

logo

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NTDB id 1028 TT RS08215 WP 011173991.1		MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
NTDB id 1168 A1552VC RS11070 WP 000957200.1			0
NTDB id 1401 DSB67 RS12665 WP 010643259.1			0
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NTDB id 1293 VP RS12240 WP 005479695.1			0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1			0
NTDB id 1018 ACIAD RS01685 WP 004920473.1			0
consensus			

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		SLTPAEKAKELRAIPFALDEEAGVVRVAFLNPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLEIVVPSGA	
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NTDB id 1018 ACIAD RS01685 WP 004920473.1			0
consensus			

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consensus			

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NTDB id 1028 TT RS08215 WP 011173991.1	LLLLRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSR	317
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consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	EALKEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLED TLVQ..SGKLRPEALAA.....VATQL	388
NTDB id 1168 A1552VC RS11070 WP 000957200.1	MLTNLVA ILRQAE LISATQE QAVVTQVSASGTSVPEALLE..LSIFHAQELTEQ.....LSHIF	57
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NTDB id 168327 AL552 RS13105 WP 104969782.1	MVTNLST VLRQSG LLSLAQEESLLEQVKASGISMPEALLS..SGFFTSSSELVEH.....LSSLF	57
NTDB id 1293 VP RS12240 WP 005479695.1	MHSNLST ILRQKGLLTFSQEESLIEQVKASGISMPEALLS..SGFFTSSSELAEH.....LSSIF	57
NTDB id 1110 NGFG RS09235 WP 003689817.1	MSVGLLR ILVQNQ VVTVEQA EHY..NESQAGKEVLPMLFS..DGVISP KSLAAL.....IARVF	56
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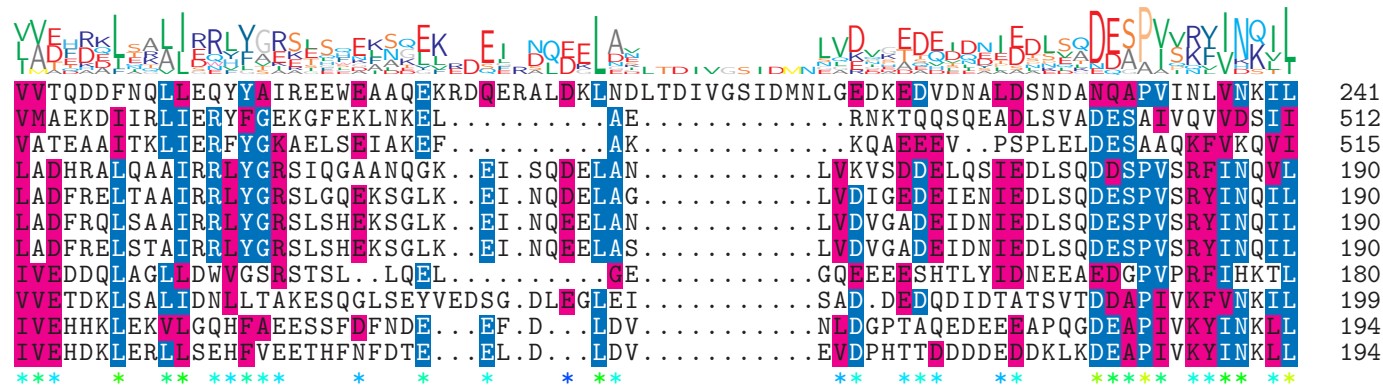
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consensus	* * * * *	

