

logo

NTDB id 1410 SGL RS13950 WP 010873519.1	MSXLSIGDRRLGAALLDQGYLDNQLQRALERHSEVGGRLAELVLYDSGYSERRIASXIESEFGIPLVNLHSEVETSPKXK	0
NTDB id 1306 DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAL	79
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSGLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
NTDB id 1401 DSB67 RS12665 WP 010643259.1		0
NTDB id 1064652 ACHFJ9 RS20010 WP 395356092.1		0
NTDB id 1293 VP RS12240 WP 005479695.1		0
NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		0
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	SETPAEKAKELRAEPFALDEEADYRVAEYNPLDSESEYEDSGHYEPYQTTREAYMAATHYPELGLEYMPSCSK	0
NTDB id 1306 DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGA	157
NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
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NTDB id 1028 TT RS08215 WP 011173991.1	SGEGQKDKLKGELLQKGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAA	237
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consensus		

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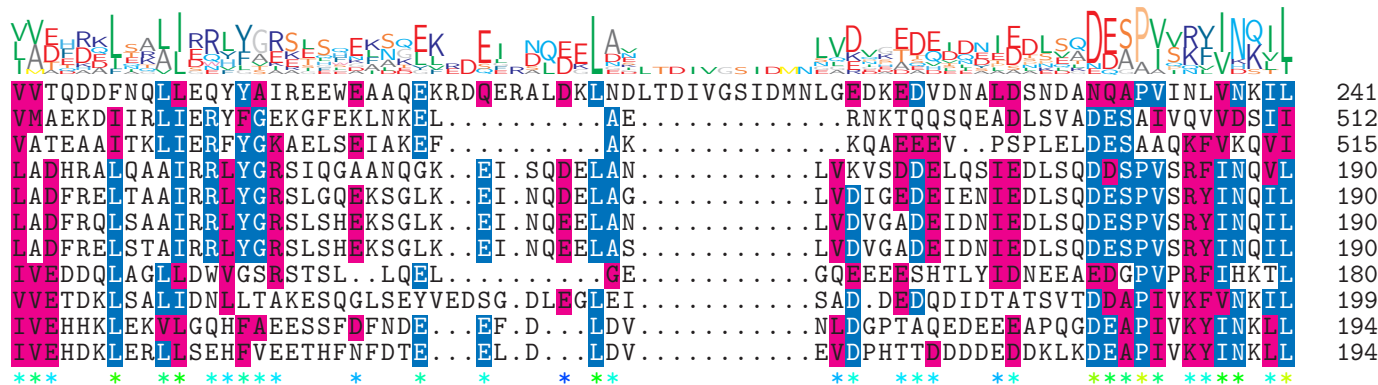
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NTDB id 1401	DSB67 RS12665 WP 010643259.1	MLTNLSTVLRQAGILTFSQEESLLEQVKASGISMPEALLS..SGFFQSHELTEH.....LSSVF	57
