

NTDB id	32	MW	RS07995	WP	000472256.1	MFKFL..KKTQAFTLIEMLVLLIIISLILLILIIPNIAKQTAHIQSTGCNAQVKVMVNSQIEAYALKHNRNP	68
NTDB id	16	SA	RS07765	WP	000472256.1	MFKFL..KKTQAFTLIEMLVLLIIISLILLILIIPNIAKQTAHIQSTGCNAQVKVMVNSQIEAYALKHNRNP	68
NTDB id	1055082	ACE8TC	RS01055	WP	029487321.1	MRKLKHQGFTLVEMLVLLVISILLLFVPNLQAQRIVIDEKGNAATIVKVVEITQIELQLNEN.RT	65
NTDB id	588	KW2	RS10845	WP	051303741.1	MSKALSIVKIHGRLCQQKKQKAFTLIEMLVLAISILLLFFVPNLKEKAQVCKTGEEAAVVKVVESQAELYELDHDNDK	80
NTDB id	460	GPW51	RS00790	WP	011921720.1	MKKLIVEMKVGFTLVEMLVVLGIISLILLFFVPNLSQQKDAIQKKGDAAVVKVSVESQMELYELEHDEE	68
NTDB id	502	SM12261	RS00955	WP	000738641.1	MKKMMTFLKKAQVKAFTLVEMLVLLIIISVLFFFVPNLTQKQEAVNDKGKAAAVVKVSVESQAELYSLEKN.ED	72
NTDB id	162	SP	RS10370	WP	000738627.1	MKKMMTFLKKAQVKAFTLVEMLVLLIIISVLFFFVPNLTQKQEAVNDKGKAAAVVKVSVESQAELYSLEKN.ED	72
NTDB id	272	KZH43	RS09255	WP	000738626.1	MKKMMTFLKKAQVKAFTLVEMLVLLIIISVLFFFVPNLTQKQEAVNDKGKAAAVVKVSVESQAELYSLEKN.ED	72
NTDB id	231	SPD	RS09885	WP	000738626.1	MKKMMTFLKKAQVKAFTLVEMLVLLIIISVLFFFVPNLTQKQEAVNDKGKAAAVVKVSVESQAELYSLEKN.ED	72
NTDB id	197	SPR	RS09420	WP	000738626.1	MKKMMTFLKKAQVKAFTLVEMLVLLIIISVLFFFVPNLTQKQEAVNDKGKAAAVVKVSVESQAELYSLEKN.ED	72
NTDB id	530	SMSK321	RS09230	WP	000738643.1	MKKMMTFLKKAQVKAFTLVEMLVLLIIISVLFFFVPNLTQKQEAVNDKGKAAAVVKVSVESQAELYSLEKN.ED	72
NTDB id	431	SGO	RS09410	WP	012130940.1	MNKLKKLRVKAFTLVEMLVLLIIISVLMLLFVPNLTQKQEAVSDTGNAAAVVKVSVESQAELYELKNTGDQ	69
NTDB id	396	FSA28	RS09395	WP	002263437.1	MKKIKRVSVRAFTLIEMLVLLIIISLILLFFVPNLTQKQDKVKDDTGGSAVVKVSVESQAELYELNKTT.DS	68
NTDB id	387	SMU	RS09025	WP	002263437.1	MKKIKRVSVRAFTLIEMLVLLIIISLILLFFVPNLTQKQDKVKDDTGGSAVVKVSVESQAELYELNKTT.DS	68
consensus						* * * * *	68

NTDB	id	32	MW	RS07995	WP	000472256.1	SSIEDLIADGFIKEAQKTKCSGETITISN.GEAVAN	103
NTDB	id	16	SA	RS07765	WP	000472256.1	SSIEDLIADGFIKEAQKTKCSGETITISN.GEAVAN	103
NTDB	id	1055082	ACE8TC	RS01055	WP	029487321.1	PTRQELLAGNYVTEEQYAIYLAHRNE.....	91
NTDB	id	588	KW2	RS10845	WP	051303741.1	PNLSELLSAGMITCKQVTAAYDDYDQNKNEQRNFDD	116
NTDB	id	460	GPW51	RS00790	WP	011921720.1	ATVADLQAKGYITEKQAKQYATAKK.....	93
NTDB	id	502	SM12261	RS00955	WP	000738641.1	ASLSKLQADGRITEEQAKAYKEYHAKQNTSQTVD.	107
NTDB	id	162	SP	RS10370	WP	000738627.1	ASLRKLQADGRITEEQAKAYKEYNDKNGGANRKVND	108
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	530	SMSK321	RS09230	WP	000738643.1	ASLSKLQADGRITEEQAKAYNEYAKNGGANRKVND	108
NTDB	id	431	SGO	RS09410	WP	012130940.1	ATLSKLVAAGNISQKQADS YKAYYGKNNSETQAVAN	105
NTDB	id	396	FSA28	RS09395	WP	002263437.1	ATLSKLVASGDITQEANSYKDYYAKHPSKTRHVAN	104
NTDB	id	387	SMU	RS09025	WP	002263437.1	ATLSKLVASGDITQEANSYKDYYAKHPSKTRHVAN	104
consensus							*** *! *** **!*! **** * * *	

- X non conserved
- X similar
- X $\geq 50\%$ conserved