

logo

NTDB id 586 KW2 RS10855 WP 011836048.1	LKNGLEAVVYQRLIAGKGVLDIAKSKFDTWSPKKWNEKIENLYANGHLTAIEAEREKISINQASKIDTTDGESFE	326
NTDB id 429 SGO RS09420 WP 012130942.1	LKIVLQGGICVYQRLIGGGGVIDFASDNYQEHEPTVWNQIDQLLAAGHIHPEQAEAEKIRNQQAKTSY.....	319
NTDB id 500 SM12261 RS00945 WP 000249545.1	LAVVLQGVVCYQRLIGGGGIVDFANKDYQEHQSTSWNAQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 270 KZH43 RS09265 WP 000249564.1	LAVVLQGVVCYQRLIGGGGIVDFASRDYQEHQAAKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 229 SPD RS09895 WP 000249564.1	LAVVLQGVVCYQRLIGGGGIVDFASRDYQEHQAAKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 195 SPR RS09430 WP 000249564.1	LAVVLQGVVCYQRLIGGGGIVDFASRDYQEHQAAKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 160 SP RS10380 WP 000249564.1	LAVVLQGVVCYQRLIGGGGIVDFASRDYQEHQAAKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 584 DLJ51 RS00890 WP 019777327.1	LDNCLAGIAYQRLIGGGGLIDFAQDHFQNHQAECWNEKIDQLLAAGHLTEEEAAAEKIKD.....	315
NTDB id 22485 GBS RS11090 WP 000250808.1	LENSLKLIAVQRLIGGGSLIDFETGNFKKHSSDKWNRQVDILAEEGHISKKKQAQVEKIIPQETTESPTF.....	323
NTDB id 385 SMU RS09035 WP 002263435.1	LQNALRGIVYQRLIGGGGIIDFAKGDYQNHSAKKWNKQIYDLFTAGYITEGEAQAEIIED.....	313
NTDB id 394 FSA28 RS09405 WP 002291652.1	LQNALCGIVYQRLIGGGGIIDFAKGDYQNHSAKKWNKQIYDLFTAGYITEGEAQAEIIED.....	313
consensus	! **!*!!!!!!*!*!!!!* *!!!!* *!!*!!!!!* *!*** **!* !!!*	

- ☐ non conserved
- ☒ similar
- ☒ ≥ 50% conserved