logo

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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 583 DLJ51 RS09560 WP 002962590.1     MYNKVILIGRLTAAPEMVKTPSNKSVTHITLAVNRRFKTQDGERQADFVNVVFWGKLAESLVSYAGKGSLVSLDGEIRTR  80
NTDB id 409 SMU RS08940 WP 002263454.1       MYNKVILIGRLVATPEMVKTPKDKSVTRVTLAVNRRFKSQDGEKQADFVNVVLWGRLAETIASYGSKGSLISLDGELRTR  80
NTDB id 100782 QUD80 RS14060 WP 078127112.1  MINNVTLVGRLTKDCDLRYTSSGVAVAAFTLAVNRNFTNQNGEREADYIQCVIWRKPAETLANYTRKGTLIGLVGRLQTR  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                                    !*!*!***!!!* **** *! ****!** **!!!!*** **!***!!******!***!!*******!!*!****!***!!
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NTDB id 498 SM12261 RS01980 WP 000282464.1   RFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAGQDL.....................................  119
NTDB id 526 SMSK321 RS08825 WP 000282440.1   RFEK.NGQMNYVAEVLVTGFQLLESRAQRAMRENNAGQDL.....................................  119
NTDB id 158 SP RS09595 WP 000282442.1        RFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAGQDL.....................................  119
NTDB id 268 KZH43 RS08520 WP 000282467.1     RFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAGQDL.....................................  119
NTDB id 227 SPD RS09110 WP 000282467.1       RFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAGQDL.....................................  119
NTDB id 193 SPR RS08685 WP 000282467.1       RFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAGQDL.....................................  119
NTDB id 583 DLJ51 RS09560 WP 002962590.1     HYEK.DGQTHYVTEILAQSFQLLESRAQRALRENNGASDL.....................................  119
NTDB id 409 SMU RS08940 WP 002263454.1       KYEK.DGVTHFVTEVLGHSFQLLESRAQRAMRENNSENDL.....................................  119
NTDB id 100782 QUD80 RS14060 WP 078127112.1  NYENQQGQRIYVTEVVVESFQLLESKEVSQQRGVNASNPT.....................PVGQ..QESNTQQPF  133
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                                    **!* *!*  *!*!***  ********************
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NTDB id 498 SM12261 RS01980 WP 000282464.1   .....ADLVLEEEEELPF  131
NTDB id 526 SMSK321 RS08825 WP 000282440.1   .....ADLVLEEEEELPF  131
NTDB id 158 SP RS09595 WP 000282442.1        .....ADLVLEEEEELPF  131
NTDB id 268 KZH43 RS08520 WP 000282467.1     .....ADLVLEEEEELPF  131
NTDB id 227 SPD RS09110 WP 000282467.1       .....ADLVLEEEEELPF  131
NTDB id 193 SPR RS08685 WP 000282467.1       .....ADLVLEEEEELPF  131
NTDB id 583 DLJ51 RS09560 WP 002962590.1     .....TDLVLEEEEELPF  131
NTDB id 409 SMU RS08940 WP 002263454.1       .....ADLVLEEEEELPF  131
NTDB id 100782 QUD80 RS14060 WP 078127112.1  PDLEGSFIEISDHDLPF   150
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 ≥ 50% conserved