
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 364147 FE326 RS07070 WP 208954032.1 .MNNVSIVGRLTREPELKYTNSNTAVVSFTVAVNRPYKDKQTGEYEADFIRCQAWRKTAELISQYGYKGMRVGVEGRIQT 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
logo
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 364147 FE326 RS07070 WP 208954032.1 RTYEGQNGQ.VFVTEVIADQFHFLESKSERSNNQYQNA NRQQNNNFN.D 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
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 409 SMU RS08940 WP 002263454.1 NDLADLVLEEEELPF 131
NTDB id 583 DLJ51 RS09560 WP 002962590.1 SDLTDLVLEEEELPF 131
NTDB id 364147 FE326 RS07070 WP 208954032.1 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
non conserved
similar
≥ 50% conserved