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logo

NTDB id 542 H702 RS02345 WP 039696227.1        MLDIFILEDNITQQFRIERQIETIMERNHWEYQRLEAFAASKDIIKKADGQGNHQVFFLDLEIKGDEKQGLDVARSIREK   80
NTDB id 1127411 FGK97 RS02605 WP 037590320.1   .MNIFILEDDLTQQYYMETIVEKIIEKHRLQYHHFEVFGKPKQLLEAISEKGSHQVFFLDIEIKTEEKKGLDVAKKIREI   79
NTDB id 55 SMU RS08710 WP 002262114.1          MISIFVLEDDFLQQGRLETTIAAIMKEKNWSYKELTIFGKPQQLIDAIPEKGNHQIFFLDIEIKKEEKKGLEVANQIRQH   80
consensus                                      ** !!*!!!**!!***!* ** !***  * !  ***!********** **!*!!*!!!!*!!! *!!*!!*!!* !!*

logo

NTDB id 542 H702 RS02345 WP 039696227.1        DPTAIIVFVTTHSEFMLLTYQSLVGAIDFIDKNMNDEAFGGRIELCLKEALKY.QTGSFGENSYLFETAKARIRVAYDDI   159
NTDB id 1127411 FGK97 RS02605 WP 037590320.1   DPYAIIVFVTSHSELMPAAFKLQVGALDYLDKSLSKETFKKRIETALLHT.ENLTGKGLADDSLLFETPHTQIQVPFKDI   158
NTDB id 55 SMU RS08710 WP 002262114.1          NPSAVIVFVTTHSEFMPLTFQYQVSALDFIDKSLNPEEFSHRIESALYYAMENSQKNGQSEELFIFHSSETQFQVPFAEI   160
consensus                                      *!*!*!!!!!*!!!*!***** *!*!*!**!!*** ! !  !!!**!  **** *  * ******!**  ****!** *!

logo

NTDB id 542 H702 RS02345 WP 039696227.1        LYFETSPTVHRVILHTKTGQTEFYGTIAEVAKSDKRLLKCHRSFVINVNNVTEFDKSTRTAYFENGSYCQVSRDKVKKLM   239
NTDB id 1127411 FGK97 RS02605 WP 037590320.1   LYIETSSRPHRLVLYTHRGRTEFTANLSDMLKKEKRFYQCHRSFVVNPANVHQVDKVNHILHFQNGATCLVSRSKIKGLT   238
NTDB id 55 SMU RS08710 WP 002262114.1          LYFETSSTAHKLCLYTYDERIEFYGSMTDIVKMDKRLFQCHRSFIVNPANITRIDRKKRLAYFRNNKSCLISRTKLTKLR   240
consensus                                      !!*!!!** !***!*!  ***!!********! *!!**!!!!!**!**!** *!*  ****! !* *!**!!*!***!

logo

NTDB id 542 H702 RS02345 WP 039696227.1        AEMR......   243
NTDB id 1127411 FGK97 RS02605 WP 037590320.1   EAIAALHRRE   248
NTDB id 55 SMU RS08710 WP 002262114.1          AVIADQRRAK   250
consensus                                      * **   *
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X non conserved

X similar

X ≥ 50% conserved