

NTDB	id 32 MW RS07995 WP 000472256.1		MFKFL..KKTQAFTLIEMLLVLLIISLLLILIIIPNIAKQTAHIQSTGCNAQVKMNVNSQIEAYALKHNRP	68
NTDB	id 16 SA RS07765 WP 000472256.1		MFKFL..KKTQAFTLIEMLLVLLIISLLLILIIIPNIAKQTAHIQSTGCNAQVKMNVNSQIEAYALKHNRP	68
NTDB	id 588 KW2 RS10845 WP 051303741.1	MSKALSLV	KIHGRKLCKKKKAFTLIEMLIVLAIIISTILLFVPVNLIKKEAKVQKGTGEAAVVKVVSQAQLYELDHDNDK	80
NTDB	id 460 GPW51 RS00790 WP 011921720.1		MKKLVEMKVGCTFLVEMLVLVIIISLLLLFVPNLSSQQKDATQKKGDAAVVKVVSQMELYELEH.D.EE	68
NTDB	id 502 SM12261 RS00955 WP 000738641.1		MKKMMTFLKKAKVKAFTLVEMLVLLIISVLFLLFVPNLTKQEAEVNDKGKAAVVKVVSQAELYSLDKN.ED	72
NTDB	id 272 KZH43 RS09255 WP 000738626.1		MKKMMTFLKKAKVKAFTLVEMLVLLIISVLFLLFVPNLTKQEAEVNDKGKAAVVKVVSQAELYSLDKN.ED	72
NTDB	id 231 SPD RS09885 WP 000738626.1		MKKMMTFLKKAKVKAFTLVEMLVLLIISVLFLLFVPNLTKQEAEVNDKGKAAVVKVVSQAELYSLDKN.ED	72
NTDB	id 197 SPR RS09420 WP 000738626.1		MKKMMTFLKKAKVKAFTLVEMLVLLIISVLFLLFVPNLTKQEAEVNDKGKAAVVKVVSQAELYSLDKN.ED	72
NTDB	id 162 SP RS10370 WP 000738627.1		MKKMMTFLKKAKVKAFTLVEMLVLLIISVLFLLFVPNLTKQEAEVNDKGKAAVVKVVSQAELYSLDKN.ED	72
NTDB	id 530 SMSK321 RS09230 WP 000738643.1		MKKMMTFLKKAKVKAFTLVEMLVLLIISVLFLLFVPNLTKQEAEVNDKGKAAVVKVVSQAELYSLDKN.ED	72
NTDB	id 431 SGO RS09410 WP 012130940.1		MNKLKLRVKAFTLVEMLVLLIISVMLLFVPNLTKQEAEVSDTGNAAVVKVVSQAELYELKN TG D Q	69
NTDB	id 240576 CG712 RS05265 WP 002946126.1		MKLM LKKLN AVK LR AFTL I E M LV V LL II SI L L L L F VP N L SK Q KD SV K ET G NA A V VK V VD SQ AE LY EM KN . KT	72
NTDB	id 396 FSA28 RS09395 WP 002263437.1		MKKIKRVSVRAFTLIEMLVLLIISILLFFVPNLTKQKDKVKDTGG SAVVKVVSQAELYELNKT DS	68
NTDB	id 387 SMU RS09025 WP 002263437.1		MKKIKRVSVRAFTLIEMLVLLIISILLFFVPNLTKQKDKVKDTGG SAVVKVVSQAELYELNKT DS	68
consensus		** * *	* ** * * * * * * * * * * * * * * * * * * * * * * * * * * * *	

NTDB id	32 MW RS07995 WP 000472256.1	SSTIEDLIADGFIKEAQKTKSGETIT.I.SNGEAVAN	103
NTDB id	16 SA RS07765 WP 000472256.1	SSTIEDLIADGFIKEAQKTKSGETIT.I.SNGEAVAN	103
NTDB id	588 KW2 RS10845 WP 051303741.1	PNLSELLSACMITQKQVTA YDD YDQNKNEQRNFDD	116
NTDB id	460 GPW51 RS00790 WP 011921720.1	ATVADLQAKGYITEKQAKQYATAKK.....	93
NTDB id	502 SM12261 RS00955 WP 000738641.1	ASLSKLQADGRITEEQAKAYKEYHYAKNNTSQTVAD.	107
NTDB id	272 KZH43 RS09255 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVN	108
NTDB id	231 SPD RS09885 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVN	108
NTDB id	197 SPR RS09420 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVN	108
NTDB id	162 SP RS10370 WP 000738627.1	ASLRKLQADGRITEEQAKAYKEYNDKNGGANRKVN	108
NTDB id	530 SMSK321 RS09230 WP 000738643.1	ASLSKLQADGRITEEQAKAYNEYAKNGGANRKVN	108
NTDB id	431 SGO RS09410 WP 012130940.1	ATLSKLVAAGNISQKQADSYPKAYYGKNNSETQAVAN	105
NTDB id	240576 CG712 RS05265 WP 002946126.1	ASLAALVSEGGITQKQADSINDYYAKHGGSRSVAN	108
NTDB id	396 FSA28 RS09395 WP 002263437.1	ATLSKLVASGDITQEANSYKDYYAKHPSKTRHVAN	104
NTDB id	387 SMU RS09025 WP 002263437.1	ATLSKLVASGITQEANSYKYDYAKHPSKTRHVAN	104
consensus		*** * ! ** ! ! **** ! ***** ** *	

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