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NTDB id 1410 SGL RS13950 WP 010873519.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	0
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
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NTDB id 1293 VP RS12240 WP 005479695.1		0
NTDB id 1401 DSB67 RS12665 WP 010643259.1		0
NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 299282 DPV67 RS15815 WP 001274987.1		0
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 1410 SGL RS13950 WP 010873519.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	0
NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	160
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consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	EGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDPSAALLL	0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
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consensus		

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NTDB id 1251 GC085 RS07725 WP 011946523.1		0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

NTDB id	1410	SGL RS13950 WP 010873519.1	SKKSRVAVALRNYFSPFGNKL [*] VAGCHVD ^{*!} AEQLRQALVQVKKTGRSLPEELKAVTGRELSPELLRQYKKNQLFELKVLYGVD	86
NTDB id	1028	TT RS08215 WP 011173991.1	KEAIV [*] E [*] VQKGLPFAKPVGETLVELCLARPEDVEEALQKQRGGRRLEDTLVQ..SGKLRPEALAQA.....VATQLGYP	391
NTDB id	1168	A1552VC RS11070 WP 000957200.1	MLTNLVALLRQAELISATQECAVVTVQSASGTSVPEALLE..LSIFHAQELTEQ.....LSHIFGLP	60
NTDB id	1293	VP RS12240 WP 005479695.1	MHSNLSTILRLRKQGILTFSSQEESLIEQVKASGISMPALLS..SGFFTSSELAEH.....LSSIIFGLS	60
NTDB id	1401	DSB67 RS12665 WP 010643259.1	MLTINLTSTRVLRAQIGILTFSQEESLIHQVKASGISMEALLS..SGFFGSHELTEH.....LSSFVGLS	60
NTDB id	1110	NGFG RS09235 WP 003689817.1	MSVGLLRLLVQNQQVITVEQAAEHYY.NESQACKEVLPMFLFS..DGVISPKS LAAL.....IARVFYSYS	59
NTDB id	1251	GCO85 RS07725 WP 011946523.1	...MALATEEYRLQTIGQLLVLEKLLDKTKAIELHKLA-AAEKISLLQYIVK..NKILSAEQIALT.....AAQNFQGP	68
NTDB id	299282	DPV67 RS15815 WP 001274987.1	...MSALHTSPKFTGFFFRRLVEEKHVSAATMQTALDAAKRAKQDTVAYLIE..EVHLSPSLLAET.....ISAEEFAEP	68
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NTDB id	1018	ACIAD RS01685 WP 004920473.1	...MSAF [*] TTPPKFKSFIRRLVEEGYNANQMMAALEKAKKFKDIPYLID..NFSISPLTI AEI.....ISLEFGEP	68
consensus			* **** !* ***** *** ***** *	

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NTDB	id	1028	TT	RS08215	WP	011173991.1	Y	V	D	P	E	E	D	P	P	D	P	.	.	.	.	.	.	G	A	P	L	L	P	E	D	L	C	R	R	Y	G	V	F	P	H	R	L	E	.	.	G	N	R	L	V	L	L	M	K	D	P	R	N	I	L	A	L	D	D	V	R	L	A	L	K	R	K	G	L	N	Y	E	V	A	P	A	V	A	T	463		
NTDB	id	1168	A1552VC	RS11070	WP	000957200.1	E	T	D	L	S	R	Y	D	Y	A	N	.	.	.	.	.	.	L	C	Q	Q	L	G	L	R	E	L	I	T	R	Y	D	A	L	P	I	A	K	Q	.	.	G	N	L	L	L	A	V	S	D	P	T	L	L	Q	A	E	E	E	F	R	F	A	T	G	.	.	.	.	.	.	L	Q	V	E	L	A	L	A	D	127	
NTDB	id	1293	VP	RS12240	WP	005479695.1	Q	P	E	L	S	Q	Y	E	Y	A	S	.	.	.	.	.	.	L	C	Q	Q	L	G	L	R	E	L	I	T	R	H	N	A	L	P	L	H	R	T	.	.	P	S	T	L	L	L	A	V	A	D	P	T	N	Q	Q	A	E	D	D	F	R	F	A	T	G	.	.	.	.	.	.	L	Q	V	E	L	V	L	A	D	127
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NTDB	id	1058	ABD1	RS01615	WP	001274986.1	Y	F	D	L	D	V	Y	D	T	S	Q	.	.	.	.	.	.	I	P	K	D	L	V	D	Q	K	L	I	L	K	H	R	I	L	P	L	I	Q	R	.	.	G	Q	I	L	Y	V	A	T	S																																

