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consensus		

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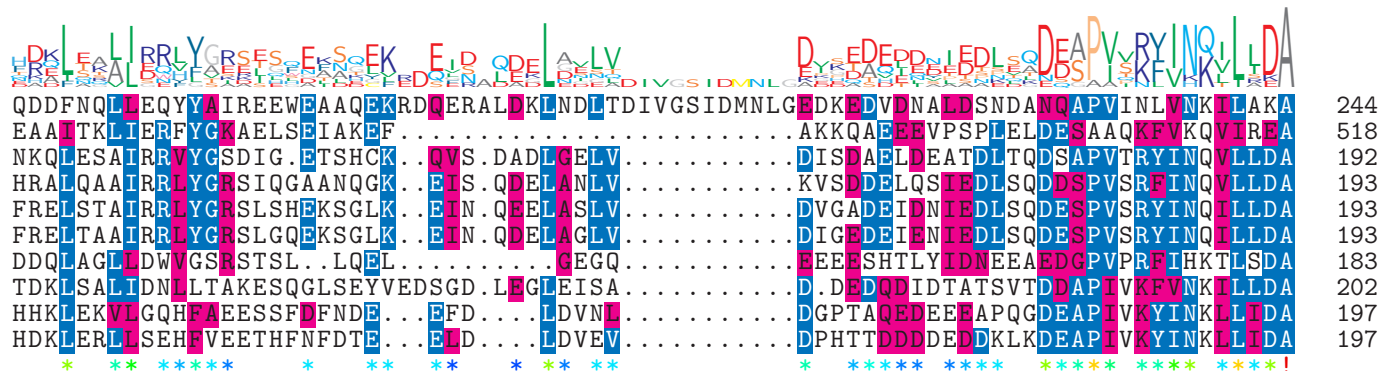
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consensus		

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NTDB id	1028	TT RS08215 WP 011173991.1	KEALEVQKGLPRAKPLGEITL	VELGLARPEDVEEALQKQRGGGRLEDTLVQ	.SGKL RPEALACA.....VATQLGY P	391
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NTDB id	1168	A1552VC RS11070 WP 000957200.1	MLTNLAIVAILRQA	ELISATQEQAVTQVSASGTSPPEALIE	.LSIFHAQELTEQ.....LSHIFGLP	60
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NTDB id	1058	ABD1 RS01615 WP 001274986.1	...MSALHTSPKFIFGFRR	LVEEKHVSAA TMQTALDAAKRAKQDTVAYLIE	.EVHLSPSLLAET.....ISAFAEP	68
NTDB id	1018	ACIAD RS01685 WP 004920473.1	...MSAFTTPPKFSGFIRRL	LV EECYNNAQNMQQA L EKAKFKQDIVPYLID	.NFSISPLTI A EI.....ISLEFGEP	68
consensus			* * *	*! * * * * *	* * * * * * * * * * * * * *	* * * * *

NTDB id	1410	SGL RS13950 WP 010873519.1	SVDPEVAPIATQQMAEIGRFFPLDTCRRFKFLPLAQEGEDPPSVLVAIVDPDNAAQDELNRILRVKGF..ELRRLVVT	164
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NTDB id	1293	VP RS12240 WP 005479695.1	QPQLSQVEYA.....SLCQQLGLRELITRHNALPLHRTPT...STLLLAVDPTNQQAEDDFRFATG....LQVELVLAD	127
NTDB id	1401	DSB67 RS12665 WP 010643259.1	CTSLNQVEYA.....SLCQTLLGLRDILTRHNALPLNRTS...STLLLAVDPTNLQAEDDFRFATG....LQVELVLAD	127
NTDB id	1110	NGFG RS09235 WP 003689817.1	IIDLRLHYPRH.....RVLMGVLTTEEQWVEFHCVPVVFRRG...DKVFFFAVSDDPTQMPQIQKTVSAAG....IATELVIVE	126
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NTDB id	1058	ABD1 RS01615 WP 001274986.1	YFDLDVYDTS.....QIPKDLVDQKLILKHRILPLIQRG...QILYVATSNSPNIIEAIDAIRFNASK....LLVEPVIVE	135
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consensus			* * * * *	

