

logo

NTDB id 1323 RA0C RS07335 WP 004916218.1	MRQPLEDR	EV	TISRAKFT	VNYPASF	MLVASMNP	SPSG	FFPDDPN	.NTSS	SFEM	QRYLN	KL	SGP	LLDR	ID	TH	IEV	QK	VEFD	397		
NTDB id 1271 LPP RS03220 WP 011213198.1	LREPLESG	HIC	CISRAA	QIEFPA	KFQLIA	AMNP	CPGQ	WGN	SQANCM	CTPD	RISRYL	AKLS	APLL	DR	ID	MQVT	IH	ALS	SQE	394	
NTDB id 1248 GC085 RS03150 WP 011213198.1	LREPLESG	HIC	CISRAA	QIEFPA	KFQLIA	AMNP	CPGQ	WGN	SQANCM	CTPD	RISRYL	AKLS	APLL	DR	ID	MQVT	IH	ALS	SQE	394	
NTDB id 1034833 AB9V60 RS00645 WP 003413980.1	LREPLESG	HIV	ISRARD	RVRFP	PARFQL	VAA	MNP	PCPG	YLGE	PTGR	CRCS	STEQ	VQRY	RNKL	SGP	LLDR	ID	LHLTVAREAT	A	393	
NTDB id 1153 A1552VC RS00115 WP 000521788.1	LREPLESG	EIV	ISRAQ	GKTRF	PARFQL	V	GALNP	SPTG	YEG	SQ	..ARAN	PQSIL	RYLS	R	LSG	P	LLDR	FDM	STIEIPAL	PKG	392
NTDB id 1397 DSB67 RS15690 WP 005535577.1	LREPLESG	EII	ISRAQ	GKTRF	PARFQ	I	V	GALNP	SPTG	YEG	NQ	..ARTN	PQA	ILRYL	GRL	SGP	LLDR	FDM	SLEIPAL	PKG	392
NTDB id 1387 A4U84 RS06900 WP 020457569.1	LRQPLESG	EII	ISRA	N	AKVQ	F	PACFQL	I	A	MNP	SPTG	H	Y	Q	G	T	H	..NRT	SPQQLMRYLNRLSGPFLDRFDLSIEVPLLP	Q	395
NTDB id 1349 HI 1117 AAC22771.1	LRQPLESG	EII	ISRA	N	AKIQ	F	PARFQL	VAA	MNP	SPTG	H	Y	Q	G	T	H	..NRT	SPQQLMRYLNRLSGPFLDRFDLSIEVPLLP	Q	396	
consensus	*****	!!!!	*****	!!!!	*****	!!!!	*****	!!!!	*****	!!!!	*****	!!!!	*****	!!!!	*****	!!!!	*****	!!!!	*****	!!!!	

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NTDB id 1323 RA0C RS07335 WP 004916218.1	QLS	..DKR	KGESSE	ITRQ	RVLK	AREI	Q	QERY	QDLA	ISYNAQ	MGP	KEIE	HFC	CEL	DEV	SLLL	IKNA	MEK	LNLS	SARAYDRILK	475					
NTDB id 1271 LPP RS03220 WP 011213198.1	ELIKPN	THLEKQ	SLA	IREK	VTKM	HEI	Q	MARQD	...	SLNAN	LNSK	TC	EMV	CEL	GSEE	Q	LFL	REV	MSK	LKLS	SARGYHRLLK	470				
NTDB id 1248 GC085 RS03150 WP 011213198.1	ELIKPN	THLEKQ	SLA	IREK	VTKM	HEI	Q	MARQD	...	SLNAN	LNSK	TC	EMV	CEL	GSEE	Q	LFL	REV	MSK	LKLS	SARGYHRLLK	470				
NTDB id 1034833 AB9V60 RS00645 WP 003413980.1	..LNP	DP	TTSEN	TASAA	VVAE	AARD	Q	RRQ	Q	...	CANAF	L	DL	PGL	REY	CAL	SSV	DES	W	LET	ACER	RLTSLRS	SAHRLLK	467		
NTDB id 1153 A1552VC RS00115 WP 000521788.1	TLANG	.GDR	G	ESTAA	V	RQ	R	VW	ARER	M	L	ARSG	...	KVN	ALLQ	SREIE	QY	CP	LLK	ADAE	FLES	A	I	HRLGLSIRAYHRI	IK	467
NTDB id 1397 DSB67 RS15690 WP 005535577.1	TLSEG	.GDR	G	EPTAI	V	KERV	NQ	ARDC	M	L	ARSG	...	KVN	ALLQ	SREIE	QY	CP	LQKS	ADAE	FLEN	A	I	HRLGLSIRAYHRI	IK	467	
NTDB id 1387 A4U84 RS06900 WP 020457569.1	ALQN	NTDQ	RGETTE	Q	ARKRV	LE	ARHL	Q	F	NRRG	...	KINA	QLT	TKEIE	QDCK	L	TDR	DAL	FLE	NAL	T	KLGLS	VRAYHRILK	471		
NTDB id 1349 HI 1117 AAC22771.1	SLQNT	.GDR	G	ETSA	V	REK	V	LKV	REI	Q	MERAG	...	KINAY	L	NSKEI	ERDCK	L	NDK	DAF	FLE	KAL	N	KLGLS	VRAYHRILK	471	
consensus	*		***	*		***	!	*		*****	!	*		!!!!	*	*****	!	!	*		***	*	!	!!!!	*****	!!!!

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NTDB id 1323 RAOC RS07335 WP 004916218.1	VSRTIADLERETNIQSHHIAEAIQYRSLDRDFWKV.....	510
NTDB id 1271 LPP RS03220 WP 011213198.1	VSRTIADMNSSKKVLLNHLQQALS YKQNLHLPK.....	503
NTDB id 1248 GC085 RS03150 WP 011213198.1	VSRTIADMNSSKKVLLNHLQQALS YKQNLHLPK.....	503
NTDB id 1034833 AB9V60 RS00645 WP 003413980.1	VARTLADLEQVDAINRSHLAEALQYRPSTN.....	497
NTDB id 1153 A1552VC RS00115 WP 000521788.1	VARTIADLQGEAQIARPHLAEALGYRAMDRLKQLSAQNV	507
NTDB id 1397 DSB67 RS15690 WP 005535577.1	VARTIADLEGSPQIERTHLAEALGYRAMDRLKQLTAQAV	507
NTDB id 1387 A4U84 RS06900 WP 020457569.1	VSRTIADLAGEAQIQQPHLAEALGYRAMDRLQLRQLGES.	510
NTDB id 1349 HI 1117 AAC22771.1	VSRTIADLQGEQIQIFQPHLAEALGYRAMVRLQLKLSNM..	509
consensus	!!!!!!* * !***** !* * ** *	

- ☒ non conserved
- ☒ similar
- ☒ ≥ 50% conserved