

logo

NTDB id 1105948 V3C22 RS05395 WP 226706963.1

NTDB id 372 SMU RS05280 WP 002262198.1

NTDB id 413 AAK55817.1 141..845()

NTDB id 374 SMU RS06885 WP 002262930.1

consensus

Sequence logo for the first region. The y-axis represents information content, with values 80, 80, 75, and 75. The consensus sequence is shown at the bottom with color-coded markers: blue for highly conserved, red for similar, and green for non-conserved.

logo

NTDB id 1105948 V3C22 RS05395 WP 226706963.1

NTDB id 372 SMU RS05280 WP 002262198.1

NTDB id 413 AAK55817.1 141..845()

NTDB id 374 SMU RS06885 WP 002262930.1

consensus

Sequence logo for the second region. The y-axis represents information content, with values 150, 150, 154, and 155. The consensus sequence is shown at the bottom with color-coded markers: blue for highly conserved, red for similar, and green for non-conserved.

logo

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NTDB id 372 SMU RS05280 WP 002262198.1

NTDB id 413 AAK55817.1 141..845()

NTDB id 374 SMU RS06885 WP 002262930.1

consensus

Sequence logo for the third region. The y-axis represents information content, with values 229, 229, 234, and 235. The consensus sequence is shown at the bottom with color-coded markers: blue for highly conserved, red for similar, and green for non-conserved.

<0

logo

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NTDB id 372 SMU RS05280 WP 002262198.1

NTDB id 413 AAK55817.1 141..845()

NTDB id 374 SMU RS06885 WP 002262930.1

consensus

Sequence logo for the fourth region. The y-axis represents information content, with values 229, 229, 234, and 235. The consensus sequence is shown at the bottom with color-coded markers: blue for highly conserved, red for similar, and green for non-conserved.

- non conserved
- similar
- ≥ 50% conserved