

NTDB id	1250 GC085 RS13065 WP 027219989.1	. . MNNLQPLLA LNRMKRIG PRTVLKLQKRWPDNLNMFQLSSVDLEEAGLPSSLAQTIKNFDPDFIKTDLNWLSSENHHI	78
NTDB id	599 KW2 RS05940 WP 021037268.1	. MITNFD LFRWKKGAMTNLGVNKLLKFFRKYDRKISLRQM . . GGVAQVKSIPNFIEQYKNQDVVKILRADYKKFS S	72
NTDB id	525 SMSK321 RS06295 WP 000705298.1	MKITNYE IYKLKKSGLTNQQILKVLEYGENVDQELLGDI . . ADISGCNPAPVFMEFYFQIDDAHLEKEFKKTIP S	73
NTDB id	267 KZH43 RS05590 WP 000705306.1	MKITNYE IYKLKKSGLTNQQILKVLEYGENVDQELLGDI . . ADISGCNPAPVFMEFYFQIDDAHLSKEFQKFP S	73
NTDB id	226 SPD RS05990 WP 000705306.1	MKITNYE IYKLKKSGLTNQQILKVLEYGENVDQELLGDI . . ADISGCNPAPVFMEFYFQIDDAHLSKEFQKFP S	73
NTDB id	192 SPR RS05715 WP 000705306.1	MKITNYE IYKLKKSGLTNQQILKVLEYGENVDQELLGDI . . ADISGCNPAPVFMEFYFQIDDAHLSKEFQKFP S	73
NTDB id	157 SP RS06205 WP 000705306.1	MKITNYE IYKLKKSGLTNQQILKVLEYGENVDQELLGDI . . ADISGCNPAPVFMEFYFQIDDAHLSKEFQKFP S	73
NTDB id	497 SM12261 RS05565 WP 000705318.1	MKITNYE IYKLKKSGLTNQQVLA VLEYGENVDQELLGDI . . AEISGCNPAPVFMEFYFQIDDAHLEKEFQKFP S	73
NTDB id	66106 M3B RS05435 WP 011054501.1	. . MNHFE LYKLKKA GLTNLN ILNI LDY QKHQE KSLSRDM . . AVVSGCKHP SHFT EAYKQLD IQKLKMEFKQFP S	71
NTDB id	407 SMU RS04605 WP 002262865.1	. . MDNFQ LFKLKKAGLTNL NII NI IDYEERTQSLSLRDM . . AVVS KNKKP LI FMEHYKNLD SKAL RKEFNRF S	71
consensus		* * * * *	*

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NTDB id	1250	GC085	RS13065	WP	027219989.1	DVLAIPGSIHNPLARGCHYLQGGAKLVTSVADVLDELKIEHHQFTSNKPIFSLASGKENLVKFIFGETTTIDQIIDRSGYGM EQ	333
NTDB id	599	KW2	RS05940	WP	021037268.1	DIFAIPGNIA DGTSDGCNHLIQGGAKLVYQAQDILEEYLYN.....	282
NTDB id	525	SMSK321	RS06295	WP	000705298.1	DVFAIPGSILDGLSDGCHH LIQE GAKLVTSGQDV LAEF EF	282
NTDB id	267	KZH43	RS05590	WP	000705306.1	DVFAIPGSILDGLSDGCHH LIQE GAKLVTSGQDV LAEF EF	282
NTDB id	226	SPD	RS05990	WP	000705306.1	DVFAIPGSILDGLSDGCHH LIQE GAKLVTSGQDV LAEF EF	282
NTDB id	192	SPR	RS05715	WP	000705306.1	DVFAIPGSILDGLSDGCHH LIQE GAKLVTSGQDV LAEF EF	282
NTDB id	157	SP	RS06205	WP	000705306.1	DVFAIPGSILDGLSDGCHH LIQE GAKLVTSGQDV LAEF EF	282
NTDB id	497	SM12261	RS05565	WP	000705318.1	DVFAIPGSILDGLSDGCHH LIQE GAKLVTSGQDV LAEF EF	282
NTDB id	66106	M3B	RS05435	WP	011054501.1	DIFAVPGNI LDGKS EGC LQLIK EGATC VTSGMD ILSEY	278
NTDB id	407	SMU	RS04605	WP	002262865.1	DVFVVPGNI LDGQS EGC HHL IQEG AK C ITSGFD I LNEFN F	280
consensus						!*****! !*****! !*****! !*****! !*****! !*****!	

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NTDB id	Accession	Sequence	Length
1250	GC085 RS13065 WP 027219989.1	VTSGLAELKGVIAVPGGYIRCEYER	361
599	KW2 RS05940 WP 021037268.1		282
525	SMSK321 RS06295 WP 000705298.1		282
267	KZH43 RS05590 WP 000705306.1		282
226	SPD RS05990 WP 000705306.1		282
192	SPR RS05715 WP 000705306.1		282
157	SP RS06205 WP 000705306.1		282
497	SM12261 RS05565 WP 000705318.1		282
66106	M3B RS05435 WP 011054501.1		278
407	SMU RS04605 WP 002262865.1		280
consensus			

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