

logo

NTDB id 311090 D1F63 RS01405 WP 021299351.1

NTDB id 475 HSISS4 RS07005 WP 021143809.1

NTDB id 324 STU RS16140 WP 011226306.1

NTDB id 292 STER RS06940 WP 011681419.1

consensus

Sequence logo for position 1-100. The logo shows the conservation of amino acids across four sequences. The consensus sequence is indicated by exclamation marks at the bottom.

80

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logo

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NTDB id 324 STU RS16140 WP 011226306.1

NTDB id 292 STER RS06940 WP 011681419.1

consensus

Sequence logo for position 101-200. The logo shows the conservation of amino acids across four sequences. The consensus sequence is indicated by exclamation marks at the bottom.

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logo

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NTDB id 475 HSISS4 RS07005 WP 021143809.1

NTDB id 324 STU RS16140 WP 011226306.1

NTDB id 292 STER RS06940 WP 011681419.1

consensus

Sequence logo for position 201-300. The logo shows the conservation of amino acids across four sequences. The consensus sequence is indicated by exclamation marks at the bottom.

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logo

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NTDB id 324 STU RS16140 WP 011226306.1

NTDB id 292 STER RS06940 WP 011681419.1

consensus

Sequence logo for position 301-400. The logo shows the conservation of amino acids across four sequences. The consensus sequence is indicated by exclamation marks at the bottom.

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logo

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consensus

Sequence logo for position 401-500. The logo shows the conservation of amino acids across four sequences. The consensus sequence is indicated by exclamation marks at the bottom.

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logo

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NTDB id 292 STER RS06940 WP 011681419.1

consensus

Sequence logo for position 501-600. The logo shows the conservation of amino acids across four sequences. The consensus sequence is indicated by exclamation marks at the bottom.

477

476

476

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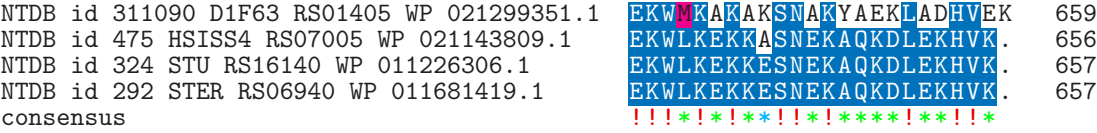
logo



logo



logo



X non conserved
X similar
X ≥ 50% conserved