

logo

NTDB id 336988 EQH41 RS08860 WP 000742235.1

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 asterisks below the logo.

80

77

77

77

logo

NTDB id 336988 EQH41 RS08860 WP 000742235.1

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 101-200. The logo shows the conservation of amino acids across four sequences. The consensus sequence is indicated by asterisks below the logo.

160

157

157

157

logo

NTDB id 336988 EQH41 RS08860 WP 000742235.1

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 201-300. The logo shows the conservation of amino acids across four sequences. The consensus sequence is indicated by asterisks below the logo.

240

237

237

237

logo

NTDB id 336988 EQH41 RS08860 WP 000742235.1

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 301-400. The logo shows the conservation of amino acids across four sequences. The consensus sequence is indicated by asterisks below the logo.

320

317

317

317

logo

NTDB id 336988 EQH41 RS08860 WP 000742235.1

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 401-500. The logo shows the conservation of amino acids across four sequences. The consensus sequence is indicated by asterisks below the logo.

400

397

397

397

logo

NTDB id 336988 EQH41 RS08860 WP 000742235.1

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 501-600. The logo shows the conservation of amino acids across four sequences. The consensus sequence is indicated by asterisks below the logo.

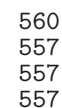
480

477

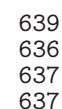
477

477

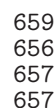
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consensus



-  non conserved
-  similar
-  > 50% conserved