

NTDB id 588 KW2 RS10845 WP 051303741.1	MSKALSIVKIHGRKIQCKKQKQFTLIEMLIVLAIIISILLFVFPNLIKQKQAVQKKTGEAAVVKVSVESQAQLYELDHDNDK	80
NTDB id 396 FSA28 RS09395 WP 002263437.1	MKKKIKRVSVRAFTLIEMLVLLIISILLLLFVFPNLTQKQDKVKDTGGSAAVVKVSVESQAELYELNKTDS	68
NTDB id 387 SMU RS09025 WP 002263437.1	MKKKIKRVSVRAFTLIEMLVLLIISILLLLFVFPNLTQKQDKVKDTGGSAAVVKVSVESQAELYELNKTDS	68
NTDB id 502 SM12261 RS00955 WP 000738641.1	MKKMMTFLEKKAQKVAFTLIEMLVLLIISVLLLLFVFPNLTQKQEAVNDKGKAAAVVKVSVESQAELYSLDKNE.D	72
NTDB id 272 KZH43 RS09255 WP 000738626.1	MKKMMTFLEKKAQKVAFTLIEMLVLLIISVLFLLFVFPNLTQKQEAVNDKGKAAAVVKVSVESQAELYSLDKNE.D	72
NTDB id 231 SPD RS09885 WP 000738626.1	MKKMMTFLEKKAQKVAFTLIEMLVLLIISVLFLLFVFPNLTQKQEAVNDKGKAAAVVKVSVESQAELYSLDKNE.D	72
NTDB id 197 SPR RS09420 WP 000738626.1	MKKMMTFLEKKAQKVAFTLIEMLVLLIISVLFLLFVFPNLTQKQEAVNDKGKAAAVVKVSVESQAELYSLDKNE.D	72
NTDB id 162 SP RS10370 WP 000738627.1	MKKMMTFLEKKAQKVAFTLIEMLVLLIISVLFLLFVFPNLTQKQEAVNDKGKAAAVVKVSVESQAELYSLDKNE.D	72
NTDB id 530 SMSK321 RS09230 WP 000738643.1	MKKMMTFLEKKAQKVAFTLIEMLVLLIISVLLLLFVFPNLTQKQEAVNDKGKAAAVVKVSVESQAELYSLDKNE.D	72
NTDB id 98 BSU 24710 NP 390351.1	MNEKGFTLIEMLIVLFIISILLITTPNVTKHNQTIQKKGCEGLQNMVKAQMTAFELDHEGQT	63
NTDB id 1094223 L1F34 RS06690 WP 070045385.1	MKNIKFFKKEEGFTLIEMLIVLLIISILLIIITPNISKQSETVKSTGCCQAQVKMVAGQVEAYTLKHDAPK	70
NTDB id 32 MW RS07995 WP 000472256.1	MFKFLKKTQAFTLIEMLVLLIISILLIITIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id 16 SA RS07765 WP 000472256.1	MFKFLKKTQAFTLIEMLVLLIISILLIITIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
consensus	* * * * *	

NTDB	id	588	KW2	RS10845	WP	051303741.1	PNLSELLSAGMITKQVTA	YDYYDQNKNEQ	RNFDD	116
NTDB	id	396	FSA28	RS09395	WP	002263437.1	ATLSKLVASGDITQE	QANSYKDYYAKHPSK	TRHVN	104
NTDB	id	387	SMU	RS09025	WP	002263437.1	ATLSKLVASGDITQE	QANSYKDYYAKHPSK	TRHVN	104
NTDB	id	502	SM12261	RS00955	WP	000738641.1	ASLSKLQADGRITEE	QAKAYKEYHAKQNTS	SQTVD	107
NTDB	id	272	KZH43	RS09255	WP	000738626.1	ASLRKLQADGRITEE	QAKAYKEYHDKNGG	ANRKVD	108
NTDB	id	231	SPD	RS09885	WP	000738626.1	ASLRKLQADGRITEE	QAKAYKEYHDKNGG	ANRKVD	108
NTDB	id	197	SPR	RS09420	WP	000738626.1	ASLRKLQADGRITEE	QAKAYKEYHDKNGG	ANRKVD	108
NTDB	id	162	SP	RS10370	WP	000738627.1	ASLRKLQADGRITEE	QAKAYKEYNDKNGG	ANRKVD	108
NTDB	id	530	SMSK321	RS09230	WP	000738643.1	ASLSKLQADGRITEE	QAKAYNEYAKNGG	ANRKVD	108
NTDB	id	98	BSU	24710	NP	390351.1	PSLADLQSEGYVKKD	AVCPNGKRIITG	GEVKEH	98
NTDB	id	1094223	L1F34	RS06690	WP	070045385.1	NSINDLVSDGFIKEN	QLTCKSGQNIVLNN	CEVVAQ	105
NTDB	id	32	MW	RS07995	WP	000472256.1	SSIEDLIADGFIKEA	QKTCKSGETITISN	GEAVAN	103
NTDB	id	16	SA	RS07765	WP	000472256.1	SSIEDLIADGFIKEA	QKTCKSGETITISN	GEAVAN	103
consensus							*** * ! * * ! * * * * * * * * * * * * *			

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