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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 010888596.1
NTDB id 1086 NMV RS00245 WP 002218487.1
NTDB id 1116 NFGF RS10615 WP 003690200.1
NTDB id 1185 GTF74 RS11940 WP 000350195.1
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NTDB id 1259 GCO85 RS10305 WP 010947729.1
NTDB id 1020 ACIAD RS04210 WP 011182164.1
NTDB id 1081 FDQ49 RS05335 WP 006581387.1
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NTDB id 22735 PSPTO RS26095 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 1194 PAKAF RS01995 WP 003084552.1
NTDB id 1202 PSJM300 01600 AFN76400.1
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EKNERFRVSAFQQRDMPGMVIRRIETIKTIPSPDDLHLPLLELAMSKRGITIFSGATGTGKSTSLASMIQHRNQHSKGI	156
D. KARFRVNAFMQRGNVGCVMRLIPTITVRITAEMLGPKQTVIDIANPRLGLVLTGPTGSGKSTTLASIMIDHINTTKLHI	159
N. VARFRVNAFNGRGVAAVFRPTIPSTVLSLEELKAPSIQKIAESPGRMVLVTGPTGSGKSTTLAAMINYINETQPAHI	155
N. VARFRVNAFNTGRGPAAVFRPTIPSTVLSLEELKAPSIQKIAESPGRMVLVTGPTGSGKSTTLAAMINYINETQPAHI	155
N. VGRFRVNAFHQARGCSAVFRPTIPTVIPTLEQLDAPEIFSKIANYEKGLVLVTGPTGSGKSTTLAAMVNYVNAHHNKHI	155
N. VGRFRVNAFHQARGCSAVFRPTIPTVIPTLEQLDAPEIFSKIANYEKGLVLVTGPTGSGKSTTLAAMVNYVNAHHNKHI	155
N. LARFRVNAFNQSRGAAAVFRPTIPSEILSMEDLGLPPIFKEMASFSRGLVLVTGPTGSGKSTTLAAVIDYINSSRYEHI	155
N. LARFRVNAFNQSRGAAAVFRPTIPSEILSMEDLGLPPIFKEMASFSRGLVLVTGPTGSGKSTTLAAVIDYINSSRYEHI	155
N. LARFRVNVFNQNRGAGAVFRPTIPSKVLTLEELGLGKIFRDI CDYPRGLVLVTGPTGSGKSTTLAAMDYINNHRYDHI	155
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G. VARFRVNAFNQNRGAGAVFRPTIPSKILSMEDLCMGSVFRKITDVARGLTILVTGPTGSGKSTTLAAMDYLNCKNHHHI	155
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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										
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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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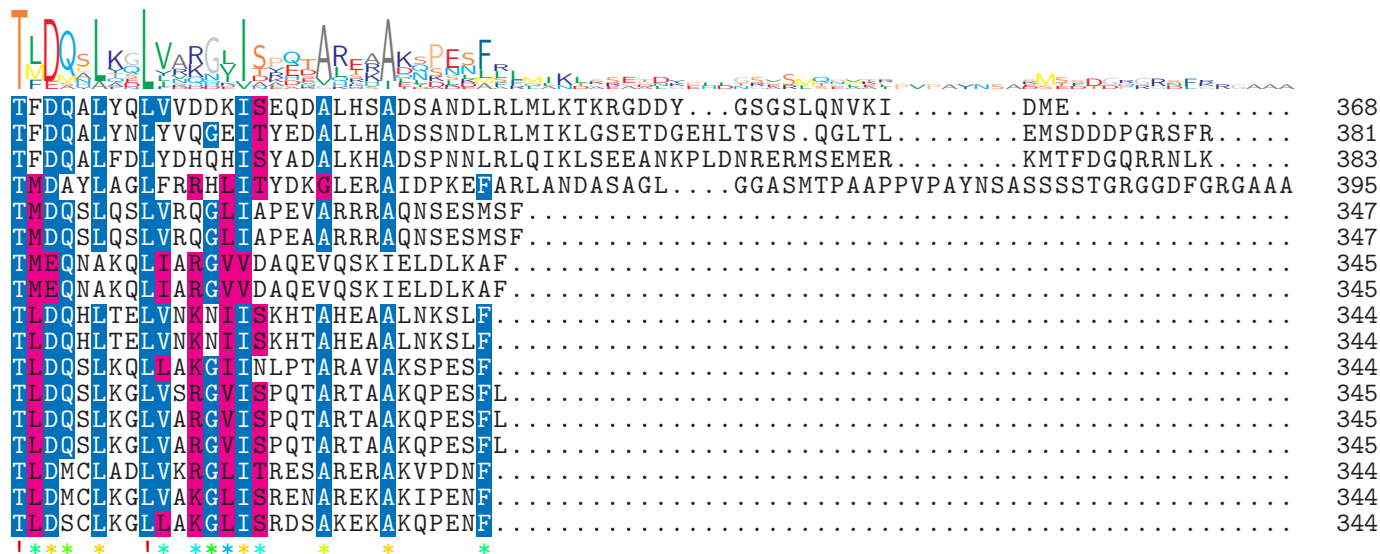
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consensus



logo

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consensus

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 PATPAAPAAAGRSEPGRAGGFGRR
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 344

-  non conserved
-  similar
-  >50% conserved