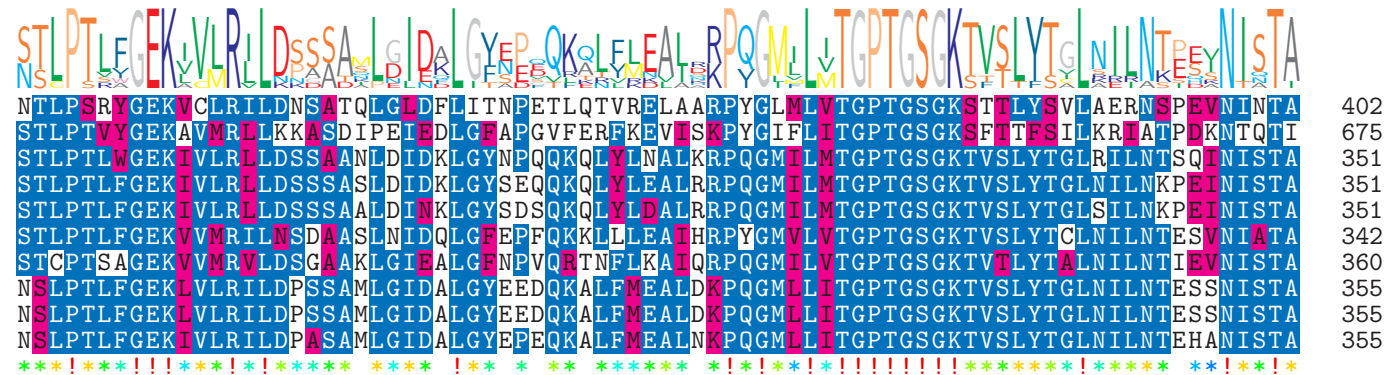
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consensus
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consensus

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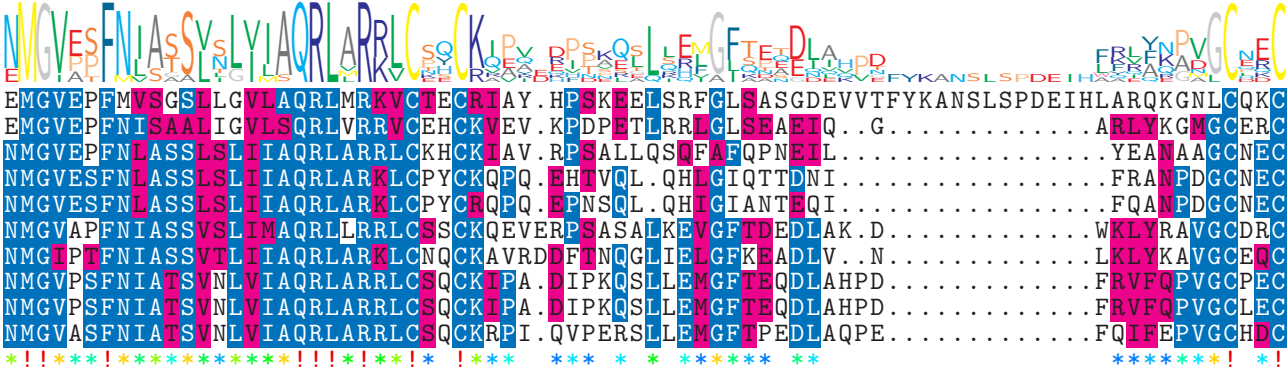
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consensus



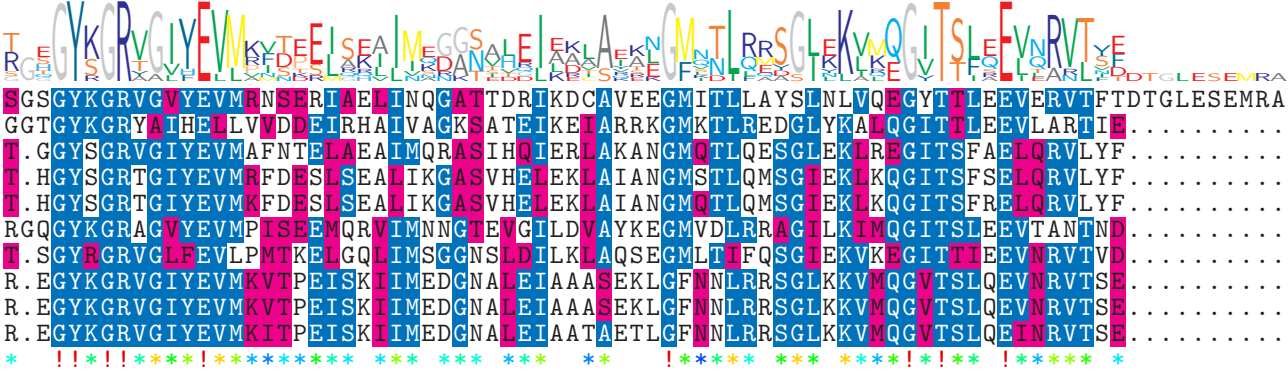
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consensus



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NTDB id 1410 SGL RS13950 WP 010873519.1
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consensus



logo

NTDB id 1410 SGL RS13950 WP 010873519.1	KRKSALTCRTCAAELQQEWLDCPYCMTSRFS	672
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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved