logo
NTDB id 23932 STU RS17095 WP 011226464.1 MANKTKSKSLLEKMWRIKWWLLSIFTVLFLLFALFFPLNNYYVELPGGAFDTKEVLTVNKKADDSKGSYNFVAVAQTKATLALML 85
NTDB id 360 SMU RS02495 WP 002262039.1 .....MKTNKKFKWWLISGISLILLLMVFFFPL.PYYLEMPGGAYDIRSVVTVNHKEDKEKGSYNFVAVSLSKATPIQVL 74
consensus ********** * !!!!*! *** !! !!!!**!!*!*!!!!*! * !*!!!*! ! !!!!!!!!! *!!!* *!
logo
NTDB id 23932 STU RS17095 WP 011226464.1 YTQFNDFAKLQTAEEATGNYSDEDFMRINQFYMETSQNQAVYQGLTLAGKEVSLEYMGVYVLQVADDSSFKGVLNIADTVTAVNG 170
NTDB id 360 SMU RS02495 WP 002262039.1 YAWLTPFTEITSAKETTGGVSDADYMRINQFYMETSQNGAIYKALSLAHKKASFNYMGVYVLQVSRNSTFKGVLNIADTVTGVND 159
consensus ! *! * *! ! !! !! !*!!!!!!!!!!!!!! !*! *!*!!*! ! !!!!!!!!! !*!!!!!!!!!!!!*!!
logo
NTDB id 23932 STU RS17095 WP 011226464.1 KTFDNSTDMIKYVQGLKLGSKVKVTYMRDGKEKTATGKIIKIANGKNGIGIGLTDHTEIKSPENVKFKLDGVGGPSAGLMFTLAI 255
NTDB id 360 SMU RS02495 WP 002262039.1 KTFHSSKELIKYVSGLKLGSKVSVQYTSENKKKTAKGKVIKLSNGKNGIGIGLVDHTKVTSDDKIEFSTNGIGGPSAGLMFTLDI 244
consensus !!!* ! **!!!! !!!!!!!! ! ! * ! !!! !!*!!* !!!!!!!!!! !!! * !** * ! !*!!!!!!!!!!! !
logo
NTDB id 23932 STU RS17095 WP 011226464.1 YDQVSGQDLKAGRKIAGTGTIEKDGAVGDIGGAYLKVKSAADSGADIFFVPNNLVTKEMKKADPDAKTNYQEAKEAAEKLGTKMK 340
NTDB id 360 SMU RS02495 WP 002262039.1 YDQLVKEDLRKGRIIAGTGTIGEDGSVGDIGGADMKVAAADRINADIFFVPNNPVSKATLKKNPKALTNYQEAKQAAKKLGTKMK 329
consensus !!!* !!* !! !!!!!!! !! !!!!!!! *!! ! !!!!!!!!!*!*! ! ! ! !!!!!!! !! !!!!!!!
logo
NTDB id 23932 STU RS17095 WP 011226464.1 IVPVKTAQEAIDYLKKTK 358
NTDB id 360 SMU RS02495 WP 002262039.1 IVPVKNVQEAIDYLRKH. 346
consensus !!!!! !!!!!!!*!** X non conserved X similar X ≥50% conserved