

logo

NTDB id 1410 SGL RS13950 WP 010873519.1	MSXLTIGDRRLGAALLDAGLLDQGYLDSQLRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	0
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
NTDB id 1306 DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAL	79
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NTDB id 1355 HI 0298 AAC21962.1		0
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NTDB id 1468 ASU2 RS05090 WP 014991734.1		0
NTDB id 73535 AU33 RS00730 WP 000650595.1		0
NTDB id 1436 BW25113 RS00530 WP 001025145.1		0
NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
NTDB id 1293 VP RS12240 WP 005479695.1		0
NTDB id 1401 DSB67 RS12665 WP 010643259.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	RYTFKAKELRAFPFALDEEAGVVRVAFNLPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLEIVVPSGA	0
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NTDB id 1306 DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPFALE..GDRRLVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGA	157
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consensus		

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consensus			

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NTDB id 1306	DR RS10060 WP 027479822.1	GSMRLRADALRLLAVPVDETEQGVTVLTS DPRKRDPIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGR LGEQLVQEGTLSR	317	
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NTDB id 1468	ASU2 RS05090 WP 014991734.1		0	
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NTDB id 1436	BW25113 RS00530 WP 001025145.1		0	
NTDB id 1168	A1552VC RS11070 WP 000957200.1		0	
NTDB id 1293	VP RS12240 WP 005479695.1		0	
NTDB id 1401	DSB67 RS12665 WP 010643259.1		0	
NTDB id 1251	GCO85 RS07725 WP 011946523.1		0	
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consensus				

