

logo

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NTDB id 1028	TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
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NTDB id 1058	ABD1 RS01615 WP 001274986.1		0
NTDB id 1018	ACIAD RS01685 WP 004920473.1		0
consensus			

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NTDB id 1058	ABD1 RS01615 WP 001274986.1		0
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consensus			

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NTDB id 1058	ABD1 RS01615 WP 001274986.1		0
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consensus			

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consensus		

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NTDB id 1251	GC085 RS07725 WP 011946523.1	MALATEEYRLQGLLGLLVLEKLLDKTKAIELHKLAAAEEKISLLQYIV..KNKILSAEQIALT.....AAQNF	65
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consensus		* * * * *	

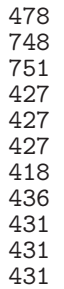
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NTDB id 1293 VP RS12240 WP 005479695.1	GLSQPE	LSQYBYA	SLCQQL	LGLRELIT	RHNALPLHRT	...STLLAVADPTN	QQAEDDFRFATG	LQVELV	124	
NTDB id 1401 DSBB7 RS12665 WP 010643259.1	GLSCTSL	MSQYBYA	SLCQT	LGLRDLIT	RHNALPLNR	TS...STLLAVADPTN	LQAEEDFRFATG	LQVELV	124	
NTDB id 1110 NGFG RS09235 WP 003689817.1	SYSILDL	RHYPRH	RVLMG	VLTEEQM	VEFHCVPV	FRRG...DKVFFAVSDPT	QMPQIQKTVS	AAG....I	AVELV	123
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	AEPIYFD	LDVYDTS	QIPKD	LVDQKL	LILKHRIL	PLIQRG...QILYVATS	NPTNIEAIDAIR	FNSK....LLVEP	132	
NTDB id 1018 ACIAD RS01685 WP 004920473.1	GEPLLD	LGVFDPA	LFLKDK	ID	EKLQIKYR	IMPLVHRG...HVLVYATS	NPTNIEAMDAIR	FNSK....LKVEPI	132	
consensus	*****	*****	*****	*	*****	*****	*****	*****	*****	

NTDB	id	1410	SGL	RS13950	WP	010873519.1	VVTQDDFNQ	LLEQYYA	AIREEW	EAAQEKRD	QERALDK	LNLDLTD	IVGSI	ID.MN	LG	EDK	EDVDNAL	DSNDA	NQAPVINL	VNKI	240									
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NTDB	id	1028	TT	RS08215	WP	011173991.1	VATEAA	ITK	LIERFY	YGKAELS	EIA						KEFA		KKQA	EEEV		PSPLEL	DESAAQ	KFVKQV	514					
NTDB	id	1168	A	1552VC	RS11070	WP	000957200.1	LADHRA	LQAA	IRRLY	GRSISQA	ANQGK	EI				SQ	DEL	AN	LKVKS	DDELQSI	EDLSQ	DDSPV	SRYINQV	189					
NTDB	id	1293	VP	RS12240	WP	005479695.1	LADFREL	STA	IRRLY	GRSLSH	EKSGLK	EI						NQ	EEL	ASLV	DVGAE	IDNIED	LSQ	DESPV	SRYINQI	189				
NTDB	id	1401	DSB67	RS12665	WP	010643259.1	LADFREL	TAA	IRRLY	GRSLGQ	EKSGLK	EI						NQ	DEL	AGLV	DIG	EDEIENI	EDLSQ	DESPV	SRYINQI	189				
NTDB	id	1110	NGFG	RS09235	WP	003689817.1	IVEDDQ	LAG	LD	WVGS	RSTSL	LQE	LGE							GQ		EEEE	SHTLYI	DNEEA	EDGPV	PRFIHKT	179			
NTDB	id	1251	GC085	RS07725	WP	011946523.1	VVETDK	LSA	LIDN	LLTAKES	QGLSEY	VED								SG	DIEGLE	ISADD	EDQDID	TATSVT	DDAPI	IVKVF	VNKI	198		
NTDB	id	270922	C3Y93	RS11345	WP	004975865.1	VVEHDK	LVK	LIEQN	FASESA	F	EFE	DD							DF	DLN	FED	GNS	EPEQAAE	DEPQG	DESP	IVKY	INKL	193	
NTDB	id	1058	ABD1	RS01615	WP	001274986.1	IVEHHK	L	EKVL	GQHFA	EESSFD	N	DE							EF	DL	VNL	DGPTAQ	EDEE	EAPQG	DEAP	IVKY	INKL	193	
NTDB	id	1018	ACIAD	RS01685	WP	004920473.1	IVEHDK	L	ERL	L	SEHF	VEETHF	NFD	TE							EL	DL	VEV	DPHTT	DDDED	DDKLK	DEAP	IVKY	INKL	193
consensus							***	*	***	**	*	*					**	*	*	***	*		*****							

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NTDB id 1110	NGFG RS09235 WP 003689817.1	LSDALRS GASDIHFEFYEHNAIRIFRVVDGQLREV VQ.PPIAVRGQLASRIKVM SRLDISEKRIPQDGRMQLTFQKGGKPV	258
NTDB id 1251	GC085 RS07725 WP 011946523.1	LLDAIRQGASDIHFEPYEREYRIRYRQDGI LHE VAT.PPASLSSRITARIKVM SNLDISERRIPQDGGFMMKISK.SRAI	276
NTDB id 270922	C3Y93 RS11345 WP 004975865.1	LIDAIRM GASDLHFEPYEKT YRVRYRV DGV LR Q IAT.PPLQLANRLA SRLKVM SQMDISEKRV PPDGRIKLKLSK.NKSI	271
NTDB id 1058	ABD1 RS01615 WP 001274986.1	LIDAIRM GASDLHFEPYEKS YRVRYRV DGV LR Q IAN.PPLQLANRLA SRLKVM SQMDISEKRV PPDGRIKLKLSK.SKAI	271
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consensus		***** *!!!!!! ***** ****!*!!!!!!*!**** * * ** ***** *!!!!!!*!!!!!!*!!!!!!*! *	

NTDB	id	1410	SGL	RS13950	WP	010873519.1	D	F	R	V	N	T	L	P	S	R	Y	G	E	K	V	C	L	R	I	L	D	N	S	A	T	Q	L	G	L	D	F	L	I	T	N	P	E	T	L	Q	T	V	R	E	L	A	A	R	P	Y	G	L	M	V	T	G	P	T	G	S	G	K	S	T	T	L	Y	S	V	L	A	E	R	N	S	P	E	V	N	398
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consensus
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consensus

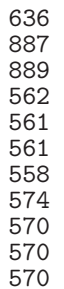
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

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logo

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NTDB id 1058 ABD1 RS01615 WP 001274986.1		570
NTDB id 1018 ACIAD RS01685 WP 004920473.1		570
consensus		

- X non conserved
- X similar
- X ≥ 50% conserved