

logo

NTDB id 336799 EQH43 RS02030 WP 000842571.1

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 asterisks below the logo.

80

78

78

78

logo

NTDB id 336799 EQH43 RS02030 WP 000842571.1

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 asterisks below the logo.

160

158

158

158

logo

NTDB id 336799 EQH43 RS02030 WP 000842571.1

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 asterisks below the logo.

240

238

238

238

logo

NTDB id 336799 EQH43 RS02030 WP 000842571.1

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 asterisks below the logo.

320

318

318

318

logo

NTDB id 336799 EQH43 RS02030 WP 000842571.1

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

400

398

398

398

logo

NTDB id 336799 EQH43 RS02030 WP 000842571.1

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 asterisks below the logo.

480

478

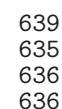
478

478

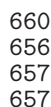
NTDB id 336799 EQH43 RS02030 WP 000842571.1
NTDB id 475 HSISS4 RS07005 WP 021143809.1
NTDB id 324 STU RS16140 WP 011226306.1
NTDB id 292 STER RS06940 WP 011681419.1
consensus



NTDB id 336799 EQH43 RS02030 WP 000842571.1
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



NTDB id 336799 EQH43 RS02030 WP 000842571.1
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



-  non conserved
-  similar
-  > 50% conserved