

logo

			MSXLIIGDRRLGALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVK	
NTDB id 1028	TT	RS08215	WP 011173991.1	80
NTDB id 1306	DR	RS10060	WP 027479822.1	79
NTDB id 1110	NGFG	RS09235	WP 003689817.1	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	GCO85	RS07725	WP 011946523.1	0
NTDB id 43948	SBAL678	RS43955	WP 006084368.1	0
NTDB id 1018	ACIAD	RS01685	WP 004920473.1	0
NTDB id 1058	ABD1	RS01615	WP 001274986.1	0
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			RSTFRKELAKLRAIPFALDEEAGVVRVAFNLPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLEIVVPSGA	
NTDB id 1028	TT	RS08215	WP 011173991.1	158
NTDB id 1306	DR	RS10060	WP 027479822.1	157
NTDB id 1110	NGFG	RS09235	WP 003689817.1	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	GCO85	RS07725	WP 011946523.1	0
NTDB id 43948	SBAL678	RS43955	WP 006084368.1	0
NTDB id 1018	ACIAD	RS01685	WP 004920473.1	0
NTDB id 1058	ABD1	RS01615	WP 001274986.1	0
consensus				

logo

			SGEGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGFLESTEGIVPDPSSAA	
NTDB id 1028	TT	RS08215	WP 011173991.1	237
NTDB id 1306	DR	RS10060	WP 027479822.1	237
NTDB id 1110	NGFG	RS09235	WP 003689817.1	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	GCO85	RS07725	WP 011946523.1	0
NTDB id 43948	SBAL678	RS43955	WP 006084368.1	0
NTDB id 1018	ACIAD	RS01685	WP 004920473.1	0
NTDB id 1058	ABD1	RS01615	WP 001274986.1	0
consensus				

logo

NTDB id 1028 TT RS08215 WP 011173991.1 LLLLRS DALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSR 317

NTDB id 1306 DR RS10060 WP 027479822.1 GSMLRADALRLLAVPVDETEQGVTVLTSDPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPQRGRLGEQLVQEGTLSR 317

NTDB id 1110 NGFG RS09235 WP 003689817.1

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 1251 GCO85 RS07725 WP 011946523.1

NTDB id 43948 SBAL678 RS43955 WP 006084368.1

NTDB id 1018 ACIAD RS01685 WP 004920473.1

NTDB id 1058 ABD1 RS01615 WP 001274986.1

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NTDB id 1028 TT RS08215 WP 011173991.1 EALKEAL EVQKGLPRAKPL GEI LVELGLARPEDVEALQKQRRCGRIEDTLVQSGKLRPEALAGAVATQLGYPYVDPEE 397

NTDB id 1306 DR RS10060 WP 027479822.1 DQLREAL QVQAREGKVKPL GEVITELGFASPDVEVDSALQKQNVGCGRIEDTLVQSGKLSPEMLARSLAAQLGYEFLDPIQ 397

NTDB id 1110 NGFG RS09235 WP 003689817.1 MSVGLLRILVQNVVIVEQAHHYINES.QACKEVLPMLFSDGVI SPKSLAAL IARVFSYSILDLRH 65

NTDB id 1168 A1552VC RS11070 WP 000957200.1 MLTINLVAILRQAE LISATQEQAVVTVQVSASCTSVPEALIELSIFHAQELTEQLSHIFGLPETDLSR 66

NTDB id 1293 VP RS12240 WP 005479695.1 MHSNLSTILRQKGLTFSQEE SLIEQVMASCISMPEALLSSCGFFTSSELAEHLSSIFGLSQPELSQ 66

NTDB id 1401 DSB67 RS12665 WP 010643259.1 MLTINLSTVLRQAQILTFSQEE SLIEQVMASCISMPEALLSSCGFFQSHELTEHLSSVFCLSCTSLNQ 66

NTDB id 1251 GCO85 RS07725 WP 011946523.1 MALATEEYRLQGIGQLLVLEKLLDKTKATELHKLAAAEEKISLLQYIVKNKILSAEQIALTAANQFGVPMLDINC 74

NTDB id 43948 SBAL678 RS43955 WP 006084368.1 MPTTGLHLGLSTLFI RKGLINEEQMANA ITKSRQNKHALVSTLVQSQLV SARAI AELCYEYEGTPLL DINE 71

NTDB id 1018 ACIAD RS01685 WP 004920473.1 MSAFTTPPKFSGFIRRLVEEGYVNAQNMQQALEKAKKFKQDIVPYLDNFST SPLTIAETISLEFGEPPLD LGV 74

NTDB id 1058 ABD1 RS01615 WP 001274986.1 MSALHTSPKFTGFFRRLVEEKHVS AATMQTALDAAKRAKQDTWAYLIEEVHLSPSLLAETISAEFAEPYFDL DV 74

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NTDB id 1028 TT RS08215 WP 011173991.1 DPPDPGAPL LIPEDLCRRYGVFPHRLEGNRLVLLMKDPRNLLAADDVRLALKRKGNYEVAPAVATEAAITKLIERFYGA 477

NTDB id 1306 DR RS10060 WP 027479822.1 NPPDPKVALMIP EATARRYVWVVR LQGNLSLVAMKDPRNVFALDDLKLTG..... KEILPAVMAEKDIRLIERFYGE 472

NTDB id 1110 NGFG RS09235 WP 003689817.1 YPRHRVLMGVLT EEQMVVEFHCVPVFRGDKVFFAVSDPTQMPQLQKTVSAAG..... IAVELVIVEDDQLAGLDDWVGSR 140

NTDB id 1168 A1552VC RS11070 WP 000957200.1 YDYANLCQQLGLRELITRYDALPIAKQGNL LLLAVSDPTLLQAE EEFRFATG..... LQVELALADHRA LQAATRRLYGR 141

NTDB id 1293 VP RS12240 WP 005479695.1 YEYASLCQQLGLRELITRHNALPLHRTPTSTLLAVADPTNQQAE DDFRFATG..... LQVELVLADFRELSTAIRLYGR 141

NTDB id 1401 DSB67 RS12665 WP 010643259.1 YEYASLCQTLGLRDLITRHNALPLNRTSSSTLILAVADPTNLQAE DDFRFATG..... LQVELVLADFRELTAATRRLYGR 141

NTDB id 1251 GCO85 RS07725 WP 011946523.1 IDVGTIPVNLVNEKLIKRHAMVPLFSRGTNLYLATDDPSKQASLKEIQFHTG..... LNTHAIVETDKLSALIDNLLTA 149

NTDB id 43948 SBAL678 RS43955 WP 006084368.1 FDLASIPEDFLNKKLIEKHRCLPLFKRCNRLYICTSDPTNIAALED FQFSAG..... LHAFAIIVVEDKLALEKILEE 146

NTDB id 1018 ACIAD RS01685 WP 004920473.1 FDPALFLKDKIDEKLIQKYRIMPLVHRGHVLYVATSNPTNTEAMDARFN SK..... LKVEPIIVEHDKLERLSEHFVE 149

NTDB id 1058 ABD1 RS01615 WP 001274986.1 YDTSQIPKDLVDGKILKHLRLLPLIQRGGILYVATSNPSNTEAMDARFN SK..... LLVEPVIIVEHKKLEKVLGQHFAE 149

consensus

