

NTDB id 1250 GC085 RS13065 WP 027219989.1	..MNNLQLPLLA	LNRMKRIGP....RTVLKLQKRPDLNLMFQLSSV	DLEEAGLPSWLAQTIKNFPDPDFIKTDLNLWLSSS	73
NTDB id 599 KW2 RS05940 WP 021037268.1	.MITNFD....	LFRWKKAGMTNLGVNKLKLFRRKYDRKISLRQM..	GQVAQVKKSIPNFIEQYKNQDVKKLRADYKKFS..	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....	IYKLKKSGLTNQQVLAVLEYGENVDQELLGDI..	AEISGCRNPAVFMERYFQIDDAHLEKEFQKFP..	72
NTDB id 341171 ETT69 RS05040 WP 014407585.1	..MNHFE....	LYKLKKAAGLTNKNILNILDYQEHQEKSLSLRDM..	AVVSGCKHPSHFIEAYKQLDIQKLKMEFKQFP..	70
NTDB id 407 SMU RS04605 WP 002262865.1	..MDNFG....	LFLKKAAGLTNKNILNIIIDYEERTQKSLSLRDM..	AVVSKNKKPLIFMEHYKNLDSKALRKEFNRF..	70
consensus	*****	*****	*****	

NTDB id 1250 GC085 RS13065 WP 027219989.1	ENHHILTWDSPYYPALKEIADPPFITYAKGELSALTQPSLATIVGSRNA SVTG NENARA FSR EISRHGVS IVSGLA LGID	153
NTDB id 599 KW2 RS05940 WP 021037268.1	 SFS ILDD L YPERL REIYN PPVLI FYQGNI DL LKNPKLA FVG SRLAG QSG I KSV QKIV TELNQ . SFT IVSGLAKGID	146
NTDB id 525 SMSK321 RS06295 WP 000705298.1	 SFS ILDD CYPWDL SEIYDAPVLLFYKG NL D LLKF PKVA VVG SR SCSKQGA KSVEKI IQGLE N. ELVIVSGLAKGID	147
NTDB id 267 KZH43 RS05590 WP 000705306.1	 SFS ILDD CYPWDL SEIYDAPVLLFYKG NL D LLKF PKVA VVG SRACSKQA KSVEKVIQGLE N. ELVIVSGLAKGID	147
NTDB id 226 SPD RS05990 WP 000705306.1	 SFS ILDD CYPWDL SEIYDAPVLLFYKG NL D LLKF PKVA VVG SRACSKQA KSVEKVIQGLE N. ELVIVSGLAKGID	147
NTDB id 192 SPR RS05715 WP 000705306.1	 SFS ILDD CYPWDL SEIYDAPVLLFYKG NL D LLKF PKVA VVG SRACSKQA KSVEKVIQGLE N. ELVIVSGLAKGID	147
NTDB id 157 SP RS06205 WP 000705306.1	 SFS ILDD CYPWDL SEIYDAPVLLFYKG NL D LLKF PKVA VVG SRACSKQA KSVEKVIQGLE N. ELVIVSGLAKGID	147
NTDB id 497 SM12261 RS05565 WP 000705318.1	 SFS ILDD CYPWDL SEIYDAPVLLFYKG NL D LLKF PKVA VVG SRACSKQA KSVEKVIQGLE N. ELVIVSGLAKGID	147
NTDB id 341171 ETT69 RS05040 WP 014407585.1	 STSILD KHYPMALKEIYNPPVLLFFYQGNDLLEKPKLATIVGSRRSSDTGVKSVRKIKELGN.RFVIVSGLARGID	145
NTDB id 407 SMU RS04605 WP 002262865.1	 SL SILD KEPL ELKNIYNPVLLFFYQGDLLARPKLAVVGSRNASQM GVA AVKKIIQDLSK.QFVIISGLARGID	145
consensus	* * * * *	

[illegible]

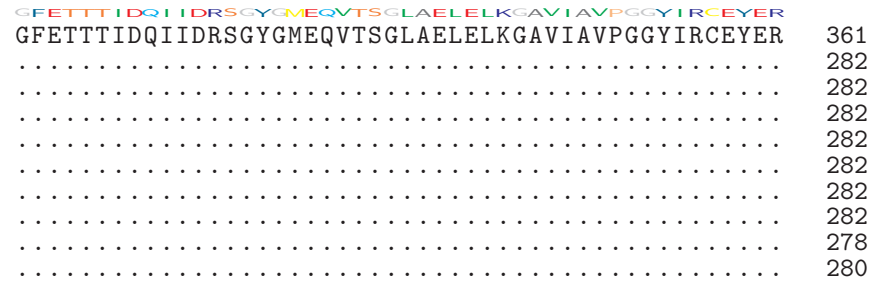
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consensus
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- X non conserved
- X similar
- X $\geq 50\%$ conserved