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logo
NTDB id 591 KW2 RS10830 WP 014735174.1        ..........MKLKIRAFTLLECLVALLAISGSVLVISGLTRMIEEQMKISQNDSRKDWQIFCEQMRSELSGAKLDNVNQ   70
NTDB id 275 KZH43 RS09240 WP 000250534.1      .MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKVEG   79
NTDB id 234 SPD RS09870 WP 000250534.1        .MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKVEG   79
NTDB id 200 SPR RS09405 WP 000250534.1        .MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKVEG   79
NTDB id 165 SP RS10355 WP 000250534.1         .MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKVEG   79
NTDB id 505 SM12261 RS00970 WP 000250509.1    .MVQNSCWLSKSHKIKAFTLLESLIALIVISGSLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEAELDRSQFEKVEG   79
NTDB id 533 SMSK321 RS09215 WP 000250512.1    .MVQNSCWLSKSHKVKAFTLLESLIALIVISGSLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEAELDRSQFEKIEG   79
NTDB id 315455 D5R90 RS06490 WP 119773875.1   MKSIKKSLALRKISLKAFTLLECMLALVILSGAILVYQGMTKSLFFNMSYLVENDQDKWLLFSQQLRAELTGARFEKVEN   80
NTDB id 399 FSA28 RS09380 WP 002264245.1      .........MLKSKLKAFTLLECLLALLVIAGSVSVYNGLTKAISSNIHYLSKNQEDAWLLFCQQFRSELEGTTLQKLDS   71
NTDB id 390 SMU RS09010 WP 002263440.1        .........MLKSKLKAFTLLECLLALLVIAGSVSVYNGLTKAISSNIHYLSKNQEDAWLLFCQQFRSELEGTTLQKLDS   71
consensus                                      ******* *******!!!!!!**!!****!*************************!**!**!** !!**********

logo
NTDB id 591 KW2 RS10830 WP 014735174.1        NFLYVTKDK.KLRFGLV.GDDFRKSDDKGQGYQPMLYDLKGAKIQAEENLIKITIDFDNGGERVFIYRFTDTK.   141
NTDB id 275 KZH43 RS09240 WP 000250534.1      NRLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS   153
NTDB id 234 SPD RS09870 WP 000250534.1        NRLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS   153
NTDB id 200 SPR RS09405 WP 000250534.1        NRLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS   153
NTDB id 165 SP RS10355 WP 000250534.1         NRLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS   153
NTDB id 505 SM12261 RS00970 WP 000250509.1    DRIYMKQDGKEIAIGKSKSDDFRKTDASGRGYQPMVYGLKSAQITEENQVVRFRFQFQKGLEREFIYRVEKAKS   153
NTDB id 533 SMSK321 RS09215 WP 000250512.1    NRIYMKQDGKDISIGKLKSDDFRKTDASGRGYQPMVYGLKSAQITEDNQLVRFRFQFQKGLEREFIYRVEKAKS   153
NTDB id 315455 D5R90 RS06490 WP 119773875.1   NKLYVAKEGKQIAFGLSKQGDFRKTAANGHGYHPMLLDLKDAKISHHNKEITVSITWKSGLERVFLYAF.....   149
NTDB id 399 FSA28 RS09380 WP 002264245.1      NKLYVQKNNQSLAFGKSKASDFRKTNSDGRGYQPMLTEIKAANFSQSGKIIKLDLTFKDGLERTFVYAFHEKN.   144
NTDB id 390 SMU RS09010 WP 002263440.1        NKLYIQKNNQSLAFGKSKASDFRKTNSDGRGYQPMLTEIKAANFSQSGKIIKLDLTFKDGLERTFVYAFHEKN.   144
consensus                                     ***!**********!*****!!!!****!*!!*!!****!************* *****!*!!*!*!**** **
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X non conserved

X similar

X ≥ 50% conserved