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NTDB id	977524	MGAS37843	RS00670	WP	002986552.1	MINQWNNLRHKLLKGF T LLEMLLVILVLSVLMLLFVPNLSKKQDRVETGNAAVVKLVENQAELYELSGG.SK	72
NTDB id	588	KW2	RS10845	WP	051303741.1	MSKALSLV K IHGRLCQC K KAFTLIEMLIVLAIIISILLFVPNLIK E KAQVQKTGEAAVVKVVESQAQLYELDHDNK	80
NTDB id	460	GPW51	RS00790	WP	011921720.1	M K KLVE M KVG F TLEVEMLVVLGIISLLLLFVPNLSQQKD A IQKKGDAAVVKVVESQMELYEL H D.EE	68
NTDB id	502	SM12261	RS00955	WP	000738641.1	M K KMMTFL K KA V KAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGA A AVVKVVESQAELYSLDKN.ED	72
NTDB id	272	KZH43	RS09255	WP	000738626.1	M K KMMTF L KKAKVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGA A AVVKVVESQAELYSL E KN.ED	72
NTDB id	231	SPD	RS09885	WP	000738626.1	M K KMMTFL L KKAKVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGA A AVVKVVESQAELYSL E KN.ED	72
NTDB id	197	SPR	RS09420	WP	000738626.1	M K KMMTF L KKAKVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGA A AVVKVVESQAELYSL E KN.ED	72
NTDB id	162	SP	RS10370	WP	000738627.1	M K KMMTFL L KKAKVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGA A AVVKVVESQAELYSL E KN.ED	72
NTDB id	530	SMSK321	RS09230	WP	000738643.1	M K KMMTFL L KKAKVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGA A AVVKVVESQAELYSLDKN.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 KRVSVRAFTLIEMLVLLIISILLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DS	68
NTDB id	387	SMU	RS09025	WP	002263437.1	MKK I KRVSVRAFTLIEMLVLLIISILLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DS	68
consensus						** * **** ** ** ** **	

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NTDB id	977524 MGAS37843 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 G 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 K	93
NTDB id	502 SMS12261 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 Q N T S Q 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 SMSK321 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 Q 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 Q 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