

NTDB id	32	MW	RS07995	WP	000472256.1		MFKF..LKKTQAFTLIEMLLVLLIISLLLILIIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id	16	SA	RS07765	WP	000472256.1		MFKF..LKKTQAFTLIEMLLVLLIISLLLILIIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id	588	KW2	RS10845	WP	051303741.1	MSKALSLV	KIHGRKLCKKKQAFTLIELVLVAIISILILLFPVNLIKKEKAQVQKTGEAAVVKVVSQAQLYELDHDNDK	80
NTDB id	460	GPW51	RS00790	WP	011921720.1		MKKLVEMKVKGFTLVEMLVLVGIIISLLLLFVPNLSSQQDKAIQKGDAAVVKVVESQMELYELEHD.EE	68
NTDB id	404963	GOM47	RS00915	WP	235080778.1		MNKLMTNFKKAKVQAFTLVEMLVLLIISVLLLFVPNLTKQKDVDVDDKGKAAAVKVVESQAELYSLDKN.ED	72
NTDB id	502	SM12261	RS00955	WP	000738641.1		MKKMMTFLKKAHVKAFTLVEMLVLLIISVLLLFVPNLTKQKEAVNDKGKAAAVKVVESQAELYSLEKN.ED	72
NTDB id	272	KZH43	RS09255	WP	000738626.1		MKKMMTFLKKAHVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAAVKVVESQAELYSLEKN.ED	72
NTDB id	231	SPD	RS09885	WP	000738626.1		MKKMMTFLKKAHVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAAVKVVESQAELYSLEKN.ED	72
NTDB id	197	SPR	RS09420	WP	000738626.1		MKKMMTFLKKAHVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAAVKVVESQAELYSLEKN.ED	72
NTDB id	162	SP	RS10370	WP	000738627.1		MKKMMTFLKKAHVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAAVKVVESQAELYSLEKN.ED	72
NTDB id	530	SMSK321	RS09230	WP	000738643.1		MKKMMTFLKKAHVKAFTLVEMLVLLIISVLLLFVPNLTKQKEAVNDKGKAAAVKVVESQAELYSLEKN.ED	72
NTDB id	431	SGO	RS09410	WP	012130940.1		MNKLKLRVKAFTLVEMLVLLIISVLMLLFVPNLTKQKEAVSDTGNAAVKVVESQAELYELKNTGDQ	69
NTDB id	396	FSA28	RS09395	WP	002263437.1		MKKIKRVSVRAFTLIEMLVLLIISILLFFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DS	68
NTDB id	387	SMU	RS09025	WP	002263437.1		MKKIKRVSVRAFTLIEMLVLLIISILLFFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DS	68
consensus						*	* * * * *	

NTDB	id	32	MW	RS07995	WP	000472256.1	SSIEDLIADGFIKEAQKTKSGETITISNG.EAVAN	103
NTDB	id	16	SA	RS07765	WP	000472256.1	SSIEDLIADGFIKEAQKTKSGETITISNG.EAVAN	103
NTDB	id	588	KW2	RS10845	WP	051303741.1	PNLSELLSACMITQKQVTAYDDYDQNKNEQRNFDD	116
NTDB	id	460	GPW51	RS00790	WP	011921720.1	ATVADLQAKGYITEKQAKQYATAKK.....	93
NTDB	id	404963	GOM47	RS00915	WP	235080778.1	ASLSKLQADGRITAEQVKAYDKDYAKQKTSQTVAD.	107
NTDB	id	502	SM12261	RS00955	WP	000738641.1	ASLSKLQADGRITEEQAKAYKEYHAKQNTSQTVAD.	107
NTDB	id	272	KZH43	RS09255	WP	000738626.1	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRKKVND	108
NTDB	id	231	SPD	RS09885	WP	000738626.1	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRKKVND	108
NTDB	id	197	SPR	RS09420	WP	000738626.1	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRKKVND	108
NTDB	id	162	SP	RS10370	WP	000738627.1	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRKKVND	108
NTDB	id	530	SMSK321	RS09230	WP	000738643.1	ASLSKLQADGRITEEQAKAYNEYAKNGGANRKKVND	108
NTDB	id	431	SGO	RS09410	WP	012130940.1	ATLSKLVAAGNISQKKQADSYKAYYGGNNSETQAVAN	105
NTDB	id	396	FSA28	RS09395	WP	002263437.1	ATLSKLVASGDITQEQANSYKDYAKHPSKTRHVAN	104
NTDB	id	387	SMU	RS09025	WP	002263437.1	ATLSKLVASGDITQEQANSYKDYAKHPSKTRHVAN	104
consensus							*** * ! **** ! ! **** ! **** * * * * *	

- X non conserved
- X similar
- X ≥ 50% conserved