

logo

NTDB id 1186	GTF74	RS01320	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA	AVLV	VELASPS	AVVDVQKVQ	VEGLSI	76
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKA	AVLV	VELASPS	AVVDVQKVQ	VEGLSI	76
NTDB id 1408	DSB67	RS13980	WP 010643500.1	MKMRHGLTLTPRKLTGLFSLWFSLLAP...	FVAAEEVQ	SNSLKS	SIDFR	TNKDKDAVIVVELVSPAAIVDIKRV	77
NTDB id 1347	HI 0435	AAC22094.1							0
NTDB id 1373	NTHI	RS02645	WP 011272046.1						0
NTDB id 1465	ASU2	RS06170	WP 014991934.1						0
NTDB id 1381	A4U84	RS02010	WP 021113424.1						0
consensus									

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDY	YTL	LK	GK	Y	L	V	V	I	S	K	L	K	A	D	E	K	P	K	P	K	S	V	L	E	K	E	G	156			
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDY	YTL	LK	GK	Y	L	V	V	I	S	K	L	K	A	D	E	K	P	K	P	K	S	V	L	E	K	E	G	156			
NTDB id 1408	DSB67	RS13980	WP 010643500.1	DLLNTSVKDDQLYLLDVKDFSTTVESVEVFRDTSTTRLVAYIDEDYSHD	YRL	T	G	R	Y	L	E	V	K	V	G	K	L	K	A	N	E	K	A	P	D	K	S	I	L	D	K	E	G	157	
NTDB id 1347	HI 0435	AAC22094.1			M	K	K	Y	F	L	K	C	G	Y	F	L	V	C	F	C	L	P	L	I	V	F	A		N	P	K	T	D	N	29
NTDB id 1373	NTHI	RS02645	WP 011272046.1		M	K	K	Y	F	L	K	C	G	Y	F	L	V	C	F	C	L	P	L	I	V	F	A		N	P	K	T	D	N	29
NTDB id 1465	ASU2	RS06170	WP 014991934.1		M	K	R	M	I	L	L	L	F	F	I		L	P	P	V	I	A												17	
NTDB id 1381	A4U84	RS02010	WP 021113424.1		M	R	Y	L	F	L	L	F	F	A	T		F	P	V	L	A	N												17	
consensus													*	*	*	*		*	*	*	*					*	*	*	*	*	*				

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVVS	SDSV	EGNLT	RLDNV	PWQQVLDI	ILQVKGLDKRVDGNVIL	IAPKEELD	DLRE	236
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVVS	SDSV	EGNLT	RLDNV	PWQQVLDI	ILQVKGLDKRVDGNVIL	IAPKEELD	DLRE	236
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KLISINFQDIPVRNVLQLIADYNGFNLVVS	SDSV	EGNLT	RLDNV	PWQQVLDI	ILQVKGLDKRVDGNVIL	IAPKEELD	DLRE	237
NTDB id 1347	HI 0435	AAC22094.1		ERFFIRLSQAPLAQTLEQLAFQQDVNLVIGDILENKISLKLNNIDMPRL	LQIIAKSKHL	TLNKDDG	IYYL	NGSQS	G			105
NTDB id 1373	NTHI	RS02645	WP 011272046.1	ECFFIRLSQAPLAQTLEQLAFQQDVNLVIGDILENKISLKLNNIDMPRL	LQIIAKSKHL	TLNKDDG	VYYL	NGSQS	G			105
NTDB id 1465	ASU2	RS06170	WP 014991934.1	HAFSISILKNAPTAKILSYLAEHKGKNI	IVLS	DNITETNT	TLRIENS	NFDSVLKS	ITRANKLTSAYENQIYF	IGHKKD	D	93
NTDB id 1381	A4U84	RS02010	WP 021113424.1	QQISLSIRNAPTAETISYLAETGKNITISDEIKDTKNFRVEKSHFDEILNSLIKTHQLNLK	KENGIYYIHQAQ	H						93
consensus										*** * !!***** *!* ** !***** ***** * * *! ***** * ! * * ***** * *		

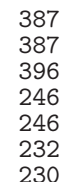
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NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KQALEKARLAEELGDLKSEI	IKINFAKASDIAAM	IGGEGNVNMLSERGSIS	IDERTNSLLIRELPDNI	IAVIREII	IESLDI	316
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KQALEKARLAEELGDLKSEI	IKINFAKASDIAAM	IGGEGNVNMLSERGSIS	IDERTNSLLIRELPDNI	IAVIREII	IESLDI	317
NTDB id 1347	HI 0435	AAC22094.1		KGQV.AGNLTTNEPHLV	SHTVKLHF	FAKASELMKSLT	.TGSGSLLSPAGSIT	FDDRSNLLVIQDEPRSVQ	NIKKLIAEMDK	183
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NTDB id 1465	ASU2	RS06170	WP 014991934.1	KDPNRGLNLELLKPKLIT	KTIKLDYAKAAE	EVIESLT	.KSGNFLSENGYLH	FDDRSNLSIIKDSPE	SMKNIVKLIRN	172
NTDB id 1381	A4U84	RS02010	WP 021113424.1	KQHT.TAQLVNALPKLIT	KTIKLDYAKAAE	EVIESLT	.KGQGNLLSESGYLH	FDDRSNLSIIKDSPE	SMKNIVKLIRN	171
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 1347 HI 0435 AAC22094.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
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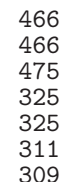
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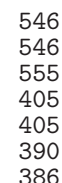
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NTDB id 1373 NTHI RS02645 WP 011272046.1
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consensus

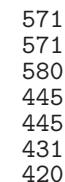
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NTDB id 1186 GTF74 RS01320 WP 000788426.1
NTDB id 1175 A1552VC RS12105 WP 000788426.1
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NTDB id 1347 HI Q435 AAC22094.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
NTDB id 1465 ASU2 RS06170 WP 014991934.1
NTDB id 1381 A4U84 RS02010 WP 021113424.1
consensus

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