

logo



NTDB id 586 KW2 RS10855 WP 011836048.1	LKNGLEAVVYQRLIAGKGVLDIAKSKFDTWSPKKNWNEKIENLYANGHLTAIEAEREKISINQASKIDTTDGESFE	326
NTDB id 429 SGO RS09420 WP 012130942.1	LKIVLQGITCYQLIGGGGVIDFASDNYQEHEPTVWVQQIDQLLAAGHIHPEQAEAEKIRNQQAKTSY.....	319
NTDB id 500 SM12261 RS00945 WP 000249545.1	LAVVLQGVVYQRLIGGGGIVDFANKDYQEHQSTSWNAQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 270 KZH43 RS09265 WP 000249564.1	LAVVLQGVVYQRLIGGGGIVDFASRDYQEHQAAKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 229 SPD RS09895 WP 000249564.1	LAVVLQGVVYQRLIGGGGIVDFASRDYQEHQAAKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 195 SPR RS09430 WP 000249564.1	LAVVLQGVVYQRLIGGGGIVDFASRDYQEHQAAKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 160 SP RS10380 WP 000249564.1	LAVVLQGVVYQRLIGGGGIVDFASRDYQEHQAAKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 584 DLJ51 RS00890 WP 019777327.1	LDNCLAGIAYQRLIGGGGLIDFAQDHFQNHQAECWNEKIDQLLAAGHLTEEEAAAEEKIKD.....	315
NTDB id 121862 EN71 RS11255 WP 000250805.1	LENSLKLIAVYQRLIGGGSLIDFETGNFKKHSSDKWNRQVDILAEEGHISKKQAQVEKIIPQETTESPTF.....	323
NTDB id 385 SMU RS09035 WP 002263435.1	LQNALRGIVYQRLIGGGGVIDFAKGDYQNHSAKKWVQIYDLFTAGYITEGEAQAEETIED.....	313
NTDB id 394 FSA28 RS09405 WP 002291652.1	LQNALCGIVYQRLIGGGGVIDFAKGDYQNHSAKKWVQIYDLFTAGYITEGEAQAEETIED.....	313
consensus	! **!*!!!!!!*!*****!*** ***** *!*****!* *!*** **!* !!!*	

non conserved
 similar
 ≥ 50% conserved