```
logo
NTDB id 39731 TSC RS12310 WP 015718358.1  MSGPRAFFPKPPLSTWGPGVGLGLGGLLGAWGLFHPWVLLLAPLLLPFLRLPFALGLVFVLLRGLLFPVPEPPYGTRLEGVFTLH   85
NTDB id 1035 TT RS08105 WP 011173974.1    ...........MRPSWTWGVGLGAGALVAAWGLFAPWAFLG.LGLLPFLRLPFALGFGLVLARALFFPLPEPPWGAAVEGVFAVR   73
consensus                                 *********** **! *!!!!! !*!**!!!!!*!!   ! ** !!!!!!!!!!!!    !! !*! !!*!!!!*!   *!!!! **

logo
NTDB id 39731 TSC RS12310 WP 015718358.1  QGTILWQGHRLWVQHYPGLEDGRYRLRGYLAPPQGKRNPGGFDQRTWLLSRGIRGVFHVERAEALAPLPDPRAPWRERLTAGLSP   170
NTDB id 1035 TT RS08105 WP 011173974.1    GGFTRAEGHLLWVSHYPPLPDGRYRLRGRIAPPQGRTNPGGFDQRAWLLSRGAKGVLRAESAEPLGPLPDWREGFRERLGAGLSL   158
consensus                                  !     !! !!! !!!*!*!!!!!!!! *!!!!!* !!!!!!!! !!!!!! *!! * ! !!*!*!!!!*! **!!!! !!!!*

logo
NTDB id 39731 TSC RS12310 WP 015718358.1  QVGEVVEGLVLGDKRGLEDAYPLFQKAGLAHLLAVSGQNVGYLA..ATLALLPLGRWRYLLALLLLPAYLWLAGPSPSLLRASLM   253
NTDB id 1035 TT RS08105 WP 011173974.1    EAREVVEGLVLGDKGGLETAYPLFQRAGLAHLLALSGLHVGVLVGFAVLGLYPLGRWRYLLALALLPFYLLLAGPSPSLVRASLM   243
consensus                                    !!!!!!!!!!! !!! !!!!!!*!!!!!!!!*!! *!! ! **! !*! !!!!!!!!!!! !!! !! !!!!!!!!*!!!!!

logo
NTDB id 39731 TSC RS12310 WP 015718358.1  AGLSLLGLFLGLGAAGVFQALGLALFLQLLLRPEALLGLGFQLSYLAVLGLALVLPALALPSGARGWLLGGLAASLAAQLPLIPL   338
NTDB id 1035 TT RS08105 WP 011173974.1    AGLSLLGLFLGLGGAGVVQALGLALFLQLLLRPEALLGLGFQLSYLAVLGLALVLPALRLPPGPRGYLLGGVAASLAVQAFLLPL   328
consensus                                 !!!!!!!!!!!!!*!!! !!!!!!!!!!!!!
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logo

NTDB id 39731 TSC RS12310 WP 015718358.1  GPKALLLADVPGEVEERLTPPQVAVVKVSHHGSRTGTDEDLLERTRPQVALIGVGRNPHGHPHPEVLERLDQHGVRVYRTDQNGA  678
NTDB id 1035 TT RS08105 WP 011173974.1    RGRALLLADLPREVEARLPVGEVEALKVSHHGSQSGTSEVLLERTRPRVALIGVGPNPFGHPHPEVLERLERHGAEIRRTDQDGA  668
consensus                                  **!!!!!!*! !!! !!*** !  *!!!!!!! *!! ! !!!!!!! !!!!!!!*!!*!!!!!!!!!!!* !!   * !!!! !!

logo

NTDB id 39731 TSC RS12310 WP 015718358.1  VRVLFGYAW   687
NTDB id 1035 TT RS08105 WP 011173974.1    VRVLFGYAW   677
consensus                                 !!!!!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved