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logo
NTDB id 591 KW2 RS10830 WP 014735174.1          ............MKLKIRAFTLLECLVALLAISGSVLVISGLTRMIEEQMKISQNDSRKDWQIFCEQMRSELSGAKLDNV   68
NTDB id 275 KZH43 RS09240 WP 000250534.1        ...MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKV   77
NTDB id 234 SPD RS09870 WP 000250534.1          ...MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKV   77
NTDB id 200 SPR RS09405 WP 000250534.1          ...MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKV   77
NTDB id 165 SP RS10355 WP 000250534.1           ...MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFEKV   77
NTDB id 505 SM12261 RS00970 WP 000250509.1      ...MVQNSCWLSKSHKIKAFTLLESLIALIVISGSLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEAELDRSQFEKV   77
NTDB id 533 SMSK321 RS09215 WP 000250512.1      ...MVQNSCWLSKSHKVKAFTLLESLIALIVISGSLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEAELDRSQFEKI   77
NTDB id 109589 ACGHT2 RS00765 WP 058692632.1   MFMRGKRKYYTLKKTSLKAFTLIECLVSLLIISGAILVYNGLTQSISANVHYLSENQEENWLLFSQQLRAELANCQLDKV   80
NTDB id 399 FSA28 RS09380 WP 002264245.1        ...........MLKSKLKAFTLLECLLALLVIAGSVSVYNGLTKAISSNIHYLSKNQEDAWLLFCQQFRSELEGTTLQKL   69
NTDB id 390 SMU RS09010 WP 002263440.1          ...........MLKSKLKAFTLLECLLALLVIAGSVSVYNGLTKAISSNIHYLSKNQEDAWLLFCQQFRSELEGTTLQKL   69
consensus                                          ******* *******!!!!*!*!**!**!*!*************************!**!**!** !!********
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logo
NTDB id 591 KW2 RS10830 WP 014735174.1          NQNFLYVTKDK.KLRFGL.VGDDFRKSDDKGQGYQPMLYDLKGAKIQAEENLIKITIDFDNGGERVFIYRFTDTK.   141
NTDB id 275 KZH43 RS09240 WP 000250534.1        EGNRLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS   153
NTDB id 234 SPD RS09870 WP 000250534.1          EGNRLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS   153
NTDB id 200 SPR RS09405 WP 000250534.1          EGNRLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS   153
NTDB id 165 SP RS10355 WP 000250534.1           EGNRLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEKEKS   153
NTDB id 505 SM12261 RS00970 WP 000250509.1      EGDRIYMKQDGKEIAIGKSKSDDFRKTDASGRGYQPMVYGLKSAQITEENQVVRFRFQFQKGLEREFIYRVEKAKS   153
NTDB id 533 SMSK321 RS09215 WP 000250512.1      EGNRIYMKQDGKDISIGKLKSDDFRKTDASGRGYQPMVYGLKSAQITEDNQLVRFRFQFQKGLEREFIYRVEKAKS   153
NTDB id 109589 ACGHT2 RS00765 WP 058692632.1   ENNKLYVTKSSQKLAFGQSKADDFRKTNASGQGYQPMIFGVKSSAISRDGQKVTMTLNLENGLERTFVYTFETAS.   155
NTDB id 399 FSA28 RS09380 WP 002264245.1        DSNKLYVQKNNQSLAFGKSKASDFRKTNSDGRGYQPMLTEIKAANFSQSGKIIKLDLTFKDGLERTFVYAFHEKN.   144
NTDB id 390 SMU RS09010 WP 002263440.1          DSNKLYIQKNNQSLAFGKSKASDFRKTNSDGRGYQPMLTEIKAANFSQSGKIIKLDLTFKDGLERTFVYAFHEKN.   144
consensus                                       *****!**********!*****!!!!****!*!!!!!****!************* *****!*!!*!*!**** **
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X non conserved

X similar

X ≥ 50% conserved