logo		
NTDB id 591 KW2 RS10830 WP 014735174.1	.........MKLKIRAFTLLECLVALLAISGSVLVISGLTRMIEEQMKISQNDSRKDWQIFCEQMRSELSGAKLDNVNQN	71
NTDB id 275 KZH43 RS09240 WP 000250534.1	MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKVEGN	80
NTDB id 234 SPD RS09870 WP 000250534.1	MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKVEGN	80
NTDB id 200 SPR RS09405 WP 000250534.1	MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKVEGN	80
NTDB id 165 SP RS10355 WP 000250534.1	MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKVEGN	80
NTDB id 505 SM12261 RS00970 WP 000250509.1	MVQNSCWLSKSHKIKAFTLLESLIALIVISGSLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEAELDRSQFEKVEGD	80
NTDB id 533 SMSK321 RS09215 WP 000250512.1	MVQNSCWLSKSHKVKAFTLLESLIALIVISGSLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEAELDRSQFEKIEGN	80
NTDB id 240574 CG712 RS05250 WP 022096897.1	........MYLSSKLRAFTLLECLVALLVIAGSVQVYQGLTTVLVSNVKQVRQQENQDWLLFVQQMEAELEGCQLVKVEGN	73
NTDB id 399 FSA28 RS09380 WP 002264245.1	........MLKSKLKAFTLLECLLALLVIAGSVSVYNGLTKAISSNIHYLSKNQEDAWLLFCQQFRSELEGTTLQKLDSN	72
NTDB id 390 SMU RS09010 WP 002263440.1	........MLKSKLKAFTLLECLLALLVIAGSVSVYNGLTKAISSNIHYLSKNQEDAWLLFCQQFRSELEGTTLQKLDSN	72
consensus	******* ****!**!!!!!!*!*!!**!*!**************************!**!**!** !!**********	

logo		
NTDB id 591 KW2 RS10830 WP 014735174.1	FLYVTKDK.KLRFGL.VGDDFRKSDDKGQGYQPMLYDLKGAKIQAEENLIKITIDFDNGGERVFIYRFTDTK.	141
NTDB id 275 KZH43 RS09240 WP 000250534.1	RLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS	153
NTDB id 234 SPD RS09870 WP 000250534.1	RLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS	153
NTDB id 200 SPR RS09405 WP 000250534.1	RLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS	153
NTDB id 165 SP RS10355 WP 000250534.1	RLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS	153
NTDB id 505 SM12261 RS00970 WP 000250509.1	RIYMKQDGKEIAIGKSKSDDFRKTDASGRGYQPMVYGLKSAQITEENQVVRFRFQFQKGLEREFIYRVEKAKS	153
NTDB id 533 SMSK321 RS09215 WP 000250512.1	RIYMKQDGKDISIGKLKSDDFRKTDASGRGYQPMVYGLKSAQITEDNQLVRFRFQFQKGLEREFIYRVEKAKS	153
NTDB id 240574 CG712 RS05250 WP 022096897.1	KLYVKKDNQDLAFGLSSASDFRKTNADGRGYQPMLFDVKSSKVSQKNQIVTIQVTLKNELQRSFVYAFEKTG.	145
NTDB id 399 FSA28 RS09380 WP 002264245.1	KLYVQKNNQSLAFGKSKASDFRKTNSDGRGYQPMLTEIKAANFSQSGKIIKLDLTFKDGLERTFVYAFHEKN.	144
NTDB id 390 SMU RS09010 WP 002263440.1	KLYIQKNNQSLAFGKSKASDFRKTNSDGRGYQPMLTEIKAANFSQSGKIIKLDLTFKDGLERTFVYAFHEKN.	144
consensus	**!**********!*****!!!!****!*!!!!!****!************* ********!*!*!**** **	

X non conserved

X similar

X $\geq$ 50% conserved