

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
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NTDB id 1251 GCO85 RS07725 WP 011946523.1		0
NTDB id 996484 ABDE90 RS17485 WP 001274990.1		0
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NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	0
NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	160
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consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL	0
NTDB id 1028 TT RS08215 WP 011173991.1	EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL	240
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NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

logo

NTDB id	Accession	Sequence	Score
NTDB id 1410	SGL RS13950 WP 010873519.1	LRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL	6
NTDB id 1028	TT RS08215 WP 011173991.1	LRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL	320
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NTDB id 1401	DSB67 RS12665 WP 010643259.1		0
NTDB id 1110	NGFG RS09235 WP 003689817.1		0
NTDB id 1251	GC085 RS07725 WP 011946523.1		0
NTDB id 996484	ABDE90 RS17485 WP 001274990.1		0
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	SKKSRAVALRNYFS	SPFGNKLV	AGGHVDAE	QLRQALVQ	VKKTKGR	SLPEELK	AVTGRELS	PELLRQ	YKKNQL	FELK	LV	GYD	86
NTDB id 1028 TT RS08215 WP 011173991.1	KEAL	EVQKGLP	RAKPL	GETL	VELGLAR	PEDEEAL	QKQR	GGRL	EDTL	VQ	SGKLR	PEALQA	391
NTDB id 1168 A1552VC RS11070 WP 000957200.1		MLTN	LVAILR	QAEELIS	ATQE	QAVVTQ	VSASGT	SVPEAL	LE	..	LSIFHA	QELTEQ	60
NTDB id 1293 VP RS12240 WP 005479695.1		MHSN	LSTILR	QKGLLT	FSQEES	LIEQVK	ASGISM	PEALLS	..	SGFF	TSSELA	EH	60
NTDB id 1401 DSB67 RS12665 WP 010643259.1		MLTN	LSTVLR	QAGILT	FSQEES	LIEQVK	ASGISM	PEALLS	..	SGFF	QSHELTEH	..	60
NTDB id 1110 NGFG RS09235 WP 003689817.1		MSVG	LLRLV	QNVV	TVEQA	EHYY	NESQAC	KEVL	PMLFS	..	DGVIS	PKSLAAL	59
NTDB id 1251 GC085 RS07725 WP 011946523.1	...	MALATEEY	RLQCT	IGQLLV	LEKLLD	DKTKA	IELHKL	AAAEKI	SLLQY	IVK	..	NKILSAEQ	68
NTDB id 996484 ABDE90 RS17485 WP 001274990.1	...	MSALHT	SPKFT	GFFRRL	VEEKHV	SAATM	QTALD	AAKRAK	QDTVAY	LIE	..	EVHLSP	68
NTDB id 1058 ABD1 RS01615 WP 001274986.1	...	MSALHT	SPKFT	GFFRRL	VEEKHV	SAATM	QTALD	AAKRAK	QDTVAY	LIE	..	EVHLSP	68
NTDB id 1018 ACIAD RS01685 WP 004920473.1	...	MSAFTT	PPKFS	GFFIRRL	VVEEGY	VNAQNM	QQALE	KAKK	FKQD	IVPY	LID	..	68
consensus	*	*	***	!**	*****	*	*****	*	*****	*	*****	***	***

logo

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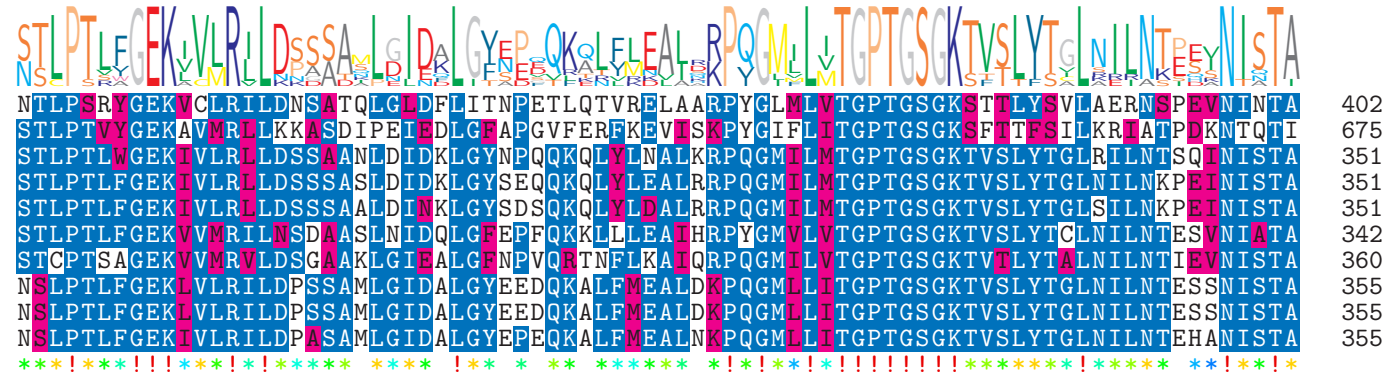
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus
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consensus

