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logo

NTDB id 498 SM12261 RS01980 WP 000282464.1   MYNKVIMIGRLTSTPELHKTNNDKSVARATIAVNRRYKD.QNGEREADFVNMVLWGKLAESLASYATKGSLISVDGELRT   79
NTDB id 526 SMSK321 RS08825 WP 000282440.1   MYNKVILIGRLTSTPELHKTNNDKSVARATIAVNRRYKD.QNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRT   79
NTDB id 158 SP RS09595 WP 000282442.1        MYNKVILIGRLTSTPELHKTNNDKSVARATIAVNRRYKD.QNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRT   79
NTDB id 268 KZH43 RS08520 WP 000282467.1     MYNKVIMIGRLTSTPELHKTNNDKSVARATIAVNRRYKD.QNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRT   79
NTDB id 227 SPD RS09110 WP 000282467.1       MYNKVIMIGRLTSTPELHKTNNDKSVARATIAVNRRYKD.QNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRT   79
NTDB id 193 SPR RS08685 WP 000282467.1       MYNKVIMIGRLTSTPELHKTNNDKSVARATIAVNRRYKD.QNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRT   79
NTDB id 409 SMU RS08940 WP 002263454.1       MYNKVILIGRLVATPEMVKTPKDKSVTRVTLAVNRRFKS.QDGEKQADFVNVVLWGRLAETIASYGSKGSLISLDGELRT   79
NTDB id 583 DLJ51 RS09560 WP 002962590.1     MYNKVILIGRLTAAPEMVKTPSNKSVTHITLAVNRRFKT.QDGERQADFVNVVFWGKLAESLVSYAGKGSLVSLDGEIRT   79
NTDB id 364037 FE331 RS00440 WP 208967762.1 .MNNVSIVGRLTRDPELKYTNSNTAVVSFTVAVNRPYKDKQTGEYEADFIRCQAWRKTAELISQYGYKGMRVGVEGRIQT   79
NTDB id 624 LCA RS00040 WP 011373726.1       MINRVVLVGRLTRDVDLRYTSSGAAVGTFSMAVNRQFTN.ANGDREADFINCVIWRKSAENFANFTKKGSLVGVDGRLQT   79
NTDB id 113 BSU 40900 NP 391970.1            MLNRVVLVGRLTKDPELRYTPNGAAVATFTLAVNRTFTN.QSGEREADFINCVTWRRQAENVANFLKKGSLAGVDGRLQT   79
NTDB id 112 BSU 36310 NP 391512.2            MFNQVMLVGRLTKDPDLRYTSAGAAVAHVTLAVNRSFKN.ASGEIEADYVNCTLWRKTAENTALYCQKGSLVGVSGRIQT   79
consensus                                    **!*!***!!!* **** *!*****!** **!!!!**** **!***!!*****!***!!*******!!******!***!

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NTDB id 498 SM12261 RS01980 WP 000282464.1   RRFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAG.........................................   116
NTDB id 526 SMSK321 RS08825 WP 000282440.1   RRFEK.NGQMNYVAEVLVTGFQLLESRAQRAMRENNAG.........................................   116
NTDB id 158 SP RS09595 WP 000282442.1        RRFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAG.........................................   116
NTDB id 268 KZH43 RS08520 WP 000282467.1     RRFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAG.........................................   116
NTDB id 227 SPD RS09110 WP 000282467.1       RRFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAG.........................................   116
NTDB id 193 SPR RS08685 WP 000282467.1       RRFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAG.........................................   116
NTDB id 409 SMU RS08940 WP 002263454.1       RKYEK.DGVTHFVTEVLGHSFQLLESRAQRAMRENNSE.........................................   116
NTDB id 583 DLJ51 RS09560 WP 002962590.1     RHYEK.DGQTHYVTEILAQSFQLLESRAQRALRENNGA.........................................   116
NTDB id 364037 FE331 RS00440 WP 208967762.1 RSYEGQNGQ.VFVTEVIADQFHFLESKSERSNNQQPNTSR.................................QRNNNF.ND   126
NTDB id 624 LCA RS00040 WP 011373726.1       RNYENQQGQRVYVTEVVVDNFSLLESRTTTEQRQGDGASQNFNSNQSNGSQ.......QSGFTSPQQTGNAPAANNTQAD   152
NTDB id 113 BSU 40900 NP 391970.1            RNYENQQGQRVFVTEVQAESVQFLEPKNGGGSGSG.....GYNEGNSGGGQYFGGGQNDNPFGGNQNNQRRNQGNSFNDD   154
NTDB id 112 BSU 36310 NP 391512.2            RSYENEEGVNVYVTEVLADTVRFMDPKPREKAAD.........................................   113
consensus                                    !**!* *!*  *!*!***  *****************

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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 409 SMU RS08940 WP 002263454.1       ...NDLADLVLEEEELPF   131
NTDB id 583 DLJ51 RS09560 WP 002962590.1     ...SDLTDLVLEEEELPF   131
NTDB id 364037 FE331 RS00440 WP 208967762.1  SF...QDDIHIDDGDLPF   141
NTDB id 624 LCA RS00040 WP 011373726.1       PFANNGQAIDISDDDLPF   170
NTDB id 113 BSU 40900 NP 391970.1            PFANDGKPIDISDDDLPF   172
NTDB id 112 BSU 36310 NP 391512.2            ..................   113
consensus                                       ***************
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X non conserved

X similar

X ≥ 50% conserved