

logo

NTDB id 1410 SGL RS13950 WP 010873519.1	MSVLTITCDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	0
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
NTDB id 1293 VP RS12240 WP 005479695.1		0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

logo

NTDB id 1410 SGL RS13950 WP 010873519.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	0
NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	160
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NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

logo

NTDB id 1410 SGL RS13950 WP 010873519.1	EGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL	0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
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consensus		

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NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

NTDB id 1410	SGL RS13950 WP 010873519.1	SKKSRAVALRNYFSPFGNKL	VAGGHVDAEQ	LRQALVQVK	KTGRSLPEEL	KAVTGRELSPELLRQYKKNQLFEL	KVLYGVD	86						
NTDB id 1028	TT RS08215 WP 011173991.1	KEA	EVQKGLP	RAKPL	GEILVELGLARPEDVEEAL	QKQRGGGRLEDTLVQ	SGKLRPEALACA	VATQLGY	391					
NTDB id 1168	A1552VC RS11070 WP 000957200.1		ML	NLVAL	LRQAE	LISATQE	QAVVTQVSASGTS	PEALIE	LSIFHAQELTEQ	LSHIFGLP	60			
NTDB id 1293	VP RS12240 WP 005479695.1		MHSN	LSTLR	QKGL	LTFSGEES	LIEQVKASGISMPEAL	LS	SGFFTSS	ELAEH	LSSTFGLS	60		
NTDB id 1401	DSB67 RS12665 WP 010643259.1		ML	NLSTVLR	QAQGL	LTFSGEES	LIEQVKASGISMPEAL	LS	SGFFTSS	ELAEH	LSSTFGLS	60		
NTDB id 1110	NGFG RS09235 WP 003689817.1		MSVGL	LRLLV	QNNQV	TV	EQAEHYY	NESQACKEV	IPMLFS	DGV	SPKSLAAL	IARVFSYS	59	
NTDB id 1251	GCO85 RS07725 WP 011946523.1	...	MALATEEY	RLQ	IGQL	LVLEK	LIDKTKA	IELHKLAAAEKI	SL	LQYIVK	NKILSAEQ	IALT	AAQNFGVP	68
NTDB id 110493	ACAK33 RS02705 WP 001274987.1	...	MSALHTSP	KFT	GFFRR	LVEEKHV	SAATMQT	ALDAAKRAKQDT	VAYLIE	EVHL	SPSLLAET	SAEFAEP	68	
NTDB id 1058	ABD1 RS01615 WP 001274986.1	...	MSALHTSP	KFT	GFFRR	LVEEKHV	SAATMQT	ALDAAKRAKQDT	VAYLIE	EVHL	SPSLLAET	SAEFAEP	68	
NTDB id 1018	ACIAD RS01685 WP 004920473.1	...	MSAFTTPP	KFS	GFI	RLV	EEGYVNAQNM	QQALEKAK	FKQDIV	PLYLID	NFS	SPLTIAEI	SLEFGEP	68
consensus		*	*****	*!	*****	*	*****	*	*****	*	*****	*	*****	*

[illegible]

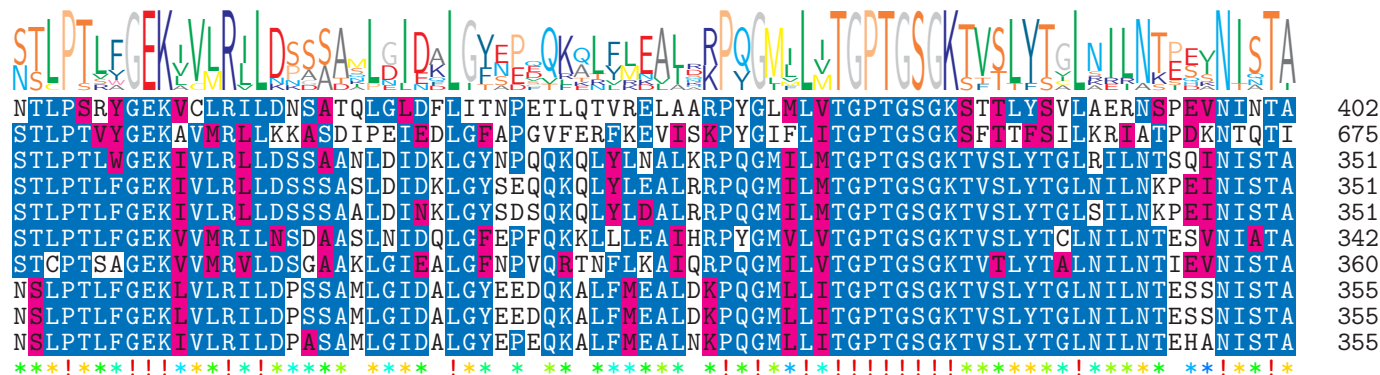
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus
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NTDB id 1410 SGL RS13950 WP 010873519.1
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