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logo

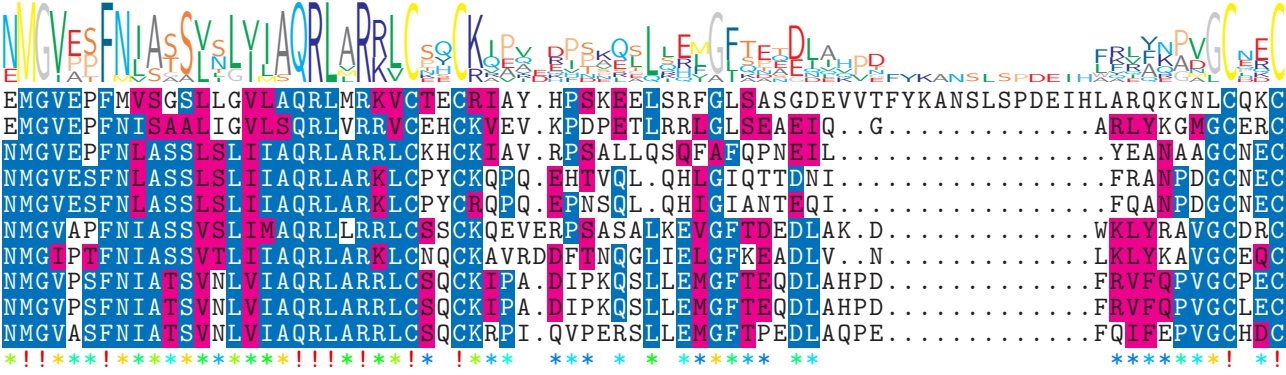
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NTDB id 996484 ABDE90 RS17485 WP 001274990.1
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus



482
755
431
431
431
422
440
435
435
435

logo

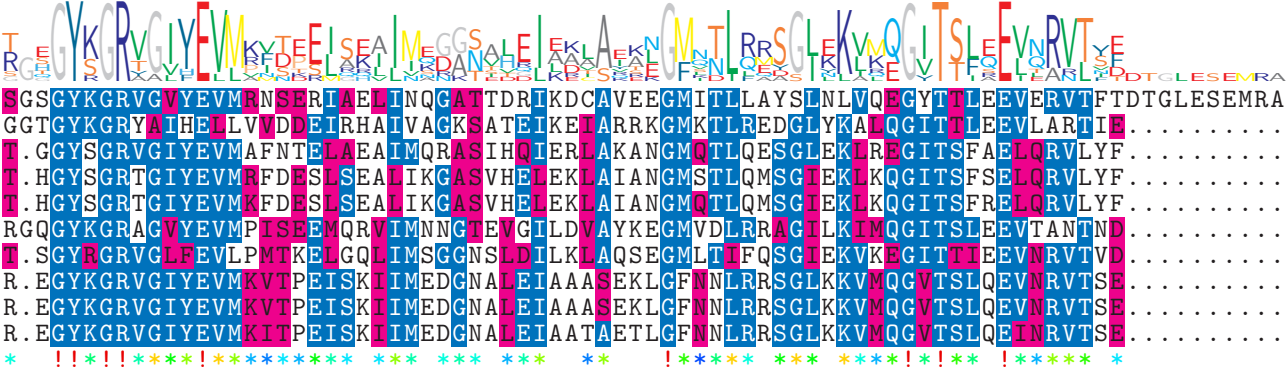
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NTDB id 996484 ABDE90 RS17485 WP 001274990.1
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus



561
819
493
492
492
488
505
501
501
501

logo

NTDB id 1410 SGL RS13950 WP 010873519.1
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NTDB id 996484 ABDE90 RS17485 WP 001274990.1
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus



641
889
562
561
561
558
574
570
570
570

logo

NTDB id 1410 SGL RS13950 WP 010873519.1	KRKSALTCTCCTCAAELQQEWLDCPYCMTSRFS	672
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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
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		570
NTDB id 1018 ACIAD RS01685 WP 004920473.1		570
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved