

NTDB id 32	MW	RS07995	WP	000472256.1	MFKFL..KKTQAFTLIEMLLVLLIISLLLILIIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id 16	SA	RS07765	WP	000472256.1	MFKFL..KKTQAFTLIEMLLVLLIISLLLILIIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id 588	KW2	RS10845	WP	051303741.1	MSKALSLVKIHGRKLCKQKQKAFTLIEMLLVLAIIISILILLFVPNLIKEAKVQKTEGAADVVKVVESQAQLYELDHDNDK	80
NTDB id 422299	G3T13	RS09895	WP	174519819.1	MKKKIVRAFTLVEMLVLLVLSVLILFFIPNLSKQKETVKTNGGAVVKKVNSQAELYRLDHT.DD	65
NTDB id 460	GPW51	RS00790	WP	01921720.1	MKKLVEMKVKGFTLVEMLVLLGIISILLFFVPNLSQKKDAIQKKGDGAADVVKVVESQMELEYELHD.EE	68
NTDB id 502	SM12261	RS00955	WP	000738641.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLLFFVPNLTQKKEAVNDKGKAAVVKVVESQAELYSLDKN.ED	72
NTDB id 272	KZH43	RS09255	WP	000738626.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLLFFVPNLTQKKEAVNDKGKAAVVKVVESQAELYSLDKN.ED	72
NTDB id 231	SPD	RS09885	WP	000738626.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLLFFVPNLTQKKEAVNDKGKAAVVKVVESQAELYSLDKN.ED	72
NTDB id 197	SPR	RS09420	WP	000738626.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLLFFVPNLTQKKEAVNDKGKAAVVKVVESQAELYSLDKN.ED	72
NTDB id 162	SP	RS10370	WP	000738627.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLLFFVPNLTQKKEAVNDKGKAAVVKVVESQAELYSLDKN.ED	72
NTDB id 530	SMSK321	RS09230	WP	000738643.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLLFFVPNLTQKKEAVNDKGKAAVVKVVESQAELYSLDKN.ED	72
NTDB id 431	SGO	RS09410	WP	012130940.1	MNKLLKLLRVKAFTLVEMLVLLIISVLLMFFVPNLTQKKEAVSDTGNAAVVKVVESQAELYELKNTGDQ	69
NTDB id 396	FSA28	RS09395	WP	002263437.1	MKKIKRVSVRAFTLIEMLVLLIISILLFFVPNLTQKQDKVKDTGGSAVVKVVESQAELYELNKT.DS	68
NTDB id 387	SMU	RS09025	WP	002263437.1	MKKIKRVSVRAFTLIEMLVLLIISILLFFVPNLTQKQDKVKDTGGSAVVKVVESQAELYELNKT.DS	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	PNLSELLSACGMITQKQVTAYDDYDQNKNEQRNFDD	116
NTDB	id	422299	G3T13	RS09895	WP	174519819.1	VTLGKLVSNGNITQKQADAYNEYKKNTTESREIPS	101
NTDB	id	460	GPW51	RS00790	WP	011921720.1	ATVADLQAKGYITEKQAKQYATAKK.....	93
NTDB	id	502	SM12261	RS00955	WP	000738641.1	ASLSKQLADGRITEEQAKAYKEYHAKQNTSQTIVAD.	107
NTDB	id	272	KZH43	RS09255	WP	000738626.1	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRKVND	108
NTDB	id	231	SPD	RS09885	WP	000738626.1	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRKVND	108
NTDB	id	197	SPR	RS09420	WP	000738626.1	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRKVND	108
NTDB	id	162	SP	RS10370	WP	000738627.1	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRKVND	108
NTDB	id	530	SMSK321	RS09230	WP	000738643.1	ASLSKQLADGRITEEQAKAYNEYEYAKNGGANRKVND	108
NTDB	id	431	SGO	RS09410	WP	012130940.1	ATLSKLVAAGNISQKQADSYKAYYGKNNSETQAVAN	105
NTDB	id	396	FSA28	RS09395	WP	002263437.1	ATLSKLVASGDITQEQANSYKDYYAKHPSKTRHVN	104
NTDB	id	387	SMU	RS09025	WP	002263437.1	ATLSKLVASGDITQEQANSYKDYYAKHPSKTRHVN	104
consensus							*** *! ***! !****!***** ** *	

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