

NTDB id 1186	GTF74	RS01320	WP	000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA	AAVLIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1175	A1552VC	RS12105	WP	000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA	AAVLIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1408	DSB67	RS13980	WP	010643500.1	MKMRHGLTLTPRKLTGLFSLWFSLLAP...	FVAAEEVQSNLSKSIDFRTNKKDKDAVIVVELVSPAAIVDIKRVQEGLSI	77
NTDB id 1381	A4U84	RS02010	WP	021113424.1			0
NTDB id 219832	BV121	RS02595	WP	011272046.1			0
NTDB id 1373	NTHI	RS02645	WP	011272046.1			0
NTDB id 1347	HI	0435	AAC22094.1				0
consensus							0

NTDB id	1186	GTF74	RS01320	WP	000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDYTLKGKYLEVVISK.LKADEKPKPKSVLEKE	155
NTDB id	1175	A1552VC	RS12105	WP	000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDYTLKGKYLEVVISK.LKADEKPKPKSVLEKE	155
NTDB id	1408	DSB67	RS13980	WP	010643500.1	DLLNTSVKDDQLYLLDVKDFSTVVESVEVFRDSTSTRRLVAYIDEDYSHDYRLTGRYLEVKKVGK.LKAN EKAPDKSILDKE	156
NTDB id	1381	A4U84	RS02010	WP	021113424.1	MRYLFLFFATF.....PVL A	16
NTDB id	219832	BV121	RS02595	WP	011272046.1	MKKYFLKCGYFLVCFCLPLIVFAN.....PKTD	28
NTDB id	1373	NTHI	RS02645	WP	011272046.1	MKKYFLKCGYFLVCFCLPLIVFAN.....PKTD	28
NTDB id	1347	HI	0435	AAC22094.1	MKKYFLKCGYFLVCFCLPLIVFAN.....PKTD	28	
consensus						Y*G**!*G**G*	* * *

[illegible]

NTDB id	1186	GTF74	RS01320	WP	000788426.1	EKQALEKARLAEEELGDLKSEIKINFAKASDIAAMIGGEGNVNMLSERGSISIDERTNSLLIRELPDNIAVIREIIESLD	315
NTDB id	1175	A1552VC	RS12105	WP	000788426.1	EKQALEKARLAEEELGDLKSEIKINFAKASDIAAMIGGEGNVNMLSERGSISIDERTNSLLIRELPDNIAVIREIIESLD	315
NTDB id	1408	DSB67	RS13980	WP	010643500.1	EKQQLEKKQLKEIKELSELSEIKVINFAKASDIAEMINGEGRISMILSRDRSGMTIDERTNSLLIRELPDNIEVIREIIESLD	316
NTDB id	1381	A4U84	RS02010	WP	021113424.1	..KQHHTAQLVNALPKLITTKIKLVHYSKASEVIESLT.KGGGNLLLESGLYLFHDDRNSNIIVKDSAASVKNFQTQLIESLD	170
NTDB id	219832	BV121	RS02595	WP	011272046.1	..KGQVAGNLTTNEPHLVSHTVKLHFPAKASELMKSLT.TGSGSLLSSAGSITFDDRSNLLVIQDEPRFVQNIXKLIAMD	182
NTDB id	1373	NTHI	RS02645	WP	011272046.1	..KGQVAGNLTTNEPHLVSHTVKLHFPAKASELMKSLT.TGSGSLLSPAGSITFDDRSNLLVIQDEPRFVQNIXKLIAMD	182
NTDB id	1347	HI	0435	AAC22094.1		..KGQVAGNLTTNEPHLVSHTVKLHFPAKASELMKSLT.TGSGSLLSPAGSITFDDRSNLLVIQDEPRS	182
consensus						* * * * *	

