```
logo
NTDB id 583 DLJ51 RS09560 WP 002962590.1   MYNKVILIGRLTAAPEMVKTPSNKSVTHITLAVNRRFKTQDGERQADFVNVVFWGKLAESLVSYAGKGSLVSLDGEIRTRHYEK.   84
NTDB id 624 LCA RS00040 WP 011373726.1     MINRVVLVGRLTRDVDLRYTSSGAAVGTFSMAVNRQFTNANGDREADFINCVIWRKSAENFANFTKKGSLVGVDGRLQTRNYENQ   85
NTDB id 65504 ANG RS02415 WP 003026323.1   MINNVVLVGRMTRDAELRYTPQNQAVATFTLAVNRNFKNQSGEREADFINVVIWRQQAENLANWAKKGALIGITGRIQTRNYENQ   85
NTDB id 113 BSU 40900 NP 391970.1          MLNRVVLVGRLTKDPELRYTPNGAAVATFTLAVNRTFTNQSGEREADFINCVTWRRQAENVANFLKKGSLAGVDGRLQTRNYENQ   85
NTDB id 112 BSU 36310 NP 391512.2          MFNQVMLVGRLTKDPDLRYTSAGAAVAHVTLAVNRSFKNASGEIEADYVNCTLWRKTAENTALYCQKGSLVGVSGRIQTRSYENE   85
consensus                                  !*!*!*!*!!*!*******!* ***!*****!!!! !****!***!!**!***!** !!****** *!!*!****!***!!*!!**

logo
NTDB id 583 DLJ51 RS09560 WP 002962590.1   DGQTHYVTEILAQSFQLLESRAQRALRENNGASDL.........................................TDLVLEEEE   128
NTDB id 624 LCA RS00040 WP 011373726.1     QGQRVYVTEVVVDNFSLLESRTTTEQRQGDGAS.QNFNSNQS..NGSQQSGFTSPQQTGNAPAANNTQADPFANNGQAIDISDDD   167
NTDB id 65504 ANG RS02415 WP 003026323.1   QGQRVYVTEVVADNFQLLESRASREGQSSGGYTSNNFGNSS........SNFGNST.SANQVPNFSRDENPFG.NTNPMDISDDD   160
NTDB id 113 BSU 40900 NP 391970.1          QGQRVFVTEVQAESVQFLEPKN.GGGSGSGGYNEGNSGGGQYFGGGQNDNPFGGNQNNQRRNQGNSFNDDPFANDGKPIDISDDD   169
NTDB id 112 BSU 36310 NP 391512.2          EGVNVYVTEVLADTVRFMDPKP.REKAAD...............................................   113
consensus                                  *!****!!!************  **      * *  *                *                  ***     *******

logo
NTDB id 583 DLJ51 RS09560 WP 002962590.1   LPF   131
NTDB id 624 LCA RS00040 WP 011373726.1     LPF   170
NTDB id 65504 ANG RS02415 WP 003026323.1   LPF   163
NTDB id 113 BSU 40900 NP 391970.1          LPF   172
NTDB id 112 BSU 36310 NP 391512.2          ...   113
consensus                                  ***
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X non conserved
X similar
X $\geq$ 50% conserved