

logo

NTDB id 1410 SGL RS13950 WP 010873519.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	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
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 361948 FDN00 RS03270 WP 001274990.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
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
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 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 361948 FDN00 RS03270 WP 001274990.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	EGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDPSAALLL	0
NTDB id 1028 TT RS08215 WP 011173991.1	EGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDPSAALLL	240
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 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 361948 FDN00 RS03270 WP 001274990.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

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consensus

[illegible]

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[illegible]

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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

Sequence	Position
SVDPEVAPVATQQMAELIGRFFPLDTCRRFKFLPLAQEGDPPSVLVAIVDPDNAAQDEINRILRVKGF...EIRRLVVT	164
YVDPEEDPPD.....GAPILPEDLCRRYGFPHRLE...GNRLVLLMKDPNRNILADVRLALKRGLNYEVAPAVAT	463
ETDLIRYDYAN.....CQQGLGLRELITRYDALPQIAKQ...GNLLLLAVSDPTLLQAEFFRFATG.....LQVELALAD	127
QPELSQYEYAS.....LCQQGLGLRELITRHNALPLHRT...PSTLLAVADPTNQQAEDDFRFATG.....LQVELVLAD	127
CTSLNQYEYAS.....CQTLGLRLDITRHNALPLNRT...SSTLLAVADPTNQQAEDDFRFATG.....LQVELVLAD	127
ITDLRHYPRHR.....VLMGVLTEEQVEFHCVFVR...GDKVFFAVSDPTQMPQIKQTVSAAQ.....IAVELVIVE	126
MDTNCIDVGT.....IPVNLVNEKLIKRHAMVPLFSR...GTNLLYLATDDPSKQASKEIQFHTG.....LNTHAIVVE	135
YFDLDVYDTSQ.....IPKDLVDQKLLILKHRILPLIQR...GQILYVATSNPSNIEAIDAIRFNSK.....LLVEPVIVE	135
YFDLDVYDTSQ.....IPKDLVDQKLLILKHRILPLIQR...GQILYVATSNPSNIEAIDAIRFNSK.....LLVEPVIVE	135
LIDLGVFDPAL.....FLDKKIDEKLIQKYRIMPLVHR...GHVLYVATSNPTNIEAMDIAIRFNSK.....LKVEPTIVE	135

NTDB id 1410	SGL RS13950 WP 010873519.1	QDDFNQLLEQYVAIREEWAAAEKRDQERALDKLNDLTDIVGSIDMNLG	EDKEEDVDNALDSNDA	NQAPVILNVNKILLAKA	244
NTDB id 1028	TT RS08215 WP 011173991.1	EAAITTKLIERFYGKAELSIIAKEF.....	AKKQAEFEVPSPLEL	DESAAQKFVKQVIREA	518
NTDB id 1168	A1552VC RS11070 WP 000957200.1	HRAIQAAIRRLYGRSIQAANQGK..EI.SQDELANL.....	VKVSDDELQSIEDLSQ	DSPVSRFINQVLLDA	193
NTDB id 1293	VP RS12240 WP 005479695.1	FRELSTAIRRLYGRSLSHEKSGLK..EI.NQEELASL.....	VDVGADEIDNIEDLSQ	DESPVSRYINQILLDA	193
NTDB id 1401	DSB67 RS12665 WP 010643259.1	FRELTAAIRRLYGRSLGSEKSGLK..EI.NQDELAGL.....	VDIGEDEIEINEDLSQ	DESPVSRYINQILLDA	193
NTDB id 1110	NGFG RS09235 WP 003689817.1	DDQLAGLLDWVGSRSTSL.LQEL.....GEG.....	QEEESHTLYIDNEEA	EDGPVPRFIHKTLSDA	183
NTDB id 1251	GCO85 RS07725 WP 011946523.1	TDKLSALIDNLLTAKESQGLSEYVEDSG.DLEGLEIS.....	AD.DEDQDIDTATSVT	DDAPIVKFVNKILLDA	202
NTDB id 361948	FDN00 RS03270 WP 001274990.1	HHKLEKVLGGHF AEESSFDFNDE...EF.D...LDVN.....	LDGPTTQEDEE EAPQG	DEAPIVKYINKLLIDA	197
NTDB id 1058	ABD1 RS01615 WP 001274986.1	HHKLEKVLGGHF AEESSFDFNDE...EF.D...LDVN.....	LDGPTAQEDEE EAPQG	DEAPIVKYINKLLIDA	197
NTDB id 1018	ACIAD RS01685 WP 004920473.1	HDKLERLLSEHFEVEETHFNFDTE...EL.D...LDVE.....	VDPHTTDDDDEDKCLK	DEAPIVKYINKLLIDA	197
consensus		* ** **** * *	** ***** **	***** !	

[illegible]

NTDB id 1410	SGL RS13950 WP 010873519.1	N	T	L	P	S	R	Y	G	E	K	V	C	L	R	I	L	D	N	S	A	T	Q	L	G	L	D	F	L	I	T	N	P	E	T	L	Q	T	V	R	E	L	A	A	R	P	Y	G	L	M	V	T	G	P	T	G	S	G	K	S	T	T	L	Y	S	V	L	A	E	R	N	S	P	E	V	N	I	N	T	A	
NTDB id 1028	TT RS08215 WP 011173991.1	S	T	L	P	T	V	Y	G	E	K	A	V	M	R	I	L	K	K	A	S	D	I	P	E	I	E	D	L	G	F	A	P	G	V	F	E	R	F	K	E	V	I	S	K	P	Y	G	I	F	L	I	T	G	P	T	G	S	G	K	S	F	T	T	F	S	I	L	K	R	I	A	T	P	D	K	N	T	Q	T	I
NTDB id 1168	A1552VC RS11070 WP 000957200.1	S	T	L	P	T	L	W	G	E	K	I	V	L	R	L	D	S	S	A	A	N	L	D	I	D	K	L	G	L	Y	N	P	Q	Q	K	Q	L	Y	L	N	A	L	K	R	P	Q	G	M	I	L	M	T	G	P	T	G	S	G	K	T	V	S	L	Y	T	G	L	R	I	L	N	T	S	Q	I	N	I	S	T	A
NTDB id 1293	VP RS12240 WP 005479695.1	S	T	L	P	T	L	F	G	E	K	I	V	L	R	L	D	S	S	A	L	D	I	D	K	L	G	Y	S	E	Q	Q	K	Q	L	Y	L	E	A	L	R	R	P	Q	G	M	I	L	M	T	G	P	T	G	S	G	K	T	V	S	L	Y	T	G	L	N	I	L	N	K	P	E	I	N	I	S	T	A			
NTDB id 1401	DSB67 RS12665 WP 010643259.1	S	T	L	P	T	L	F	G	E	K	I	V	L	R	L	D	S	S	A	L	D	I	D	K	L	G	Y	S	D	S	E	Q	Q	K	Q	L	Y	L	D	A	L	R	R	P	Q	G	M	I	L	M	T	G	P	T	G	S	G	K	T	V	S	L	Y	T	G	L	S	I	L	N	K	P	E	I	N	I	S	T	A	
NTDB id 1110	NGFG RS09235 WP 003689817.1	S	T	L	P	T	L	F	G	E	K	V	M	R	I	L	N	S	D	A	A	S	L	N	I	D	Q	L	G	F	E	P	F	Q	K	K	L	L	E	A	T	H	R	P	Y	G	M	V	L	T	G	P	T	G	S	G	K	T	V	S	L	Y	T	C	L	N	I	L	N	T	E	S	V	N	I	A	T	A			
NTDB id 1251	GC085 RS07725 WP 011946523.1	S	T	C	P	T	S	A	G	E	K	V	M	R	V	L	D	S	G	A	A	K	L	G	I	E	A	L	G	F	N	P	V	Q	R	T	N	F	L	K	A	I	Q	R	P	Q	G	M	I	L	V	T	G	P	T	G	S	G	K	T	V	T	L	Y	T	A	L	N	I	L	N	T	I	E	V	N	I	S	T	A	
NTDB id 361948	FDM00 RS03270 WP 001274990.1	N	S	L	P	T	L	F	G	E	K	I	V	L	R	I	L	D	P	S	S	A	M	L	G	I	D	A	L	G	Y	E	E	D	Q	K	A	L	F	M	E	A	L	D	K	P	Q	G	M	L	I	T	G	P	T	G	S	G	K	T	V	S	L	Y	T	G	L	N	I	L	N	T	E	S	S	N	I	S	T	A	
NTDB id 1058	ABD1 RS01615 WP 001274986.1	N	S	L	P	T	L	F	G	E	K	I	V	L	R	I	L	D	P	S	S	A	M	L	G	I	D	A	L	G	Y	E	E	D	Q	K	A	L	F	M	E	A	L	D	K	P	Q	G	M	L	I	T	G	P	T	G	S	G	K	T	V	S	L	Y	T	G	L	N	I	L	N	T	E	S	S	N	I	S	T	A	
NTDB id 1018	ACIAD RS01685 WP 004920473.1	N	S	L	P	T	L	F	G	E	K	I																																																																					

