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NTDB id	978587	MGAS37637	RS00670	WP	002986552.1	MINQWNNLRHKKLLKGFTLLEMLLVILVISVLMFLFVPNLSKKQDRVETGNAAAVVKLVENQAELYELSQG.SK	72
NTDB id	588	KW2	RS10845	WP	051303741.1	MSKALSLVKIIGHGRKLCCQKKQKAFTLIEMLIVLAITISILILLFVPNLIKKEKAQVCKTGEEAAVVKVVSQAQLYELDHDNK	80
NTDB id	460	GPW51	RS00790	WP	011921720.1	MKKLVEMKVKGFTEVLVLLIIISLLLLLFVPNLSQQKDAIQKKGDAAVVKVVSQMELEYELHDEE	68
NTDB id	502	SM12261	RS00955	WP	000738641.1	MKKMMTFLKKAKVKAFTLVEMLVLLIIISVLLFLLFVPNLTKQKEAVNDKGKAADVVKVVSQAELYSLDKN.ED	72
NTDB id	272	KZH43	RS09255	WP	000738626.1	MKKMMTFLKKAQVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGKAADVVKVVSQAELYSLKN.ED	72
NTDB id	231	SPD	RS09885	WP	000738626.1	MKKMMTFLKKAQVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGKAADVVKVVSQAELYSLKN.ED	72
NTDB id	197	SPR	RS09420	WP	000738626.1	MKKMMTFLKKAQVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGKAADVVKVVSQAELYSLKN.ED	72
NTDB id	162	SP	RS10370	WP	000738627.1	MKKMMTFLKKAQVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGKAADVVKVVSQAELYSLKN.ED	72
NTDB id	530	SMSK321	RS09230	WP	000738643.1	MKKMMTFLKKAQVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGKAADVVKVVSQAELYSLDKN.ED	72
NTDB id	431	SGO	RS09410	WP	012130940.1	MNKLLKL RVKAFTLVEMLVLLIIISVLMFLFVPNLTKQKEAVSDTGNAADVVKVVSQAELYELKNTGDQ	69
NTDB id	396	FSA28	RS09395	WP	002263437.1	MKKIKRVSVRAFTLIEMLVLLIIISILLLLFPNLTQKDKVKVDGTGSAAVVKVVSQAELYELNKT.DS	68
NTDB id	387	SMU	RS09025	WP	002263437.1	MKKIKRVSVRAFTLIEMLVLLIIISILLLLFPNLTQKDKVKVDGTGSAAVVKVVSQAELYELNKT.DS	68
consensus						** * **** ** ** *	

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[illegible]

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