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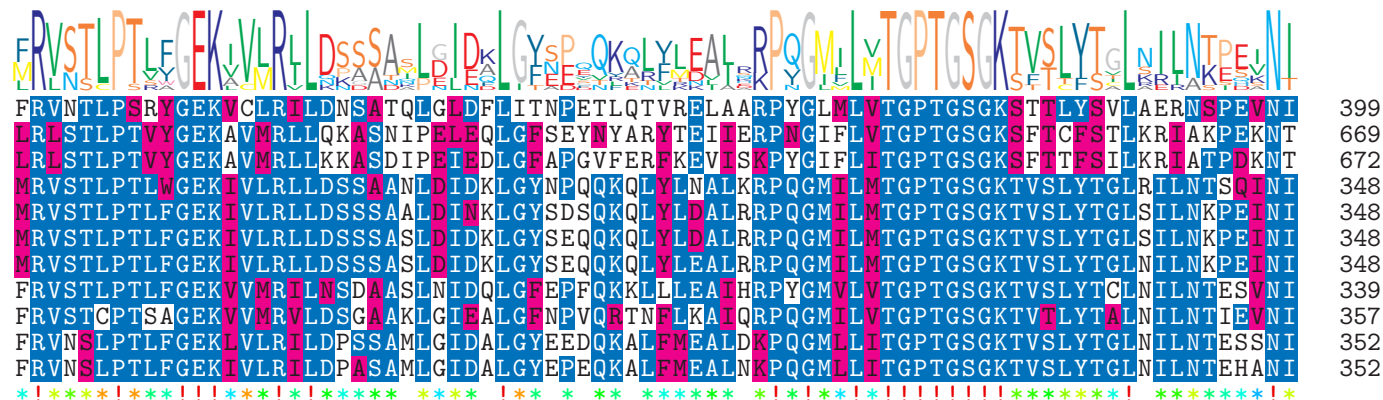
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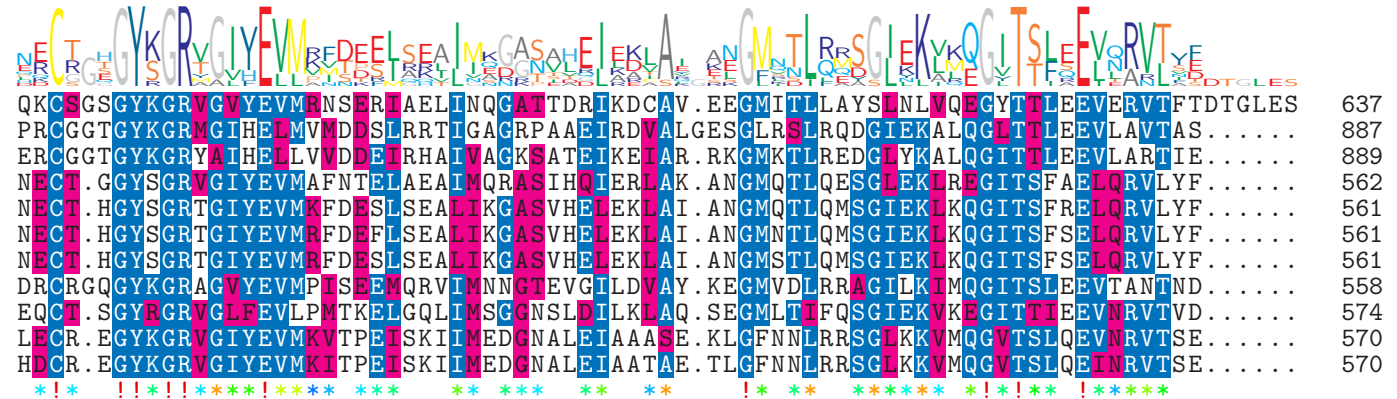
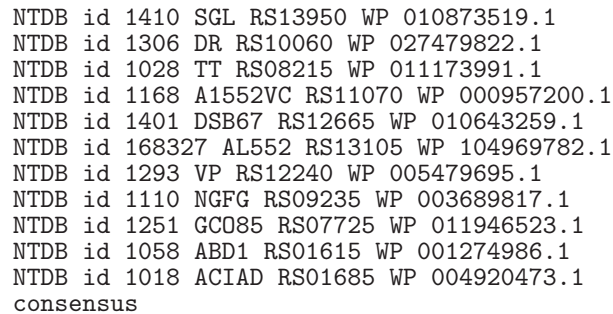
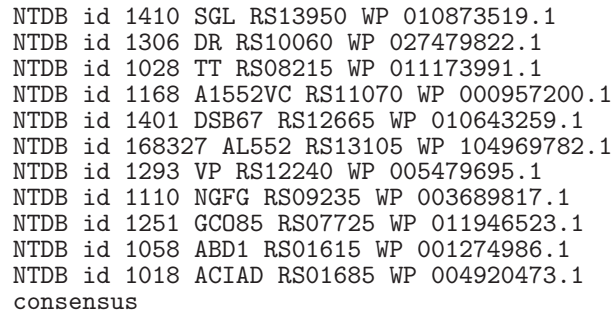
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consensus

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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved