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NTDB id 1203 PSJM300 01605 AFN76401.1
NTDB id 1019 ACIAD RS04205 WP 004922051.1
NTDB id 1307 DR RS10055 WP 01088596.1
NTDB id 1086 NMV RS00245 WP 002218487.1
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NTDB id 1185 GTF74 RS11940 WP 000350195.1
NTDB id 1176 A1552VC RS01160 WP 000350195.1
NTDB id 1283 ACE17W RS02630 WP 010947729.1
NTDB id 1259 GCO85 RS10305 WP 010947729.1
NTDB id 1020 ACIAD RS04210 WP 011182164.1
NTDB id 1081 FDQ49 RS05335 WP 006581387.1
NTDB id 1060 ABD1 RS04380 WP 000355489.1
NTDB id 1040 HON27 RS13360 WP 000355489.1
NTDB id 218161 B1R35 RS27395 WP 003377212.1
NTDB id 1194 PAKAF RS01995 WP 003084552.1
NTDB id 1202 PSJM300 01600 AFN76400.1

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[illegible]

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NTDB id 1177 A1552VC RS01165 WP 000422572.1
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LT	ED	PI	EF	VH	Q	SKK	CL	IN	QR	EV	H	R	D	T	S	F	E	A	L	R	S	A	L	R	E	D	P	D	Y	I	L	V	G	E	R	D	E	T	I	R	L	A	L	T	A	A	E	T	G	H	L	V	F	G	T	L	H	T	S	A	A	K										
L	T	V	E	D	P	I	E	N	S	S	V	T	L	C	G	H	I	G	L	R	S	A	L	R	E	D	P	D	Y	I	L	V	G	E	R	D	E	T	I	R	L	A	L	T	A	A	E	T	G	H	L	V	F	G	T	L	H	T	S	A	A	K										
LT	VED	PI	EF	VH	EH	K	R	C	I	V	T	Q	R	E	V	G	L	D	T	S	E	V	E	A	L	K	N	S	L	R	Q	A	P	M	I	L	I	G	E	I	R	S	R	E	T	M	E	Y	A	M	T	F	A	E	T	G	H	L	C	M	A	T	L	H	A	N	N	A	N	Q		
IS	IED	PI	EF	I	H	Q	H	Q	N	C	I	V	T	Q	R	E	V	G	I	D	T	S	F	E	V	A	L	K	N	T	L	R	Q	A	P	V	I	L	I	G	E	V	R	T	R	E	T	M	D	H	A	V	A	F	A	E	T	G	H	L	C	L	A	T	L	H	A	N	N	A	N	Q
IT	IED	PI	EF	I	E	H	A	G	C	I	T	Q	R	E	V	G	I	D	T	S	F	E	I	A	L	K	N	T	L	R	Q	A	P	V	I	L	I	G	E	I	R	S	R	E	V	M	D	Y	A	L	A	F	A	E	T	G	H	L	V	F	A	T	L	H	A	N	N	A	N	Q		
LT	IED	PI	EF	M	H	P	N	K	A	S	I	N	Q	R	E	V	G	S	D	T	S	F	D	A	L	R	A	A	R	Q	A	P	V	I	L	V	G	E	M	R	D	Y	E	T	I	K	A	A	V	T	A	A	E	T	G	H	L	V	M	G	T	L	H	T	S	A	P					
LT	IED	PI	EF	VH	Q	SKK	S	L	I	N	Q	R	E	L	H	Q	H	T	L	S	F	A	N	A	L	R	S	A	L	R	E	D	P	V	I	L	V	G	E	M	R	D	P	E	T	I	G	L	A	L	T	A	A	E	T	G	H	L	V	F	G	T	L	H	T	G	A	A	K			
LT	IED	PI	EF	VH	Q	SKK	S	L	I	N	Q	R	E	L	H	Q	H	T	L	S	F	A	N	A	L	R	S	A	L	R	E	D	P	V	I	L	V	G	E	M	R	D	P	E	T	I	G	L	A	L	T	A	A	E	T	G	H	L	V	F	G	T	L	H	T	G	A	A	K			
LT	IED	PI	EF	VH	S	N	N	K	C	L	I	N	Q	R	E	V	H	R	D	T	H	S	F	K	N	A	L	R	S	A	L	R	E	D	P	V	I	L	V	G	E	L	R	D	Q	E	T	I	S	L	A	L	T	A	A	E	T	G	H	L	V	F	G	T	L	H	T	S	A	A	K	
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LT	V	E	D	PI	EF	VH	Q	SKK	CL	V	N	Q	R	E	V																																																									

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TLERIIHFFDADRHQQIFMDLSLNLKAVISQQLIPTHDQGSRAAVEVLINTPLSDYIRGETYKIKELMARSSELGMQ	316
SIDRIIDVFPPEEQEQRIRVQLANNLVAVMTQQLPRADGGQRLAYEILLANPAVRALIREKGTFTQISVMQTGAREGMV	319
TIDRIIDVFPAGEKEMVRSMLSESLTAVISQNLKTHDGNGRVASHEILLIANPAVRNLIRENKITQINSVLQTGGASGMQ	315
TVDRIIDVFPAGEKEMVRSMLSESLTAVISQNLKTHDGNGRVASHEILLIANPAVRNLIRENKITQINSVLQTGGASGMQ	315
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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
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NTDB id 1307 DR RS10055 WP 010888596.1	TMDAYLAGLFRHHLITYDKGLERAIDPKFARLANDASAGL....GGASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	395
NTDB id 1086 NMV RS00245 WP 002218487.1	TMDQSLQSLVRQGLIAPEVARRRAQNSESMSF.....	347
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NTDB id 1185 GTF74 RS11940 WP 000350195.1	TMEQNAKQLIARGVVDAQEVQSKIELDLKAF.....	345
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NTDB id 218161 B1R35 RS27395 WP 003377212.1	TLDMCLADLVKRGLITRESARERAKVPDNF.....	344
NTDB id 1194 PAKAF RS01995 WP 003084552.1	TLDMCLKGLVAKGLISRAREKAKIPENF.....	344
NTDB id 1202 PSJM300 01600 AFN76400.1	TLDSCCLKGLIAKGLISRDSAKEKAKQPENF.....	344
consensus	!*** * !* ***** * *	

logo

NTDB id 1177 A1552VC RS01165 WP 000422572.1	PATPAAPAAACRSEPCRRACCFRR	368
NTDB id 1203 PSJM300 01605 AFN76401.1		381
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NTDB id 1020 ACIAD RS04210 WP 011182164.1		344
NTDB id 1081 FDQ49 RS05335 WP 006581387.1		345
NTDB id 1060 ABD1 RS04380 WP 000355489.1		345
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NTDB id 1202 PSJM300 01600 AFN76400.1		344
consensus		

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