

NTDB id 1250 GC085 RS13065 WP 027219989.1	..MNNLQLPLLA	LNRMKRIGP....RTVLKLQKRPDLNLMFQLSSV	DLEEAGLPSWLAQTIKNFPDPDFIKTDLNLWLSSS	73
NTDB id 599 KW2 RS05940 WP 021037268.1	.MITNFD....	LFRWKKAGMTNLGVNKLKLFRRKYDRKISLRQM..	GQVAQVKISIPNFIEQYKNQDVKKLRADYKKFS..	71
NTDB id 525 SMSK321 RS06295 WP 000705298.1	MKITNYE....	IYKLKKSGLTNQQILKVLEYGENVDQELLGDI..	ADISGCRNPAVFMERYFQIDDAHLEKEFKKIP..	72
NTDB id 267 KZH43 RS05590 WP 000705306.1	MKITNYE....	IYKLKKSGLTNQQILKVLEYGENVDQELLGDI..	ADISGCRNPAVFMERYFQIDDAHLSKEFQKFP..	72
NTDB id 226 SPD RS05990 WP 000705306.1	MKITNYE....	IYKLKKSGLTNQQILKVLEYGENVDQELLGDI..	ADISGCRNPAVFMERYFQIDDAHLSKEFQKFP..	72
NTDB id 192 SPR RS05715 WP 000705306.1	MKITNYE....	IYKLKKSGLTNQQILKVLEYGENVDQELLGDI..	ADISGCRNPAVFMERYFQIDDAHLSKEFQKFP..	72
NTDB id 157 SP RS06205 WP 000705306.1	MKITNYE....	IYKLKKSGLTNQQILKVLEYGENVDQELLGDI..	ADISGCRNPAVFMERYFQIDDAHLSKEFQKFP..	72
NTDB id 497 SM12261 RS05565 WP 000705318.1	MKITNYE....	IYKLKKSGLTNQQVLA	AVLEYGENVDQELLGDI..AEISGCRNPAVFMERYFQIDDAHLEKEFQKFP..	72
NTDB id 308319 DZV55 RS05025 WP 002995009.1	..MNHFE....	LYKLKKAAGLTNKNILN	IILDYQEHQEKSLSLRDM..AVVSGCKHPSHFIEAYKQLDIQKLKIEFKQFP..	70
NTDB id 407 SMU RS04605 WP 002262865.1	..MDNFG....	LFLKKAAGLTNKNILN	IIDIYEERTQKSLSLRDM..AVVSKNKKPLIFMEHYKNLDSKALRKEFNRF..	70
consensus	*****	*****!*****	*****	*****

NTDB id	1250 GC085 RS13065 WP 027219989.1	ENHHIL	TWDSPLYPALKEIADPPFI	LTKELG	LAIVGSRNASVTGNENARAFSREISRHGV	SIVSGLALGID	153
NTDB id	599 KW2 RS05940 WP 021037268.1		SFSILDDLYPERLREIYNPPVLI	FYQGNIDLKPNKLAFVGSRLAGQS	GIKSVQKIVTELNQ	.SFTIVSGLAKGID	146
NTDB id	525 SMSK321 RS06295 WP 000705298.1		SFSILDDCYPWDLSEIYDAPVLL	LFYKGNLDLLKFPKVAVVGSR	SCSKQGAKSVEKI	IQQLEN.ELVIVSGLAKGID	147
NTDB id	267 KZH43 RS05590 WP 000705306.1		SFSILDDCYPWDLSEIYDAPVLL	LFYKGNLDLLKFPKVAVVGSRACSK	QGAKSVEKVIQGLE	N.ELVIVSGLAKGID	147
NTDB id	226 SPD RS05990 WP 000705306.1		SFSILDDCYPWDLSEIYDAPVLL	LFYKGNLDLLKFPKVAVVGSRACSK	QGAKSVEKVIQGLE	N.ELVIVSGLAKGID	147
NTDB id	192 SPR RS05715 WP 000705306.1		SFSILDDCYPWDLSEIYDAPVLL	LFYKGNLDLLKFPKVAVVGSRACSK	QGAKSVEKVIQGLE	N.ELVIVSGLAKGID	147
NTDB id	157 SP RS06205 WP 000705306.1		SFSILDDCYPWDLSEIYDAPVLL	LFYKGNLDLLKFPKVAVVGSRACSK	QGAKSVEKVIQGLE	N.ELVIVSGLAKGID	147
NTDB id	497 SM12261 RS05565 WP 000705318.1		SFSILDDCYPWDLSEIYDAPVLL	LFYKGNLDLLKFPKVAVVGSRACSK	QGAKSVEKVIQGLE	N.ELVIVSGLAKGID	147
NTDB id	308319 DZV55 RS05025 WP 002995009.1		STILDKHYPMALKEIYNPPVLL	FFQGNLDLLEKPKLAIVGSRRSSDT	GVKSVRKILKELGN	.RFVIVSGLARGID	145
NTDB id	407 SMU RS04605 WP 002262865.1		SLSIDKEYPLFKNIYNPPVLL	FYQGDLLARPKLAVVGSRNASQM	GVAVKKKIIQDLSK	.QFVIIISGLARGID	145
consensus			* * * * *	! ! ! ! !	* * * * *	! ! ! ! !	* * * * *

[illegible]

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NTDB id 1250 GC085 RS13065 WP 027219989.1
NTDB id 599 KW2 RS05940 WP 021037268.1
NTDB id 525 SMSK321 RS06295 WP 000705298.1
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NTDB id 192 SPR RS05715 WP 000705306.1
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NTDB id 308319 DZV55 RS05025 WP 002995009.1
NTDB id 407 SMU RS04605 WP 002262865.1
consensus



X non conserved
X similar
X ≥ 50% conserved