

NTDB id	32	MW	RS07995	WP	000472256.1	MFKFL..KKTQAFTLIEMLVLLLIISLLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id	16	SA	RS07765	WP	000472256.1	MFKFL..KKTQAFTLIEMLVLLLIISLLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id	588	KW2	RS10845	WP	051303741.1	MSKALSLV ^V KIHGRKLCKKKQAFTL IEMLVLAIIISILILLFVPNLKEKAQVKGTGEAAVVKVVSQAQLYELDHDNDK	80
NTDB id	460	GPW51	RS00790	WP	011921720.1	MKKLIVEMKVKCFTLVEMLVLVGIIISLLLLFVPNLSQQKDIAIKKGDAAVVKVVESQMELYELEHDEE	68
NTDB id	502	SM12261	RS00955	WP	000738641.1	MKKMMTFLKKAKVKAFTLVEMLVLLLIISVLLLFVPNLTKQKEAVNDKGKAAAVKVVSQAELYSLDKNEED	72
NTDB id	272	KZH43	RS09255	WP	000738626.1	MKKMMTFLKKAKVKAFTLVEMLVLLLIISVLFLLFVPNLTKQKEAVNDKGKAAAVKVVSQAELYSLDKNEED	72
NTDB id	231	SPD	RS09885	WP	000738626.1	MKKMMTFLKKAKVKAFTLVEMLVLLLIISVLFLLFVPNLTKQKEAVNDKGKAAAVKVVSQAELYSLDKNEED	72
NTDB id	197	SPR	RS09420	WP	000738626.1	MKKMMTFLKKAKVKAFTLVEMLVLLLIISVLFLLFVPNLTKQKEAVNDKGKAAAVKVVSQAELYSLDKNEED	72
NTDB id	162	SP	RS10370	WP	000738627.1	MKKMMTFLKKAKVKAFTLVEMLVLLLIISVLFLLFVPNLTKQKEAVNDKGKAAAVKVVSQAELYSLDKNEED	72
NTDB id	530	SMSK321	RS09230	WP	000738643.1	MKKMMTFLKKAKVKAFTLVEMLVLLLIISVLLLFVPNLTKQKEAVNDKGKAAAVKVVSQAELYSLDKNEED	72
NTDB id	431	SGO	RS09410	WP	012130940.1	MNKLLKLRVKAFTLVEMLVLLLIISVLMLLFVPNLTKQKEAVSDTGNAAVVKVVSQAELYELKNITGDQ	69
NTDB id	302450	DTA40	RS08820	WP	002946126.1	MKLMLKKLNNAVKLRAFTL IEMLVLLLIISILLLLFPVNLSSQKQDSVKETGNAAVVKVVDQAELYEMKNNK.T	72
NTDB id	396	FSA28	RS09395	WP	002263437.1	MKKI ^R KRVSVRAFTL IEMLVLLLIISILLLLFPVNLTKQKDKVKDTGGSAVVKVVSQAELYELNKT.DS	68
NTDB id	387	SMU	RS09025	WP	002263437.1	MKKI ^R KRVSVRAFTL IEMLVLLLIISILLLLFPVNLTKQKDKVKDTGGSAVVKVVSQAELYELNKT.DS	68
consensus						* * * * *	

NTDB id	32 MW RS07995 WP 000472256.1	SSI E DLIADGFIKEAQKTKSGETIT I .SNGEAVAN	103
NTDB id	16 SA RS07765 WP 000472256.1	SSI E DLIADGFIKEAQKTKSGETIT I .SNGEAVAN	103
NTDB id	588 KW2 RS10845 WP 051303741.1	PNLSELLSACMITQKVTA YDDYDQ NKNEQRNFDD	116
NTDB id	460 GPW51 RS00790 WP 011921720.1	ATVADLQAKGYITEKQAKKYATAKK.....	93
NTDB id	502 SM12261 RS00955 WP 000738641.1	ASLSKLQADGRITEEQAKAYKEYHAKNNTSQTVAD.	107
NTDB id	272 KZH43 RS09255 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVN D	108
NTDB id	231 SPD RS09885 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVN D	108
NTDB id	197 SPR RS09420 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVN D	108
NTDB id	162 SP RS10370 WP 000738627.1	ASLRKLQADGRITEEQAKAYKEYNDKNGGANRKVN D	108
NTDB id	530 SMSK321 RS09230 WP 000738643.1	ASLSKLQADGRITEEQAKAYNEYAKNGGANRKVN D	108
NTDB id	431 SGO RS09410 WP 012130940.1	ATLSKLVAAGNISQKQADSYPKAYYGKNNSETQAVAN	105
NTDB id	302450 DTA40 RS08820 WP 002946126.1	ASLAALVSEGGITQKQADSYN D YAKHGGSRSVAN	108
NTDB id	396 FSA28 RS09395 WP 002263437.1	ATLSKLVASGDITQEANSYKDYYAKHPSKTRHVAN	104
NTDB id	387 SMU RS09025 WP 002263437.1	ATLSKLVASGITQEANSYKDYAKHPSKTRHVAN	104
consensus		*** *! *** ! ****! ***** ** *	

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