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NTDB id 624 LCA RS00040 WP 011373726.1	MINRVVLVGRLLTRDVLRYTSSGAAVCTFSMAVNRQFTNANGDREADFINCVIWRKSAENFANFTKKGSLVGVDGRLQTR	80
NTDB id 113 BSU 40900 NP 391970.1	MLNRVVLVGRLLTKDPELRYTPNGAAVATFTLAVNRITFTNQSGEREADFINCVITWRRAENVANFLKKGSLAGVDGRLQTR	80
NTDB id 597 KW2 RS01935 WP 021036727.1	.MNKTMILIGRLTNAPEISKTTNNKSYVRVTLAVNRRFKNEKGEREADFISITLWCKSAETLVSYAKKGSLISVEGETRTR	79
NTDB id 498 SM12261 RS01980 WP 000282464.1	MYNKVIMIGRLTSTPELHKTNNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGKLAESLASYATKGSLISVDGELRTR	80
NTDB id 526 SMSK321 RS08825 WP 000282440.1	MYNKVILIGRLTSTPELHKTNNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRTR	80
NTDB id 53363 HMPREF1038 RS09320 WP 000282467.1	MYNKVIMIGRLTSTPELHKTNNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRTR	80
NTDB id 268 KZH43 RS08520 WP 000282467.1	MYNKVIMIGRLTSTPELHKTNNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRTR	80
NTDB id 227 SPD RS09110 WP 000282467.1	MYNKVIMIGRLTSTPELHKTNNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRTR	80
NTDB id 193 SPR RS08685 WP 000282467.1	MYNKVIMIGRLTSTPELHKTNNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRTR	80
NTDB id 158 SP RS09595 WP 000282442.1	MYNKVILIGRLTSTPELHKTNNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRTR	80
NTDB id 409 SMU RS08940 WP 002263454.1	MYNKVILIGRLVATPEVVKTPKDKSVTRVTLAVNRRFKSQDGEKQADFVNVVLWGRLAETIASYGSKGSLISLDGELRTR	80
NTDB id 583 DLJ51 RS09560 WP 002962590.1	MYNKVILIGRLTAAPENVKTPSNKSVTHITLAVNRRFKTQDGERQADFVNVVFWGKLAESLVSYAGKGSLVSLDGEIRTR	80
consensus	***!*****!!!!*****!!*****!!!!*****!*****!!!!*****!*****!!!!*****!*****!!!!	

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NTDB id 624 LCA RS00040 WP 011373726.1	RYEKNQGMYVTEVLVTEVQLLESRAQRAMRENNAGQ	
NTDB id 113 BSU 40900 NP 391970.1	NYENQGGQRVYVTEVVVDNFSLLESRTTTEQRQGDGASQNFNSNQSNQSQ.....QSGFTSPQQTGNAPAAANTQADP	153
NTDB id 597 KW2 RS01935 WP 021036727.1	NYENQGGQRVFVTEVQAESVQFLEPKNGGSGS....GGYNEGNSGGGQYFGGGQNDNPFGGNQNNQRRNQGNSFNDDP	155
NTDB id 498 SM12261 RS01980 WP 000282464.1	NYTDKNEQKHYTEILGLSYDLESRAATLALRESAV.....	115
NTDB id 526 SMSK321 RS08825 WP 000282440.1	RFEKNGQMNYVTEVLVTGTFQLLESRAQRAMRENNAGQ.....	117
NTDB id 53363 HMPREF1038 RS09320 WP 000282467.1	RFEKNGQMNYVAEVLVTGTFQLLESRAQRAMRENNAGQ.....	117
NTDB id 268 KZH43 RS08520 WP 000282467.1	RFEKNGQMNYVTEVLVTGTFQLLESRAQRAMRENNAGQ.....	117
NTDB id 227 SPD RS09110 WP 000282467.1	RFEKNGQMNYVTEVLVTGTFQLLESRAQRAMRENNAGQ.....	117
NTDB id 193 SPR RS08685 WP 000282467.1	RFEKNGQMNYVTEVLVTGTFQLLESRAQRAMRENNAGQ.....	117
NTDB id 158 SP RS09595 WP 000282442.1	RFEKNGQMNYVTEVLVTGTFQLLESRAQRAMRENNAGQ.....	117
NTDB id 409 SMU RS08940 WP 002263454.1	KYELKDGVTHTFVTEVLGHSEFQLLESRAQRAMRENNSEN.....	117
NTDB id 583 DLJ51 RS09560 WP 002962590.1	HYELKDGQTHYVTEVTLAQSFQLLESRAQRAIRENNGAS.....	117
consensus	*** *****!!*****!!**********	

logo

NTDB id 624 LCA RS00040 WP 011373726.1	FANRGQATIDISDDDLPF	170
NTDB id 113 BSU 40900 NP 391970.1	FANDGKPIDISDDDLPF	172
NTDB id 597 KW2 RS01935 WP 021036727.1	...KTEELLEADELPF	129
NTDB id 498 SM12261 RS01980 WP 000282464.1	...DLADLVLEEEELPF	131
NTDB id 526 SMSK321 RS08825 WP 000282440.1	...DLADLVLEEEELPF	131
NTDB id 53363 HMPREF1038 RS09320 WP 000282467.1	...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 158 SP RS09595 WP 000282442.1	...DLADLVLEEEELPF	131
NTDB id 409 SMU RS08940 WP 002263454.1	...DLADLVLEEEELPF	131
NTDB id 583 DLJ51 RS09560 WP 002962590.1	...DLTDLVLEEEELPF	131
consensus	*****!!!!	

-  non conserved
-  similar
-  ≥ 50% conserved