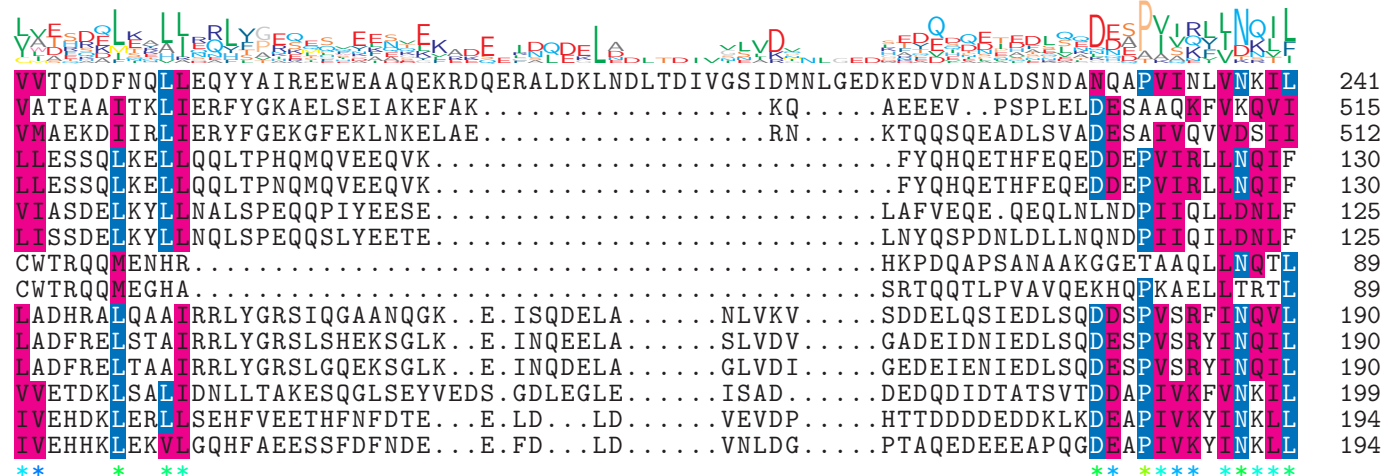
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NTDB	id	1410	SGL RS13950 WP 010873519.1	GVDSDVDEPVAPIATQQMAELIGRFFPLDTCRRF	KFLPLAQQEGDPSPVLVAMVDPDNLAQDELNRILRVKGF..ELRRL	161
NTDB	id	1028	TT RS08215 WP 011173991.1	GYPYVDPEEDPPD....PGAPLLLPEDLCRRYGVFPHRLEG...NRLVLMLMKDPRNILALDDVRRLAKRKGLNYEVAPA		460
NTDB	id	1306	DR RS10060 WP 027479822.1	GYEFLDPIQNPPD....PKVALMIPEATARRVVVPVRLQG...NSLVVAMKDPRNVFALDDLKLITG.....KEILPA		455
NTDB	id	1366	NTHI RS01950 WP 011271972.1	TISPD.....LWERNQQQSLLRLRYFALPLKEEN...NRLWLGVDSLSNLACETIAFITG.....KPVEPI		79
NTDB	id	1355	HI 0298 AAC21962.1	TISPD.....LWERNQQQSLLRLRYFALPLKEEN...NRLWLGVDSLSNLACETIAFITG.....KPVEPI		79
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NTDB	id	73535	AU33 RS00730 WP 000650595.1	MKDAQLNTLCQRHQAVLIN SAS...NSTTVAVVDAPSHALLDAIHFAFTQ.....KQIDIIV		52
NTDB	id	1436	BW25113 RS00530 WP 001025145.1	MNIPQLTALCLRYHGVLLDASE...EVVHVAVVDAPSHHELLDALHFATT.....KRTEIT		52
NTDB	id	1168	A1552VC RS11070 WP 000957200.1	CLPETDLSTRYDYA....NLCQQQLGLRELITRYDALPTIAKQG...NLLLLAVSDPTTLQAEEFRFATG....LQVELA		124
NTDB	id	1293	VP RS12240 WP 005479695.1	GLSQPELSQYEYA....SLCQQLGLRELITRHNALPLHRTTP...STLLLAVADPTNQQAEDDFRFATG....LQVELV		124
NTDB	id	1401	DSB67 RS12665 WP 010643259.1	GLSCTSLNQYEYA....SLCQTGLGRDLITRHNALPLNRTS...STLLLAVADPTNLQAEDDFRFATG....LQVELV		124
NTDB	id	1251	GC085 RS07725 WP 011946523.1	GVPMLDINCIDVG....TIPVNLVNEKLIKRRHAMVPLFSRG...TNLYLATDPSKQASLKEIQFHTG....LNTHAI		132
NTDB	id	1018	ACIAD RS01685 WP 004920473.1	GEPLDLLGVFDPA....LFLKDKIDEKLIQKKYRIMPLVHRG...HVLVYVATSNPNTTEAMDAIRFN SK....LKVEPI		132
NTDB	id	1058	ABD1 RS01615 WP 001274986.1	AEPYFDLDVYDTS....QIPKDLVDQKILLKHLRILPLIQRG...QILYVATSNPNTTEAMDAIRFN SK....LLVEPV		132
consensus				*	* * * *	* * * *

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NTDB id 1410 SGL RS13950 WP 010873519.1
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
consensus

