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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
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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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NTDB id 1410 SGL RS13950 WP 010873519.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	0
NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	160
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NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	EGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDPSAALLL	0
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consensus		

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NTDB id 422116	G0027 RS06465 WP 016659655.1		0
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NTDB id 1018	ACIAD RS01685 WP 004920473.1		0
consensus			

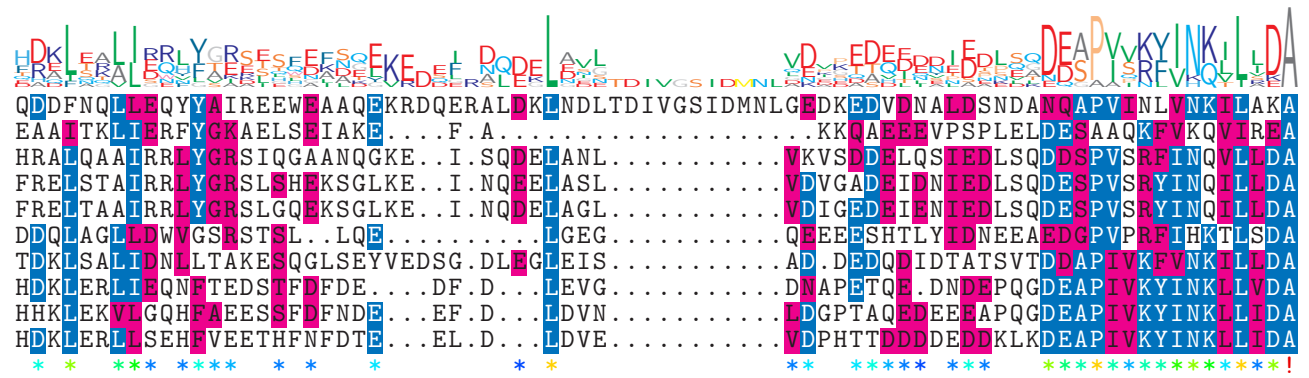
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NTDB id 1251	GCO85 RS07725 WP 011946523.1	...MALATEEYRLQGTGLLVLEKLLDKTKAIELHKLA AA EKI SLLQYIVK..NKILSAEQIALT.....AAQNFQVP	68
NTDB id 422116	G0027 RS06465 WP 016659655.1	...MSVQQAMPRTFTGFIRRLVEEGLVTAENMQQAIVAAKKAETDIVPYLIQ..QLQISPLTIAEK.....ISYEFGEF	68
NTDB id 1058	ABD1 RS01615 WP 001274986.1	...MSALHTSPKFTGFFRRLVEEKHVSAATMQTALDAAKRAKQDTVAYLIE..EVHLSPSLLAET.....ISAEFAEP	68
NTDB id 1018	ACIAD RS01685 WP 004920473.1	...MSAFTTPPKFS GFIRRLVEEGYVNANMQQALEKAKKFKQDIVPYLID..NFSISPLTIAEI.....ISLEFGEF	68
consensus		* * * * *	

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NTDB id 1168 A1552VC RS11070 WP 000957200.1	ETDLRSYDYA.....NLCCQLGLRELI TRYDALPTAKQ...GNLILLAVSDPTLLQAEEEFRAFATG.....LQVELALAD	127
NTDB id 1293 VP RS12240 WP 005479695.1	QPELSQYEYA.....SLCQQLGLRELITRHNA LPHRT...PSTLLAVADPTNQAEEDDFRFATG.....LQVELVLAD	127
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NTDB id 1110 NGFG RS09235 WP 003689817.1	ILDLRHYPRH.....RVLMGVLTTEEQMVEFHCVPVFRR...GDKVFFAVSDPTQMPQIQKTVSAAAG.....I AVELVIVE	126
NTDB id 1251 GC085 RS07725 WP 011946523.1	MLDINCIDVG.....TIPVNLVNEKLIKRHAMVPLFSR...GTNLYLATDDPSKQASLKEIQFHTG.....LNTHAIVE	135
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	YFDLDVYDT S.....QLPKDLVDQKLILKHRI LPLIQR...GQILYVATSNPSNIEAIDAIRFN SK.....LLVEPVIVE	135
NTDB id 1018 ACIAD RS01685 WP 004920473.1	LLEDLGVEDPA.....LFLKD KIDEKL IQKYRI MPLVHR...GHVLYVATSNPSNIEAMD AIRFN SK.....LKVEPIIVE	135
consensus	* * * * *	

