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logo

NTDB id 158 SP RS09595 WP 000282442.1   MYNKVILIGRLTSTPELHKTNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRTRRFEK.   84
NTDB id 624 LCA RS00040 WP 011373726.1  MINRVVLVGRLTRDVDLRYTSSGAAVGTFSMAVNRQFTNANGDREADFINCVIWRKSAENFANFTKKGSLVGVDGRLQTRNYENQ   85
NTDB id 75295 GOZ47 RS03985 WP 048589852.1 MINNVVLVGRLTKDPDLRYTSGDSAVATFTLAVNRNFKNQNGDREADFVNCVIWRKPAETLANYACKGTLLGVVGRIQSRSYENQ   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 158 SP RS09595 WP 000282442.1   NGQMNYVTEVLVTGFQLLESRAQRAMRENNAGQD.L.....................................ADLVLEEE   127
NTDB id 624 LCA RS00040 WP 011373726.1  QGQRVYVTEVVVDNFSLLESRTTTEQRQGDGASQNFNSNQSN....GSQQSGFTSPQQTGNAPAANNTQADPFANNGQAIDISDD   166
NTDB id 75295 GOZ47 RS03985 WP 048589852.1 QGQRVYVTEVVCDNFQLLESKNTRESQNKQDR.DNLNKNESNYIQEQ..NNVF......KTQNSEQLFNRNNNQSFGNQIDISDD   161
NTDB id 113 BSU 40900 NP 391970.1       QGQRVFVTEVQAESVQFLEPKNGGGSGSGGYN.EGN.SGGGQYFGGGQNDNPFGGNQNNQRRNQGNSFNDDPFANDGKPIDISDD   168
NTDB id 112 BSU 36310 NP 391512.2       EGVNVYVTEVLADTVRFMDPKPREKAAD.....................................................   113
consensus                               *!****!!!!* * *******          *       *           *           *  *       *  ******

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NTDB id 158 SP RS09595 WP 000282442.1   ELPF   131
NTDB id 624 LCA RS00040 WP 011373726.1  DLPF   170
NTDB id 75295 GOZ47 RS03985 WP 048589852.1 DLPF   165
NTDB id 113 BSU 40900 NP 391970.1       DLPF   172
NTDB id 112 BSU 36310 NP 391512.2       ....   113
consensus                               ****
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X non conserved
X similar
X ≥ 50% conserved