```
logo
NTDB id 275 KZH43 RS09240 WP 000250534.1     .MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKVEGNRLYM   84
NTDB id 234 SPD RS09870 WP 000250534.1       .MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKVEGNRLYM   84
NTDB id 200 SPR RS09405 WP 000250534.1       .MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKVEGNRLYM   84
NTDB id 165 SP RS10355 WP 000250534.1        .MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKVEGNRLYM   84
NTDB id 505 SM12261 RS00970 WP 000250509.1  .MVQNSCWLSKSHKIKAFTLLESLIALIVISGSLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEAELDRSQFEKVEGDRIYM   84
NTDB id 533 SMSK321 RS09215 WP 000250512.1  .MVQNSCWLSKSHKVKAFTLLESLIALIVISGSLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEAELDRSQFEKIEGNRIYM   84
NTDB id 591 KW2 RS10830 WP 014735174.1      ..........MKLKIRAFTLLECLVALLAISGSVLVISGLTRMIEEQMKISQNDSRKDWQIFCEQMRSELSGAKLDNVNQNFLYV   75
NTDB id 64585 STRIC RS08545 WP 050897399.1 MKIKKKFLEWSKSKVTAFTLLECLLTLLIISGSLIVYQGLTKSLFANCHYLAKNDQDQWLLFAQQLRTELAESHLPKVVNNKLYV   85
NTDB id 399 FSA28 RS09380 WP 002264245.1    .........MLKSKLKAFTLLECLLALLVIAGSVSVYNGLTKAISSNIHYLSKNQEDAWLLFCQQFRSELEGTTLQKLDSNKLYV   76
NTDB id 390 SMU RS09010 WP 002263440.1      .........MLKSKLKAFTLLECLLALLVIAGSVSVYNGLTKAISSNIHYLSKNQEDAWLLFCQQFRSELEGTTLQKLDSNKLYI   76
consensus                                    ******* ****!**!!!!!!*!**!**!*!*************************!**!**!** !!**************!*

logo
NTDB id 275 KZH43 RS09240 WP 000250534.1     KQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS   153
NTDB id 234 SPD RS09870 WP 000250534.1       KQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS   153
NTDB id 200 SPR RS09405 WP 000250534.1       KQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS   153
NTDB id 165 SP RS10355 WP 000250534.1        KQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS   153
NTDB id 505 SM12261 RS00970 WP 000250509.1  KQDGKEIAIGKSKSDDFRKTDASGRGYQPMVYGLKSAQITEENQVVRFRFQFQKGLEREFIYRVEKAKS   153
NTDB id 533 SMSK321 RS09215 WP 000250512.1  KQDGKDISIGKLKSDDFRKTDASGRGYQPMVYGLKSAQITEDNQLVRFRFQFQKGLEREFIYRVEKAKS   153
NTDB id 591 KW2 RS10830 WP 014735174.1      TKDKK.LRFGL.VGDDFRKSDDKGQGYQPMLYDLKGAKIQAEENLIKITIDFDNGGERVFIYRFTDTK.   141
NTDB id 64585 STRIC RS08545 WP 050897399.1 EKGKKSLAFGQYKSHDFRKTSQTGRGYQPMLFHLTTSKLTLEGERLHIHFKWESGLERIFIYDFQEKS.   153
NTDB id 399 FSA28 RS09380 WP 002264245.1    QKNNQSLAFGKSKASDFRKTNSDGRGYQPMLTEIKAANFSQSGKIIKLDLTFKDGLERTFVYAFHEKN.   144
NTDB id 390 SMU RS09010 WP 002263440.1      QKNNQSLAFGKSKASDFRKTNSDGRGYQPMLTEIKAANFSQSGKIIKLDLTFKDGLERTFVYAFHEKN.   144
consensus                                   *********!*****!!!!****!*!!!!!*************************!*!!*!*!**** **
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X non conserved

X similar

X ≥ 50% conserved