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NTDB id 1410 SGL RS13950 WP 010873519.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	0
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
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 1018 ACIAD RS01685 WP 004920473.1		0
NTDB id 668414 MOW14 RS01525 WP 168370330.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	0
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consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	EGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL	0
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consensus		

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NTDB id 1028	TT RS08215 WP 011173991.1	LRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARYLALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL	320
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NTDB id 1058	ABD1 RS01615 WP 001274986.1		0
consensus			

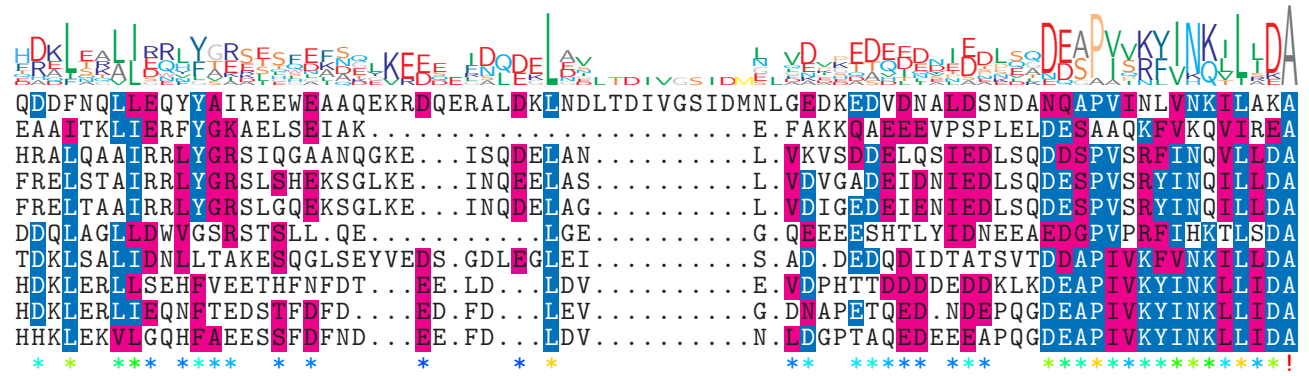
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NTDB id 1293	VP RS12240 WP 005479695.1	MHS	NLST	ILRQGLLTFSQ	EESLIEQVKASGISMEALLS..SGFFTSSELAEH.....	LSSIFGLS	60		
NTDB id 1401	DSB67 RS12665 WP 010643259.1	MLT	NLST	ILVRQAGILTFS	QEESSLIEQVKASGISMEALLS..SGFFQSHELTEH.....	LSSVFGLS	60		
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NTDB id 1251	GC085 RS07725 WP 011946523.1	...MALATEEY	RLQGIGQL	LVLEKLLDKTKAIELHKLA	AAEKISLLQYIVK..NKILSAEQIALT.....	AAQNFGVP	68		
NTDB id 1018	ACIAD RS01685 WP 004920473.1	...MSAFTTPPK	FSGFIRRLVEEGYVNAQNMQQALEKA	KKFKQDIVPYLID..NFSISPLTIAEI.....	ISLEFGEP	68			
NTDB id 668414	MOW14 RS01525 WP 168370330.1	...MSVQQAMP	RFTGFIRRLVEEGLVTAENMQQATIVA	AKKAETDIVPYLIQ..QLQISPLTIAEK.....	ISYEFGEP	68			
NTDB id 1058	ABD1 RS01615 WP 001274986.1	...MSALHTSP	KFTGFFRR	LVEEKHVSAATMTALDAAK	RKAKQDTVAYLIE..EVHLSPSLIAET.....	ISAFAEP	68		
consensus		*	* * * *	* ! *	* * * * *	* * * *	* * * *	* * * * *	

NTDB id 1410	SGL RS13950 WP 010873519.1	SVDPEVAPIATQQMAELIGRFFPLDTCRRFKFLPLAQEGEDPPSVLVAMVPDPNLAQAQDELNRILRVKGF..ELRRLLVV	164
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NTDB id 1293	VP RS12240 WP 005479695.1	QP E LSQY E YA.....SLCQTGLRL E LITHNALPLHRT...PSTLLLAVADPTNQQAEDDFRFATG.....LQVELVLAD	127
NTDB id 1401	DSB67 RS12665 WP 010643259.1	CTSLNQYEYA.....SLCQTGLRL E LITHNALPLNRT...SSTLILAVADPTNIQAEDDFRFATG.....LQVELVLAD	127
NTDB id 1110	NGFG RS09235 WP 003689817.1	I D DLRHYP R H.....RVLMGVLT E EQMVEFHCVPVFR...GDKVFFAVSDPTQMPQIQKTVSAAAG.....I A VELVIVE	126
NTDB id 1251	GCO85 RS07725 WP 011946523.1	M L DINCIDVG.....TIPVNLVNEKLIKRHAMVPLFSR...GTNLYLATDDPSKQASLKEIQFHTG.....LNTHAIVIVE	135
NTDB id 1018	ACIAD RS01685 WP 004920473.1	L L DLGVFDPA.....LFLKDKIDEKLIQKYRIMPLVHR...GHVLYVATSNPNTIEAMD A IRFN S K.....LK V EP I IVE	135
NTDB id 668414	MOW14 RS01525 WP 168370330.1	IFDLAVYDSR.....LIVREGLD E KLF T KYRI L P F FR...HNQLYIATSNPTNIEAIDAIRFN S K.....LNIEV V IVE	135
NTDB id 1058	ABD1 RS01615 WP 001274986.1	YFDLDVYDTS.....QIPKDLVDQKLLIKHL R LPLIQR...GQILYVATSNPN N IEAIDAIRFN S K.....LLVEPVIVE	135
consensus		*** * * * * * * * * ! * * * * * * * * * * *	