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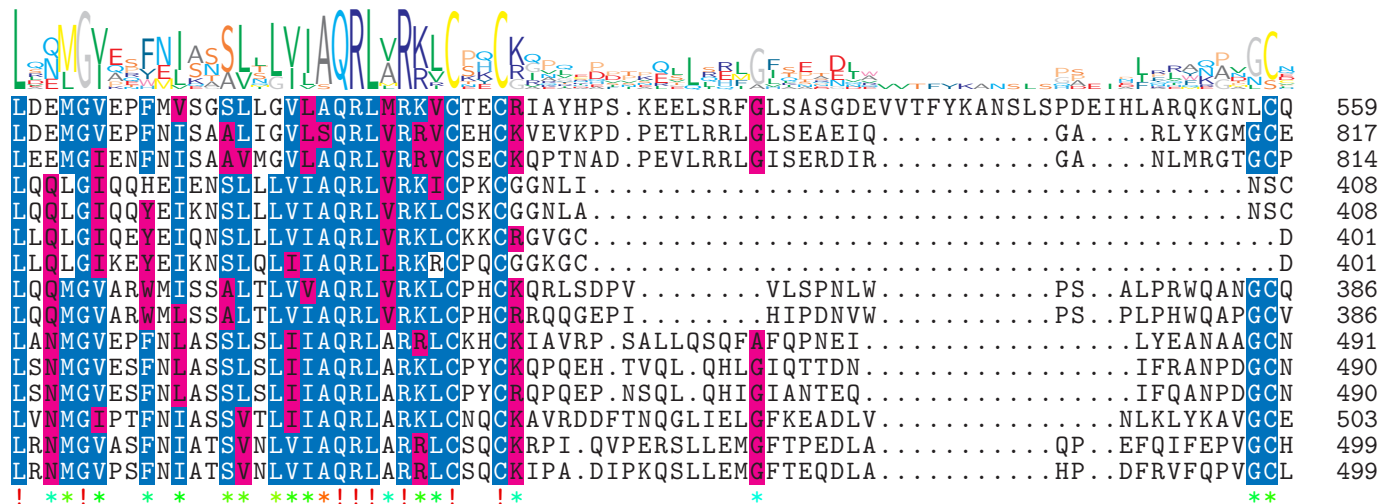
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consensus

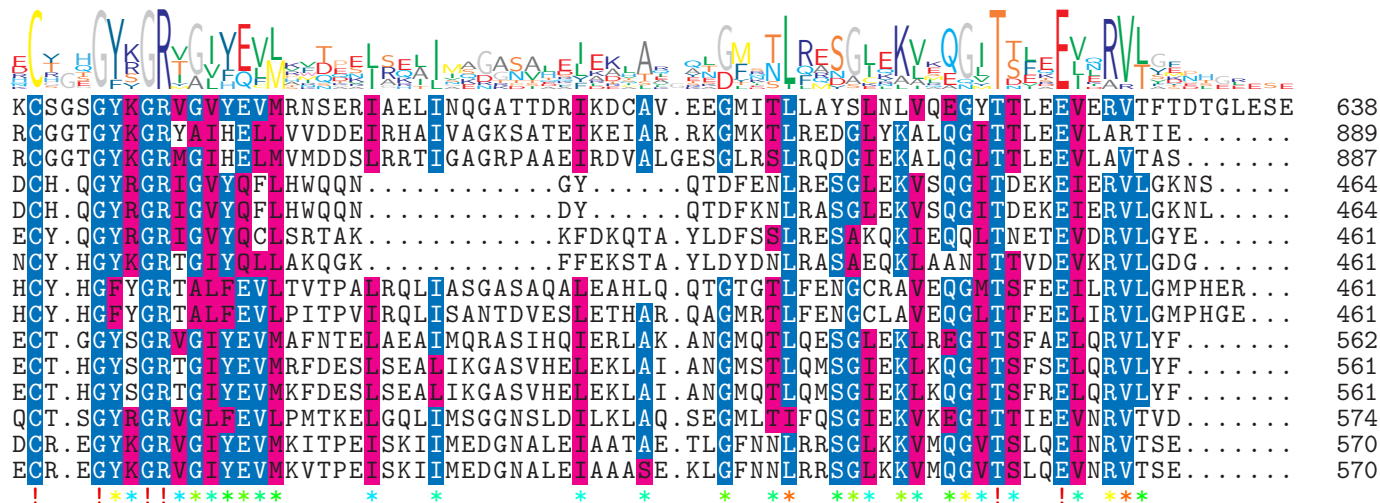
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
consensus

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NTDB id 1410 SGL RS13950 WP 010873519.1	MR A KRKSALTCRTCAAEL Q QEWLDCPYCMTSRFS	672
NTDB id 1028 TT RS08215 WP 011173991.1		889
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NTDB id 1355 HI 0298 AAC21962.1		464
NTDB id 1384 A4U84 RS08525 WP 010787006.1		461
NTDB id 1468 ASU2 RS05090 WP 014991734.1		461
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consensus		

- X non conserved
- X similar
- X ≥ 50% conserved