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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 583 DLJ51 RS09560 WP 002962590.1       MYNKVILIGRLTAAPEMVKTPSNKSVTHITLAVNRRFKTQDGERQADFVNVVFWGKLAESLVSYAGKGSLVSLDGEIRTR   80
NTDB id 409 SMU RS08940 WP 002263454.1         MYNKVILIGRLVATPEMVKTPKDKSVTRVTLAVNRRFKSQDGEKQADFVNVVLWGRLAETIASYGSKGSLISLDGELRTR   80
NTDB id 1155205 QML75 RS05880 WP 047338779.1  MINSVSLTGRLIREVDLRYSKTGKPVAQFTLAVNRKYPDQNGERQADFIRCIIWGKSAEALEKYTQKGSLIGVEGRIQTR   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.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 583 DLJ51 RS09560 WP 002962590.1       HYEK.DGQTHYVTEILAQSFQLLESRAQRALRENNGA.......................................   116
NTDB id 409 SMU RS08940 WP 002263454.1         KYEK.DGVTHFVTEVLGHSFQLLESRAQRAMRENNSE.......................................   116
NTDB id 1155205 QML75 RS05880 WP 047338779.1  NYTNSQGKRVYLTEVVAQDFVFLESKGTAESRNGAGIATDYEDDLPD.............................   127
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     ..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 583 DLJ51 RS09560 WP 002962590.1       ..SDLTDLVLEEEELPF   131
NTDB id 409 SMU RS08940 WP 002263454.1         ..NDLADLVLEEEELPF   131
NTDB id 1155205 QML75 RS05880 WP 047338779.1  ..LDMGVEEFSDDDLPF   142
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