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NTDB id 1410 SGL RS13950 WP 010873519.1	MALSIIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	0
NTDB id 1306 DR RS10060 WP 027479822.1	MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	80
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NTDB id 1468 ASU2 RS05090 WP 014991734.1		0
NTDB id 50949 K5616 RS18905 WP 033567069.1		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
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	SIRPQTALNLQAFPPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	0
NTDB id 1306 DR RS10060 WP 027479822.1	SIRPQTALNLQAFPPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	160
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consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	GRTGGKLGGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFFHPGEEVLGSM	0
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consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	LRADALRLLAVPVDETEQGVTVLTSDPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGGEQLVQEGT	6
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consensus		

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NTDB id 1355 HI 0298 AAC21962.1	MTSYA.....LLHTQRVTAQ.....NGEVF...	20
NTDB id 1384 A4U84 RS08525 WP 010787006.1	DISP.....MSQYT.....ICEQQ.....TERIF...	15
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NTDB id 1401 DSB67 RS12665 WP 010643259.1	MLTNLSTVLRQAGILTFSQEESLLEQVKASGISMPEALLS..SGFFQSHELTEH.....LSSVFGLS	60
NTDB id 1251 GC085 RS07725 WP 011946523.1	...MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEKISILLQYIVK..NKILSAEQIALT.....AAQNFGVP	68
NTDB id 1018 ACIAD RS01685 WP 004920473.1	...MSAFTTPPKFSGFIRRLVEEGYVNAQNMQALEKAKKFKQDIVPYLID..NFSISPLTIAEI.....ISLEFGEP	68
NTDB id 1058 ABD1 RS01615 WP 001274986.1	...MSALHTSPKFTGFFRRLVEEKHVSAATMQTALDAAKRAKQDTVAYLIE..EVHLSPSLLAET.....ISAEFAEP	68
consensus	* * * *	

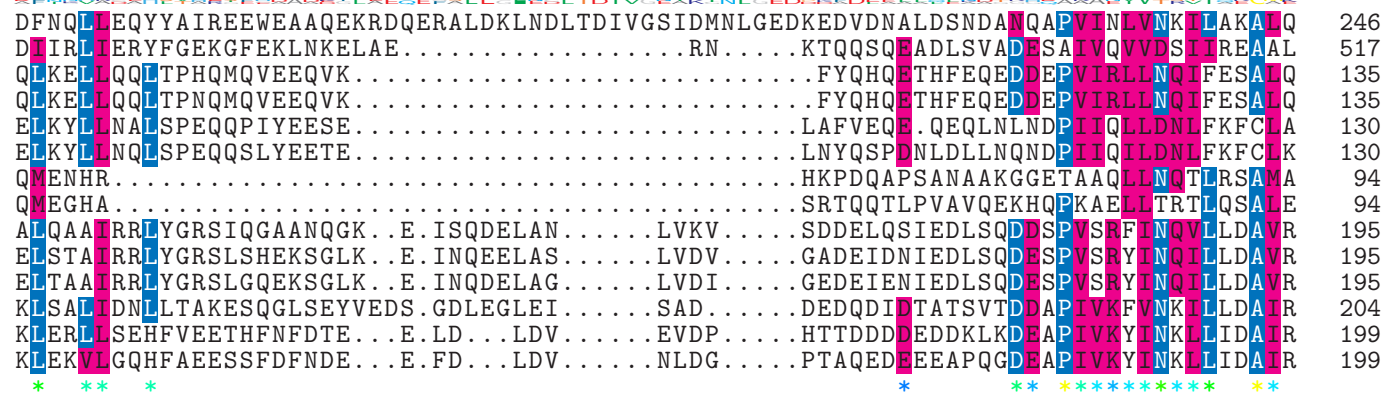
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NTDB id 50949 K5616 RS18905 WP 033567069.1	MKDAQNLNTLCQRHQAVLINSAS...NSITVAVVDAPSHALLDALHFA...TQKQIDIVCWTRQ	57
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NTDB id 1251 GC085 RS07725 WP 011946523.1	MLDINCIDV....GTIPVNLVNEKLIKRRHAMVPLFSRG...TNLYLATDPSKQASLKEITQFH...TGLNTHAIVVETD	137
NTDB id 1018 ACIAD RS01685 WP 004920473.1	LLDLGVFDP....ALFLKDKIDEKLIQKYRIMPLVHRG...HVLVYVATSNPTNIEAMDAIRFN...SKLKVPIIVEHD	137
NTDB id 1058 ABD1 RS01615 WP 001274986.1	YFDLDVYDT....SQIPKDLVDQKLILKHRILPLIQRG...QILYVATSNPSNIEAIDAIRFN...SKLLVEPVIVEHH	137
consensus	* * * * * * * * * * * * * * * *	

