

NTDB	id	32	MW	RS07995	WP	000472256.1		MFKFL..KKTQAFTLIEMLVLLIIISLLLILIIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB	id	16	SA	RS07765	WP	000472256.1		MFKFL..KKTQAFTLIEMLVLLIIISLLLILIIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB	id	588	KW2	RS10845	WP	051303741.1		MSKALSLVKIHGRKLCKKKQKAFTLIELVLVLAIIISILILLFVPNLIKEKAQVKQTGEAAVVKVVSQAQLYELDHDNDK	80
NTDB	id	174486	A2G56	RS06510	WP	237334442.1		MLLVLLIISVLMFLFVPNLISKQKAADVDEKGAAVVKVVSQMELYENED.VK	52
NTDB	id	460	GPW51	RS00790	WP	01921720.1		MKKLIVEMKVKGFTLEVMLVVLGIISLLLLFVPNLSQQKDAIQKKGDAAAVVKVVSQMELYELED.EE	68
NTDB	id	502	SM12261	RS00955	WP	000738641.1		MKKMMTFLKKAKVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVSQAELYSLDKN.ED	72
NTDB	id	272	KZH43	RS09255	WP	000738626.1		MKKMMTFLKKAKVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVSQAELYSLDKN.ED	72
NTDB	id	231	SPD	RS09885	WP	000738626.1		MKKMMTFLKKAKVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVSQAELYSLDKN.ED	72
NTDB	id	197	SPR	RS09420	WP	000738626.1		MKKMMTFLKKAKVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVSQAELYSLDKN.ED	72
NTDB	id	162	SP	RS10370	WP	000738627.1		MKKMMTFLKKAKVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVSQAELYSLDKN.ED	72
NTDB	id	530	SMSK321	RS09230	WP	000738643.1		MKKMMTFLKKAKVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVSQAELYSLDKN.ED	72
NTDB	id	431	SGO	RS09410	WP	012130940.1		MNKLLKLLRVKAFTLVEMLVLLIISVLMFLFVPNLTKQKEAVSDTGNAAVVKVVSQAELYELKNKTGDQ	69
NTDB	id	396	FSA28	RS09395	WP	002263437.1		MKKITKRVSVRRAFTLIEMLVLLIIISILLLFVPNLTKQKDVKVDGTGSAAVVKVVSQAELYELNKTD.SS	68
NTDB	id	387	SMU	RS09025	WP	002263437.1		MKKITKRVSVRRAFTLIEMLVLLIIISILLLFVPNLTKQKDVKVDGTGSAAVVKVVSQAELYELNKTD.SS	68
							consensus	* * * * *	

NTDB id	32 MW RS07995 WP 000472256.1	SSI E DLIADGFIKEAQKTKSGETITISN.GEAVAN	103
NTDB id	16 SA RS07765 WP 000472256.1	SSI E DLIADGFIKEAQKTKSGETITISN.GEAVAN	103
NTDB id	588 KW2 RS10845 WP 051303741.1	PNLSELLSACMITQKQVTA YD DYYDQNKNEQRNFDD	116
NTDB id	174486 A2G56 RS06510 WP 237334442.1	PSIQQLVGEIGYITQKQADYA A KTPK.....	77
NTDB id	460 GPW51 RS00790 WP 01921720.1	ATVADLQA K G I YITEKQAKQAYATAKK.....	93
NTDB id	502 SM12261 RS00955 WP 000738641.1	ASLSKLQADGRITEEQAKAYKE Y HAKQNTSQTVAD.	107
NTDB id	272 KZH43 RS09255 WP 000738626.1	ASLRKLQADGRITEEQAKAYKE Y HDKNGGANRKVN	108
NTDB id	231 SPD RS09885 WP 000738626.1	ASLRKLQADGRITEEQAKAYKE Y HDKNGGANRKVN	108
NTDB id	197 SPR RS09420 WP 000738626.1	ASLRKLQADGRITEEQAKAYKE Y HDKNGGANRKVN	108
NTDB id	162 SP RS10370 WP 000738627.1	ASLRKLQADGRITEEQAKAYKE Y HDKNGGANRKVN	108
NTDB id	530 SMSK321 RS09230 WP 000738643.1	ASLSKLQADGRITEEQAKAYNE Y YAKNGGANRKVN	108
NTDB id	431 SGO RS09410 WP 012130940.1	ATLSKLVAAGNISQKQADSYKAYYGKNNSETQAVAN	105
NTDB id	396 FSA28 RS09395 WP 002263437.1	ATLSKLVASGDITQE Q ANSYKDYYAKHPSKTRHVAN	104
NTDB id	387 SMU RS09025 WP 002263437.1	ATLSKLVASGDITQE Q ANSYKDYYAKHPSKTRHVAN	104
consensus		*** * ! ** ! ! *** ! **** * * *	

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