

logo

		LSMRLRSDALRYEAVPYGEENQVEVYZSDPRRFRERYRRLNRPXSBMLTPEPBYEELEHBBYYPQRNRLGELVQEGQLSR	
NTDB id 1306	DR RS10060 WP 027479822.1	GSMLRADALRLLAVPVDETEQQVTVLTSDPRKRPDIDALIGRPVQMLTRPRDIERLIEQFYPQRGRLGELVQEGTLSR	317
NTDB id 1028	TT RS08215 WP 011173991.1	LLLLRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSR	317
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 1124044	EL305 RS10900 WP 002221967.1	0
NTDB id 1110	NGFG RS09235 WP 003689817.1	0
NTDB id 1251	GCO85 RS07725 WP 011946523.1	0
NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
NTDB id 1058	ABD1 RS01615 WP 001274986.1	0
consensus		0

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		MxYQTTTEPKENLFKREARYASERHKKYKESKKBYSMTTEPFFHAFETTCDAEELSSEPTETES	
		MLTGLCRIVLVEGLYTAQEEALQKASGHSYPEALYSSGLSPLEELAEILSYFQYPLDLS	
NTDB id 1306	DR RS10060 WP 027479822.1	DQLREALQVQAREGKVKPLGEVITELGFASPDVDSALQKQNVCCGRIEDTLVQSCKLRPEMLARSAAQLGYEFLDPIQ	397
NTDB id 1028	TT RS08215 WP 011173991.1	EALKEALEVQKGLPRAKPLGETLVLEGLARPEDVDEALQKQRCCGRIEDTLVQSCKLRPEALACAVATQIGYPYVDPEE	397
NTDB id 1168	A1552VC RS11070 WP 000957200.1	66
NTDB id 1293	VP RS12240 WP 005479695.1	66
NTDB id 1401	DSB67 RS12665 WP 010643259.1	66
NTDB id 1124044	EL305 RS10900 WP 002221967.1	65
NTDB id 1110	NGFG RS09235 WP 003689817.1	65
NTDB id 1251	GCO85 RS07725 WP 011946523.1	74
NTDB id 1018	ACIAD RS01685 WP 004920473.1	74
NTDB id 1058	ABD1 RS01615 WP 001274986.1	74
consensus		

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		YDYLALCOGLLELLRYAYLERRNLYLAVSDPTNLAADDRFATG	
		ELFDEYKAYDRKXKSEFGLVSRGKRVYFLVNRSGEQQEERFVNAKRRKCLNKFAALADERSAFALRQVSE	
		LEVELVIVEHDKLIRLIRLYGR	
NTDB id 1306	DR RS10060 WP 027479822.1	NPPDPKVALMPEATARRVWVVRRLQGNLSLVAMKDRNVFALDDDLKLITG.....KEILPAVMAEKDIRLIERVYGE	472
NTDB id 1028	TT RS08215 WP 011173991.1	DPPDPGAPLLLPEDLCRRYGVVPHRLEGNRLVLLMKDPRNIALDDVRLALKRKGNYEVAPAVATEAATKLIBRFYK	477
NTDB id 1168	A1552VC RS11070 WP 000957200.1	YDYANLCQQLGLRELITRYDALPTAKQCNMLLLAVSDPTLLQAEEDFRFATG.....LQVELALADHRAAQAIRRLYGR	141
NTDB id 1293	VP RS12240 WP 005479695.1	YEYASLCQQLGLRELITRHNAIPLHRTPTSTLLAVADPTNQAEDDFRFATG.....LQVELVLADFRELSTAIRRLYGR	141
NTDB id 1401	DSB67 RS12665 WP 010643259.1	YEYASLCQQLGLRELITRHNAIPLHRTPTSTLLAVADPTNQAEDDFRFATG.....LQVELVLADFRELSTAIRRLYGR	141
NTDB id 1124044	EL305 RS10900 WP 002221967.1	YPRHRVLMGVLTTEEQMVEFHCVVFRGDKVFFAVSDPTQMPQIQKTVSAAG.....TEVELVIVEDDQLAGLLDWWGSR	140
NTDB id 1110	NGFG RS09235 WP 003689817.1	YPRHRVLMGVLTTEEQMVEFHCVVFRGDKVFFAVSDPTQMPQIQKTVSAAG.....TAVELVIVEDDQLAGLLDWWGSR	140
NTDB id 1251	GCO85 RS07725 WP 011946523.1	IDVGTIPVNLVNEKLIKRHAMVPLFSRGTNLYLATDDPSKQASIKETQFHTG.....LNTHAVVETDKLSALIDNLLTA	149
NTDB id 1018	ACIAD RS01685 WP 004920473.1	FDPALFLKDKIDEKLIQKYRIMPLVHRGHVLYVATSNPTNTEAMDAIRFNISK.....LKVEPTIVEHDKLERLSEHFVE	149
NTDB id 1058	ABD1 RS01615 WP 001274986.1	YDTSQIPKDLVDQKLLKLRILPLIQRGQILYVATSNPSNTEADAIRFNISK.....LLVEPVIVEHKKLEKVLGQHFAE	149
consensus		