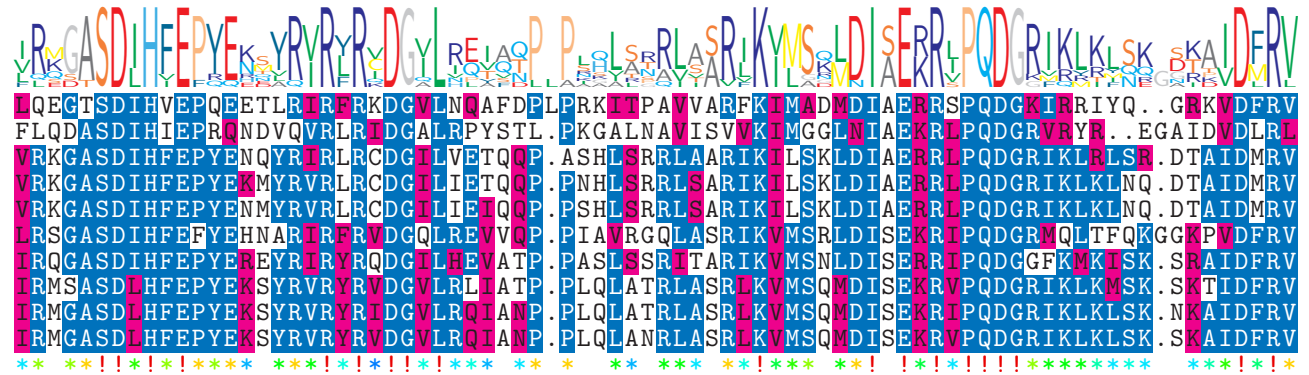
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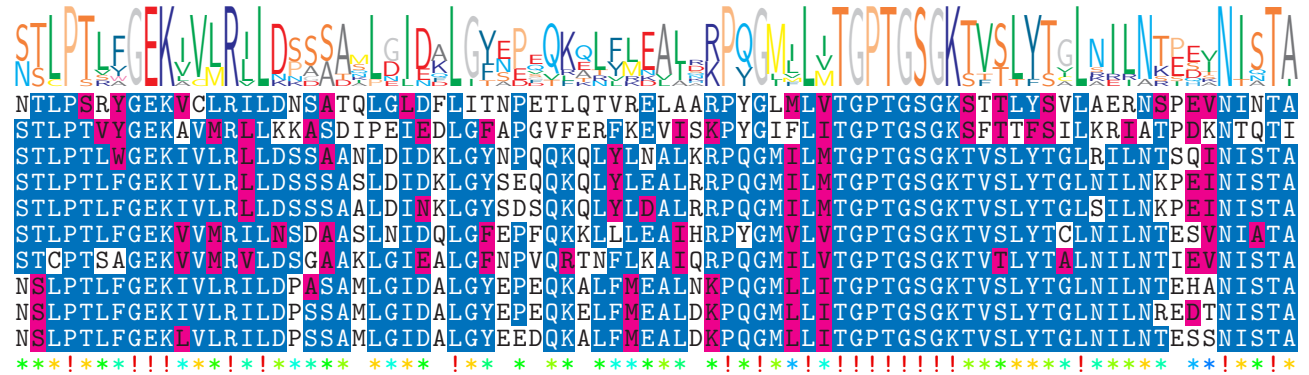
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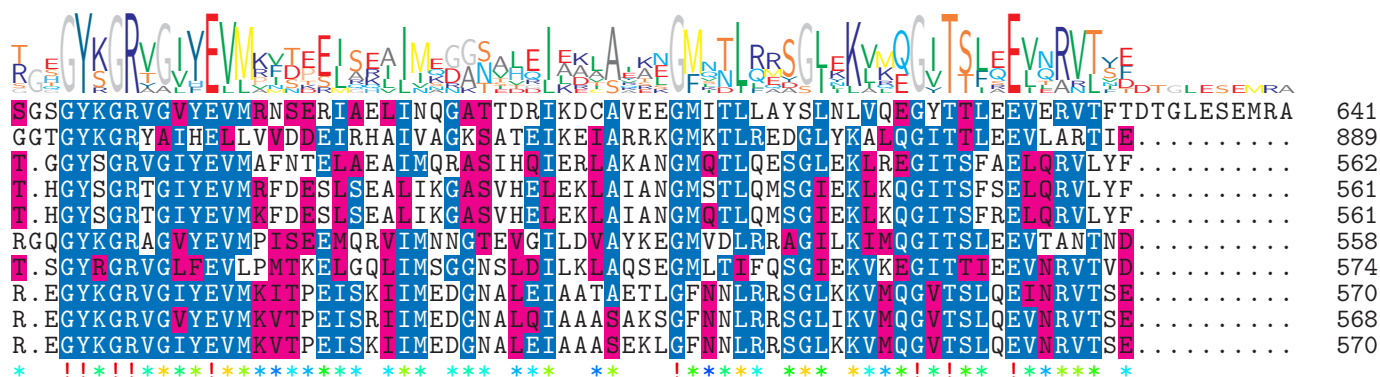
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consensus



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consensus



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consensus



logo

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NTDB id 1110 NGFG RS09235 WP 003689817.1		558
NTDB id 1251 GC085 RS07725 WP 011946523.1		574
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		570
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved