NGFG RS11455 WP 003696286.1     MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVE.SLAESVKGRAESAVE........TVSGAVEQVKET   71
consensus                                    !!!!!**!* *  * * *    *****  *! ! !** ** ! ** !**  ***! *! **********  !*    **

logo

NTDB id 205805 BO996 RS08570 WP 013803937.1  PPETPPAPAAAPASTIAP.....ADFIA..APEPTAKPEAEGASRQNWMQRLKTGLRKT....GSSIATVFTGTQINDAL  140
NTDB id 1118 NGFG RS11455 WP 003696286.1     VAEMPSE.AGEAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDL  150
consensus                                    **! !* *!* *!  * ******!* **** * *   *     **  !  !!! !! !****** !*! !! ! !! * !

logo

NTDB id 205805 BO996 RS08570 WP 013803937.1  YEELEDALLMADTGVKATEHLLADLRRRVKEAKATDPAAVKALLADALADLLRPLEKPLVIGE.HQPTVIMVAGVNGAGK  219
NTDB id 1118 NGFG RS11455 WP 003696286.1     YEELETVLITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGK  230
consensus                                    !!!!!  !* *! !* !!!*!* !*! !!      !*  *** ! *!! !!**!!!!!!!**!** ! !!!*!!*!!!!!

logo

NTDB id 205805 BO996 RS08570 WP 013803937.1  TTSIGKLTRHLADEGQSVLLAAADTFRAAAREQLGVWANRNTVEIVSQEGGDPAAVSFDAVTAGRARKKDVVLVDTAGRL  299
NTDB id 1118 NGFG RS11455 WP 003696286.1     TTSIGKLAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRL  310
consensus                                    !!!!!!! **    ! !!!!!!*!!!!!!!!!!  !* !! ! **!!  !!*!!!*!!!! !**!!  !*!! !!!!!!

logo

NTDB id 205805 BO996 RS08570 WP 013803937.1  PTQLHLMEELRKIRRVVSKADATAPHEVLLVIDGNTGQNALAQVRAFDDALQLTGLIVTKLDGTAKGGVLAAIAQERPIP  379
NTDB id 1118 NGFG RS11455 WP 003696286.1     PTQLHLMEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVP  390
consensus                                    !!!!!!!!!**!**!!* !! * !!!!***!*!*! !!!!* !!*!!!!!! !!!!!!!!!!!!!!!!*!!!*! *!!*!

logo

NTDB id 205805 BO996 RS08570 WP 013803937.1  VYFIGVGEKVEDLETFNAREFAQALLS   406
NTDB id 1118 NGFG RS11455 WP 003696286.1     VRYIGVGEGIDDLRPFDARAFVDALLD   417
consensus                                    ! *!!!!! **!! *! !! !  !!!
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X non conserved

X similar

X ≥ 50% conserved