NTDB id 1064652	ACHFJ9 RS20010 WP 395356092.1	MQTNLSTILRQKGFLTFSQEESLLEQVKASGISMPEALLS..SGFFTSSSELAEH.....LSSIF	57
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NTDB id 1251	GC085 RS07725 WP 011946523.1	MALATEEYRLQGIQLLVLEKLLDKTKAIELHKLAAAEKISLLQYIVK..NKILSAEQIALT.....AAQNF	65
NTDB id 1058	ABD1 RS01615 WP 001274986.1	MSALHTSPKFTIGFFRRLVVEEKHVSAAATMQTALDAAKRAKQDVTAYLIE..EVHLSPSLLAET.....ISAEF	65
NTDB id 1018	ACIAD RS01685 WP 004920473.1	MSAFTTPPKFISGFIRRLVEEGYVNAQNMQQALEKAKKFKQDIPVYLID..NFSISPLTIAEI.....ISLEF	65
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NTDB id 1410	SGL RS13950 WP 010873519.1	GVDSVDPEVAPIATQQMAELIGRFFPLDTCRRFKFLPLAQEGDPPSVLVAMVDPDNLAQAQDELNRILRVKGF..ELRRL	161
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NTDB id 1028	TT RS08215 WP 011173991.1	GYPYVDPEEDPPD....PGAPLILPEDLCRRYGVFPHRLE...GNRLVLMLMKDPRNILALDDVRLALKRKGLNYEVAPA	460
NTDB id 1168	A1552VC RS11070 WP 000957200.1	GLPETDLSTRYDYA....NLCCQLGLRELITRYDALPIAKQ...GNLLLLAVSDPTLLQAEFEFRFATG.....LQVELA	124
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NTDB id 1064652	ACHFJ9 RS20010 WP 395356092.1	GLSQPELSQYEYA....SLCQQLGLRELITRHNAALPLQRT...PSTLLAVADPTNQQAEEDDFRFATG.....LQIELV	124
NTDB id 1293	VP RS12240 WP 005479695.1	GLSQPELSQYEYA....SLCQQLGLRELITRHNAALPLHRT...PSTLLAVADPTNQQAEEDDFRFATG.....LQVELV	124
