```
logo
NTDB id 498 SM12261 RS01980 WP 000282464.1     MYNKVIMIGRLTSTPELHKTNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGKLAESLASYATKGSLISVDGELRTR   80
NTDB id 526 SMSK321 RS08825 WP 000282440.1     MYNKVILIGRLTSTPELHKTNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRTR   80
NTDB id 158 SP RS09595 WP 000282442.1          MYNKVILIGRLTSTPELHKTNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRTR   80
NTDB id 268 KZH43 RS08520 WP 000282467.1       MYNKVIMIGRLTSTPELHKTNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRTR   80
NTDB id 227 SPD RS09110 WP 000282467.1         MYNKVIMIGRLTSTPELHKTNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRTR   80
NTDB id 193 SPR RS08685 WP 000282467.1         MYNKVIMIGRLTSTPELHKTNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRTR   80
NTDB id 153846 MOTHE RS00675 WP 011391677.1    MLNRVILIGRLTRDPELRYTASGVAMTTFTLAVDRNFVNQQGERGTDFIRITVWRKLAETCANHLGKGRLVAVDGRLQTR   80
NTDB id 624 LCA RS00040 WP 011373726.1         MINRVVLVGRLTRDVDLRYTSSGAAVGTFSMAVNRQFTNANGDREADFINCVIWRKSAENFANFTKKGSLVGVDGRLQTR   80
NTDB id 113 BSU 40900 NP 391970.1              MLNRVVLVGRLTKDPELRYTPNGAAVATFTLAVNRTFTNQSGEREADFINCVTWRRQAENVANFLKKGSLAGVDGRLQTR   80
NTDB id 112 BSU 36310 NP 391512.2              MFNQVMLVGRLTKDPDLRYTSAGAAVAHVTLAVNRSFKNASGEIEADYVNCTLWRKTAENTALYCQKGSLVGVSGRIQTR   80
consensus                                      !*!*!***!!!!****!**!***********!!*!******!****!******!***!!**!****!!*!**!*!***!!

logo
NTDB id 498 SM12261 RS01980 WP 000282464.1     RFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAG.............................................   116
NTDB id 526 SMSK321 RS08825 WP 000282440.1     RFEK.NGQMNYVAEVLVTGFQLLESRAQRAMRENNAG.............................................   116
NTDB id 158 SP RS09595 WP 000282442.1          RFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAG.............................................   116
NTDB id 268 KZH43 RS08520 WP 000282467.1       RFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAG.............................................   116
NTDB id 227 SPD RS09110 WP 000282467.1         RFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAG.............................................   116
NTDB id 193 SPR RS08685 WP 000282467.1         RFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAG.............................................   116
NTDB id 153846 MOTHE RS00675 WP 011391677.1    TYETPDGQKRSVTDVVAEDVRFLDWPKEGRGPAGSGPVAGP.........................GTGGGMETAGSGFNQD   137
NTDB id 624 LCA RS00040 WP 011373726.1         NYENQQGQRVYVTEVVVDNFSLLESRTTTEQRQGDGASQNFNSNQSNGSQ.......QSGFTSPQQTGNAPAANNTQADP   153
NTDB id 113 BSU 40900 NP 391970.1              NYENQQGQRVFVTEVQAESVQFLEPKNGGGSGSG.....GYNEGNSGGGQYFGGGQNDNPFGGNQNNQRRNQGNSFNDDP   155
NTDB id 112 BSU 36310 NP 391512.2              SYENEEGVNVYVTEVLADTVRFMDPKPREKAAD.................................................   113
consensus                                      **!* *!****!**!**********************

logo
NTDB id 498 SM12261 RS01980 WP 000282464.1     ..QDLADLVLEEEELPF   131
NTDB id 526 SMSK321 RS08825 WP 000282440.1     ..QDLADLVLEEEELPF   131
NTDB id 158 SP RS09595 WP 000282442.1          ..QDLADLVLEEEELPF   131
NTDB id 268 KZH43 RS08520 WP 000282467.1       ..QDLADLVLEEEELPF   131
NTDB id 227 SPD RS09110 WP 000282467.1         ..QDLADLVLEEEELPF   131
NTDB id 193 SPR RS08685 WP 000282467.1         ..QDLADLVLEEEELPF   131
NTDB id 153846 MOTHE RS00675 WP 011391677.1    FSDLGTEVEIGEDDLPF   154
NTDB id 624 LCA RS00040 WP 011373726.1         FANNGQAIDISDDDLPF   170
NTDB id 113 BSU 40900 NP 391970.1              FANDGKPIDISDDDLPF   172
NTDB id 112 BSU 36310 NP 391512.2              ................    113
consensus                                        ***************
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X non conserved

X similar

X $\geq$ 50% conserved