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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	hEhIhEhAKhLRAhPFALhEhEhRYVAhYhPLhEhStEhEDhGhYhEPYQhRERhYhAhAhHYPELGLEhYhPsh	0
NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
NTDB id 1306 DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGA	157
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consensus		

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NTDB id 1306	DR RS10060 WP 027479822.1	SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	237
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consensus			

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NTDB id 1306	DR RS10060 WP 027479822.1	GSMRLADALRLLAVPVDETEQGVTVLTSDPRKRPDIDALIGRPVQLMLLTRPRDIERLIEQFYPPQRGRLGEQLVQEGTLSR	317
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NTDB id 1468	ASU2 RS05090 WP 014991734.1		0
NTDB id 78474	IY65 RS00745 WP 000650595.1		0
NTDB id 1436	BW25113 RS00530 WP 001025145.1		0
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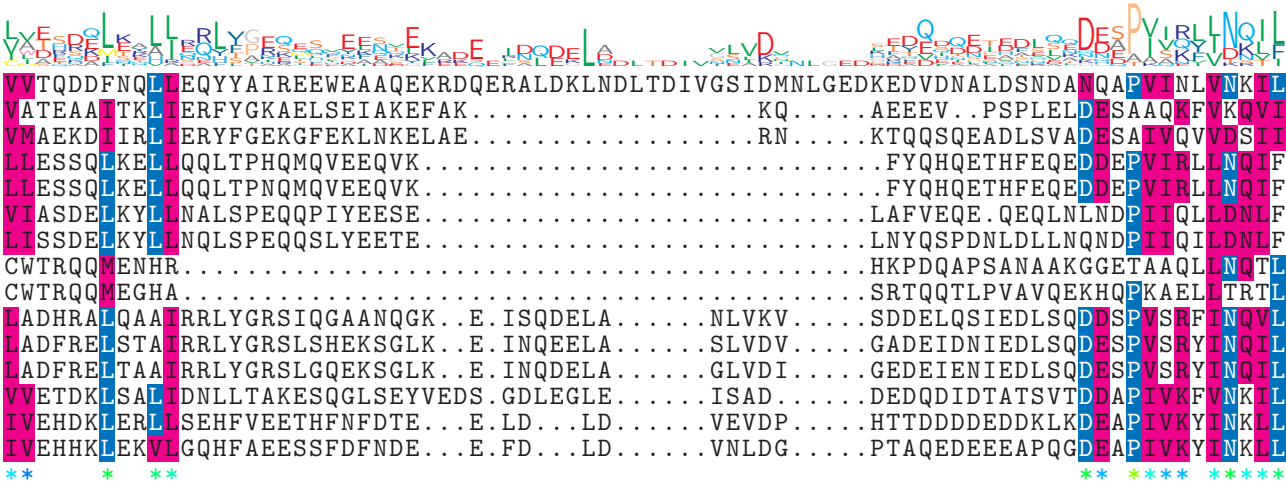
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NTDB	id	1436	BW25113	RS00530	WP	001025145.1											0						
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NTDB	id	1293	VP	RS12240	WP	005479695.1			MHSNLSTIL	R	QKGLLTFS	QEE SLIEQVKASGISM	P	EAALL	..SSGFFTSS	E	LAEH		LSSI	F	57		
NTDB	id	1401	DSB67	RS12665	WP	010643259.1			MLTNLSTVL	R	QAGILTFS	QEE SLLEQVKASGISM	P	EAALL	..SSGFFQSHE	L	TEH		LSSV	F	57		
NTDB	id	1251	GC085	RS07725	WP	011946523.1			MALATEEYRL	Q	GIGQLLV	EKLLDKTKAIELHKLAAA	E	KISLLQYIV	..KNKILSAEQ	I	ALT		AAQN	F	65		
NTDB	id	1018	ACIAD	RS01685	WP	004920473.1			MSAFTTPPKF	S	GFIRRLV	E EGYVNAQNM	Q	QALEKAKKFKQD	I	VPYLI	..DNFSISPLT	I	AEI		ISLE	F	65
NTDB	id	1058	ABD1	RS01615	WP	001274986.1			MSALHTSPKFT	G	FFRRILV	E EKHVSAATM	Q	TALDAAKRAKQD	T	VAYLI	..EEVHLSPSL	I	AEI		ISAE	F	65
consensus								*				*		*		*		*					

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NTDB	id	1028	TT RS08215 WP 011173991.1	GYPYVDPEEDPPD....PGAPLLLPEDLCRRYGVFPHRLEG...NRLVLMLMKDPRNILALDDVRLALKRKGLNYEVAPA			460
NTDB	id	1306	DR RS10060 WP 027479822.1	CYEFLDPIQMPPD....PKVALMIPEATARRVVVPVRLQG...NSLTVAMKDPRNVFALDDLKLITG.....KEILPA			455
NTDB	id	1366	NTHI RS01950 WP 011271972.1	TISPD....LWERNQQQS	LRLRYFALPLKEEN...NRLWLGVDSLSNL	SACETIAFITG....KPVEPI	79
NTDB	id	1355	HI 0298 AAC21962.1	TISPD....LWERNQQQS	LRLRYFALPLKEEN...NRLWLGVDSLSNL	SACETIAFITG....KPVEPI	79
NTDB	id	1384	A4U84 RS08525 WP 010787006.1	DISPS....RWQKNCEEKQL	LRLRYLAVPVQETE...HKLWLAVDDENNLTACEIFAFMTH....KQIEPV		74
NTDB	id	1468	ASU2 RS05090 WP 014991734.1	DISES....LWQKNCDEKH	LRLRYLAIPIQEND...DTLWLAIIDDMNLSACEIFAFIAH....KTIEPV		73
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NTDB	id	1168	A1552VC RS11070 WP 000957200.1	CLPETDLTRYDYA....NLCQQQLGLREL	LTRYDALPTAKQG...NLLLLAVSDPTTLQAEEFRFATG....LQVELV		124
NTDB	id	1293	VP RS12240 WP 005479695.1	CLSQPESLSQEYA....SLCQQLGLREL	LTRHNALPLHRTTP...STLLLAVADPTNQQAEDDFRFATG....LQVELV		124
NTDB	id	1401	DSB67 RS12665 WP 010643259.1	GLSCTSLNQYEYA....SLCQTGLGRDL	LTRHNALPLNRTS...STLLLAVADPTNLQAEDDFRFATG....LQVELV		124
NTDB	id	1251	GC085 RS07725 WP 011946523.1	GVPMLDINCIDVG....TIPVNLVNEKL	LKRHAMVPLFSRG...TNLYLATDPSKQASLKEIQFHTG....LNTHAI		132
NTDB	id	1018	ACIAD RS01685 WP 004920473.1	GEPLLDLGVFDPA....LFLKDKIDEKL	LQKYRIMPLVHRG...HVLVYVATSNPNTTEAMDAIRFNSK....LKVEPI		132
NTDB	id	1058	ABD1 RS01615 WP 001274986.1	AEPYFDLDVYDTS....QIPKDLVDQKL	LKLKHLRILPLIQRG...QILYVATSNPNTTEAMDAIRFNSK....LLVEPV		132
consensus				*	** * *	* ** * *	** *

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consensus		



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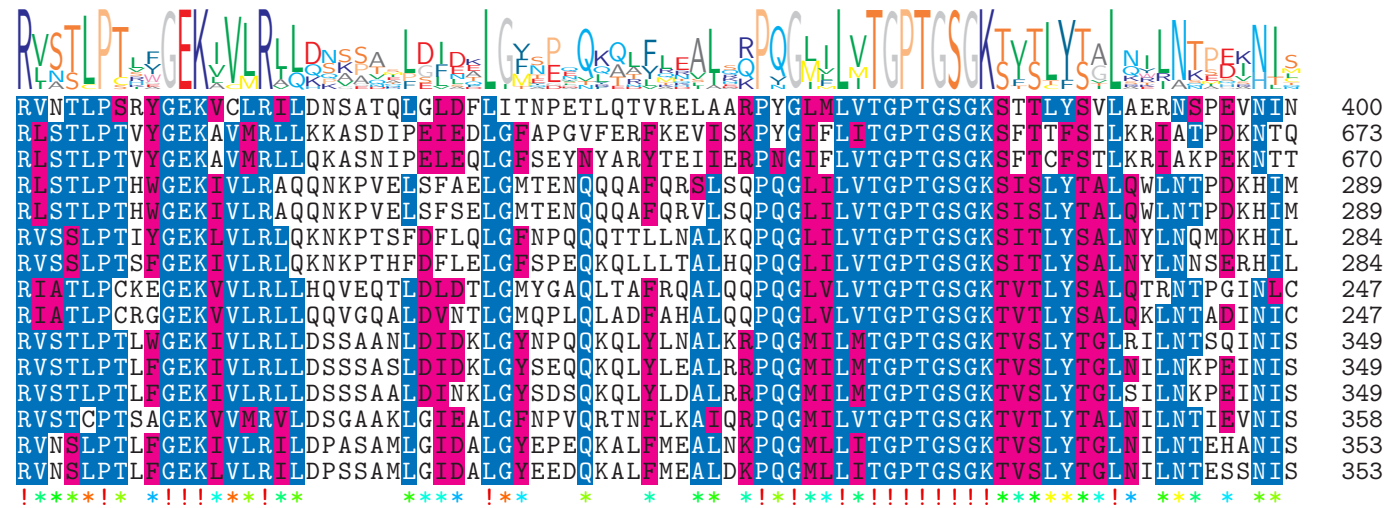


