

logo

NTDB id 345405 EXW31 RS05930 WP 215572532.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



639
634

logo

NTDB id 345405 EXW31 RS05930 WP 215572532.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



719
712

logo

NTDB id 345405 EXW31 RS05930 WP 215572532.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



799
792

logo

NTDB id 345405 EXW31 RS05930 WP 215572532.1
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consensus



879
872

logo

NTDB id 345405 EXW31 RS05930 WP 215572532.1
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consensus



959
949

logo

NTDB id 345405 EXW31 RS05930 WP 215572532.1
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1036
1029

logo

NTDB id 345405 EXW31 RS05930 WP 215572532.1
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1115
1109

logo

NTDB id 345405 EXW31 RS05930 WP 215572532.1

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[illegible]

logo

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consensus

Sequence logo for the 1241 and 1232 motifs. The logo shows the conservation of amino acids at each position. The top sequence (1241) is AKPTLKKRVKVLQSLYTKAVEQSLEKTPVKGKCLYFFDGNHWTL. The bottom sequence (1232) is AKPTLEDTRYKVLQSLYAKALEKSLKHPVKEKCLYFFDGNHVNIEE. The sequences are color-coded by amino acid type: A (green), K (blue), P (orange), T (red), L (purple), K (blue), R (red), V (green), K (blue), V (green), L (purple), Q (blue), S (green), L (purple), Y (red), T (red), K (blue), A (green), V (green), E (red), Q (blue), S (green), L (purple), K (blue), P (orange), V (green), K (blue), G (green), K (blue), C (blue), L (purple), Y (red), F (red), F (red), D (red), G (green), N (blue), H (blue), W (green), T (red), L (purple).

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