

logo

NTDB id 1186	GTF74	RS01320	WP 000788426.1	..MKNGLKT..YVAQ	TLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA	AAVLIVELASPSAVVDVQKVQEGLSI	76			
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	..MKNGLKT..YVAQ	TLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA	AAVLIVELASPSAVVDVQKVQEGLSI	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 47252	R2866	RS00705	WP 005649302.1							0
NTDB id 1373	NTHI	RS02645	WP 011272046.1							0
consensus										

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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	LKAN	EAPDKS	ILDK	EG	157					
NTDB id 1381	A4U84	RS02010	WP 021113424.1											17					
NTDB id 1347	HI 0435	AAC22094.1												29					
NTDB id 47252	R2866	RS00705	WP 005649302.1											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	VAPKDELD	DLRE	237															
NTDB id 1381	A4U84	RS02010	WP 021113424.1	QQISLSIRNAPTAEIISYLAEEETGKNITISDEIK	DTKNFRVEKSHFDEILNSLIKTHQLNLK	KENGITYYIHQAQEH....		93															
NTDB id 1347	HI 0435	AAC22094.1		ERFFIRLSQAPLAQTLEQLAFQQQDVNLVIGDILENK	ISLKLNNIDMPRL	LQIIAKSKH	LTNLKDDG	IYYLNGSQSG....	105														
NTDB id 47252	R2866	RS00705	WP 005649302.1	ERFFIRLSQAPLAQTLEQLAFQQQDVNLVIGDILENK	ISLKLNNIDMPRL	LQIIAKSKH	LTNLKDDG	IYYLNGSQSG....	105														
NTDB id 1373	NTHI	RS02645	WP 011272046.1	ECFFIRLSQAPLAQTLEQLAFQQQDVNLVIGDILENK	ISLKLNNIDMPRL	LQIIAKSKH	LTNLKDDG	IYYLNGSQSG....	105														
consensus																							
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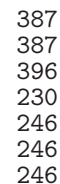
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NTDB id 1186	GTF74	RS01320	WP 000788426.1	KQALEKARLAEELGDLKSEI	IKINF	AKASDIAAM	IGGEGNVN	MLSERGSIS	IDERTNS	LLIRELP	DNIAVIRE	IIIESLDI	316				
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KQALEKARLAEELGDLKSEI	IKINF	AKASDIAAM	IGGEGNVN	MLSERGSIS	IDERTNS	LLIRELP	DNIAVIRE	IIIESLDI	316				
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KQALEKARLAEELGDLKSEI	IKINF	AKASDIAAM	IGGEGNVN	MLSERGSIS	IDERTNS	LLIRELP	DNIAVIRE	IIIESLDI	317				
NTDB id 1381	A4U84	RS02010	WP 021113424.1	.KQHTTAQLVNALPKLITKT	IKLHYSK	ASEVIESLT	.KQQGNLL	SES	GYLHFD	DRSNS	IIIVK	DSAASV	KNFTQL	IESLDK	171		
NTDB id 1347	HI 0435	AAC22094.1		.KQQVAGNLT	TNEPHLV	SHTVKL	HFAKASELMKSLT	.TGSGSLL	SPAGSIT	FDDRSN	LLVIQ	DEPRSV	QNIKKLIA	EMDK	183		
NTDB id 47252	R2866	RS00705	WP 005649302.1	.KQQVAGNLT	TNEPHLV	SHTVKL	HFAKASELMKSLT	.TGSGSLL	SPAGSIT	FDDRSN	LLVIQ	DEPRSV	QNIKKLIA	EMDK	183		
NTDB id 1373	NTHI	RS02645	WP 011272046.1	.KQQVAGNLT	TNEPHLV	SHTVKL	HFAKASELMKSLT	.TGSGSLL	SPAGSIT	FDDRSN	LLVIQ	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 47252 R2866 RS00705 WP 005649302.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
consensus

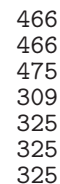
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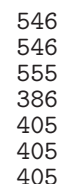
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consensus

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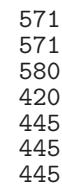
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NTDB id 1381 A4U84 RS02010 WP 021113424.1
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NTDB id 47252 R2866 RS00705 WP 005649302.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
consensus

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