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consensus

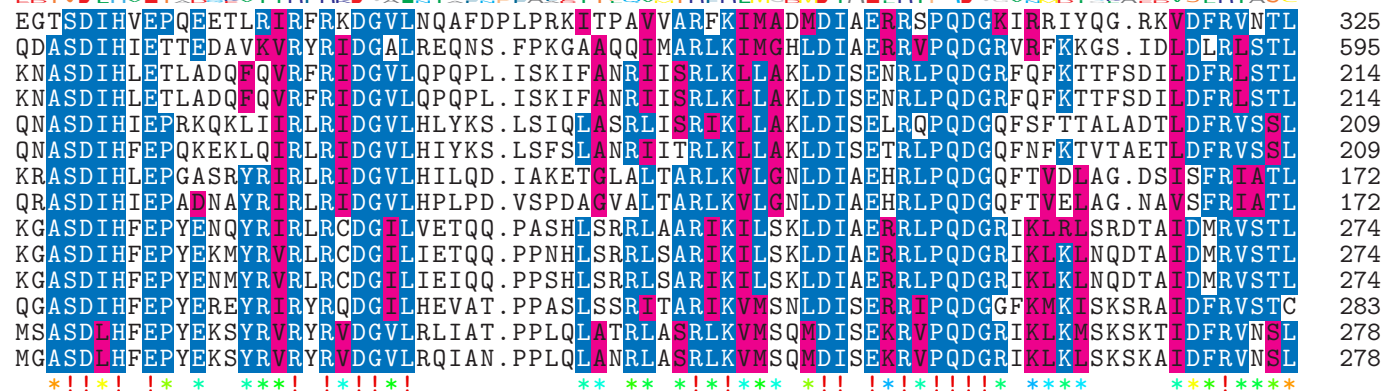
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consensus

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consensus

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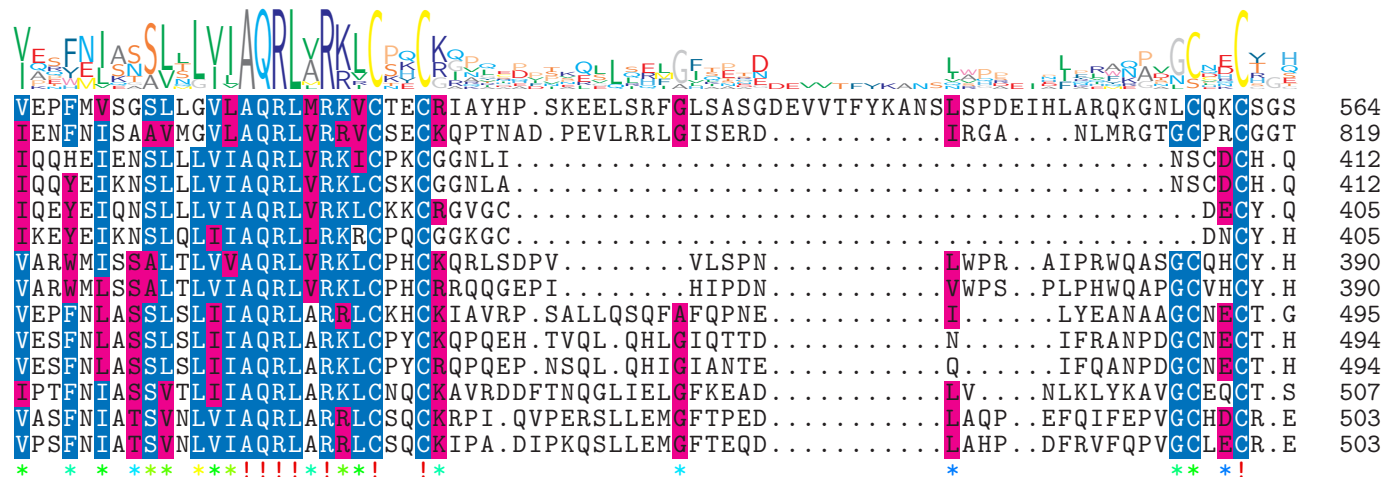
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 consensus



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 consensus



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 consensus



logo

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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved