

logo

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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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NTDB id 1018 ACIAD RS01685 WP 004920473.1			0
consensus			

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NTDB id 1028 TT RS08215 WP 011173991.1		SGEGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAA	237
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consensus			

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consensus		

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LLLLRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSR	317
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NTDB id	1018	ACIAD RS01685 WP 004920473.1
consensus		

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DQLREALQVQAREGKVKPLGEVITELGFASPDVDSALQKQNVGGGRLEDTLVQ..SGKLSPEMLARS.....LAAQL	388
EALKEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQ..SGKLRPEALAA.....VATQL	388
.....MLTNLVAILRQAEILSATQEAVVTQVSASGTSVPEALLE..LSIFHAQELTEQ.....LSHIF	57
.....MLTNLSTVLRQAGILTFSQEESLLEQVKASGISMPEALLS..SGFFQSHELTEH.....LSSVF	57
.....MLTNLSTVLRQKGLLTFSQEESLIEQIKASGISMPEALLS..SGFFTSSSELAEH.....LSSIF	57
.....MHSNLTILRQKGLLTFSQEESLIEQVKASGISMPEALLS..SGFFTSSSELAEH.....LSSIF	57
.....MSVGLLRILVQNQVVTVEQAEHYY..NESQAGKEVLPMLFS..DGVISPKSALAAL.....IARVF	56
.....MALATEEYRLQGITQLLVLEKLLDKTKAIELHKLAAAEKISLLQYIVK..NKILSAEQIALT.....AAQNF	65
.....MSALHTSPKFTIGFFRRLVEEKHVSAAATMQTALDAAKRAKQDTVAYLIE..EVHLSPSLLAET.....ISAEF	65
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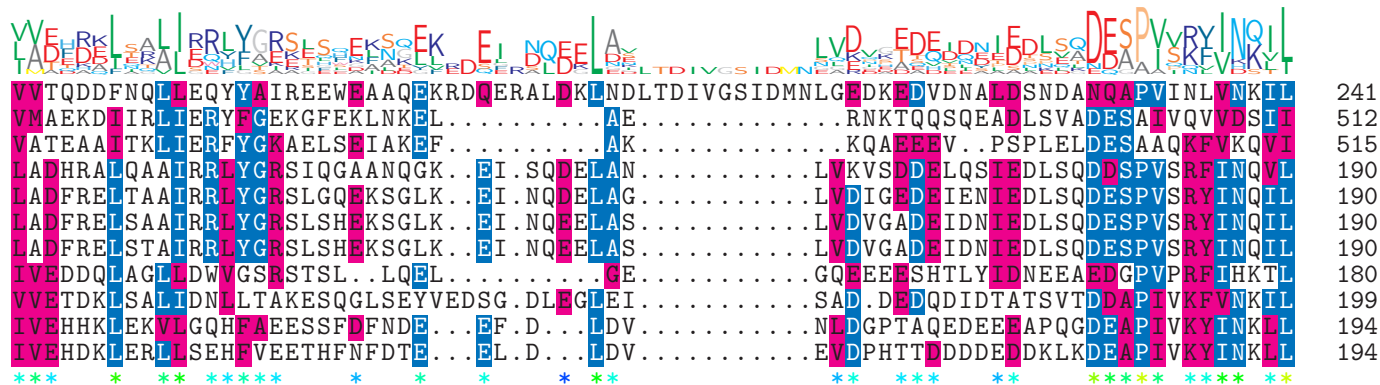
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GYEFLDPIQNPPD....PKVALMPEATARRVVVPVRLQ...GNSLVVAMKDPRNVFALDDDLKLITG.....KEILPA	455
