```
logo
NTDB id 591 KW2 RS10830 WP 014735174.1      .........MKLKIRAFTLLECLVALLAISGSVLVISGLTRMIEEQMKISQNDSRKDWQIFCEQMRSELSGAKLDNVNQNFLYVT   76
NTDB id 275 KZH43 RS09240 WP 000250534.1    MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKVEGNRLYMK   85
NTDB id 234 SPD RS09870 WP 000250534.1      MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKVEGNRLYMK   85
NTDB id 200 SPR RS09405 WP 000250534.1      MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKVEGNRLYMK   85
NTDB id 165 SP RS10355 WP 000250534.1       MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKVEGNRLYMK   85
NTDB id 505 SM12261 RS00970 WP 000250509.1 MVQNSCWLSKSHKIKAFTLLESLIALIVISGSLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEAELDRSQFEKVEGDRIYMK   85
NTDB id 533 SMSK321 RS09215 WP 000250512.1 MVQNSCWLSKSHKVKAFTLLESLIALIVISGSLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEAELDRSQFEKIEGNRIYMK   85
NTDB id 309 STER RS08975 WP 011681686.1    .......MYLSSKLRAFTLLECLVALLVIAGSVQVYQGLITVLVSNVKQVRQQENQDWLLFVQQMEAELEGCQLVKVEGNKLYVK   78
NTDB id 399 FSA28 RS09380 WP 002264245.1   ........MLKSKLKAFTLLECLLALLVIAGSVSVYNGLIKAISSNIHYLSKNQEDAWLLFCQQFRSELEGTTLQKLDSNKLYVQ   77
NTDB id 390 SMU RS09010 WP 002263440.1     ........MLKSKLKAFTLLECLLALLVIAGSVSVYNGLIKAISSNIHYLSKNQEDAWLLFCQQFRSELEGTTLQKLDSNKLYIQ   77
consensus                                  ******* ****!**!!!!!!*!*!!**!*!*****************************!**!**!** !!**************!**

logo
NTDB id 591 KW2 RS10830 WP 014735174.1      KDK.KLRFGL.VGDDFRKSDDKGQGYQPMLYDLKGAKIQAEENLIKITIDFDNGGERVFIYRFTDTK.   141
NTDB id 275 KZH43 RS09240 WP 000250534.1    QDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS   153
NTDB id 234 SPD RS09870 WP 000250534.1      QDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS   153
NTDB id 200 SPR RS09405 WP 000250534.1      QDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS   153
NTDB id 165 SP RS10355 WP 000250534.1       QDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS   153
NTDB id 505 SM12261 RS00970 WP 000250509.1 QDGKEIAIGKSKSDDFRKTDASGRGYQPMVYGLKSAQITEENQVVRFRFQFQKGLEREFIYRVEKAKS   153
NTDB id 533 SMSK321 RS09215 WP 000250512.1 QDGKDISIGKLKSDDFRKTDASGRGYQPMVYGLKSAQITEDNQLVRFRFQFQKGLEREFIYRVEKAKS   153
NTDB id 309 STER RS08975 WP 011681686.1    QDNQDLAFGLSSASDFRKTNADGRGYQPMLFDVKSSKVSQKNQIVTIQVTLKNELQRSFVYAFEKTG.   145
NTDB id 399 FSA28 RS09380 WP 002264245.1   KNNQSLAFGKSKASDFRKTNSDGRGYQPMLTEIKAANFSQSGKIIKLDLTFKDGLERTFVYAFHEKN.   144
NTDB id 390 SMU RS09010 WP 002263440.1     KNNQSLAFGKSKASDFRKTNSDGRGYQPMLTEIKAANFSQSGKIIKLDLTFKDGLERTFVYAFHEKN.   144
consensus                                  ********!*****!!!!****!*!!!!!****!************* ********!*!*!**** **
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X non conserved

X similar

X ≥ 50% conserved