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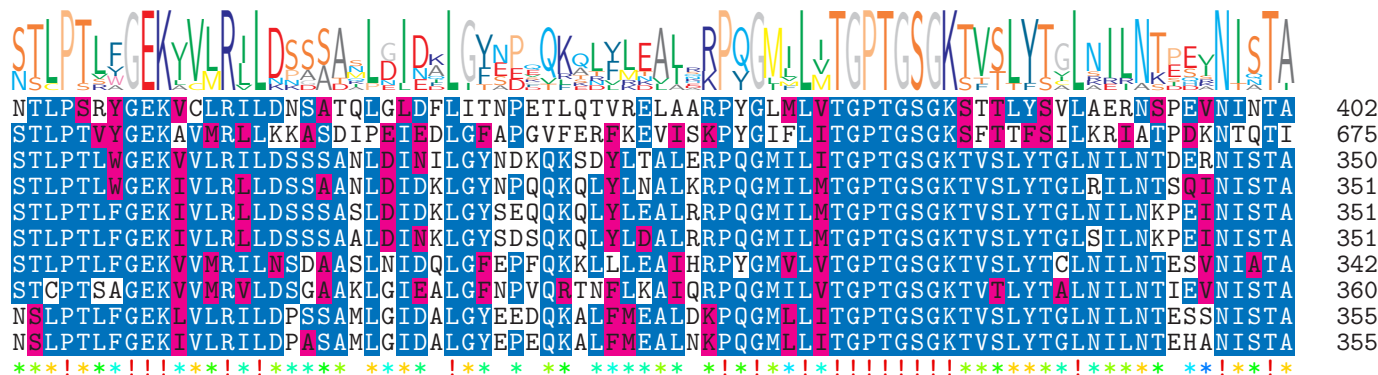
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consensus



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consensus



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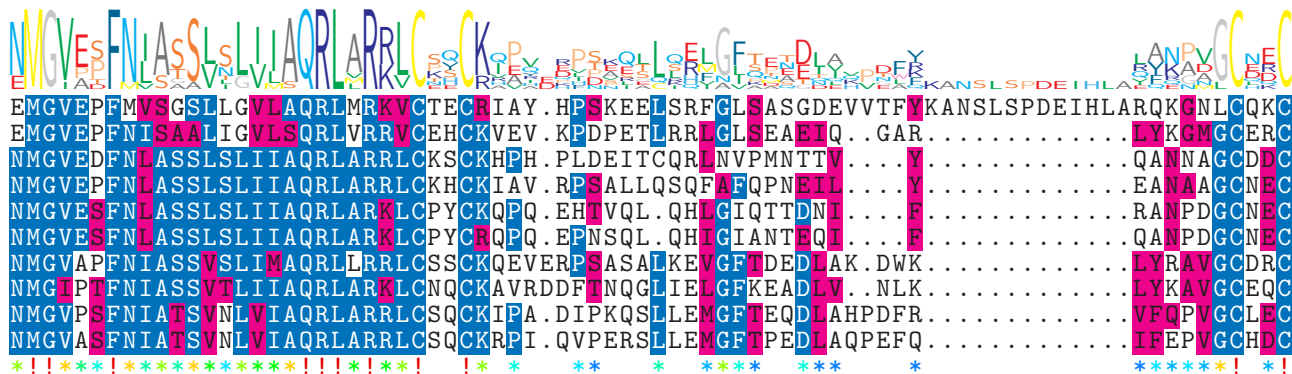
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consensus



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logo

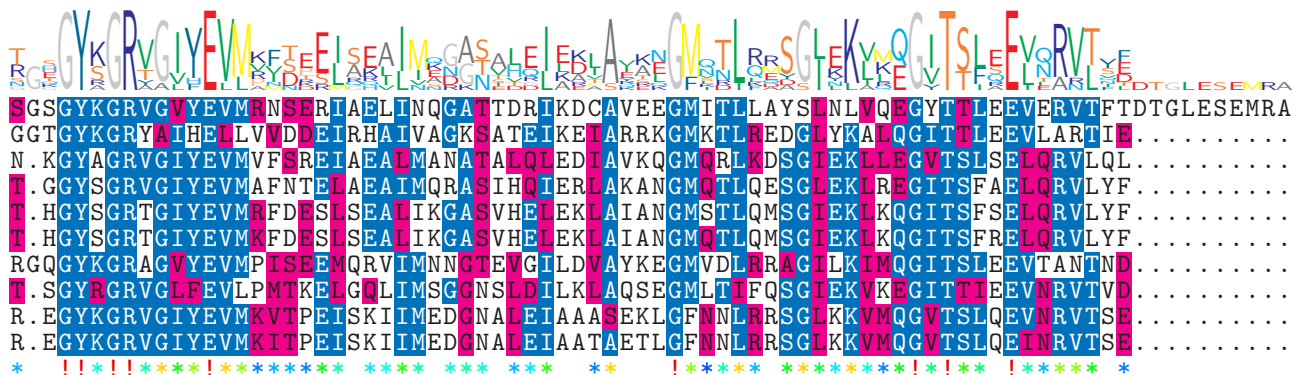
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consensus



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consensus



641
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561
562
561
561
558
574
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570

logo

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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved