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NTDB id	977468	MGAS37844	RS00670	WP	002986552.1	MINQWNNLRHKLLKGF T LLEMLLVILVLSVLMLLFVPNLSKKQDRVETGNAAVVKLVENQAELYELSQG.SK	72
NTDB id	588	KW2	RS10845	WP	051303741.1	MSKALSLV K IHGRLCQC K KAFTLIEMLIVLAIIISILLFLLFVPNLIK E KAQVCKTGEAAVVKVVESQAQLYELDHDNK	80
NTDB id	460	GPW51	RS00790	WP	011921720.1	M K KLVE M KVG F TFLVEMLVVLGGIISLLLLLFVPNLSQQKD A IQKKGDAAVVKVVESQMELYEL H D.EE	68
NTDB id	502	SM12261	RS00955	WP	000738641.1	M K KMMTF L KKAKVKAFTLVEMLVLLIIISVLLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKN.ED	72
NTDB id	272	KZH43	RS09255	WP	000738626.1	M K KMMTF L KKAKVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSL E KN.ED	72
NTDB id	231	SPD	RS09885	WP	000738626.1	M K KMMTF L KKAKVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSL E KN.ED	72
NTDB id	197	SPR	RS09420	WP	000738626.1	M K KMMTF L KKAKVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSL E KN.ED	72
NTDB id	162	SP	RS10370	WP	000738627.1	M K KMMTF L KKAKVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSL E KN.ED	72
NTDB id	530	SMSK321	RS09230	WP	000738643.1	M K KMMTF L KKAKVKAFTLVEMLVLLIIISVLLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKN.ED	72
NTDB id	431	SGO	RS09410	WP	012130940.1	MNK L KKL R VKAFTLVEMLVLLIIISVLMLLFVPNLTKQKEAVSDTGNAAVVKVVESQAELYELKNTGDQ	69
NTDB id	396	FSA28	RS09395	WP	002263437.1	MKK I KRVSVRAFTLIEMLVLLIIISILLFLLFVPNLTKQKDKVKDTGGSAAVVKVVESQAELYELNKT.DS	68
NTDB id	387	SMU	RS09025	WP	002263437.1	MKK I KRVSVRAFTLIEMLVLLIIISILLFLLFVPNLTKQKDKVKDTGGSAAVVKVVESQAELYELNKT.DS	68
consensus						** * ** * **** * ** * ** * ** * ** * ** * ** * ** * ** *	

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[illegible]

- X non conserved
- X similar
- X >50% conserved