

logo

NTDB id 342337 ETT47 RS01420 WP 063629325.1

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. The logo shows the conservation of amino acids across four sequences. The consensus sequence is indicated by exclamation marks (!) below the logo.

80

76

76

76

logo

NTDB id 342337 ETT47 RS01420 WP 063629325.1

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 2. The logo shows the conservation of amino acids across four sequences. The consensus sequence is indicated by exclamation marks (!) below the logo.

159

156

156

156

logo

NTDB id 342337 ETT47 RS01420 WP 063629325.1

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 3. The logo shows the conservation of amino acids across four sequences. The consensus sequence is indicated by exclamation marks (!) below the logo.

238

236

236

236

logo

NTDB id 342337 ETT47 RS01420 WP 063629325.1

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 4. The logo shows the conservation of amino acids across four sequences. The consensus sequence is indicated by exclamation marks (!) below the logo.

318

316

316

316

logo

NTDB id 342337 ETT47 RS01420 WP 063629325.1

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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 5. The logo shows the conservation of amino acids across four sequences. The consensus sequence is indicated by exclamation marks (!) below the logo.

397

396

396

396

logo

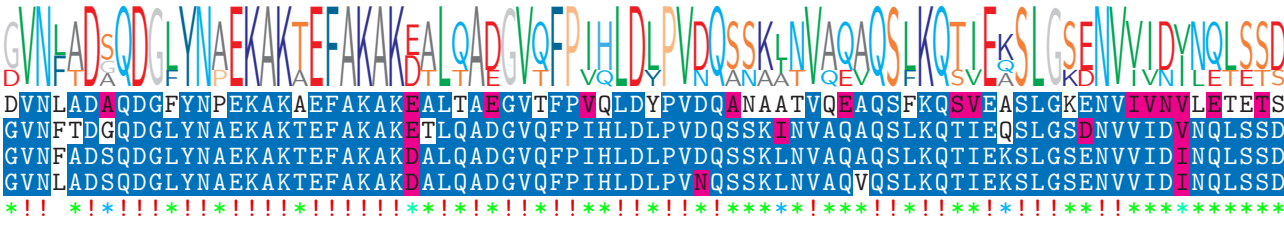
NTDB id 342337 ETT47 RS01420 WP 063629325.1

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 6. The logo shows the conservation of amino acids across four sequences. The consensus sequence is indicated by exclamation marks (!) below the logo.

477

476

476

476

logo

NTDB id 342337 ETT47 RS01420 WP 063629325.1

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

556
555
555
555

logo

NTDB id 342337 ETT47 RS01420 WP 063629325.1

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

635
633
634
634

logo

NTDB id 342337 ETT47 RS01420 WP 063629325.1

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

659
656
657
657

- non conserved
- similar
- ≥ 50% conserved