

NTDB	id	32	MW	RS07995	WP	000472256.1		MFKFL..KKTQAFTLIEMLVLLLIISLLLILIIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB	id	16	SA	RS07765	WP	000472256.1		MFKFL..KKTQAFTLIEMLVLLLIISLLLILIIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB	id	588	KW2	RS10845	WP	051303741.1	MSKALSLV	KIHGRKLCKKKQAFTLIELVLVAIISILILLFVPNLIKEAKAQVKTGEAAVVKVVSQAELYELDHDNDK	80
NTDB	id	164077	AU077	RS09230	WP	039670352.1		MKKFWKLLQKTVKCFTLVEMLIVLLIISVLMMLFVPNLSKKDKDVREKGDAAVVKVVSQMDELMEYKMGTDK	72
NTDB	id	460	GPW51	RS00790	WP	01921720.1		MKKLVEMKVKGKFTLVEMLVVLGIISILLFLFVPNLSQQKDAIQKKGDDAAVVKVVSQMELYELEHDEE	68
NTDB	id	502	SM12261	RS00955	WP	000738641.1		MKKMMTFLKKAKVKAFTLVEMLVLLIISVLLLFVPNLTQKKEAVNDKGKAAVVKVVSQAELYSLDKNEED	72
NTDB	id	272	KZH43	RS09255	WP	000738626.1		MKKMMTFLKKAKVKAFTLVEMLVLLIISVLLLFVPNLTQKKEAVNDKGKAAVVKVVSQAELYSLDKNEED	72
NTDB	id	231	SPD	RS09885	WP	000738626.1		MKKMMTFLKKAKVKAFTLVEMLVLLIISVLLLFVPNLTQKKEAVNDKGKAAVVKVVSQAELYSLDKNEED	72
NTDB	id	197	SPR	RS09420	WP	000738626.1		MKKMMTFLKKAKVKAFTLVEMLVLLIISVLLLFVPNLTQKKEAVNDKGKAAVVKVVSQAELYSLDKNEED	72
NTDB	id	162	SP	RS10370	WP	000738627.1		MKKMMTFLKKAKVKAFTLVEMLVLLIISVLLLFVPNLTQKKEAVNDKGKAAVVKVVSQAELYSLDKNEED	72
NTDB	id	530	SMSK321	RS09230	WP	000738643.1		MKKMMTFLKKAKVKAFTLVEMLVLLIISVLLLFVPNLTQKKEAVNDKGKAAVVKVVSQAELYSLDKNEED	72
NTDB	id	431	SGO	RS09410	WP	012130940.1		MNKCLKLRVKAFTLVEMLVLLIISVLMMLFVPNLTQKKEAVSDTGNAAVVKVVSQAELYELKNKTDGQ	69
NTDB	id	396	FSA28	RS09395	WP	002263437.1		MKKITKRVSVRRAFTLIELVLLIISILLFFVPNLTQKDKVKVDGTGSAAVVKVVSQAELYELNKTDSDS	68
NTDB	id	387	SMU	RS09025	WP	002263437.1		MKKITKRVSVRRAFTLIELVLLIISILLFFVPNLTQKDKVKVDGTGSAAVVKVVSQAELYELNKTDSDS	68
							consensus	** * * * * ** * * * * * * * * * * * * * * * * * * * * * * * *	

NTDB id 32	MW	RS07995	WP	000472256.1	SSIEDLIADGFIKEAQKTKSGETITIS.NGEAVAN	103
NTDB id 16	SA	RS07765	WP	000472256.1	SSIEDLIADGFIKEAQKTKSGETITIS.NGEAVAN	103
NTDB id 588	KW2	RS10845	WP	051303741.1	PNLSELLSAGMITCKQVTAYYDDYDQNKNEQRNFDD	116
NTDB id 164077	AU077	RS09230	WP	039670352.1	PTVDDLVVEVGYITSEKAQTYNAKK.....	97
NTDB id 460	GPW51	RS00790	WP	011921720.1	ATVADLQAKGYITEKQAKYATAKK.....	93
NTDB id 502	SM12261	RS00955	WP	000738641.1	ASLSKLQADGRITEEQAKAYKEYHAKCNTSQTIVAD.	107
NTDB id 272	KZH43	RS09255	WP	000738626.1	ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVND	108
NTDB id 231	SPD	RS09885	WP	000738626.1	ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVND	108
NTDB id 197	SPR	RS09420	WP	000738626.1	ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVND	108
NTDB id 162	SP	RS10370	WP	000738627.1	ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVND	108
NTDB id 530	SMSK321	RS09230	WP	000738643.1	ASLSKLQADGRITEEQAKAYNEYAKNGGANRKVND	108
NTDB id 431	SGO	RS09410	WP	012130940.1	ATLSKLVAAGNISQKQADSYPKAYYGNNSSETQAVAN	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