

logo

NTDB id 413 AAK55817.1 141..845()	MKKILIVDDEKPIISDIIFKFNMTKEGYEVVTAFA.NGREALEQFEAEQPDIIILDLMLPEIDGLEVAKTIRKTS	74
NTDB id 374 SMU RS06885 WP 002262930.1	MKKILIVDDEKPIISDIIFKFNLAKEGYDTITAF.DGREALSKEYEENPDIIILDLMLPELDGLEVAKEVRKNSHVP	74
NTDB id 1065869 ACGYMG RS19900 WP 070113453.1	MESIKDKKILVDDHKELKLMIDEILRKEGFSRVFLASSYEEAVRVFKNVKPDCAILDVVLPGDGFSLMRKIRETSKIP	80
NTDB id 372 SMU RS05280 WP 002262198.1	.MNYKEKKILIIDDNPETILEMVQESLNIAGFNNLTGVQSKKEALEQLGKRSDLAILDIMLPDGSGFVLRREIRKTSITIP	79
consensus	*****	

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NTDB id 413 AAK55817.1 141..845()	ITMLSAKDSEFDKVIQLELGADDYVTKPFSNRELQARVKALLRRSQMPVVDGQ.EADSKPQPTIQIGDL.EIVPDAYVAKK	152
NTDB id 374 SMU RS06885 WP 002262930.1	ITMLSAKDSEFDKVIQLEIGADDYVTKPFSNRELLARVKAHLRRTENIESAVAEENASGIPEIIGDL.QILPDAFVAKK	153
NTDB id 1065869 ACGYMG RS19900 WP 070113453.1	VIFLSARGEDEDRLIGLGLGADDYIVKPFLPKELTLRLVIGILNRVYVPIEE.....EELPVFKLGDSVVVNLNSACVEK	154
NTDB id 372 SMU RS05280 WP 002262198.1	VIFLSAVSDIEKQYQGFELGADDYIVKPFRPKLELRLILSILKRAYPEKDD.....TLI.....LPTCQVNFSQLITK	148
consensus	*****	

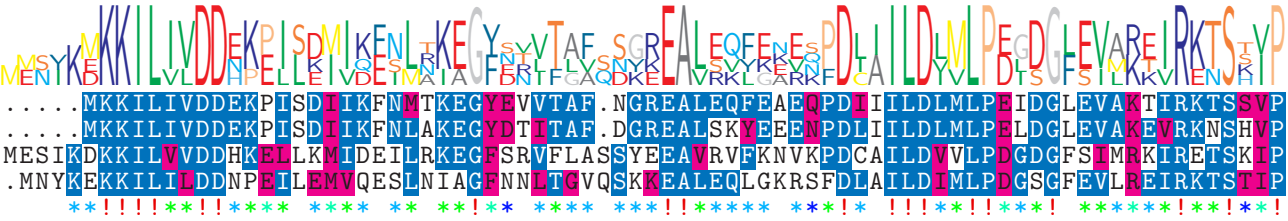
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NTDB id 413 AAK55817.1 141..845()	YGEELDLTHREFELLYHLASHTGQVITREHLLLETVWGYDYFGDVRITVDVITVRRRLREKIEDTPSRPEYILTRRGVGYMMRN	232
NTDB id 374 SMU RS06885 WP 002262930.1	RGTEVELTHREFELLHHLAHTGQVMTREHLLLETVWGYDYFGDVRITVDVITVRRRLREKIEDTPSRPEYILTRRGVGYMMKS	233
NTDB id 1065869 ACGYMG RS19900 WP 070113453.1	DGQEISLTAKEHSLLLKLYENKGRIVTSDSLCQAIWGDMDYGYENTLMVHVRRVREKIEKTPSTPKHLLTVRGLGYKLMT	234
NTDB id 372 SMU RS05280 WP 002262198.1	GELEIQLTAKESYILKVLVLYNNQNRIVTFDQLLEKVVWGLQYQGYENTMMAHIRKIRQKIEANPSKPECCLLTVKGLGYKLKV	228
consensus	*****	

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NTDB id 413 AAK55817.1 141..845()	NA*	234
NTDB id 374 SMU RS06885 WP 002262930.1	YD.	235
NTDB id 1065869 ACGYMG RS19900 WP 070113453.1	N..	235
NTDB id 372 SMU RS05280 WP 002262198.1	S..	229
consensus	*	



- X non conserved
- X similar
- X ≥ 50% conserved