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logo

NTDB id 591 KW2 RS10830 WP 014735174.1          .........MKLKIRAFTLLECLVALLAISGSVLVISGLTRMIEEQMKISQNDSRKDWQIFCEQMRSELSGAKLD   66
NTDB id 275 KZH43 RS09240 WP 000250534.1        MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFE   75
NTDB id 234 SPD RS09870 WP 000250534.1          MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFE   75
NTDB id 200 SPR RS09405 WP 000250534.1          MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFE   75
NTDB id 165 SP RS10355 WP 000250534.1           MVQNSCWQSKSHKVKAFTLLESLLALIVISGGLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEVELDRSQFE   75
NTDB id 505 SM12261 RS00970 WP 000250509.1      MVQNSCWLSKSHKIKAFTLLESLIALIVISGSLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEAELDRSQFE   75
NTDB id 533 SMSK321 RS09215 WP 000250512.1      MVQNSCWLSKSHKVKAFTLLESLIALIVISGSLLLFQAMSQLLISEVRYQQQSEQKEWLLFVDQLEAELDRSQFE   75
NTDB id 1111722 HMPREF9425 RS00700 WP 003091908.1 .......MSLNSKVRAFTLLECLVALLVIAGSVQVYQGLTTVLVGNVKQVRQQENQDWLLFVQQMEAELEGCQLV   68
NTDB id 399 FSA28 RS09380 WP 002264245.1        ........MLKSKLKAFTLLECLLALLVIAGSVSVYNGLTKAISSNIHYLSKNQEDAWLLFCQQFRSELEGTTLQ   67
NTDB id 390 SMU RS09010 WP 002263440.1          ........MLKSKLKAFTLLECLLALLVIAGSVSVYNGLTKAISSNIHYLSKNQEDAWLLFCQQFRSELEGTTLQ   67
consensus                                       ******* ****!**!!!!!!*!*!!**!*!*************************!**!**!** !!******

logo

NTDB id 591 KW2 RS10830 WP 014735174.1          NVNQNFLYVTKDK.KLRFGL.VGDDFRKSDDKGQGYQPMLYDLKGAKIQAEENLIKITIDFDNGGERVFIYRFTD   139
NTDB id 275 KZH43 RS09240 WP 000250534.1        KVEGNRLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEK   150
NTDB id 234 SPD RS09870 WP 000250534.1          KVEGNRLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEK   150
NTDB id 200 SPR RS09405 WP 000250534.1          KVEGNRLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEK   150
NTDB id 165 SP RS10355 WP 000250534.1           KVEGNRLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVYGLKSVRITEDNQLVRFHFQFQKGLEREFIYRVEK   150
NTDB id 505 SM12261 RS00970 WP 000250509.1      KVEGDRIYMKQDGKEIAIGKSKSDDFRKTDASGRGYQPMVYGLKSAQITEENQVVRFRFQFQKGLEREFIYRVEK   150
NTDB id 533 SMSK321 RS09215 WP 000250512.1      KIEGNRIYMKQDGKDISIGKLKSDDFRKTDASGRGYQPMVYGLKSAQITEDNQLVRFRFQFQKGLEREFIYRVEK   150
NTDB id 1111722 HMPREF9425 RS00700 WP 003091908.1 KVEGNKLYVKQDNQDLAFGLSSATDFRKTNADGRGYQPMLFDVKSSTVSQKNQIVTIQVTLKNGLQRSFIYAFEK   143
NTDB id 399 FSA28 RS09380 WP 002264245.1        KLDSNKLYVQKNNQSLAFGKSKASDFRKTNSDGRGYQPMLTEIKAANFSQSGKIIKLDLTFKDGLERTFVYAFHE   142
NTDB id 390 SMU RS09010 WP 002263440.1          KLDSNKLYIQKNNQSLAFGKSKASDFRKTNSDGRGYQPMLTEIKAANFSQSGKIIKLDLTFKDGLERTFVYAFHE   142
consensus                                       *******!**********!*****!!!!****!*!!!!!****!************* *****!**!*!*!****

logo

NTDB id 591 KW2 RS10830 WP 014735174.1          TK.   141
NTDB id 275 KZH43 RS09240 WP 000250534.1        EKS   153
NTDB id 234 SPD RS09870 WP 000250534.1          EKS   153
NTDB id 200 SPR RS09405 WP 000250534.1          EKS   153
NTDB id 165 SP RS10355 WP 000250534.1           EKS   153
NTDB id 505 SM12261 RS00970 WP 000250509.1      AKS   153
NTDB id 533 SMSK321 RS09215 WP 000250512.1      AKS   153
NTDB id 1111722 HMPREF9425 RS00700 WP 003091908.1 TG.   145
NTDB id 399 FSA28 RS09380 WP 002264245.1        KN.   144
NTDB id 390 SMU RS09010 WP 002263440.1          KN.   144
consensus                                        **
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X non conserved

X similar

X $\geq$ 50% conserved