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logo
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NTDB id 498 SM12261 RS01980 WP 000282464.1    MYNKVIMIGRLTSTPELHKTNNDKSVARATIAVNRRYKDQN.GEREADFVNMVLWGKLAESLASYATKGSLISVDGELRT   79
NTDB id 526 SMSK321 RS08825 WP 000282440.1    MYNKVILIGRLTSTPELHKTNNDKSVARATIAVNRRYKDQN.GEREADFVNMVLWGRLAETLASYATKGSLISVDGELRT   79
NTDB id 158 SP RS09595 WP 000282442.1         MYNKVILIGRLTSTPELHKTNNDKSVARATIAVNRRYKDQN.GEREADFVNMVLWGRLAETLASYATKGSLISVDGELRT   79
NTDB id 268 KZH43 RS08520 WP 000282467.1      MYNKVIMIGRLTSTPELHKTNNDKSVARATIAVNRRYKDQN.GEREADFVNMVLWGRLAETLASYATKGSLISVDGELRT   79
NTDB id 227 SPD RS09110 WP 000282467.1        MYNKVIMIGRLTSTPELHKTNNDKSVARATIAVNRRYKDQN.GEREADFVNMVLWGRLAETLASYATKGSLISVDGELRT   79
NTDB id 193 SPR RS08685 WP 000282467.1        MYNKVIMIGRLTSTPELHKTNNDKSVARATIAVNRRYKDQN.GEREADFVNMVLWGRLAETLASYATKGSLISVDGELRT   79
NTDB id 583 DLJ51 RS09560 WP 002962590.1      MYNKVILIGRLTAAPEMVKTPSNKSVTHITLAVNRRFKTQD.GERQADFVNVVFWGKLAESLVSYAGKGSLVSLDGEIRT   79
NTDB id 409 SMU RS08940 WP 002263454.1        MYNKVILIGRLVATPEMVKTPKDKSVTRVTLAVNRRFKSQD.GEKQADFVNVVLWGRLAETIASYGSKGSLISLDGELRT   79
NTDB id 624 LCA RS00040 WP 011373726.1        MINRVVLVGRLTRDVDLRYTSSGAAVGTFSMAVNRQFTNAN.GDREADFINCVIWRKSAENFANFTKKGSLVGVDGRLQT   79
NTDB id 299505 DP112 RS09810 WP 114866882.1  MINNVVLVGRLTRDVELRYTPSNQAVATFTLAVNRNFKNQST GEREADFINCVMWRQQAENLANWTKKGHLIGITGRIQT   80
NTDB id 113 BSU 40900 NP 391970.1             MLNRVVLVGRLTKDPELRYTPNGAAVATFTLAVNRTFTNQS.GEREADFINCVTWRRQAENVANFLKKGSLAGVDGRLQT   79
NTDB id 112 BSU 36310 NP 391512.2             MFNQVMLVGRLTKDPDLRYTSAGAAVAHVTLAVNRSFKNAS.GEIEADYVNCTLWRKTAENTALYCQKGSLVGVSGRIQT   79
consensus                                     !*!*!***!!!* **** *! ****!** **!!!!*** ** !***!!**!***!***!!*******!!*!****!***!
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logo
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NTDB id 498 SM12261 RS01980 WP 000282464.1    RRFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAGQ.........................................   117
NTDB id 526 SMSK321 RS08825 WP 000282440.1    RRFEK.NGQMNYVAEVLVTGFQLLESRAQRAMRENNAGQ.........................................   117
NTDB id 158 SP RS09595 WP 000282442.1         RRFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAGQ.........................................   117
NTDB id 268 KZH43 RS08520 WP 000282467.1      RRFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAGQ.........................................   117
NTDB id 227 SPD RS09110 WP 000282467.1        RRFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAGQ.........................................   117
NTDB id 193 SPR RS08685 WP 000282467.1        RRFEK.NGQMNYVTEVLVTGFQLLESRAQRAMRENNAGQ.........................................   117
NTDB id 583 DLJ51 RS09560 WP 002962590.1      RHYEK.DGQTHYVTEILAQSFQLLESRAQRALRENNGAS.........................................   117
NTDB id 409 SMU RS08940 WP 002263454.1        RKYEK.DGVTHFVTEVLGHSFQLLESRAQRAMRENNSEN.........................................   117
NTDB id 624 LCA RS00040 WP 011373726.1        RNYENQQGQRVYVTEVVVDNFSLLESRTTTEQRQGDGASQNFNSNQSNGS.......QQSGFTSPQQTGNAPAANNTQAD   152
NTDB id 299505 DP112 RS09810 WP 114866882.1  RSYENQQGQRVYVTEVVAESFQVFEKRDNTANHTSM.............................................EDQM   120
NTDB id 113 BSU 40900 NP 391970.1             RNYENQQGQRVFVTEVQAESVQFLEPKNGGGSGSGG.....YNEGNSGGGQYFGGGQNDNPFGGNQNNQRRNQGNSFNDD   154
NTDB id 112 BSU 36310 NP 391512.2             RSYENEEGVNVYVTEVLADTVRFMDPKPREKAAD.............................................   113
consensus                                     !**!* *!*  *!*!***  ******************
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logo
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NTDB id 498 SM12261 RS01980 WP 000282464.1    ....DLADLVLEEEELPF   131
NTDB id 526 SMSK321 RS08825 WP 000282440.1    ....DLADLVLEEEELPF   131
NTDB id 158 SP RS09595 WP 000282442.1         ....DLADLVLEEEELPF   131
NTDB id 268 KZH43 RS08520 WP 000282467.1      ....DLADLVLEEEELPF   131
NTDB id 227 SPD RS09110 WP 000282467.1        ....DLADLVLEEEELPF   131
NTDB id 193 SPR RS08685 WP 000282467.1        ....DLADLVLEEEELPF   131
NTDB id 583 DLJ51 RS09560 WP 002962590.1      ....DLTDLVLEEEELPF   131
NTDB id 409 SMU RS08940 WP 002263454.1        ....DLADLVLEEEELPF   131
NTDB id 624 LCA RS00040 WP 011373726.1        PFANNGQAIDISDDDLPF   170
NTDB id 299505 DP112 RS09810 WP 114866882.1  PPNFSGPTMDINDDGLPF   138
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