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NTDB id 1410 SGL RS13950 WP 010873519.1
NTDB id 1028 TT RS08215 WP 011173991.1
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consensus

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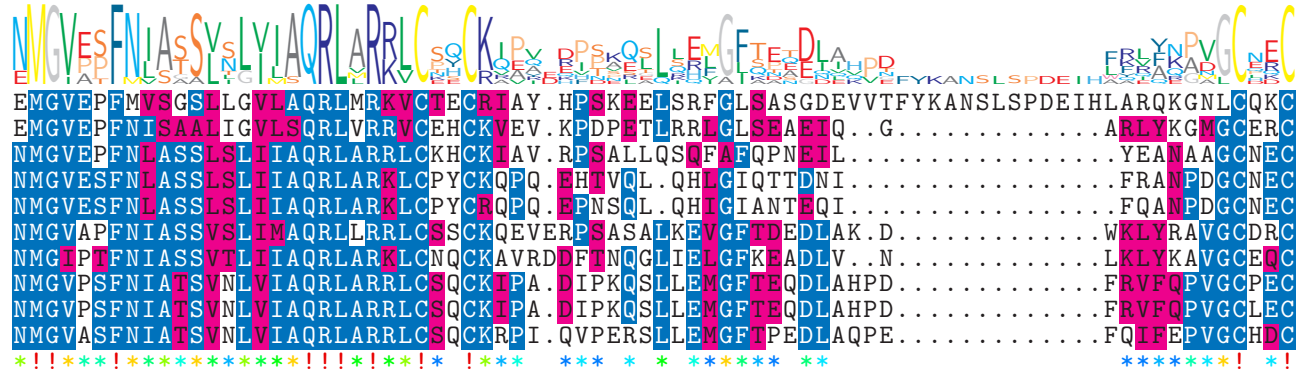
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431
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435
435

logo

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consensus

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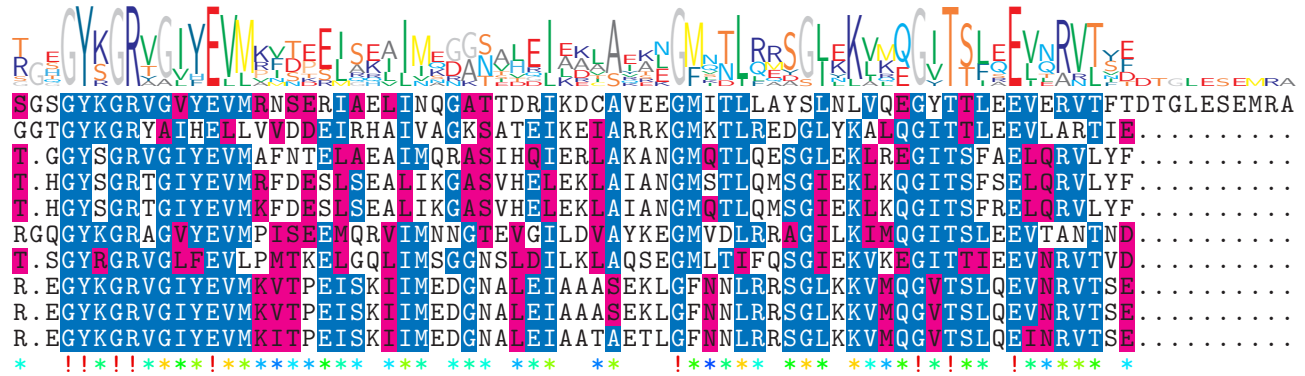
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492
488
505
501
501
501

logo

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NTDB id 1168 A1552VC RS11070 WP 000957200.1
NTDB id 1293 VP RS12240 WP 005479695.1
NTDB id 1401 DSB67 RS12665 WP 010643259.1
NTDB id 1110 NGFG RS09235 WP 003689817.1
NTDB id 1251 GCO85 RS07725 WP 011946523.1
NTDB id 110493 ACAK33 RS02705 WP 001274987.1
NTDB id 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

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641
889
562
561
561
558
574
570
570
570

logo

NTDB id 1410 SGL RS13950 WP 010873519.1	KRKSALTCRTCAAELQQEWLDCPYCMTSRFS	672
NTDB id 1028 TT RS08215 WP 011173991.1		889
NTDB id 1168 A1552VC RS11070 WP 000957200.1		562
NTDB id 1293 VP RS12240 WP 005479695.1		561
NTDB id 1401 DSB67 RS12665 WP 010643259.1		561
NTDB id 1110 NGFG RS09235 WP 003689817.1		558
NTDB id 1251 GC085 RS07725 WP 011946523.1		574
NTDB id 110493 ACAK33 RS02705 WP 001274987.1		570
NTDB id 1058 ABD1 RS01615 WP 001274986.1		570
NTDB id 1018 ACIAD RS01685 WP 004920473.1		570
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved