

logo

NTDB id 1169 A1552VC RS11075 WP 000648511.1	MKATQTLP	VSCQMLAIS	LTHRVKS	67
NTDB id 1402 DSB67 RS12670 WP 010643257.1	MKTIA	TSGNVLAMTE	ISHRVKG	66
NTDB id 1112 NGFG RS09215 WP 003689811.1	MAKNGGFS	LNGEVS	TSSKRKITQ	69
NTDB id 1113 AAA85695.1 219..1451()	MAKNGGFS	LNGEVS	TSSKRKITQ	69
NTDB id 1198 PSJM300 03950 AFN76868.1	MAQKA	IKGELS	AGKKIKP	65
NTDB id 1016 ACIAD RS01680 WP 004920476.1	MAAKKTQ	LKGELP	IFEGLLK	68
NTDB id 1059 ABD1 RS01610 WP 000279215.1	MAVKKA	IKGELP	LEGLFK	68
NTDB id 1101928 Q9271 RS15230 WP 011408067.1	MSVARSAIKKQ	MKCEQT	AGKKITP	78
NTDB id 1252 GC085 RS07730 WP 011213805.1	MDKN	MKCEQT	FDR.KN	66
consensus	*	*	*	

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NTDB id 1169 A1552VC RS11075 WP 000648511.1	KDITILTRQLATM	KDI	GEMSGN	147
NTDB id 1402 DSB67 RS12670 WP 010643257.1	KDITIFTRQIATM	KDI	GEMSGN	146
NTDB id 1112 NGFG RS09215 WP 003689811.1	EDITVFRQLSTM	EDITV	AAGETGGV	149
NTDB id 1113 AAA85695.1 219..1451()	EDITVFRQLSTM	EDITV	AAGETGGV	149
NTDB id 1198 PSJM300 03950 AFN76868.1	MDIALFTRQMATM	MDIAL	GEGQSGS	145
NTDB id 1016 ACIAD RS01680 WP 004920476.1	LDIALFTRQLATM	LDIAL	GEGQSGS	148
NTDB id 1059 ABD1 RS01610 WP 000279215.1	LDITIFTRQLATM	LDITIF	GEGQSGA	148
NTDB id 1101928 Q9271 RS15230 WP 011408067.1	KEIAFFSRQMATM	KEIAFF	GEGQSGA	158
NTDB id 1252 GC085 RS07730 WP 011213805.1	ADITVFSRQLATM	ADITV	GEGQSGS	146
consensus	*	*	*	

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NTDB id 1169 A1552VC RS11075 WP 000648511.1	LPEVFERLATYREK	LPEV	QAYS	227
NTDB id 1402 DSB67 RS12670 WP 010643257.1	LSQVFERLATYREK	LSQV	QAYS	226
NTDB id 1112 NGFG RS09215 WP 003689811.1	LESLLDKLATYKEK	LESLL	FVSYG	229
NTDB id 1113 AAA85695.1 219..1451()	LESLLDKLATYKEK	LESLL	FVSYG	229
NTDB id 1198 PSJM300 03950 AFN76868.1	LETLLDRVATYKEK	LETLL	QAW	225
NTDB id 1016 ACIAD RS01680 WP 004920476.1	LETIMLDRVATYKEK	LETIM	QAW	228
NTDB id 1059 ABD1 RS01610 WP 000279215.1	LETIMLDRVATYKEK	LETIM	QAW	228
NTDB id 1101928 Q9271 RS15230 WP 011408067.1	LETVLETVATYKEN	LETV	QAW	238
NTDB id 1252 GC085 RS07730 WP 011213805.1	LDIMLDDKVATYKEK	LDIML	QAW	226
consensus	*	*	*	

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NTDB id 1169 A1552VC RS11075 WP 000648511.1	LWAFIAIAAAIFGLKAL	LWAFI	VH	307
NTDB id 1402 DSB67 RS12670 WP 010643257.1	PFITGLSFSVSLI	PFITG	VH	306
NTDB id 1112 NGFG RS09215 WP 003689811.1	WMVLIALGFAIYGF	WMVL	VH	309
NTDB id 1113 AAA85695.1 219..1451()	WMVLIALGFAIYGF	WMVL	VH	309
NTDB id 1198 PSJM300 03950 AFN76868.1	YVVIAGFVGAAYAFK	YVVI	VH	305
NTDB id 1016 ACIAD RS01680 WP 004920476.1	FILILVIGAAIAGF	FILIL	VH	308
NTDB id 1059 ABD1 RS01610 WP 000279215.1	FIMILIAIGAVIAAF	FIMIL	VH	308
NTDB id 1101928 Q9271 RS15230 WP 011408067.1	WLMLIVTVGSSVVG	WLML	VH	318
NTDB id 1252 GC085 RS07730 WP 011213805.1	YIIFGALGGVYVS	YIIFG	VH	306
consensus	*	*	*	

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NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451( )
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
NTDB id 1101928 Q9271 RS15230 WP 011408067.1
NTDB id 1252 GC085 RS07730 WP 011213805.1
consensus

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logo



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