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NTDB id	977020	MGAS37859	RS00670	WP	002986552.1	MINQWNNLRHKLLKGF T LLEMLLVILVLSVLMFLFVPNLSKKQDRVETGNAAVVKLVENQAELYELSQG.SK	72
NTDB id	588	KW2	RS10845	WP	051303741.1	MSKALSLV K IHGRLCQKKQKAFTLIEMLIVLAIIISILLFLLFVPNLIKKAKAQVKGTGEAAVVKVVSQAQLYELDHDNDK	80
NTDB id	460	GPW51	RS00790	WP	011921720.1	M K KLVE M KVG G FTLVEMLVLVGIISLLLLLFVPNLSQQKD A IQKKGDAAVVKVVSQMELYEL E H D .EE	68
NTDB id	502	SM12261	RS00955	WP	000738641.1	M K KMMTFL K KA V KAFTLVEMLVLLIIISVLLFLLFVPNLTKQKEAVNDKGA A AVVKVVSQAELYSLDKN.ED	72
NTDB id	272	KZH43	RS09255	WP	000738626.1	M K KMMTF L KK A KVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGA A AVVKVVSQAELYSL E KN.ED	72
NTDB id	231	SPD	RS09885	WP	000738626.1	M K KMMTFL L KK A KVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGA A AVVKVVSQAELYSL E KN.ED	72
NTDB id	197	SPR	RS09420	WP	000738626.1	M K KMMTF L KK A KVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGA A AVVKVVSQAELYSL E KN.ED	72
NTDB id	162	SP	RS10370	WP	000738627.1	M K KMMTFL L KK A KVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGA A AVVKVVSQAELYSL E KN.ED	72
NTDB id	530	SMSK321	RS09230	WP	000738643.1	M K KMMTFL L KK A KVKAFTLVEMLVLLIIISVLLFLLFVPNLTKQKEAVNDKGA A AVVKVVSQAELYSLDKN.ED	72
NTDB id	431	SGO	RS09410	WP	012130940.1	MNK L KK L R V KAFTLVEMLVLLIIISVLMFLFVPNLTKQKEAVSDTGNAAVVKVVSQAELYELKNTGDQ	69
NTDB id	396	FSA28	RS09395	WP	002263437.1	MKK I KRVSVRAFTLIEMLVLLIIISILLFLLFVPNLTKQKDKVKDTGGSAVVKVVSQAELYELNK T .DS	68
NTDB id	387	SMU	RS09025	WP	002263437.1	MKK I KRVSVRAFTLIEMLVLLIIISILLFLLFVPNLTKQKDKVKDTGGSAVVKVVSQAELYELNK T .DS	68
consensus						** * ** * **** * ** * ** * ** * ** * ** * ** * ** * ** *	

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NTDB id	977020	MGAS37859	RS00670	WP	002986552.1	P S L S Q L K A D G S I T E K Q E K A Y Q D Y Y D K H K N E K A R L S N	108
NTDB id	588	KW2	RS10845	WP	051303741.1	P N L S E L L S A C M I T C K Q V T A Y D D Y Y D Q N K N E Q R N F D D	116
NTDB id	460	GPW51	RS00790	WP	011921720.1	A T V A D L Q A K G Y I T E K Q A K Q Y A T A K	93
NTDB id	502	SML12261	RS00955	WP	000738641.1	A S L S K L Q A D G R I T E E Q A K A Y K E Y H A K N T S C T V A D .	107
NTDB id	272	KZH43	RS09255	WP	000738626.1	A S L R K L Q A D G R I T E E Q A K A Y K E Y H D K N G G A N R K V N D	108
NTDB id	231	SPD	RS09885	WP	000738626.1	A S L R K L Q A D G R I T E E Q A K A Y K E Y H D K N G G A N R K V N D	108
NTDB id	197	SPR	RS09420	WP	000738626.1	A S L R K L Q A D G R I T E E Q A K A Y K E Y H D K N G G A N R K V N D	108
NTDB id	162	SP	RS10370	WP	000738627.1	A S L R K L Q A D G R I T E E Q A K A Y K E Y N D K N G G A N R K V N D	108
NTDB id	530	SMK321	RS09230	WP	000738643.1	A S L S K L Q A D G R I T E E Q A K A Y N E Y Y A K N G G A N R K V N D	108
NTDB id	431	SGO	RS09410	WP	012130940.1	A T S K L V A A G N I S Q K Q A D S Y K A Y Y G K N N S E T Q A V A N	105
NTDB id	396	FSA28	RS09395	WP	002263437.1	A T L S K L V A S G D I T Q E A N S Y K D Y Y A K H P S K T R H V A N	104
NTDB id	387	SMU	RS09025	WP	002263437.1	A T L S K L V A S G D I T Q E A N S Y K D Y Y A K H P S K T R H V A N	104
consensus						* * * * *	

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