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logo

NTDB id 583 DLJ51 RS09560 WP 002962590.1   MYNKVILIGRLTAAPEMVKTPSNKSVTHITLAVNRRFKTQDGERQADFVNVVFWGKLAESLVSYAGKGSLVSLDGEIRTRHYEK.   84
NTDB id 624 LCA RS00040 WP 011373726.1     MINRVVLVGRLTRDVDLRYTSSGAAVGTFSMAVNRQFTNANGDREADFINCVIWRKSAENFANFTKKGSLVGVDGRLQTRNYENQ   85
NTDB id 42664 TEH RS08925 WP 014125201.1   MINNVVLVGRLTKDPDLRYTQSGTAVANFTLAVNRTFTNQDGEREADFINCVIWRKAAENLANMVGKGAQIGVTGRIQTRNYENK   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   DGQTHYVTEILAQSFQLLESRAQRALRENNGASDLTD.......................................LVL   124
NTDB id 624 LCA RS00040 WP 011373726.1     QGQRVYVTEVVVDNFSLLESRTTTEQRQGDGASQNFNSNQSNGS.......QQSGFTSPQQTGNAPAANNTQADPFANNGQAIDI   163
NTDB id 42664 TEH RS08925 WP 014125201.1   EGQRVFVVEVVAENFQMLESRNKQNGQNN.........ANTGAS............NNNANEHSRNQQNSTQSNTDPFGGSSIDV   149
NTDB id 113 BSU 40900 NP 391970.1          QGQRVFVTEVQAESVQFLEPKNGGGSGSG.....GYNEGNSGGGQYFGGGQNDNPFGGNQNNQRRNQGNSFNDDPFANDGKPIDI   165
NTDB id 112 BSU 36310 NP 391512.2          EGVNVYVTEVLADTVRFMDPKPREKAAD.........................................................   113
consensus                                  *!****!*!************                  ** *                  *        *  *        *  ***

logo

NTDB id 583 DLJ51 RS09560 WP 002962590.1   EEEELPF   131
NTDB id 624 LCA RS00040 WP 011373726.1     SDDDLPF   170
NTDB id 42664 TEH RS08925 WP 014125201.1   TDDSLPF   156
NTDB id 113 BSU 40900 NP 391970.1          SDDDLPF   172
NTDB id 112 BSU 36310 NP 391512.2          .......   113
consensus                                  *******
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X non conserved
X similar
X ≥ 50% conserved