

NTDB id	586	KW2	RS10855	WP	011836048.1	LKNGLEAVVYQRLIAGKGVLIDIAKSKFDTWSPKKWNEKTENLYANGHLTAIEAREKISINQASKIDTTDGSFSE	326
NTDB id	429	SG0	RS09420	WP	012130942.1	LKIVLQGICYQRLIGGGGVIDFASDNVQEHEPTVWNQQIDQLLAAGHIHPEQAEEIKIRNQQAKTSY.....	319
NTDB id	500	SM12261	RS00945	WP	000249545.1	LAVVLQGVVCYQRLIGGGGIVDFANKDYQEHQSSTSWNAQIDQLLKDGHITSQAETEKISYS.....	313
NTDB id	270	KZH43	RS09265	WP	000249564.1	LAVVLQGVVCYQRLIGGGGIVDFASRDYQEHQAARKWNEQIDQLLKDGHITSQAETEKISYS.....	313
NTDB id	229	SPD	RS09895	WP	000249564.1	LAVVLQGVVCYQRLIGGGGIVDFASRDYQEHQAARKWNEQIDQLLKDGHITSQAETEKISYS.....	313
NTDB id	195	SPR	RS09430	WP	000249564.1	LAVVLQGVVCYQRLIGGGGIVDFASRDYQEHQAARKWNEQIDQLLKDGHITSQAETEKISYS.....	313
NTDB id	160	SP	RS10380	WP	000249564.1	LAVVLQGVVCYQRLIGGGGIVDFASRDYQEHQAARKWNEQIDQLLKDGHITSQAETEKISYS.....	313
NTDB id	584	DLJ51	RS00890	WP	019777327.1	LDNCLAGTIAYQRLIGGGGLIDFAQDHFNHNHAECWNEKIDQLLAAGHTITEEAAAEEKIKD.....	315
NTDB id	71726	EL269	RS01120	WP	000250805.1	LENLSKLTIAYQRLIGGGSLIDFETGNFKKHSSDKWNRQVDILAEFGHISKKQAQVEKIIPQETTESSPTF.....	323
NTDB id	385	SMU	RS09035	WP	002263435.1	LQNALRGIVVYQRLIGGGGIDFAKGDYQNHSAKKWNKQIYDLFTAGYITEGEAQAEEIED.....	313
NTDB id	394	FSA28	RS09405	WP	002291652.1	LQNALCGIVVYQRLIGGGGIDFAKGDYQNHSAKKWNKQIYDLFTAGYITEGEAQAEEIED.....	313
consensus						! **!* **!! ! !! *! **!* ! **! **** * ! ! ! **!* ! * ! **!* ! *	

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