

logo

NTDB	id	617	LCA	RS06480	WP	011375001.1	.MSTEEKLTTLLKHCAKQRVQDLYFTPTIGGWR	LERRQSDLVTNQELDKPTGTRYINHLKYMAGMDISERRAQTGRSELQLAD	84
NTDB	id	586	KW2	RS10855	WP	011836048.1	..MVQKKKAQELIQKAIEMEASDIYLLIASGNLYKTIYIRQQLGRITLIEEVNQEIGLALLTHFKFLAGMNTGERRRVQLGACWYELEG	83	
NTDB	id	584	DLJ51	RS00890	WP	019777327.1	MVEVQELGKGLKQAVEEQAQDIYLLIPKPAVYKTYLRFGEKRRHLIQTCPPBQAAGLISHFKFLAGMNVGEKRRQLGACDYAFAD	85	
NTDB	id	385	SMU	RS09035	WP	002263435.1	..MVQKLGKRLIIAEAVQNKAQDIYLLIPKKESEYELMRIGNERRFIDVYDHHFMANLISHFKFVAGMNVGEKRRSQLGSCDYSYDE	83	
NTDB	id	394	FSA28	RS09405	WP	002291652.1	..MVQKLGKRLIIAEAVQNKAQDIYLLIPKKESEYELMRIGNERRFIDVYDHHFMANLISHFKFVAGMNVGEKRRSQLGSCDYSYDE	83	
NTDB	id	500	SM12261	RS00945	WP	000249545.1	..MVQEIAQEIIRSARKKGAQDIYFVPKLDAYELHMRVGDERCKIGCYDFEKFAAVISHFKFVAGMNVGEKRRSQLGSCDYEYDQ	83	
NTDB	id	270	KZH43	RS09265	WP	000249564.1	..MVQEIAQEIIRSARKKGTQDIYFVPKLDAYELHMRVGDERCKIGSYDFEKFAAVISHFKFVAGMNVGEKRRSQLGSCDYAYDH	83	
NTDB	id	229	SPD	RS09895	WP	000249564.1	..MVQEIAQEIIRSARKKGTQDIYFVPKLDAYELHMRVGDERCKIGSYDFEKFAAVISHFKFVAGMNVGEKRRSQLGSCDYAYDH	83	
NTDB	id	195	SPR	RS09430	WP	000249564.1	..MVQEIAQEIIRSARKKGTQDIYFVPKLDAYELHMRVGDERCKIGSYDFEKFAAVISHFKFVAGMNVGEKRRSQLGSCDYAYDH	83	
NTDB	id	160	SP	RS10380	WP	000249564.1	..MVQEIAQEIIRSARKKGTQDIYFVPKLDAYELHMRVGDERCKIGSYDFEKFAAVISHFKFVAGMNVGEKRRSQLGSCDYAYDH	83	
NTDB	id	65556	ANG	RS10070	WP	006268058.1	..MVQNIQAQAIICQAKEERAQDIYFVPKENVYELMRIGDERRFIRNYDFEELAAVISHFKFLTAGMNVGEKRRSQLGSCDYQYSD	83	
NTDB	id	429	SGO	RS09420	WP	012130942.1	..MVQKIAQAIIVRQAKEECQAQDIYFVPKDCYELMRIGDERRFIQTYDFDQLAAVISHFKFLTAGMNVGEKRRSQLGSCDYRYDD	83	
consensus							*****	*****	*****

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NTDB	id	617	LCA	RS06480	WP	011375001.1	Q.TIYLRRLATVGDFLNRESLVIRFIYPIGAIIYHCDDQLILAQLTQMRSQSGLILFAGPTGSGKTTSLYLHLAQQSMSAKMVVAIED	168
NTDB	id	586	KW2	RS10855	WP	011836048.1	SSAKRRLRLSTVGDFEGNESLVIRLLHDKKELEFWFD.DK.LLQDFRCRKRGLYLFAGPVGSGKTSLMFDLAHRHFNSNAQVITIEE	165
NTDB	id	584	DLJ51	RS00890	WP	019777327.1	GETAALRLSTVGVDYGGRSRLVIRLLFDRKRELKFWFND.GLINALLPKPIIGRGLYLFSGPVGSCKTTLMYELVSRKYPDSPQITIED	169
NTDB	id	385	SMU	RS09035	WP	002263435.1	GKVVISRLRSSVGDYRGYESLVIRLLFSGRHLDLKFWFN.SMDLLINEIKGRGLYLFSGPVGSGKTSLMYQLVKRFETKQVITIED	167
NTDB	id	394	FSA28	RS09405	WP	002291652.1	GKVISRLRSSVGDYRGYESLVIRLLFSGRHLDLKFWFN.SMDLLINEIKGRGLYLFSGPVGSGKTSLMYQLVKRFETKQVITIED	167
NTDB	id	500	SM12261	RS00945	WP	000249545.1	K.MASRLRLSTVG DYRGHESLVIRLLHDEEQDLHFWFQ.DIEELGKQYRQRGLYLFAGPVGSGKTTLMHELAKSKSLFKCQQVM SIED	166
NTDB	id	270	KZH43	RS09265	WP	000249564.1	K.IASRLRLSTVG DYRGHESLVIRLLHDEEQDLHFWFQ.DIEELGKQYRQRGLYLFAGPVGSGKTTLMHELAKSKSLFKCQQVM SIED	166
NTDB	id	229	SPD	RS09895	WP	000249564.1	K.IASRLRLSTVG DYRGHESLVIRLLHDEEQDLHFWFQ.DIEELGKQYRQRGLYLFAGPVGSGKTTLMHELAKSKSLFKCQQVM SIED	166
NTDB	id	195	SPR	RS09430	WP	000249564.1	K.IASRLRLSTVG DYRGHESLVIRLLHDEEQDLHFWFQ.DIEELGKQYRQRGLYLFAGPVGSGKTTLMHELAKSKSLFKCQQVM SIED	166
NTDB	id	160	SP	RS10380	WP	000249564.1	K.IASRLRLSTVG DYRGHESLVIRLLHDEEQDLHFWFQ.DIEELGKQYRQRGLYLFAGPVGSGKTTLMHELAKSKSLFKCQQVM SIED	166
NTDB	id	65556	ANG	RS10070	WP	006268058.1	K.KTSLRLSTVG DYRGYESLVIRLLHDEEAELKFWFA.QIPELKERFKQRGLYLFSGPVGSGKTTLMHQLAQERFANQQIM SIED	166
NTDB	id	429	SGO	RS09420	WP	012130942.1	K.ETSIRLSTVG DYRGYESLVIRLLHDEETEELKFWFT.HFPLELREKFKDRGLYLFSGPVGSGKTTLMHQLAQLKFKCQQVM SIED	166
consensus							* * * * *	

