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NTDB id 1177 A1552VC RS01165 WP 000422572.1
NTDB id 1203 PSJM300 01605 AFN76401.1
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NTDB id 1081 FDQ49 RS05335 WP 006581387.1
NTDB id 1060 ABD1 RS04380 WP 000355489.1
NTDB id 1040 HQN27 RS13360 WP 000355489.1
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MDITELLAFSVKNGASDLHLSSAGVPPMIRVDGEVRRINLPALEHKDVHRLVYDIMNDKQRKYEEKLETFDSFEVPTMSEFEETISMAKERSFISISLPIKINLTKYCKELMPASEETKRVIVEMISELQSTFQNFVNIATLS

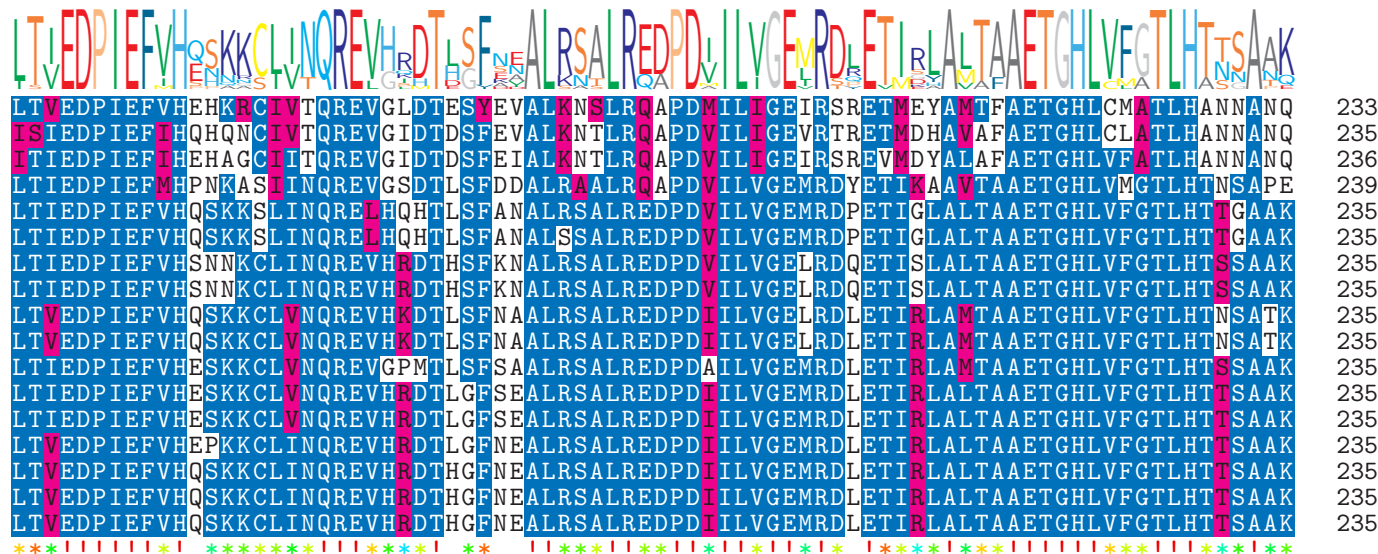
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...	ME	FE	KL	LR	LM	VE	KG	GS	DL	FI	TA	GP	PS	MK	VN	GI	LP	VN	K	TP	MS	PE	MT	RE	TV	HG	VM	NE	Q	Q	R	E	FT	EN	HE	C	N	FA	IS	AR	76																													
...	MD	FN	DL	LN	LM	AK	HK	AS	DL	FI	TA	GP	PS	LK	IN	GE	IT	VP	SK	TK	LD	GM	IA	QL	IDS	IT	TQ	KQ	R	Q	E	FI	ET	HE	C	N	FA	IL	NR	76																														
MTSGID	IT	DL	LR	LA	D	KG	AS	DL	II	LA	GL	PQ	FK	LS	GD	YA	DQ	GF	EP	LS	AD	TR	KL	MS	Y	MM	NR	Q	R	I	E	E	K	LE	VD	FS	F	EL	P	80																														
...	MQ	IT	DL	LA	FG	AK	NK	AS	DL	HL	SS	GI	SP	MIR	VH	GD	MRR	IN	LP	EM	SA	EE	VGN	MV	TS	VM	NH	Q	R	K	I	Y	Q	N	LE	VD	FS	F	EL	P	76																													
...	MQ	IT	DL	LA	FG	AK	NK	AS	DL	HL	SS	GI	SP	MIR	VH	GD	MRR	IN	LP	EM	SA	EE	VGN	MV	TS	VM	NH	Q	R	K	I	Y	Q	N	LE	VD	FS	F	EL	P	76																													
...	MD	IA	EL	LE	F	SV	K	H	N	AS	DL	HL	S	A	G	V	P	P	M	R	I	D	GE	V	R	K	L	G	V	P	A	F	T	H	S	D	V	H	R	L	I	F	E	I	M	N	D	A	Q	R	S	E	Y	E	E	K	L	E	V	D	F	S	F	E	L	P	76			
...	MD	IA	EL	LE	F	SV	K	H	N	AS	DL	HL	S	A	G	V	P	P	M	R	I	D	GE	V	R	K	L	G	V	P	A	F	T	H	S	D	V	H	R	L	I	F	E	I	M	N	D	A	Q	R	S	E	Y	E	E	K	L	E	V	D	F	S	F	E	L	P	76			
...	MD	IA	EL	LE	F	SV	K	H	N	AS	DL	HL	S	A	G	V	P	P	M	R	I	D	GE	V	R	K	L	G	V	P	A	F	T	H	S	D	V	H	R	L	I	F	E	I	M	N	D	A	Q	R	S	E	Y	E	E	K	L	E	V	D	F	S	F	E	L	P	76			
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...	MD	IT	EL	LA	FA	S	A	K	Q	G	AS	DL	HL	S	A	G	L	P	P	M	I	R	V	D	G	D	V	RR	I	N	L	P	P	L	E	H	K	Q	V	H	A	L	I	Y	D	I	M	N	D	K	Q	R	K	D	F	E	E	F	L	E	T	D	F	S	F	E	V	P	76	

