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NTDB id 275 KZH43 RS09240 WP 000250534.1	MVQNSCWQSKSHKVKAF T LL E SL L ALIVISG G LLL F QAMSQ L LISEVRYQQQSE Q KEW L LFVDQ L EV	E L D R S Q F E K V E G N	80
NTDB id 234 SPD RS09870 WP 000250534.1	MVQNSCWQSKSHKVKAF T LL E SL L ALIVISG G LLL F QAMSQ L LISEVRYQQQSE Q KEW L LFVDQ L EV	E L D R S Q F E K V E G N	80
NTDB id 200 SPR RS09405 WP 000250534.1	MVQNSCWQSKSHKVKAF T LL E SL L ALIVISG G LLL F QAMSQ L LISEVRYQQQSE Q KEW L LFVDQ L EV	E L D R S Q F E K V E G N	80
NTDB id 165 SP RS10355 WP 000250534.1	MVQNSCWQSKSHKVKAF T LL E SL L ALIVISG G LLL F QAMSQ L LISEVRYQQQSE Q KEW L LFVDQ L EV	E L D R S Q F E K V E G N	80
NTDB id 505 SM12261 RS00970 WP 000250509.1	MVQNSCWL S KSHK V KAFTLL E SL I ALIVISG S LLL F QAMSQ L LISEVRYQQQSE Q KEW L LFVDQ L EA	E L D R S Q F E K V E G D	80
NTDB id 533 SMSK321 RS09215 WP 000250512.1	MVQNSCWL S KSHK V KAFTLL E SL I ALIVISG S LLL F QAMSQ L LISEVRYQQQSE Q KEW L LFVDQ L EA	E L D R S Q F E K I E G N	80
NTDB id 591 KW2 RS10830 WP 014735174.1	MKL K I R AFTLL E C L V A L L AISG S V L VISGL T R M IEE Q M K IS N DSR K D W Q I F C E Q M R S	E L S G A K L D N V N Q N	71
NTDB id 977413 MGAS37847 RS00685 WP 010921800.1	MSK Q LSN I KAFTLL E A L I A L L VISG S LL V YQGL T R T L L IKH S H Y LARH D QDN W LL F SH Q L R E	E L S G A R F Y K V ADN	74
NTDB id 399 FSA28 RS09380 WP 002264245.1	MLK S K L KAFTLL E C L L A L L V I AG S V S VYNG L T K A I SS N I H YLSK N QEDA W LL F C Q Q F RS	E L E G I T L Q K L DSN	72
NTDB id 390 SMU RS09010 WP 002263440.1	MLK S K L KAFTLL E C L L A L L V I AG S V S VYNG L T K A I SS N I H YLSK N QEDA W LL F C Q Q F RS	E L E G I T L Q K L DSN	72
consensus	*****	*****	

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NTDB id 275 KZH43 RS09240 WP 000250534.1	RLYMKQDQGDIAIGKSKSDDFRKTNA	RGRGYQPMVYGLKS	VRITEDNQLVRFHFQFQKGLEREFYIRVEKEKS	153
NTDB id 234 SPD RS09870 WP 000250534.1	RLYMKQDQGDIAIGKSKSDDFRKTNA	RGRGYQPMVYGLKS	VRITEDNQLVRFHFQFQKGLEREFYIRVEKEKS	153
NTDB id 200 SPR RS09405 WP 000250534.1	RLYMKQDQGDIAIGKSKSDDFRKTNA	RGRGYQPMVYGLKS	VRITEDNQLVRFHFQFQKGLEREFYIRVEKEKS	153
NTDB id 165 SP RS10355 WP 000250534.1	RLYMKQDQGDIAIGKSKSDDFRKTNA	RGRGYQPMVYGLKS	VRITEDNQLVRFHFQFQKGLEREFYIRVEKEKS	153
NTDB id 505 SM12261 RS00970 WP 000250509.1	RIYMKQDQGEIAIGKSKSDDFRKTDA	RGRGYQPMVYGLKSAQ	ITEENQVVRFRFQFQKGLEREFYIRVEKA	153
NTDB id 533 SMSK321 RS09215 WP 000250512.1	RIYMKQDQGDISIGKLSKSDDFRKTDA	RGRGYQPMVYGLKSAQ	ITEDNQLVRFHFQFQKGLEREFYIRVEKA	153
NTDB id 591 KW2 RS10830 WP 014735174.1	FLYVTKDKK.LRFGL.VGDDFRKSDDK	QGQYQPMLYDLKGAK	IQAEENLIKITIDFDNGGERVFIYRFTDTK.	141
NTDB id 977413 MGAS37847 RS00685 WP 010921800.1	KLYVEKGKVLAFGQFKSHDFRKSA	NGKGYQPMLEGISRSHIHIEQSQICITLW	KSGLERTFVYAFQD...	144
NTDB id 399 FSA28 RS09380 WP 002264245.1	KLYVQKNNQSLAFGKSKASDFRKTNS	DGRGYQPMLEIKAAANFSQSGKI	IKLDLTFKDLGLERTFVYAFHEKN.	144
NTDB id 390 SMU RS09010 WP 002263440.1	KLYIQKNNQSLAFGKSKASDFRKTNS	DGRGYQPMLEIKAAANFSQSGKI	IKLDLTFKDLGLERTFVYAFHEKN.	144
consensus	*****	*****	*****	

- X non conserved
- X similar
- X $\geq 50\%$ conserved