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logo
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NTDB id 275 KZH43 RS09240 WP 000250534.1        ..MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKVE   78
NTDB id 234 SPD RS09870 WP 000250534.1          ..MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKVE   78
NTDB id 200 SPR RS09405 WP 000250534.1          ..MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKVE   78
NTDB id 165 SP RS10355 WP 000250534.1           ..MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKVE   78
NTDB id 505 SM12261 RS00970 WP 000250509.1      ..MVQNSCWLSKSHKIKAFTLLESLIALIVISGSLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEAELDRSQFEKVE   78
NTDB id 533 SMSK321 RS09215 WP 000250512.1      ..MVQNSCWLSKSHKVKAFTLLESLIALIVISGSLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEAELDRSQFEKIE   78
NTDB id 591 KW2 RS10830 WP 014735174.1          ...........MKLKIRAFTLLECLVALLAISGSVLVISGLTRMIEEQMKISQNDSRKDWQIFCEQMRSELSGAKLDNVN   69
NTDB id 1020957 ABZ559 RS11720 WP 367006524.1   MKMAELFCR.CQKIKLAAFSLLECLVALFVLSGSMLVYQGLTKSVSANVHYLSDNDNGNWLLFSQQLRSELETSHLDKVE   79
NTDB id 399 FSA28 RS09380 WP 002264245.1        ..........MLKSKLKAFTLLECLLALLVIAGSVSVYNGLTKAISSNIHYLSKNQEDAWLLFCQQFRSELEGTTLQKLD   70
NTDB id 390 SMU RS09010 WP 002263440.1          ..........MLKSKLKAFTLLECLLALLVIAGSVSVYNGLTKAISSNIHYLSKNQEDAWLLFCQQFRSELEGTTLQKLD   70
consensus                                         ******* ****!**!!*!!!*!*!!****!**********************!**!**!** !!*********
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logo
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NTDB id 275 KZH43 RS09240 WP 000250534.1        GNRLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS   153
NTDB id 234 SPD RS09870 WP 000250534.1          GNRLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS   153
NTDB id 200 SPR RS09405 WP 000250534.1          GNRLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS   153
NTDB id 165 SP RS10355 WP 000250534.1           GNRLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS   153
NTDB id 505 SM12261 RS00970 WP 000250509.1      GDRIYMKQDGKEIAIGKSKSDDFRKTDASGRGYQPMVYGLKSAQITEENQVVRFRFQFQKGLEREFIYRVEKAKS   153
NTDB id 533 SMSK321 RS09215 WP 000250512.1      GNRIYMKQDGKDISIGKLKSDDFRKTDASGRGYQPMVYGLKSAQITEDNQLVRFRFQFQKGLEREFIYRVEKAKS   153
NTDB id 591 KW2 RS10830 WP 014735174.1          QNFLYVTKDK.KLRFGL.VGDDFRKSDDKGQGYQPMLYDLKGAKIQAEENLIKITIDFDNGGERVFIYRFTDTK.   141
NTDB id 1020957 ABZ559 RS11720 WP 367006524.1   NNKLYVTKSGQTLAFGQSKADDFRKTNADGRGYQPMLYGLKSSAISQKENMIQINLTFKNGLERSLIYAFEEAV.   153
NTDB id 399 FSA28 RS09380 WP 002264245.1        SNKLYVQKNNQSLAFGKSKASDFRKTNSDGRGYQPMLTEIKAANFSQSGKIIKLDLTFKDGLERTFVYAFHEKN.   144
NTDB id 390 SMU RS09010 WP 002263440.1          SNKLYIQKNNQSLAFGKSKASDFRKTNSDGRGYQPMLTEIKAANFSQSGKIIKLDLTFKDGLERTFVYAFHEKN.   144
consensus                                       ****!**********!*****!!!!****!*!!!!!****!************* **!**!*!!***!**** **
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X non conserved
X similar
X ≥ 50% conserved