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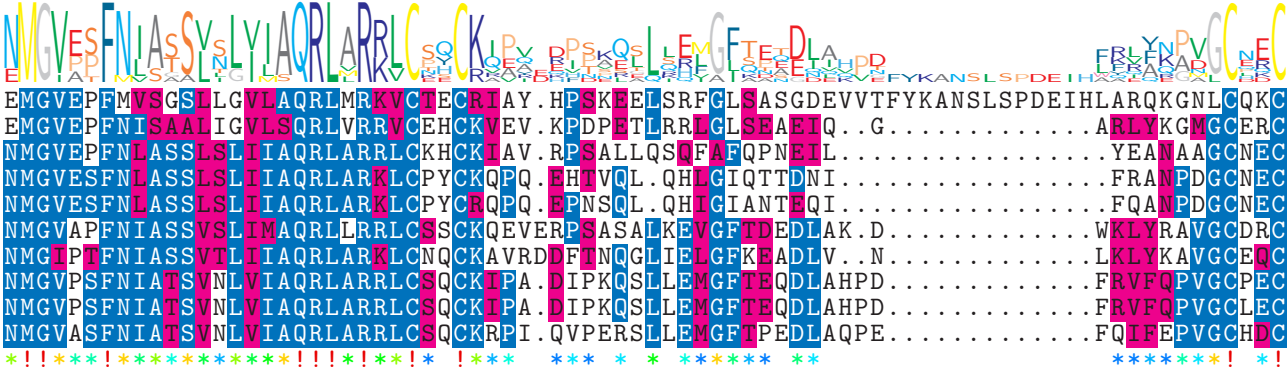
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus



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755
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logo

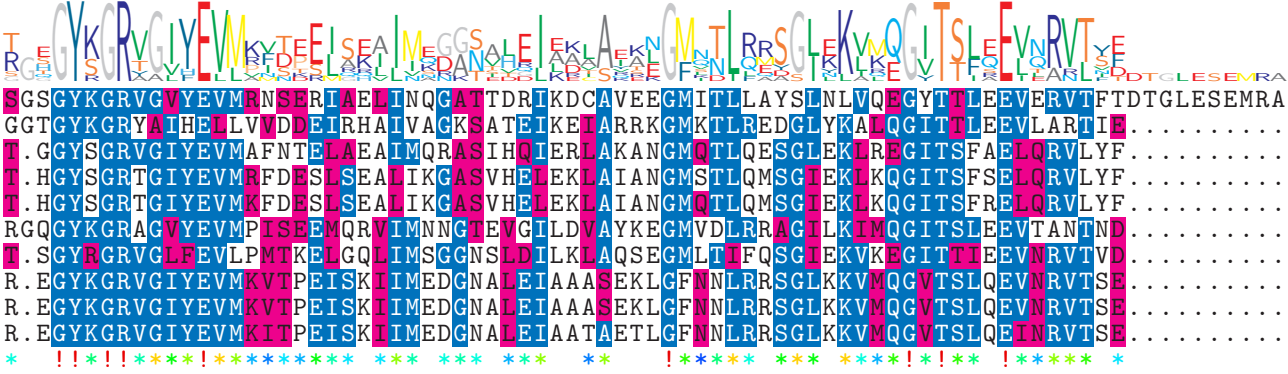
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus



561
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NTDB id 1410 SGL RS13950 WP 010873519.1
NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1168 A1552VC RS11070 WP 000957200.1
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus



641
889
562
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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 361948 FDN00 RS03270 WP 001274990.1		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