

logo

NTDB id 1186	GTF74	RS01320	WP 000788426.1	..MKNGLKT..YVAQTLWTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA	AVLV	VELASPS	AVVDVQKVQEGLSI	76		
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	..MKNGLKT..YVAQTLWTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA	AVLV	VELASPS	AVVDVQKVQEGLSI	76		
NTDB id 1408	DSB67	RS13980	WP 010643500.1	MKMRHGLTLTPRKLTGLFSLWFSLLAP...	FVAAEEVQSN	SLKSIDFR	TNKKDKDAVIVVELVSPA	AI	VDIKRVQEGLSI	77
NTDB id 1381	A4U84	RS02010	WP 021113424.1					0		
NTDB id 1347	HI 0435	AAC22094.1						0		
NTDB id 309848	CH631	RS03000	WP 116939826.1					0		
NTDB id 1373	NTHI	RS02645	WP 011272046.1					0		
consensus									0	

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDY	TLKGY	LEVVIS	KLKADEKPKPKSVLEKEG	156						
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDY	TLKGY	LEVVIS	KLKADEKPKPKSVLEKEG	156						
NTDB id 1408	DSB67	RS13980	WP 010643500.1	DLNNTSVKDDQLYLLDVKDFSTVSVESVEVFRD	TSTTRLV	AYIDEDY	SHDYRLTG	RYLEV	KVGK	KLKAN	EPDKS	ILDK	EG	157
NTDB id 1381	A4U84	RS02010	WP 021113424.1											17
NTDB id 1347	HI 0435	AAC22094.1												29
NTDB id 309848	CH631	RS03000	WP 116939826.1											29
NTDB id 1373	NTHI	RS02645	WP 011272046.1											29
consensus				* * * * *										

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLT	RLRDGVPWQQVLDIILQVKGLDKRVDGNVIL	IAPKEELD	DLRE	236							
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLT	RLRDGVPWQQVLDIILQVKGLDKRVDGNVIL	IAPKEELD	DLRE	236							
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLT	RLRDGVPWQQVLDIILQVKGLDKRVDGNVIL	IAPKEELD	DLRE	237							
NTDB id 1381	A4U84	RS02010	WP 021113424.1	QQISLSIRNAPTAEIISYLAEEETGKNITISDEIK	DTKNFRVEKSHFDEILNSLIKTHQLNLKKENG	ITYYIHQAQEH....	93								
NTDB id 1347	HI 0435	AAC22094.1	ERFFIRLSQAPLAQTLEQLAFQQQDVNLVIGDILENK	ISLKLNNIDMPRL	LQIIAKSKHLLTNLKDDGVYYLNGSQSG....	105									
NTDB id 309848	CH631	RS03000	WP 116939826.1	ERFFIRLSQAPLAQTLEQLAFQQQDVNLVMGERLEGN	ISLKLNNIDMPRL	LQIIAKSKHLLTNLKDDGVYYLNGSQSG....	105								
NTDB id 1373	NTHI	RS02645	WP 011272046.1	ECFFIRLSQAPLAQTLEQLAFQQQDVNLVMGERLEGN	ISLKLNNIDMPRL	LQIIAKSKHLLTNLKDDGVYYLNGSQSG....	105								
consensus				*** * *	***!*****	*!* *	!*****	*****	* * *	*!* *	*****	!*****	*****	*****	**

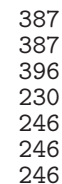
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NTDB id 1186	GTF74	RS01320	WP 000788426.1	KQALEKARLAEELGDLKSEI	IKINF	AKASDIAAM	IGGEG	NVNMLSERGSIS	IDERTNS	LLIRELP	DNIAVIRE	IIIESLDI	316												
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KQALEKARLAEELGDLKSEI	IKINF	AKASDIAAM	IGGEG	NVNMLSERGSIS	IDERTNS	LLIRELP	DNIAVIRE	IIIESLDI	316												
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KQALEKARLAEELGDLKSEI	IKINF	AKASDIAAM	IGGEG	NVNMLSERGSIS	IDERTNS	LLIRELP	DNIAVIRE	IIIESLDI	317												
NTDB id 1381	A4U84	RS02010	WP 021113424.1	.KQHTTAQLVNALPKLITKT	IKLHYSK	ASEVIESLT	.KQQGN	LLSES	GYLH	FDDRS	NSIIVK	DSAASV	KNF	QTLIESLDK	171										
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NTDB id 309848	CH631	RS03000	WP 116939826.1	.KQAVAGNLT	TNEPHLV	SHTVK	LHFAKASELMKSLT	.TGSG	LLSPAGS	ITFDDRS	NLLVIQ	DEPRSV	QNIKKLIA	EMDK	183										
NTDB id 1373	NTHI	RS02645	WP 011272046.1	.KQAVAGNLT	TNEPHLV	SHTVK	LHFAKASELMKSLT	.TGSG	LLSPAGS	ITFDDRS	NLLVIQ	DEPRSV	QNIKKLIA	EMDK	183										
consensus				**	**	!*	**	!*	**	*****	!!!!	**	**	*****	!*****	!*	*****	*****	*	*	**	**	*****	!	*

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NTDB id 1186 GTF74 RS01320 WP 000788426.1
NTDB id 1175 A1552VC RS12105 WP 000788426.1
NTDB id 1408 DSB67 RS13980 WP 010643500.1
NTDB id 1381 A4U84 RS02010 WP 021113424.1
NTDB id 1347 HI 0435 AAC22094.1
NTDB id 309848 CH631 RS03000 WP 116939826.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
consensus

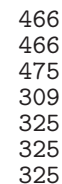
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NTDB id 309848 CH631 RS03000 WP 116939826.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
consensus

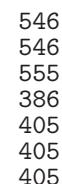
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NTDB id 1381 A4U84 RS02010 WP 021113424.1
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NTDB id 1373 NTHI RS02645 WP 011272046.1
consensus

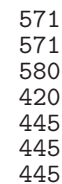
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NTDB id 1186 GTF74 RS01320 WP 000788426.1
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NTDB id 1408 DSB67 RS13980 WP 010643500.1
NTDB id 1381 A4U84 RS02010 WP 021113424.1
NTDB id 1347 HI 0435 AAC22094.1
NTDB id 309848 CH631 RS03000 WP 116939826.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
consensus

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- X non conserved
- X similar
- X >50% conserved