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NTDB id 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

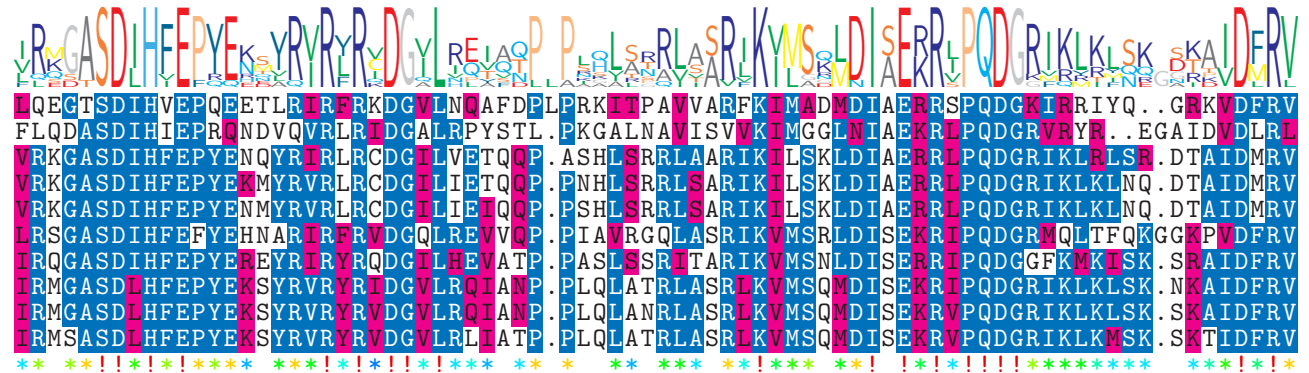
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

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consensus

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[illegible]

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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

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889	GGTGKGRVYATHELVDDEIRHAIIVAKGSATEIKETIARRKGMKTLREDGLYKALQGITTLEVLARTIE	Q95900
562	T.GGYSGRVGIYEVMAFNTELAEAIMQKASVHEIKERLAKANGMOTLQESGLEKRGITGSFAELQRVLYF	Q95900
561	T.HGYSGRGTGIYEVMRFDSELSSEALIKGASVHELEKLAIANGMSTLQMSGIEKIKQGITSFSELQRVLYF	Q95900
561	T.HGYSGRGTGIYEVMRFDSELSSEALIKGASVHELEKLAIANGMOTLQMSGIEKIKQGITSFRELQRVLYF	Q95900
558	R.GQGYKGRAGVYEVMPISEMQRVIMNGTEVGLDVAAYKEGMVDLRRAGILKIMQGITSLEVTANTND	Q95900
574	T.SGVRGRVGLFEVLPMTKEIGQLIMSGNSLDILKLAQSEGLTIFQSGIEKVKEGITITIEEVRNRTVD	Q95900
568	R.EGYKGRVGVYEVVMKVTPETISRIIMEDGNALQIAAASAKSGFNNLRRSGLIKVMQGVTSLQEVNRTSE	Q95900
570	R.EGYKGRVGIYEVVMKVTPETISKIIMEDGNALQIAAASEKLGFNINLRRSGLKKVMQGVTSLQEVNRTSE	Q95900
570	R.EGYKGRVGIYEVVMKITPETISKIIMEDGNALQIAATAETLGFNNLRRSGLKKVMQGVTSLQEVNRTSE	Q95900

logo

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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved