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NTDB id 1028 TT RS08215 WP 011173991.1		MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKK	80
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NTDB id 1018 ACIAD RS01685 WP 004920473.1			0
consensus			

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consensus			

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		SDAGRTGGKLGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFGHPGEEVL	
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consensus		

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NTDB	id 1028	TT RS08215 WP 011173991.1	EALKEALEVQKGFLPRAKP LGEILVELGIARPEDVEEALQKQRGGGRLEDTLVQ..SGKL RPEALAQA..... VATQL	388
NTDB	id 1168	A1552VC RS11070 WP 000957200.1	MLTNLVA ILRQAE LISATQEAV VTQVSASGT SVPEALLE..LSIFHAQELTEQ..... LSHIF	57
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NTDB	id 408508	GPY32 RS12915 WP 005493983.1	MLTNLST ILRQG LLTFP QEESLF EQIKASGISMPEALLS..SGFFT SSELA EH..... LSSIF	57
NTDB	id 1293	VP RS12240 WP 005479695.1	MHSNLST I LRQK GLTFS QEESL IEQVKASGISMPEALLS..SGFFT SSE LAEH..... LSSIF	57
NTDB	id 1110	NGFG RS09235 WP 003689817.1	MSVGLLR I LVQNQ VVTVE QA EHY Y.NESQAGKEVLP MLFS..DGVISP KSLA AL..... I AR VF	56
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NTDB	id 1058	ABD1 RS01615 WP 001274986.1	MSALHTSPKF T GFFRR LVEEKHV SAATMQTA LDAAKRAKQDT VAYLIE..EVHLSP SLLA ET..... TSAEF	65
NTDB	id 1018	ACIAD RS01685 WP 004920473.1	MSAF TT PPKF SG FIRRVLEE GYVNAQM NQA L EKA KKFKQ D IVPYLID..NFS IS PLTI AEI..... LSLEF	65
consensus			* * * * *	

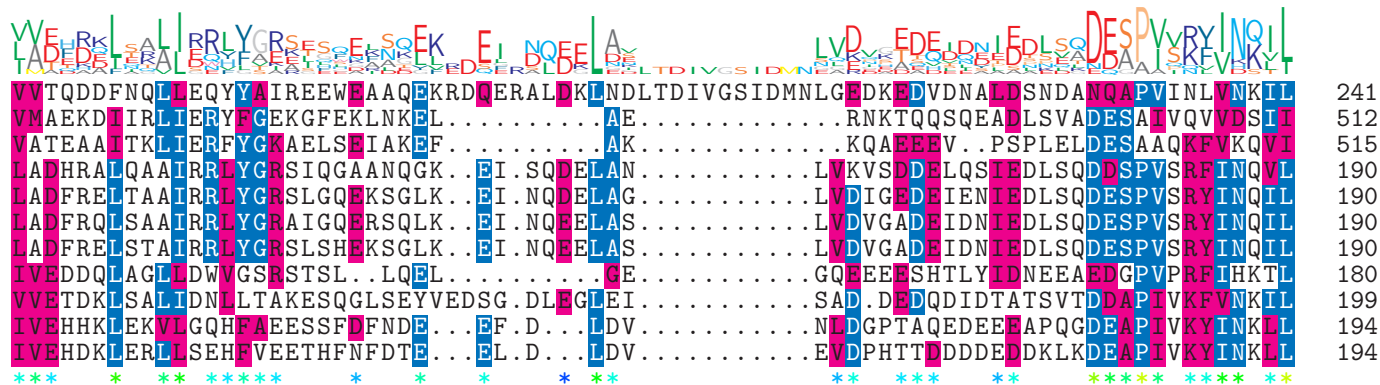
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	AEYPFDLDVYDTS....QIPKDLVDQKLILKHRIPLLIQR...GQILYVATSNPSNIEAIDAIRFNSK....LLVEPV	132
NTDB id 1018 ACIAD RS01685 WP 004920473.1	GEPLDLGVFDPA....LFLKDKIDEKLIQYRIMPLVHR...GHVLYVATSNPTNIEAMDAIRFNSK....LKVEPI	132
consensus	*** **	

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consensus

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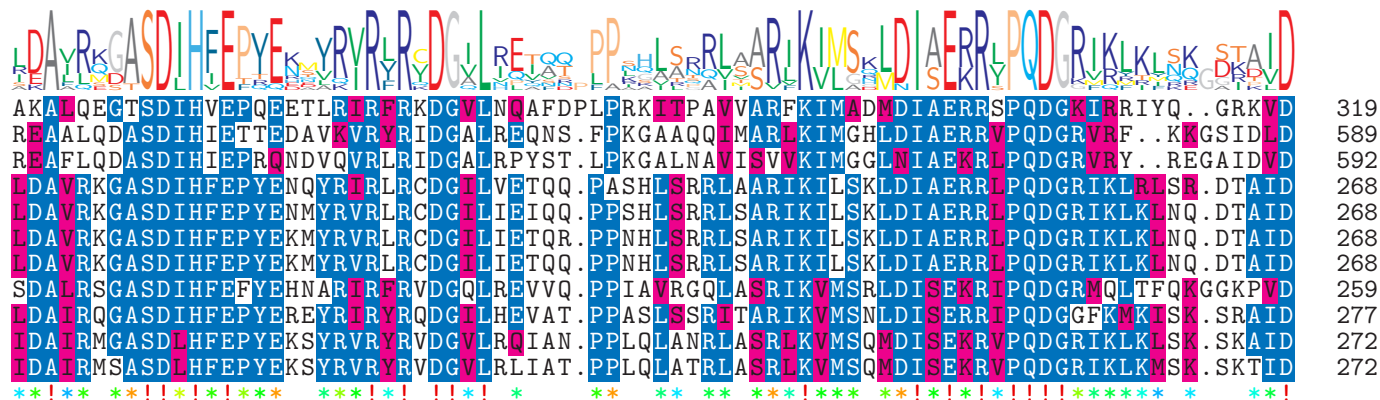


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consensus

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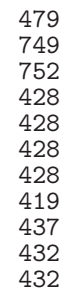
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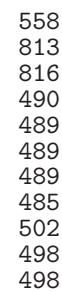

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consensus
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consensus

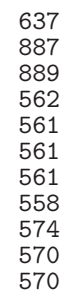
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved