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NTDB id 1410 SGL RS13950 WP 010873519.1	MSXLTIGDRRLGAALLDGYLDLQLRALERHhEVGGhLAhVLYDhGhEhRRIAhIhEhGIPLVhLhVhP	0
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHhREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
NTDB id 1306 DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAL	79
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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 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	hEhIhEhAKhLRAhPFALhEhEhRYVAhYhPLhEhStEhEhEDhGhYhEPYQhEhYhAhAhHYPELGLEhYhP	0
NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
NTDB id 1306 DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGA	157
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consensus		

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consensus			0

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NTDB id 1468	ASU2 RS05090 WP 014991734.1		0
NTDB id 73828	AV70 RS00730 WP 000650595.1		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	SSSSKKSRVAALRNYFSPFGNKLVAGGHVDAEQLRQALVQVKKTGRSLPEELKAVTGRELSPELIRQYKKNQLFELKVL	83
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NTDB	id	1058	ABD1 RS01615 WP 001274986.1	MSALHTSPKFTGFRRRLVEEKHVSAATMQ TALDAAKR AKQDTVAYLI..EEVHLSPSL LAET.....ISAE F	65
consensus				* * *	*

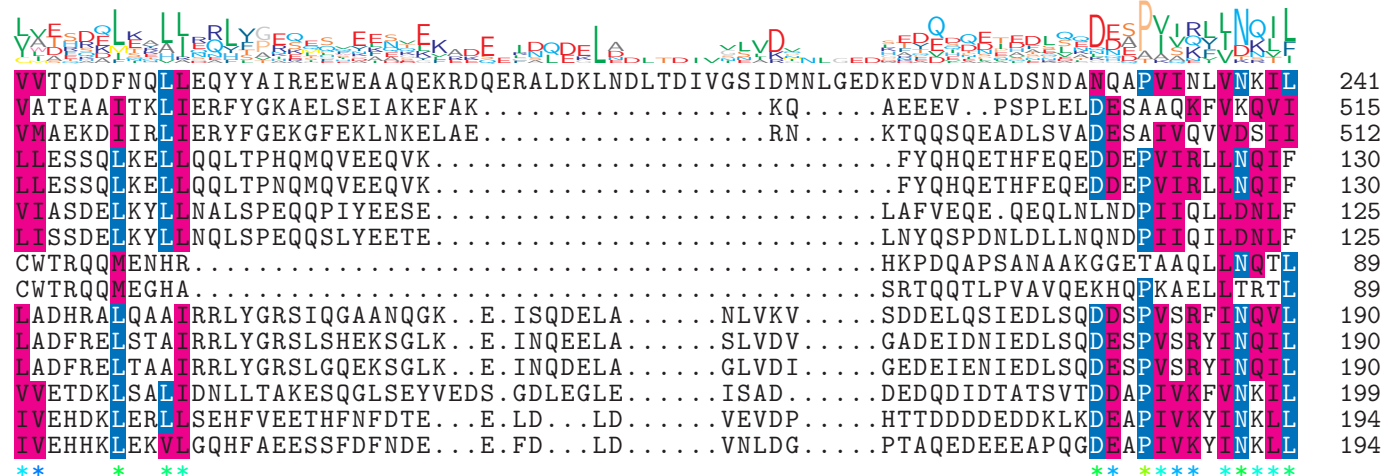
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NTDB	id	1468	ASU2 RS05090 WP 014991734.1	DISES....LWQKNCDEKHILLRLYLAIPIQEND...DTLWLAIIDDMNLSACEIFAFIAH....KTIEPV	73
NTDB	id	73828	AV70 RS00730 WP 000650595.1	MKDAQNLNTLCQRHQAVLINSAS...NSTITVAVVDAPSHALLDALHFATQ....KQIDIIV	52
NTDB	id	1436	BW25113 RS00530 WP 001025145.1	MNIPQLTALCLRYHGVLLDASE...EVVHVAVVDAPSHELLDALHFATT....KRIEIT	52
NTDB	id	1168	A1552VC RS11070 WP 000957200.1	CLPETDLSTRDYDA....NLCQQGLGRELTRYDALPTAKQG...NLLLLAVSDPTLLQAEEDFRFATG....LQVELA	124
NTDB	id	1293	VP RS12240 WP 005479695.1	GLSQPELSQYEYA....SLCQQLGLRELITRHNALPLHRTTP...STLLLAVADPTNQQAEEDFRFATG....LQVELV	124
NTDB	id	1401	DSB67 RS12665 WP 010643259.1	GLSCTSLNQYEYA....SLCQTGLGRDLITRHNALPLNRTS...STLLLAVADPTNLLQAEDDFRFATG....LQVELV	124
NTDB	id	1251	GC085 RS07725 WP 011946523.1	GVPMLDINCIDVG....TIPVNLVNEKLIKRRHAMVPLFSRG...TNLYLATDPSKQASLKEIQFHTG....LNTHAI	132
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NTDB	id	1058	ABD1 RS01615 WP 001274986.1	AEPYFDLDVYDTS....QIPKDLVDQKLLKLRHLPLIQRG...QILYVATSNPNTIEAIDAIRFNSK....LLVEPV	132
consensus				* *** ***** ***** ** *	

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consensus

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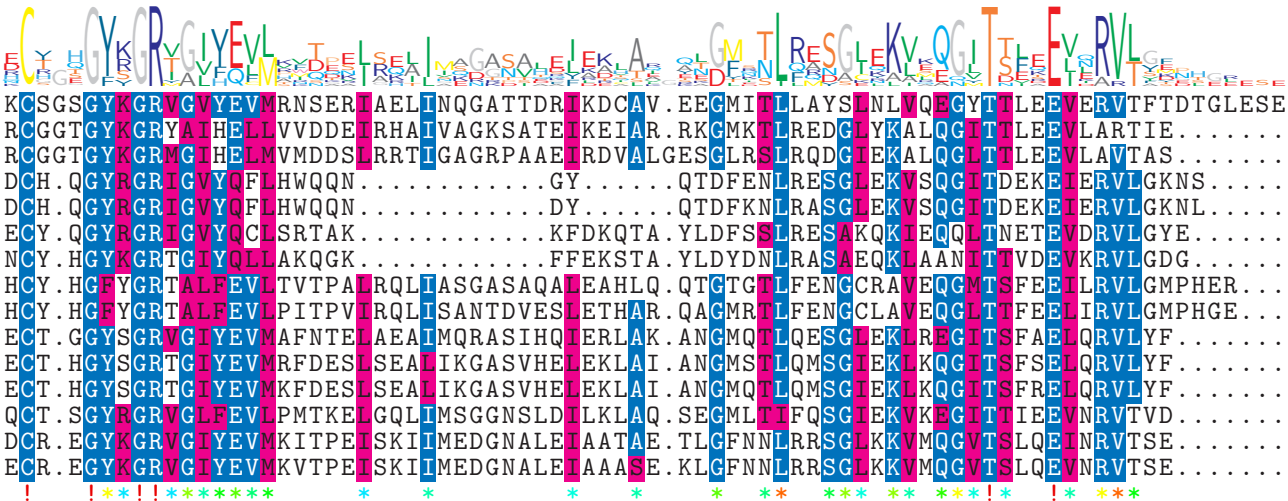
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consensus		



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consensus		



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consensus		

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