logo

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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 533 SMSK321 RS09215 WP 000250512.1    MVQNSCWLSKSHKVKAFTLLESLIALIVISGSLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEAELDRSQFEKIEGN  80
NTDB id 505 SM12261 RS00970 WP 000250509.1    MVQNSCWLSKSHKIKAFTLLESLIALIVISGSLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEAELDRSQFEKVEGD  80
NTDB id 399 FSA28 RS09380 WP 002264245.1      ........MLKSKLKAFTLLECLLALLVIAGSVSVYNGLTKAISSNIHYLSKNQEDAWLLFCQQFRSELEGTTLQKLDSN  72
NTDB id 390 SMU RS09010 WP 002263440.1        ........MLKSKLKAFTLLECLLALLVIAGSVSVYNGLTKAISSNIHYLSKNQEDAWLLFCQQFRSELEGTTLQKLDSN  72
NTDB id 183239 LLUC063 RS11025 WP 058225890.1 MERKLC..DLNFKVKAFTLLECLVALLAISGSVLVISGLTKMLKEQVAISQSDSIKDWQIFCQQMRFELSGTKLDRVEQN  78
NTDB id 591 KW2 RS10830 WP 014735174.1        ........MKLKIRAFTLLECLVALLAISGSVLVISGLTRMIEEQMKISQNDSRKDWQIFCEQMRSELSGAKLDNVNQN  71
consensus                                     ******* ****!**!!!!!!*!*!!**!*!**************************!**!**!**  !!**********
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logo

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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 533 SMSK321 RS09215 WP 000250512.1    RIYMKQDGKDISIGKLKSDDFRKTDASGRGYQPMVYGLKSAQITEDNQLVRFRFQFQKGLEREFIYRVEKAKS  153
NTDB id 505 SM12261 RS00970 WP 000250509.1    RIYMKQDGKEIAIGKSKSDDFRKTDASGRGYQPMVYGLKSAQITEENQVVRFRFQFQKGLEREFIYRVEKAKS  153
NTDB id 399 FSA28 RS09380 WP 002264245.1      KLYVQKNNQSLAFGKSKASDFRKTNSDGRGYQPMLTEIKAANFSQSGKIIKLDLTFKDGLERTFVYAFHEKN.  144
NTDB id 390 SMU RS09010 WP 002263440.1        KLYIQKNNQSLAFGKSKASDFRKTNSDGRGYQPMLTEIKAANFSQSGKIIKLDLTFKDGLERTFVYAFHEKN.  144
NTDB id 183239 LLUC063 RS11025 WP 058225890.1 FLYVTKDK.KLRFGF.MSDDFRKTDDKGQSYQPMLYDIKAAKIQATKNLITIRIDFNQGGERTFIYQFPEDT.  148
NTDB id 591 KW2 RS10830 WP 014735174.1        FLYVTKDK.KLRFGL.VGDDFRKSDDKGQGYQPMLYDLKGAKIQAEENLIKITIDFDNGGERVFIYRFTDTK.  141
consensus                                     **!**********!*****!!!!****!**!!!!****!************** **!**!*!!*!*!*******
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X non conserved
X similar
X ≥ 50% conserved