

NTDB id	586 KW2 RS10855 WP 011836048.1	LKNGLEAVVYQRLIAGKGVLIDIAKSFKDTWSPKKWNEKIENLYANGHLTAIEAREKISINQASKIDTTDGSFE	326
NTDB id	429 SGO RS09420 WP 012130942.1	LKITVLQGICYQLRIGGGGVIDFASDNYYQEHEPTVWNQQIIDQLLAAGHIHPEQAEEIKIRNQQAKTSY.....	319
NTDB id	500 SM12261 RS00945 WP 000249545.1	LAVVLQGVVCYQRLLGGGGIVDFANKDYQEHQSSTSWNAQIQDLLKDGHITSLQAETEKISYS.....	313
NTDB id	270 KZH43 RS09265 WP 000249564.1	LAVVLQGVVCYQRLLGGGGIVDFSADYQEHQAACKWNEQIDQLLKDGHITSLSQAETEKISYS.....	313
NTDB id	229 SPD RS09895 WP 000249564.1	LAVVLQGVVCYQRLLGGGGIVDFSADYQEHQAACKWNEQIDQLLKDGHITSLSQAETEKISYS.....	313
NTDB id	195 SPR RS09430 WP 000249564.1	LAVVLQGVVCYQRLLGGGGIVDFSADYQEHQAACKWNEQIDQLLKDGHITSLSQAETEKISYS.....	313
NTDB id	160 SP RS10380 WP 000249564.1	LAVVLQGVVCYQRLLGGGGIVDFSADYQEHQAACKWNEQIDQLLKDGHITSLSQAETEKISYS.....	313
NTDB id	584 DLJ51 RS00890 WP 019777327.1	LDNCLAGTIAYQRLIGGGGLIDFAQDHFNHNHAECWNEKIDQLLAAGHTITEEAAAEEKIKD.....	315
NTDB id	124402 GT95 RS01055 WP 069107363.1	LENSSLKLTIAYQRLIGGGSLIDFETGNPKKHSSDKWNRQVDILAEFGHSKKQAQVEKIIPQETTESSPTF....	323
NTDB id	385 SMU RS09035 WP 002263435.1	LQNALRCGIIVYQRLIGGGGITDAFKGDYQNHSAKKWKQIIYDLFTAGYITEGEAQAEEIED.....	313
NTDB id	394 FSA28 RS09405 WP 002291652.1	LQNALCGIIVYQRLIGGGGITDAFKGDYQNHSAKKWKQIIYDLFTAGYITEGAQAEEIED.....	313
consensus		! **!* ****!!!!!! * !***** !!** ***** !!!**** * !!*** **!* *!	

- X non conserved
- X similar
- X $\geq 50\%$ conserved