GYPYVDPEEDPPD....PGAPLLLPEDLCRRYGVFPHRLE...GNRLVILMKDPRNILALDDVRLALKRKGLNVEVAPA	460
GLPETDLSRYDYA.....NLCQQLGLRELITRYDALPIAKQ...GNLLLLAVSDPTLLQAEEDFRFATG.....LQVELA	124
GLSCTSLNQYEYA.....SLCQTLGLRDLITRHNAPLNRT...SSTLILAVADPTNLQAEDDFRFATG.....LQVELV	124
GLSQPELSQYEYA.....SLCQQLGLRELITRYNALPLHRT...PSTLLAVADPTNQAEDDFRFATG.....LQVELV	124
GLSQPELSQYEYA.....SLCQQLGLRELITRHNAPLHRT...PSTLLAVADPTNQAEDDFRFATG.....LQVELV	124
SYSILDLRHYPRH....RVLMGVITEEQMVEFHCVPVFRR...GDKVFFAVSDPTQMPQIQKTVSAAG.....IAVELV	123
GVPMLDINCDVIG....TIPVNLVNEKLIKRRHAMVPLFSR...GTNLVYLATDDPSKQASLKEIQFHTG.....LNTHAI	132
AEPYFDLDVYDTS....QIPKDLVDQKLILKHRILPLIQR...GQILYVATSNPSNIEAIDAIRFNASK....LVPEPV	132
GEPLIDLGVFDPA....LFLKDKLIDEKLIQKYRIMPLVHR...GHVLYVATSNPTNIEAMDIAIRFNASK....LKVEPI	132
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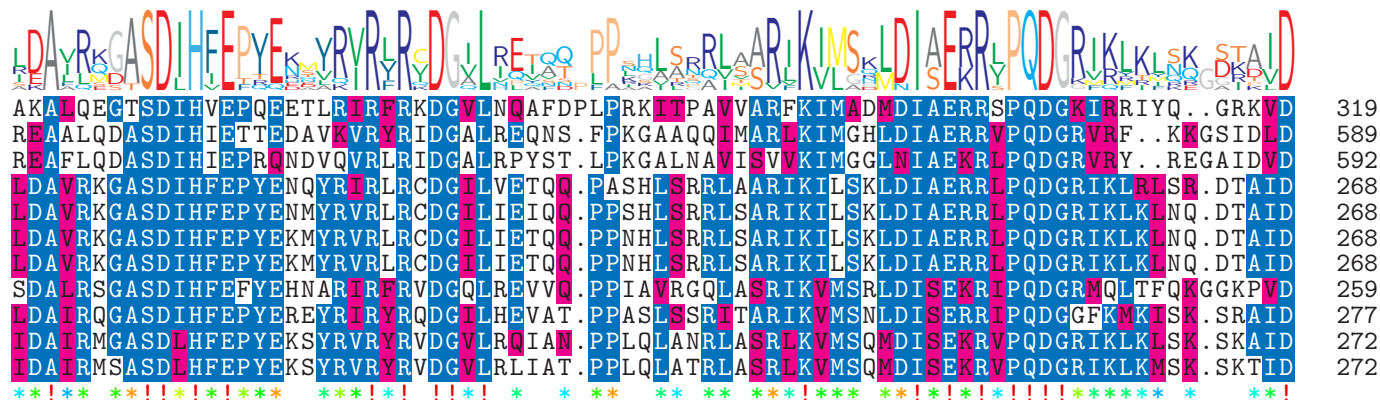
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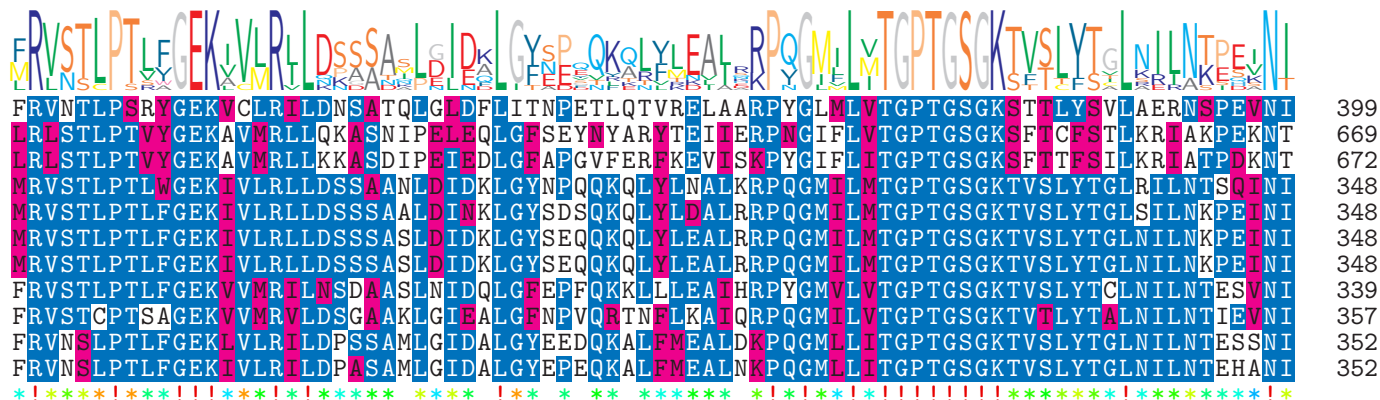
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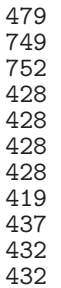
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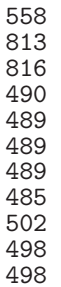
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
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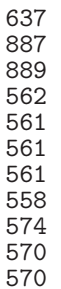
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consensus

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logo

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NTDB id 1110 NGFG RS09235 WP 003689817.1		558
NTDB id 1251 GC085 RS07725 WP 011946523.1		574
NTDB id 1058 ABD1 RS01615 WP 001274986.1		570
NTDB id 1018 ACIAD RS01685 WP 004920473.1		570
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved