

logo

		LDQSLKGLVARGYISQARAKSPESFL	
		LDQSLKGLVARGYISQARAKSPESFL	
NTDB id 1177	A1552VC	RS01165	WP 000422572.1
NTDB id 1203	PSJM300	01605	AFN76401.1
NTDB id 329483	CCS91	RS21905	WP 124866662.1
NTDB id 1307	DR	RS10055	WP 010888596.1
NTDB id 1086	NMV	RS00245	WP 002218487.1
NTDB id 1116	NGFG	RS10615	WP 003690200.1
NTDB id 1185	GTF74	RS11940	WP 000350195.1
NTDB id 1176	A1552VC	RS01160	WP 000350195.1
NTDB id 1283	ACE17W	RS02630	WP 010947729.1
NTDB id 1259	GCO85	RS10305	WP 010947729.1
NTDB id 1202	PSJM300	01600	AFN76400.1
NTDB id 1194	PAKAF	RS01995	WP 003084552.1
NTDB id 1020	ACIAD	RS04210	WP 011182164.1
NTDB id 1081	FDQ49	RS05335	WP 006581387.1
NTDB id 1060	ABD1	RS04380	WP 000355489.1
NTDB id 1040	HON27	RS13360	WP 000355489.1
	consensus		
		FDQALYQLVVDKISEQDALHSADSANDLRLMLKTKRGDD.YGSG..SLQNVKI....DME.....	368
		FDQALYNLYVQGEITYEDALLHADSSNDLRLMIKLGSETD.GEHLTSVSQGLTL....EMSDDDPGRSFR.....	381
		FQQSYQQRVGEGRLL	326
		MDAYLAGLFRHLITYDKGLERAIDPKEFARLANDASAGLGGASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAAPATPA	400
		MDQSLQSLVRQGLIAPEVARRRAQNSSEMSF.....	347
		MDQSLQSLVRQGLIAPEAARRRAQNSSEMSF.....	347
		MEQNAKQLIARGVVDAQEVQSKIELDLKAF.....	345
		MEQNAKQLIARGVVDAQEVQSKIELDLKAF.....	345
		LDQHLTELNVNKNIIISKHTAHEAALNK.SLF.....	344
		LDQHLTELNVNKNIIISKHTAHEAALNK.SLF.....	344
		LDSCCLKGLIAKGLISRDSAKEKAKQP.ENF.....	344
		LDMCLKGLVAKGLISRENAREKAKIP.ENF.....	344
		LDQSLKQLIAKGIINLPTARAVAKSP.ESF.....	344
		LDQSLKGLVSRGVISPQTARTAAKQP.ESFL.....	345
		LDQSLKGLVARGVISPQTARTAAKQP.ESFL.....	345
		LDQSLKGLVARGVISPQTARTAAKQP.ESFL.....	345
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		APAAGRSEPRACCFGR	
NTDB id 1177	A1552VC	RS01165	WP 000422572.1
NTDB id 1203	PSJM300	01605	AFN76401.1
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	consensus		
		368
		381
		326
		APAAGRSEPRACCFGR	418
		347
		347
		345
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		345
		345

X non conserved
X similar
X ≥ 50% conserved