

[illegible][illegible]

NTDB id 259 KZH43 RS03480 WP 000491790.1	LINTSQLEQASQKHEQLIVVVMASFWILSLLASLYLARVSVRPLLESQMKGQS FVENASHELRTPLAVLQNRLET LFRKPEATIM	251
NTDB id 218 SPD RS03780 WP 000491790.1	LINTSQLEQASQKHEQLIVVVMASFWILSLLASLYLARVSVRPLLESQMKGQS FVENASHELRTPLAVLQNRLET LFRKPEATIM	251
NTDB id 184 SPR RS03565 WP 000491790.1	LINTSQLEQASQKHEQLIVVVMASFWILSLLASLYLARVSVRPLLESQMKGQS FVENASHELRTPLAVLQNRLET LFRKPEATIM	251
NTDB id 148 SP RS03910 WP 000491790.1	LINTSQLEQASQKHEQLIVVVMASFWILSLLASLYLARVSVRPLLESQMKGQS FVENASHELRTPLAVLQNRLET LFRKPEATIM	251
NTDB id 34392 SDEG RS05190 WP 012766987.1	VVNVEQLDSANERYEQIIIIIVMII FWLISTI LASTYL SKWSRKPILESYEKKMFVENASHELRTPLAVLQNRLES LFRKPNETIL	253
NTDB id 363 SMU RS05195 WP 002262215.1	VINTTGLEEANTIRYVTIIISVMIVFWLISVIASVYLA KWSQKPILENYEKQKS FVENASHELRTPLAVLQNRLES LFRKPEATIL	248
consensus	* * * * *	

NTDB id 259 KZH43 RS03480 WP 000491790.1	DVSESIASSLEEVRNMRFLTTSLLNLARRDDGIKPELAEVPTSFNNFTFTNYEMIASENNVRVFRFENRIHRTIVTDQLLKQLMT	336
NTDB id 218 SPD RS03780 WP 000491790.1	DVSESIASSLEEVRNMRFLTTSLLNLARRDDGIKPELAEVPTSFNNFTFTNYEMIASENNVRVFRFENRIHRTIVTDQLLKQLMT	336
NTDB id 184 SPR RS03565 WP 000491790.1	DVSESIASSLEEVRNMRFLTTSLLNLARRDDGIKPELAEVPTSFNNFTFTNYEMIASENNVRVFRFENRIHRTIVTDQLLKQLMT	336
NTDB id 148 SP RS03910 WP 000491790.1	DVSESIASSLEEVRNMRFLTTSLLNLARRDDGIKPELAEVPTSFNNFTFTNYEMIASENNVRVFRFENRIHRTIVTDQLLKQLMT	336
NTDB id 34392 SDEG RS05190 WP 012766987.1	ENSEHLASSLDEVNRMRILTNNLLNLARRDDGINPQWTDLDATFFDAIFENYQLVAEENGKVFIFENQVNRIRMDKALLKQLMT	338
NTDB id 363 SMU RS05195 WP 002262215.1	DSSESIASSLEEVRNMRMLTANLLNLARRDDGIKPEITEIQPRFFDDTFKNYEMIARENGKKLHIINKVNRQIKTDKTLKQLMT	333
consensus	*****	

Accession	Protein	Length	Score	E-value	Identity	Positives	Gaps
NTDB id 259 KZH43 RS03480 WP 000491790.1	ILFDNAVKYTEEDGEIDFLISATDRNLYLLVSDNGIGISTEDKKKIFDRFYRVDKARTRQKGGFGLGLSLAKQIVDALKGTVTVK	421	100	0.00	100	100	0
NTDB id 218 SPD RS03780 WP 000491790.1	ILFDNAVKYTEEDGEIDFLISATDRNLYLLVSDNGIGISTEDKKKIFDRFYRVDKARTRQKGGFGLGLSLAKQIVDALKGTVTVK	421	100	0.00	100	100	0
NTDB id 184 SPR RS03565 WP 000491790.1	ILFDNAVKYTEEDGEIDFLISATDRNLYLLVSDNGIGISTEDKKKIFDRFYRVDKARTRQKGGFGLGLSLAKQIVDALKGTVTVK	421	100	0.00	100	100	0
NTDB id 148 SP RS03910 WP 000491790.1	ILFDNAVKYTEEDGEIDFLISATDRNLYLLVSDNGIGISTEDKKKIFDRFYRVDKARTRQKGGFGLGLSLAKQIVDALKGTVTVK	421	100	0.00	100	100	0
NTDB id 34392 SDEG RS05190 WP 012766987.1	ILFDNAIKYTDKEGVVDIKVKTTEKNLLISVIDNGPGIKDEKKKIFDRFYRVDKARTRQTGGFGLGLALAQQIVITSLKCSISVK	423	100	0.00	100	100	0
NTDB id 363 SMU RS05195 WP 002262215.1	ILFDNAVKYTDNDGEIWLDDVKTITDRSLIFSVADNGQGISEDKKKIFDRFYRVDKARTRQKGGFGLGLSLAKQITDSLKCSISIVK	418	100	0.00	100	100	0
consensus							

logo

NTDB id 259 KZH43 RS03480 WP 000491790.1	DNKPKGTIFEVKIAIQTPSKKKK	444
NTDB id 218 SPD RS03780 WP 000491790.1	DNKPKGTIFEVKIAIQTPSKKKK	444
NTDB id 184 SPR RS03565 WP 000491790.1	DNKPKGTIFEVKIAIQTPSKKKK	444
NTDB id 148 SP RS03910 WP 000491790.1	DNKPKGTIFEVKIAIQTPSKKKK	444
NTDB id 34392 SDEG RS05190 WP 012766987.1	DNLPKCSIFEVKL.....	436
NTDB id 363 SMU RS05195 WP 002262215.1	DKKPKQGVMEARLLANK.....	435
consensus	! ** ! * ! * * ! ! * * * * * * * * * * *	

-  non conserved
-  similar
-  ≥ 50% conserved