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NTDB id 1251	GC085 RS07725 WP 011946523.1	GVPMLDINCIDVG....TIPVNLVNEKLIKRRHAMVPLFSR...GTNLYLATDDPSKQASLKEIQFHTG.....LNTHAI	132
NTDB id 1058	ABD1 RS01615 WP 001274986.1	AEPYFDLDVYDTS....QIPKDLVDQKLILKHRILPLIQR...GQILYVATSNPSNIEAIDAIRFNSK.....LLVEPV	132
NTDB id 1018	ACIAD RS01685 WP 004920473.1	GEPLDLGVFDPA....LFLKDKLIDEKLIQKYRIMPLVHR...GHVLYVATSNPTNIEAMDAIRFNSK.....LKVEPI	132
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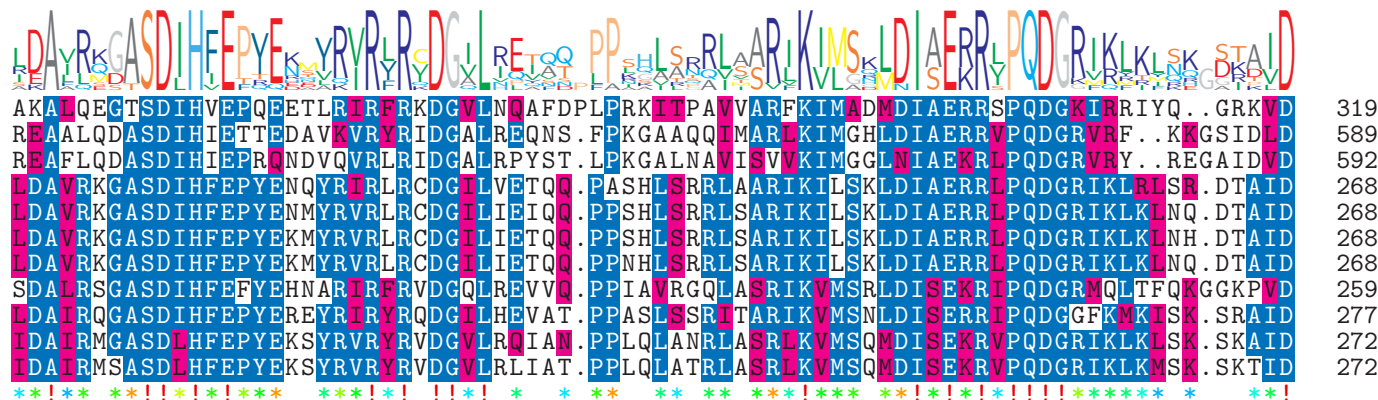
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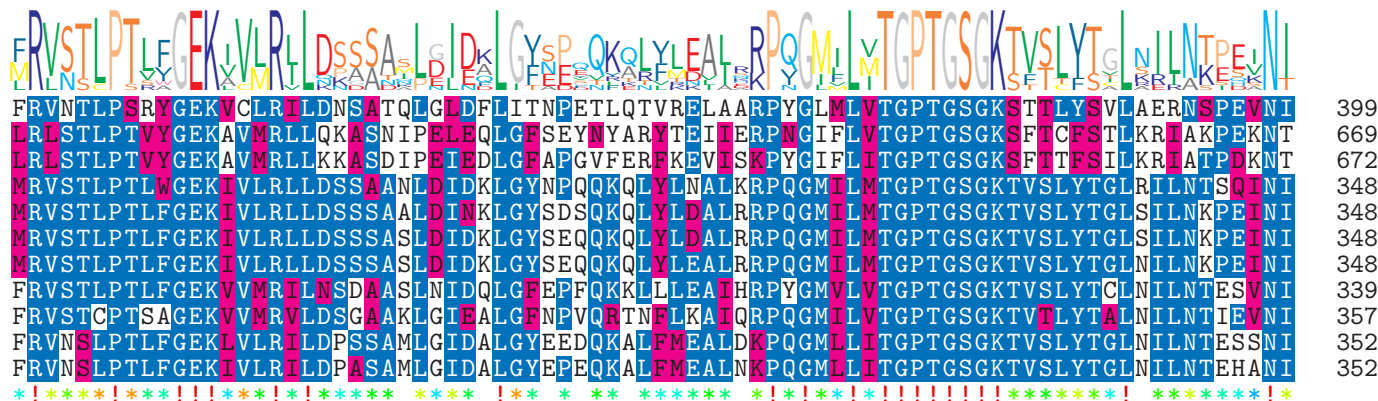
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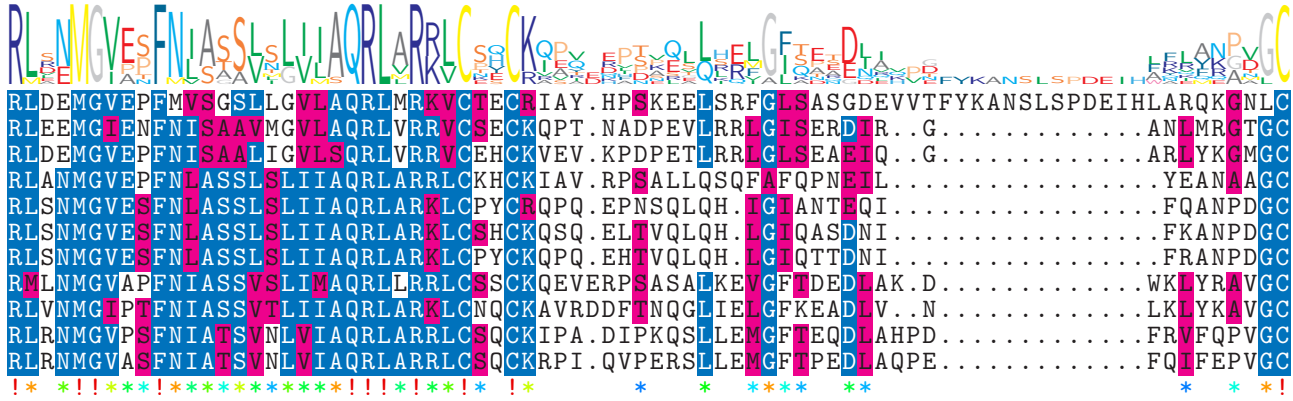
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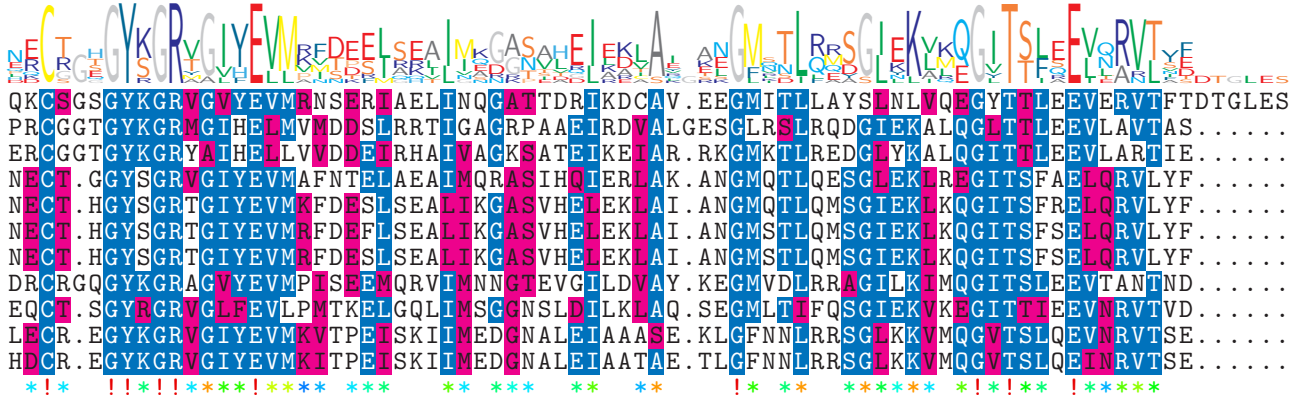
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consensus



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consensus



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NTDB id 1110 NGFG RS09235 WP 003689817.1		558
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		570
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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved