

logo

NTDB id	615	LCA	RS02545	WP	011374200.1	MGQQ.VIACGRQFTAAQLADTQN...NNYSLPQIKRRPAFLRVVKHRLVQCRCQQVV.PPQTCLPDGRHYCAQCLLFGRIVEGDW	79
NTDB id	593	KW2	RS05130	WP	021037147.1	MNLIILENLGLHILLEKDISAFNNFTDQVNNEENKLIKIGAMTSVNANKVRCNRCGTIHIKTNVKLPIGAFFCPTCLELGRVRSDEY	85
NTDB id	236	SPD	RS10765	WP	000867601.1	MKVN.LDYLGRLFTENELT.....EEERQLAEKLPMARKEKGKLFQCRCNSTI.LEEWYLPIGAYYCCECLLMKRVRSDQT	74
NTDB id	202	SPR	RS10250	WP	000867601.1	MKVN.LDYLGRLFTENELT.....EEERQLAEKLPMARKEKGKLFQCRCNSTI.LEEWYLPIGAYYCCECLLMKRVRSDQT	74
NTDB id	277	KZH43	RS10090	WP	000867601.1	MKVN.LDYLGRLFTENELT.....EEERQLAEKLPMARKEKGKLFQCRCNSTI.LEEWYLPIGAYYCCECLLMKRVRSDQT	74
NTDB id	167	SP	RS11275	WP	000867616.1	MKVN.LDYLGRLFTENELT.....EEERQLAEKLPMARKEKGKLFQCRCNSTI.LEEWYLPIGAYYCCECLLMKRVRSDQT	74
NTDB id	507	SM12261	RS09240	WP	000867722.1	MKVN.PNYLGRLFTENELT.....EEERQLAEKLPMARKEKGKLFQCRCNSTI.LEEWYLPIGTYCYCRECLLMKRVRSDQS	74
NTDB id	535	SMSK321	RS10515	WP	000867726.1	MKVN.PNYLGRLFTENELT.....KEERQLAEKLPMARKEKGKLFQCRCDSAI.LDEWYLPIGAYYCCECLLMKRVRSDQV	74
consensus						!*** * !***** *****!!!!!!!!!!!!!! **** **!!!**!*!!**!	

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NTDB	id	615	LCA	RS02545	WP	011374200.1	LYTVPECHLFENNIPKLTWEGQLTPHQEKAQAQAVVEVVQTQKRHVLTAVTGAGKTEMLEFQGITLVALQKGQRVCLAAPRVAVCLEN	164
NTDB	id	593	KW2	RS05130	WP	021037147.1	FYHLPPQDFPEK..TYLRWTGKLTENQEKTSDALCQQITNNQKLVLQAVTGAGKTEMIYQLIEQLTSHGGSVGLASPRIDVCTEL	168
NTDB	id	236	SPD	RS10765	WP	000867601.1	LYYFPQEDFPKQ..DVLKWRGQLTPFQEKVSEGLLQAVDKQKPRTLHVAVTGAGKTEMIYQVVAKVINAGGAVCLASPRIDVCLEN	157
NTDB	id	202	SPR	RS10250	WP	000867601.1	LYYFPQEDFPKQ..DVLKWRGQLTPFQEKVSEGLLQAVDKQKPRTLHVAVTGAGKTEMIYQVVAKVINAGGAVCLASPRIDVCLEN	157
NTDB	id	277	KZH43	RS10090	WP	000867601.1	LYYFPQEDFPKQ..DVLKWRGQLTPFQEKVSEGLLQAVDKQKPRTLHVAVTGAGKTEMIYQVVAKVINAGGAVCLASPRIDVCLEN	157
NTDB	id	167	SP	RS11275	WP	000867616.1	LYYFPQEDFPKQ..DVLKWRGQLTPFQEKVSEGLLVVDKKPRTLHVAVTGAGKTEMIYQVVAKVINAGGAVCLASPRIDVCLEN	157
NTDB	id	507	SM12261	RS09240	WP	000867722.1	LYYFPQEDFPKQ..DVLKWRGQLTPFQDKVSQGLLQAVDKQEPSSLHVAVTGAGKTEMIYQVVAKVINAGGAVCLASPRIDVCLEN	157
NTDB	id	535	SMSK321	RS10515	WP	000867726.1	LYYFPQEDFAKQ..DILKWRGQLTPFQEKVSEGLIQAQVDKQEPILHVAVTGAGKTEMIYQVVAKVINAGGAVCLASPRIDVCLEN	157
consensus							*!!!!***** *!!!!*! !!!!!!!*!!!!!!!!!!!!!!!!!!!!!!*!!!!*! !!!!!*! !!!!!*! !!!!!*! !!!!!*!	

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