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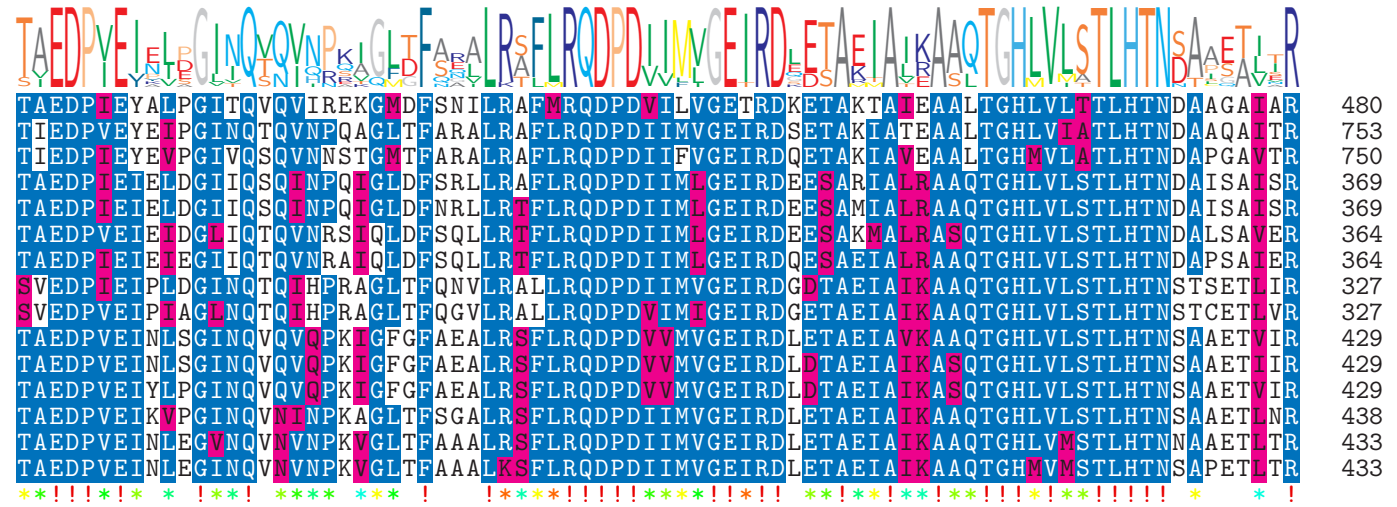
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consensus

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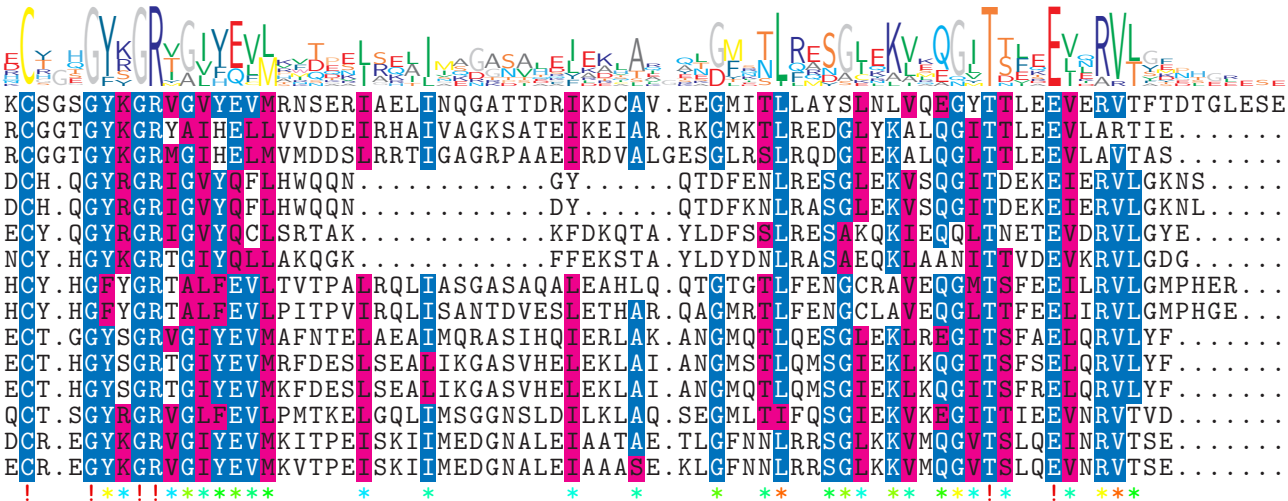
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consensus		



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consensus		



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consensus		

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