

logo

NTDB id 617 LCA RS06480 WP 011375001.1	QLRACLTGTAAYQSLVTKDDVVQA...HYELLTGSDYFSKKEG.....G.....CDV.....	296
NTDB id 586 KW2 RS10855 WP 011836048.1	ELKNGLEAVVYQRLIAGKGVLDIAKSKFDTWSPKKWNEKTENLYANGHLTAIEEAEREKISINQASKIDTTDGESFE	326
NTDB id 584 DLJ51 RS00890 WP 019777327.1	ELDNCLAGIAYQRLIGGGGLIDFAQDHFQNHQAECWNEKIDQLLAAGHLTEEEAAAEEKIKD.....	315
NTDB id 385 SMU RS09035 WP 002263435.1	ELQNALRGIVYQRLIGGGGIIDFAKGDYQNHSACKWNNKQIYDLFTAGYITEGEAQAEIIED.....	313
NTDB id 394 FSA28 RS09405 WP 002291652.1	ELQNALCGIVYQRLIGGGGIIDFAKGDYQNHSACKWNNKQIYDLFTAGYITEGEAQAEIIED.....	313
NTDB id 39995 N596 RS08115 WP 023027559.1	ELRVVLQGICYQRLIKGGGVITDFVIQDYQNHSSQKWNQQIDTLYEAGHIELDQAQAEKIIDCQARACH.....	319
NTDB id 429 SGO RS09420 WP 012130942.1	ELKIVLQGICYQRLIGGGGVITDFASDNYQEHEPTVWNNQQIDQLLAAGHIHPEQAEAEKIRNQQAKTSY.....	319
NTDB id 270 KZH43 RS09265 WP 000249564.1	ELAVVLQGVICYQRLIGGGGIVDFASRDYQEHQAACKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 229 SPD RS09895 WP 000249564.1	ELAVVLQGVICYQRLIGGGGIVDFASRDYQEHQAACKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 195 SPR RS09430 WP 000249564.1	ELAVVLQGVICYQRLIGGGGIVDFASRDYQEHQAACKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 160 SP RS10380 WP 000249564.1	ELAVVLQGVICYQRLIGGGGIVDFASRDYQEHQAACKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 500 SM12261 RS00945 WP 000249545.1	ELAVVLQGVICYQRLIGGGGIVDFANKDYQEHQSTSWNAQIDQLLKDGHITSLQAETEKISYS.....	313
consensus	*! **!*****!!*!***** **	

- non conserved
- similar
- ≥ 50% conserved