

logo

NTDB id 1250 GC085 RS13065 WP 027219989.1	..MNNLQLPLLA	LNRMKRIG....	PRTVLKLQKRPDLNLMFQLSSV	DLEEAGLPSWLAQTIKNFDPDFIKTDLNLWLSSS	73
NTDB id 599 KW2 RS05940 WP 021037268.1	.MITNFD....	LFRWKKAGMTNLGVNKLKLF	FRKYDRKISLRQM..	GQVAQVKSIPNFIEQYKNQDVVKKL	RADYKKFS.. 71
NTDB id 525 SMSK321 RS06295 WP 000705298.1	MKITNYE....	IYKLKKSGLTNQQILKVLEYGENVDQELL	LGDII..	ADISGCRNPAVFMERYFQIDDAHLEKEFKKIP..	72
NTDB id 267 KZH4 RS05590 WP 000705306.1	MKITNYE....	IYKLKKSGLTNQQILKVLEYGENVDQELL	LGDII..	ADISGCRNPAVFMERYFQIDDAHLSKEFQKFP..	72
NTDB id 226 SPD RS05990 WP 000705306.1	MKITNYE....	IYKLKKSGLTNQQILKVLEYGENVDQELL	LGDII..	ADISGCRNPAVFMERYFQIDDAHLSKEFQKFP..	72
NTDB id 192 SPR RS05715 WP 000705306.1	MKITNYE....	IYKLKKSGLTNQQILKVLEYGENVDQELL	LGDII..	ADISGCRNPAVFMERYFQIDDAHLSKEFQKFP..	72
NTDB id 157 SP RS06205 WP 000705306.1	MKITNYE....	IYKLKKSGLTNQQILKVLEYGENVDQELL	LGDII..	ADISGCRNPAVFMERYFQIDDAHLSKEFQKFP..	72
NTDB id 497 SM12261 RS05565 WP 000705318.1	MKITNYE....	IYKLKKSGLTNQQVLAVLEYGENVDQELL	LGDII..	AEISGCRNPAVFMERYFQIDDAHLEKEFQKFP..	72
NTDB id 341595 ETT61 RS04610 WP 111693218.1	..MNHFE....	LYKLKKAAGLTNKNILNILDYQKHQEKSLSLRDM..	AVVSGCKHPSHFIEAYKQLDIQNLM	MEFKQFP..	70
NTDB id 407 SMU RS04605 WP 002262865.1	..MDNFG....	LPFLKKAAGLTNKNILNIIIDYEERTQKSLSLRDM..	AVVSKNKKPLIFMEHYKNLDSKALR	KEFNRF..	70
consensus	*****	*****!*****	*****	*****!*****	*****

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NTDB id	1250 GC085 RS13065 WP 027219989.1		ENHHILTWDSPLYPALKEIADPPFIYAKGELSALTQPSLATIVGSRNA SVTG NENARAFSREISRHGVS IVSGLA LGID	153
NTDB id	599 KW2 RS05940 WP 021037268.1		SFSILD DDL YPERL L REIYN P PVLI FYQGNI DL LNPK LA FV GSRL AG QS GIKSV QK I VTELNQ . SFT IV SG LA KG ID	146
NTDB id	525 SMSK321 RS06295 WP 000705298.1		SFSILDDCYPWDLSEIYDAPVLLFYKG NL DLL KFPKV AV VGSR SC SKQG AKS VEKI IQGLEN . ELVI VS GLA KG ID	147
NTDB id	267 KZH43 RS05590 WP 000705306.1		SFSILDDCYPWDLSEIYDAPVLLFYKG NL DLL KFPKV AV VGSRACSKQGA KSVEK VIQGLEN . ELVI VS GLA KG ID	147
NTDB id	226 SPD RS05990 WP 000705306.1		SFSILDDCYPWDLSEIYDAPVLLFYKG NL DLL KFPKV AV VGSRACSKQGA KSVEK VIQGLEN . ELVI VS GLA KG ID	147
NTDB id	192 SPR RS05715 WP 000705306.1		SFSILDDCYPWDLSEIYDAPVLLFYKG NL DLL KFPKV AV VGSRACSKQGA KSVEK VIQGLEN . ELVI VS GLA KG ID	147
NTDB id	157 SP RS06205 WP 000705306.1		SFSILDDCYPWDLSEIYDAPVLLFYKG NL DLL KFPKV AV VGSRACSKQGA KSVEK VIQGLEN . ELVI VS GLA KG ID	147
NTDB id	497 SM12261 RS05565 WP 000705318.1		SFSILDDCYPWDLSEIYDAPVLLFYKG NL DLL KFPKV AV VGSRACSKQGA KSVEK VIQGLEN . ELVI VS GLA KG ID	147
NTDB id	341595 ETT61 RS04610 WP 111693218.1		STSI LD KH YP MAL KEIYN PPVL LF FY QGNLD LL EK PK LATI VGSRRSSDTGVKS VRKITL KEL GN . RFVI VS GLA RGI D	145
NTDB id	407 SMU RS04605 WP 002262865.1		SL SI LD KE YPE L KN IYN PPVL LF FY QGD L DL LAP KLAVVG SRNASQM GVAA VK KI IQDLSK . QFVI ISGL A R GI D	145
consensus			* * * * *	

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[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
NTDB id 267 KZH43 RS05590 WP 000705306.1
NTDB id 226 SPD RS05990 WP 000705306.1
NTDB id 192 SPR RS05715 WP 000705306.1
NTDB id 157 SP RS06205 WP 000705306.1
NTDB id 497 SM12261 RS05565 WP 000705318.1
NTDB id 341595 ETT61 RS04610 WP 111693218.1
NTDB id 407 SMU RS04605 WP 002262865.1
consensus

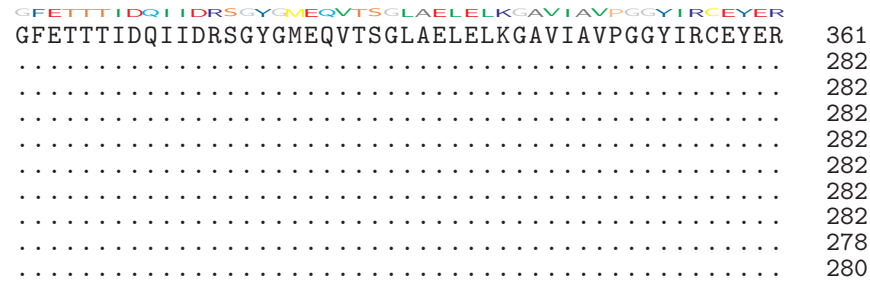
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NTDB id 1250 GC085 RS13065 WP 027219989.1
NTDB id 599 KW2 RS05940 WP 021037268.1
NTDB id 525 SMSK321 RS06295 WP 000705298.1
NTDB id 267 KZH43 RS05590 WP 000705306.1
NTDB id 226 SPD RS05990 WP 000705306.1
NTDB id 192 SPR RS05715 WP 000705306.1
NTDB id 157 SP RS06205 WP 000705306.1
NTDB id 497 SM12261 RS05565 WP 000705318.1
NTDB id 341595 ETT61 RS04610 WP 111693218.1
NTDB id 407 SMU RS04605 WP 002262865.1
consensus

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- X non conserved
- X similar
- X $\geq 50\%$ conserved