

logo

		MSXLRIGDRRLGAAALLDQGYLGDNDLQRALERHSEVGGRLAELVLYDQGYLSEERRIABXIEBEGIPLVNLAHVEITBPKK	
NTDB id 1410	SGL RS13950 WP 010873519.1	.....	0
NTDB id 1306	DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAAL	79
NTDB id 1028	TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
NTDB id 1168	A1552VC RS11070 WP 000957200.1	.....	0
NTDB id 1293	VP RS12240 WP 005479695.1	.....	0
NTDB id 1071814	ACI3L6 RS11085 WP 079400988.1	.....	0
NTDB id 1401	DSB67 RS12665 WP 010643259.1	.....	0
NTDB id 1110	NGFG RS09235 WP 003689817.1	.....	0
NTDB id 1251	GCO85 RS07725 WP 011946523.1	.....	0
NTDB id 1058	ABD1 RS01615 WP 001274986.1	.....	0
NTDB id 1018	ACIAD RS01685 WP 004920473.1	.....	0
consensus			

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		RSLRPAKALNLRALPFALDEEAGVVRVAFLNPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP	
NTDB id 1410	SGL RS13950 WP 010873519.1	.....	0
NTDB id 1306	DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLIIVVPSGA	157
NTDB id 1028	TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
NTDB id 1168	A1552VC RS11070 WP 000957200.1	.....	0
NTDB id 1293	VP RS12240 WP 005479695.1	.....	0
NTDB id 1071814	ACI3L6 RS11085 WP 079400988.1	.....	0
NTDB id 1401	DSB67 RS12665 WP 010643259.1	.....	0
NTDB id 1110	NGFG RS09235 WP 003689817.1	.....	0
NTDB id 1251	GCO85 RS07725 WP 011946523.1	.....	0
NTDB id 1058	ABD1 RS01615 WP 001274986.1	.....	0
NTDB id 1018	ACIAD RS01685 WP 004920473.1	.....	0
consensus			

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		SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	
NTDB id 1410	SGL RS13950 WP 010873519.1	.....	0
NTDB id 1306	DR RS10060 WP 027479822.1	SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	237
NTDB id 1028	TT RS08215 WP 011173991.1	SGEGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLFLESTEGIVPDPSAA	237
NTDB id 1168	A1552VC RS11070 WP 000957200.1	.....	0
NTDB id 1293	VP RS12240 WP 005479695.1	.....	0
NTDB id 1071814	ACI3L6 RS11085 WP 079400988.1	.....	0
NTDB id 1401	DSB67 RS12665 WP 010643259.1	.....	0
NTDB id 1110	NGFG RS09235 WP 003689817.1	.....	0
NTDB id 1251	GCO85 RS07725 WP 011946523.1	.....	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  
 NTDB id 1306 DR RS10060 WP 027479822.1  
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 NTDB id 1058 ABD1 RS01615 WP 001274986.1  
 NTDB id 1018 ACIAD RS01685 WP 004920473.1  
 consensus

.....MTS  
 GSMLRADALRLLAVPVDETEQGVTVLTS DPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPQGRRLGGEQLVQEGTLSR  
 LLLLRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSR  
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NTDB id 1410 SGL RS13950 WP 010873519.1  
 NTDB id 1306 DR RS10060 WP 027479822.1  
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 NTDB id 1058 ABD1 RS01615 WP 001274986.1  
 NTDB id 1018 ACIAD RS01685 WP 004920473.1  
 consensus

SSSSKKSRAVALRNYFSPFGNKL VAGGHVDAEQLRQALVQVKKTKRSIPEELKAVTGRELSPELIRQYKKNQLFELKVLV  
 DQLREALQVQAREGKVKPLGEVITELGFASPDVDSALQKQNVGGRIEDTLVQ..SGKLSPEMLARS.....LAAQL  
 EALKEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGRIEDTLVQ..SGKLRPEALAAQA.....VATQL  
 .....MLTINLVAIILRQAEILSATQEQA VVTQVSASGTSVPEALLE..LSIFHAQELTEQ.....LSHIF  
 .....MHSNLSTILRQKGLLTFSSQEEESLIEQVKASGISMPEALLS..SGFFTSSELAEH.....LSSIF  
 .....MLTINLSTVLRQAGILTFSSQEEESLIEQVKASGISMPEALLS..SGFFQSHELTEH.....LSSVF  
 .....MLTINLSTVLRQAGILTFSSQEEESLIEQVKASGISMPEALLS..SGFFQSHELTEH.....LSSVF  
 .....MSVGLLRILVQNVVITVEQA EHYH.NESQAGKEVLPMLFS..DGVITSPKSLAAL.....IARVF  
 .....MALATEEYRLQGITGLVLEKLLDKTKAIELHKLAAAEKISLLQYIVK..NKILSAEQIALT.....AAQNF  
 .....MSALHTSPKFTIGFFRRLVVEEKHVSAAATMQTALDAAKRAKQDTVAYLIE..EVHLSPSLLAET.....TSAEF  
 .....MSAFTTPPKFISGFIRRLVVEEGYVNAQNMQALEKAKKFKQDIPVYLLID..NFSISPLTIAEI.....ISLEF

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logo

NTDB id 1410 SGL RS13950 WP 010873519.1  
 NTDB id 1306 DR RS10060 WP 027479822.1  
 NTDB id 1028 TT RS08215 WP 011173991.1  
 NTDB id 1168 A1552VC RS11070 WP 000957200.1  
 NTDB id 1293 VP RS12240 WP 005479695.1  
 NTDB id 1071814 ACI3L6 RS11085 WP 079400988.1  
 NTDB id 1401 DSB67 RS12665 WP 010643259.1  
 NTDB id 1110 NGFG RS09235 WP 003689817.1  
 NTDB id 1251 GC085 RS07725 WP 011946523.1  
 NTDB id 1058 ABD1 RS01615 WP 001274986.1  
 NTDB id 1018 ACIAD RS01685 WP 004920473.1  
 consensus

GVDSVDPEVAPIATQQMAELIGRFFPLDTCRRFKFLPLAQQEGDPPSVLVAMVDPDNLAAQDELNRILRVKGF..ELRRL  
 GYFLDPIQNPDP...PKVALMIPEATARRVYVVPVRLQG...NSLVVAMKDPRNVFALDDLKLITG....KEILPA  
 GYPYVDPEEDPPD...PGAPLILPEDLCRRYGVPFHRLEG...NRLVLLMKDPRNILALDDVRLALKRKGLNYEVAPA  
 GLPETDLRSRYDYA...NLCQQLGLRELITRYDALPIAKQG...NLLLVAVSDPTLLQAEFEFRFATG.....LQVELA  
 GLSQPELSQYEYA...SLCQQLGLRELITRHNA LPHRTP...STLLAVADPTNQQAEEDFRFATG.....LQVELV  
 GLSCTSLNQY EYA...SLCQTLGLRDLITRHNA LPLNRTS...STLILAVADPTNLQAEDDFRFATG.....LQVELV  
 GLSCTSLNQY EYA...SLCQTLGLRDLITRHNA LPLNRTS...STLILAVADPTNLQAEDDFRFATG.....LQVELV  
 SYSILDLRHYP RH...RVLMGVITEEQMV EFCVVPVFRRG...DKVFFAVSDPTQMPQIQKT VSAAG.....I AVELV  
 GVPMLDINCIDVG...TIPVNLVNEKLIK RHAMVPLFSRG...TNLYLATDDPSKQASLKEIQFHTG.....LNTHAI  
 AEPYFDLDVYDTS...QIPKDLVDQKLILKHRI LPLIQRG...QILYVATSNPSNIEAIDAIRFNSK.....LLVEPV  
 GEPLLDLGVFDPA...LFLDKKIDELIKYRIMPLVHRG...HVLVATSNPTNIEAMD AIRFNSK.....LKVEPI

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NTDB id 1410 SGL RS13950 WP 010873519.1	EMRAKRKRSALTCRTCAAELQQEWLDCPYCMTSRFS	672
NTDB id 1306 DR RS10060 WP 027479822.1	.....	887
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 1071814 ACI3L6 RS11085 WP 079400988.1	.....	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 1058 ABD1 RS01615 WP 001274986.1	.....	570
NTDB id 1018 ACIAD RS01685 WP 004920473.1	.....	570
consensus		

-  non conserved
-  similar
-   $\geq 50\%$  conserved