

NTDB id	615	LCA RS02545 WP 011374200.1	.MGQQV IACGRQFTAAQLADTQNNY....SLPQIKRPAFLRVKHLVLCQRGCG.VVPPTCTCPDGRHYCAQLLFGRVEGDW	79
NTDB id	593	KW2 RS05130 WP 021037147.1	MNLILENLLCHILLEKDISAFNNFTDQVNNEKLIGIGAMTSVNANKVRGNRCGIHKTNVKLPIGAFFCPTCLELGRVRSDEY	85
NTDB id	120889	DK92 RS02635 WP 047235723.1	.MESIENSYGRILLESCLPDSA.....KQLAQPLKSVVILRGKMICQRCHY.QLD EEARLP SGAYYC RFCLVFGRNQSDKL	74
NTDB id	277	KZH43 RS10090 WP 000867601.1	.MKVNL DYLGRLFTENELTEEE.....RQLAEKLPAMRKEKGKLF CQCNCNS.TILEEWYLP IGAYYCRECLLMKRVRSDQT	74
NTDB id	236	SPD RS10765 WP 000867601.1	.MKVNL DYLGRLFTENELTEEE.....RQLAEKLPAMRKEKGKLF CQCNCNS.TILEEWYLP IGAYYCRECLLMKRVRSDQT	74
NTDB id	202	SPR RS10250 WP 000867601.1	.MKVNL DYLGRLFTENELTEEE.....RQLAEKLPAMRKEKGKLF CQCNCNS.TILEEWYLP IGAYYCRECLLMKRVRSDQT	74
NTDB id	167	SP RS11275 WP 000867616.1	.MKVNL DYLGRLFTENELTEEE.....RQLAEKLPAMRKEKGKLF CQCNCNS.TILEEWYLP IGAYYCRECLLMKRVRSDQT	74
NTDB id	507	SM12261 RS09240 WP 000867722.1	.MKVNP NYLGRLFTENELTEEE.....RQLAEKLPAMRKEKGKLF CQCNCNS.TILEEWYLP IC TYYCRECLLMKRVRSDQS	74
NTDB id	535	SMSK321 RS10515 WP 000867726.1	.MKVNP NYLGRLFTENELTK EE.....RQLAEKLPAMRKEKGKLF CQCNCNS.AILD EWYLP IGAYYCRECLLMKRVRSDQV	74
consensus			***** **!* **** ***** ! * !! * * **** ***** ! * !! * * **** ***** ! * !! * *	

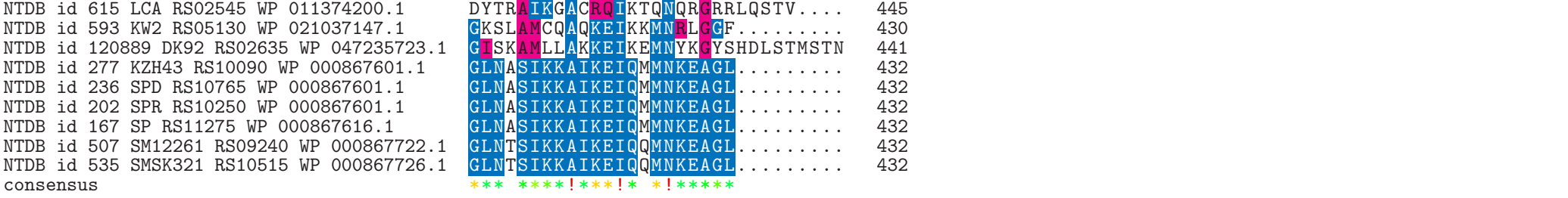
NTDB id 615 LCA RS02545 WP 011374200.1	LYTVPECHLFENNIPKLTWEGQLTPHQEAAQAVVEVVQTKQRHVLTAVTGAGKTEM	LFQGLVALQKQQRVCLAAPRVAVCLEL	164	
NTDB id 593 KW2 RS05130 WP 021037147.1	FYHLPQQDFPEKT..YLRWGTGKLTENQEKISDALCQQITNNQKL	LVQAVTGAGKTEMIYQLIEQILSHGGSVGLASPRIDVCTEL	168	
NTDB id 120889 DK92 RS02635 WP 047235723.1	LYAIPPMHFPKGN..YLVWSSGQLTAYQEILSQQLLINMQNQKT	TLVHAVTGAGKTEMIYAAAEAVINTGCWVCIASPRVDVCVEV	157	
NTDB id 277 KZH43 RS10090 WP 000867601.1	LYYFPQEDFPKQD..VLKWRGQLTPFQEKVSEGLLQAVDKQKPTLVHAVTGAGKTEMIYQVVA	KVINAGGAVCLASPRIDVCLEL	157	
NTDB id 236 SPD RS10765 WP 000867601.1	LYYFPQEDFPKQD..VLKWRGQLTPFQEKVSEGLLQAVDKQKPTLVHAVTGAGKTEMIYQVVA	KVINAGGAVCLASPRIDVCLEL	157	
NTDB id 202 SPR RS10250 WP 000867601.1	LYYFPQEDFPKQD..VLKWRGQLTPFQEKVSEGLLQAVDKQKPTLVHAVTGAGKTEMIYQVVA	KVINAGGAVCLASPRIDVCLEL	157	
NTDB id 167 SP RS11275 WP 000867616.1	LYYFPQEDFPKQD..VLKWRGQLTPFQEKVSEGLLQAVDKQKPTLVHAVTGAGKTEMIYQVVA	KVINAGGAVCLASPRIDVCLEL	157	
NTDB id 507 SM12261 RS09240 WP 000867722.1	LYYFPQEDFPKQD..VLKWRGQLTPFQDKVSGLLQAVDKQEP	SLVHAVTGAGKTEMIYQVVA	KVINAGGAVCLASPRIDVCLEL	157
NTDB id 535 SMSK321 RS10515 WP 000867726.1	LYYFPQEDFAKQD..ILKWRGQLTPFQEKVSEGLTQAVDKQEP	ILVHAVTGAGKTEMIYQVVA	KVINAGGAVCLASPRIDVCLEL	157
consensus	*****	*****		

[illegible][illegible]

logo



logo



☐ non conserved
☒ similar
☒ ≥ 50% conserved