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NTDB	id	617	LCA	RS06480	WP	011375001.1	P	T	E	I	V	A	P	E	F	L	Q	L	Q	V	N	D	N	A	G	L	S	Y	A	E	L	L	K	I	S	L	R	L	R	P	D	T	L	I	I	G	E	I	R	D	L	E	T	A	Q	Y	V	V	S	A	A	L	S	C	H	L	V	L	S	T	I	H	A	R	S	T	Q	G	V	V	A	R	L	L	D	L	G	I	S	E	L	253
NTDB	id	586	KW2	RS10855	WP	011836048.1	P	V	E	L	I	D	S	D	F	I	Q	L	Q	V	N	D	V	I	G	N	S	Y	D	E	L	I	K	L	S	L	R	H	R	P	D	L	L	I	V	G	E	I	R	D	Q	T	A	R	A	V	L	R	A	S	L	T	G	Y	T	V	F	S	T	I	H	A	S	V	K	G	V	V	A	R	L	L	E	L	G	I	S	D	W	250		
NTDB	id	584	DLJ51	RS00890	WP	019777327.1	P	V	E	I	K	Q	D	N	M	L	Q	L	Q	L	N	P	S	I	G	M	T	Y	D	H	L	I	K	L	S	L	R	H	R	P	D	V	L	I	I	G	E	I	R	D	R	E	T	A	Q	A	V	E	R	A	S	L	T	G	I	R	V	F	S	T	I	H	A	K	S	I	P	G	V	Y	A	R	L	E	L	G	V	S	E	Q	254	
NTDB	id	385	SMU	RS09035	WP	002263435.1	P	V	E	I	K	Q	D	N	M	L	Q	L	Q	L	N	P	S	I	G	M	T	Y	D	N	L	I	K	L	S	L	R	H	R	P	D	V	L	I	I	G	E	I	R	D	K	E	T	A	R	A	V	E	R	A	S	L	T	G	A	M	V	F	S	T	I	H	A	K	S	I	P	G	V	Y	A	R	L	E	L	G	T	S	E	252		
NTDB	id	394	FSA28	RS09405	WP	002291652.1	P	V	E	I	K	Q	D	N	M	L	Q	L	Q	L	N	P	S	I	G	M	T	Y	D	N	L	I	K	L	S	L	R	H	R	P	D	V	L	I	I	G	E	I	R	D	K	E	T	A	R	A	V	E	R	A	S	L	T	G	A	M	V	F	S	T	I	H	A	K	S	I	P	G	V	Y	A	R	L	E	L	G	T	S	E	252		
NTDB	id	500	SM12261	RS00945	WP	000249545.1	P	V	E	I	K	Q	D	N	M	L	Q	L	Q	L	N	E	A	I	G	L	T	Y	E	N	L	I	K	L	S	L	R	H	R	P	D	L	L	I	I	G	E	I	R	D	S	E	T	A	R	A	V	V	R	A	S	L	T	G	A	T	V	F	S	T	I	H	A	K	S	I	R	G	V	Y	E	R	L	L	E	L	G	V	S	E	251	
NTDB	id	270	KZH43	RS09265	WP	000249564.1	P	V	E	I	K	Q	D	N	M	L	Q	L	Q	L	N	E	A	I	G	L	T	Y	E	N	L	I	K	L	S	L	R	H	R	P	D	L	L	I	I	G	E	I	R	D	S	E	T	A	R	A	V	V	R	A	S	L	T	G	A	T	V	F	S	T	I	H	A	K	S	I	R	G	V	Y	E	R	L	L	E	L	G	V	S	E	251	
NTDB	id	229	SPD	RS09895	WP	000249564.1	P	V	E	I	K	Q	D	N	M	L	Q	L	Q	L	N	E	A	I	G	L	T	Y	E	N	L	I	K	L	S	L	R	H	R	P	D	L	L	I	I	G	E	I	R	D	S	E	T	A	R	A	V	V	R	A	S	L	T	G	A	T	V	F	S	T	I	H	A	K	S	I	R	G	V	Y	E	R	L	L	E	L	G	V	S	E	251	
NTDB	id	195	SPR	RS09430	WP	000249564.1	P	V	E	I	K																																																																																	

logo



X non conserved
X similar
X ≥ 50% conserved