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NTDB	id	978728	MGAS37830	RS00655	WP	002986552.1	MINQWNNLRHKKLGFTLLEMLLVILVISVLMFLFVPNLSKKQDRVETGNAAAVVKLVENQAELYELSGQ.SK	72
NTDB	id	588	KW2	RS10845	WP	051303741.1	MSKALSLVKIHRGRKLCQKKQKAFTLIEMLIVIAIISILILLFVPNLIKKAKAVVQKTGEAAAVKVVSQAQLYELDHDNDK	80
NTDB	id	460	GPW51	RS00790	WP	011921720.1	MCKLMVMKVGFTLVEMLVLVLGISLLLLFVPNLSQQKDAIQKKGDAAVVKVVSQAMELYELEHDEE	68
NTDB	id	502	SM12261	RS00955	WP	000738641.1	MKKMMTFLKKAQKVAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGAHAVVKVVSQAELYSLDKN.ED	72
NTDB	id	272	KZH43	RS09255	WP	000738626.1	MKKMMTFLKKAQKVAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGAHAVVKVVSQAELYSLDKN.ED	72
NTDB	id	231	SPD	RS09885	WP	000738626.1	MKKMMTFLKKAQKVAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGAHAVVKVVSQAELYSLDKN.ED	72
NTDB	id	197	SPR	RS09420	WP	000738626.1	MKKMMTFLKKAQKVAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGAHAVVKVVSQAELYSLDKN.ED	72
NTDB	id	162	SP	RS10370	WP	000738627.1	MKKMMTFLKKAQKVAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGAHAVVKVVSQAELYSLDKN.ED	72
NTDB	id	530	SMSK321	RS09230	WP	000738643.1	MKKMMTFLKKAQKVAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGAHAVVKVVSQAELYSLDKN.ED	72
NTDB	id	431	SGO	RS09410	WP	012130940.1	MNKLLKKLRVKAFTLVEMLVLLIIISVLMFLFVPNLTKQKEAVSDTGNAAAVVKVVSQAELYELKNTGDQ	69
NTDB	id	396	FSA28	RS09395	WP	002263437.1	MKKIKRVSVRAFTLIEMLVLLIISILILLFVPNLTKQKDKVKDTGGSAVVKVVSQAELYELNKTD.S	68
NTDB	id	387	SMU	RS09025	WP	002263437.1	MKKIKRVSVRAFTLIEMLVLLIISILILLFVPNLTKQKDKVKDTGGSAVVKVVSQAELYELNKTD.S	68
consensus							** * ** * **** * ** * ** * ** * ** * ** * ** * ** * ** *	

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NTDB	id	978728	MGAS37830	RS00655	WP	002986552.1	PSLSQLKADGS	ITEKQEAYQDYD	CHKNEKAR	SN	108	
NTDB	id	588	KW2	RS10845	WP	051303741.1	PNLSELLSAGM	ITCKQVTA	DDYDQNKNEQR	NFDD	116	
NTDB	id	460	GPW51	RS00790	WP	011921720.1	ATVADLQAKGY	ITEKQAKQY	ATAKK		93	
NTDB	id	502	SM1261	RS00955	WP	000738641.1	ASLSKLQADGR	ITEEQAKAYKEY	HA	KNTS	QTVAD	107
NTDB	id	272	KZH43	RS09255	WP	000738626.1	ASLRKLQADGR	ITEEQAKAYKEY	HD	KNGGAN	RKRVND	108
NTDB	id	231	SPD	RS09885	WP	000738626.1	ASLRKLQADGR	ITEEQAKAYKEY	HD	KNGGAN	RKRVND	108
NTDB	id	197	SPR	RS09420	WP	000738626.1	ASLRKLQADGR	ITEEQAKAYKEY	HD	KNGGAN	RKRVND	108
NTDB	id	162	SP	RS10370	WP	000738627.1	ASLRKLQADGR	ITEEQAKAYKEY	ND	KNGGAN	RKRVND	108
NTDB	id	530	SM5321	RS09230	WP	000738643.1	ASLSKLQADGR	ITEEQAKAY	NEY	YAKNGGAN	RKRVND	108
NTDB	id	431	SGO	RS09410	WP	012130940.1	ATLSKLVAAGN	ISQKQADS	YKAYYG	KNNSET	QAVAN	105
NTDB	id	396	FSA28	RS09395	WP	002263437.1	ATLSKLVASGD	ITQEANSYK	DY	YAKHPSKTR	RVAN	104
NTDB	id	387	SMU	RS09025	WP	002263437.1	ATLSKLVASGD	ITQEANSYK	DY	YAKHPSKTR	RVAN	104
consensus							*****!*****!	*****!*****!	*****!	*****!	*****!	

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