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consensus

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consensus

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logo

NTDB id 1177 A1552VC RS01165 WP 000422572.1	TFDQALYQLVVDKISEQDALHSADSANDLRLMLKTKRGDDY...GSGSLQNVKI.....DME.....	368
NTDB id 1203 PSJM300 01605 AFN76401.1	TFDQALYNLYVQGEITYEDALLHADSSNDLRLMIKLGSETDGEHLTSVS.QGLTL.....EMSDDDPGRSFR.....	381
NTDB id 1019 ACIAD RS04205 WP 004922051.1	TFDQALFDLYDHQHSYADALKHADSPNNLRLQIKLSEEANKPLDNRRERMSEMER.....KMTFDGQRRNLK.....	383
NTDB id 1307 DR RS10055 WP 010888596.1	TMDAYLAGLFRHLITYDKGLERAIDPKE.FARLANDASAG...LGGASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	395
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NTDB id 39334 ALIDE RS02460 WP 013517392.1	TLDQNLTDLVRNIIISPAEARSKAKIP.ENFPG.....	347
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NTDB id 1202 PSJM300 01600 AFN76400.1	TLDSCCLKGLIAKGLISRDSAKEKAKQP.ENF.....	344
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NTDB id 1081 FDQ49 RS05335 WP 006581387.1	TLDQSLKGLVSRGVISPQTARTAAKQP.ESFL.....	345
NTDB id 1060 ABD1 RS04380 WP 000355489.1	TLDQSLKGLVARGVISPQTARTAAKQP.ESFL.....	345
NTDB id 1040 HON27 RS13360 WP 000355489.1	TLDQSLKGLVARGVISPQTARTAAKQP.ESFL.....	345
consensus	!*** * !* ***** * *	

logo

NTDB id 1177 A1552VC RS01165 WP 000422572.1	PATPAAPAACRSEPCRCACCFGR	368
NTDB id 1203 PSJM300 01605 AFN76401.1		381
NTDB id 1019 ACIAD RS04205 WP 004922051.1		383
NTDB id 1307 DR RS10055 WP 010888596.1	PATPAAPAAGRSEPGRAGGFGR	418
NTDB id 1086 NMV RS00245 WP 002218487.1		347
NTDB id 1116 NGFG RS10615 WP 003690200.1		347
NTDB id 1185 GTF74 RS11940 WP 000350195.1		345
NTDB id 1176 A1552VC RS01160 WP 000350195.1		345
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consensus		

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