

logo



NTDB id 617 LCA RS06480 WP 011375001.1	QLRACLITGIAYQSLVTKDDVVQAH...YELLTGSDYFSKKE.....GGCDV.....	296
NTDB id 586 KW2 RS10855 WP 011836048.1	ELKNGLEAVVYQRLIAGKGVLDIAKSKFDTWSPKKWNEKIENLYANGHLTATEAEREKISINQASKIDTTDGESFE	326
NTDB id 584 DLJ51 RS00890 WP 019777327.1	ELDNCLAGIAYQRLIGGGGLIDFAQDHFQNHQAECWNEKIDQLLAAGHLTEEEAAAEEKIKD.....	315
NTDB id 385 SMU RS09035 WP 002263435.1	ELQNALRGIVYQRLIGGGGIIDFAKGDYQNHSAKKWKNQIYDLFTAGYITEGEAQAEIIEED.....	313
NTDB id 394 FSA28 RS09405 WP 002291652.1	ELQNALCGIVYQRLIGGGGIIDFAKGDYQNHSAKKWKNQIYDLFTAGYITEGEAQAEIIEED.....	313
NTDB id 429 SGO RS09420 WP 012130942.1	ELKIVLQGGICYQRLIGGGVIDFASDNYQEHSEPTVWNQQIDQLLAAGHIHPEQAEAEKIRNQQAKTSY.....	319
NTDB id 500 SM12261 RS00945 WP 000249545.1	ELAVVLQGVVVCYQRLIGGGGIVDFANKDYQEHQSTSWNAQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 98224 R8625 RS09925 WP 016398028.1	ELAVVLQGVVVCYQRLIGGGGIVDFASRDYQEHQAQAKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 270 KZH43 RS09265 WP 000249564.1	ELAVVLQGVVVCYQRLIGGGGIVDFASRDYQEHQAQAKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 229 SPD RS09895 WP 000249564.1	ELAVVLQGVVVCYQRLIGGGGIVDFASRDYQEHQAQAKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 195 SPR RS09430 WP 000249564.1	ELAVVLQGVVVCYQRLIGGGGIVDFASRDYQEHQAQAKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 160 SP RS10380 WP 000249564.1	ELAVVLQGVVVCYQRLIGGGGIVDFASRDYQEHQAQAKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
consensus	*! **!***!!**!***** **!***** **!***** **!***** **!***** **!*****	

non conserved
 similar
 ≥ 50% conserved