

logo

NTDB id 336951 EQH41 RS01710 WP 000842584.1

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 relative frequency of amino acids at this position across four sequences. The x-axis represents the amino acid, and the y-axis represents the log-odds ratio. The consensus sequence is indicated by the height of the bars.

80

78

78

78

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 2. The logo shows the relative frequency of amino acids at this position across four sequences. The x-axis represents the amino acid, and the y-axis represents the log-odds ratio. The consensus sequence is indicated by the height of the bars.

160

158

158

158

logo

NTDB id 336951 EQH41 RS01710 WP 000842584.1

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 relative frequency of amino acids at this position across four sequences. The x-axis represents the amino acid, and the y-axis represents the log-odds ratio. The consensus sequence is indicated by the height of the bars.

240

238

238

238

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 4. The logo shows the relative frequency of amino acids at this position across four sequences. The x-axis represents the amino acid, and the y-axis represents the log-odds ratio. The consensus sequence is indicated by the height of the bars.

320

318

318

318

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 5. The logo shows the relative frequency of amino acids at this position across four sequences. The x-axis represents the amino acid, and the y-axis represents the log-odds ratio. The consensus sequence is indicated by the height of the bars.

400

398

398

398

logo

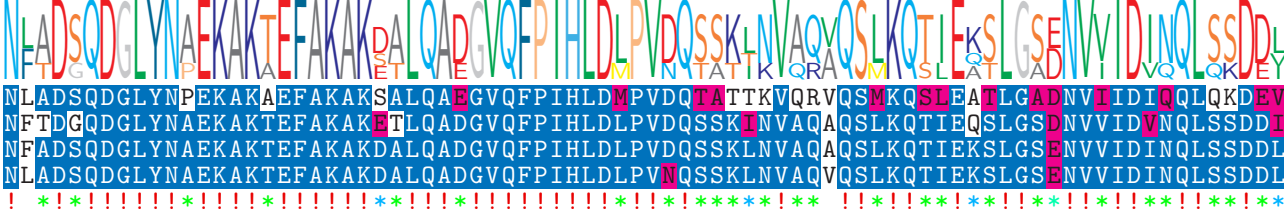
NTDB id 336951 EQH41 RS01710 WP 000842584.1

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 relative frequency of amino acids at this position across four sequences. The x-axis represents the amino acid, and the y-axis represents the log-odds ratio. The consensus sequence is indicated by the height of the bars.

480

478

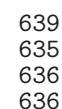
478

478

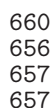
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NTDB id 336951 EQH41 RS01710 WP 000842584.1
NTDB id 475 HSISS4 RS07005 WP 021143809.1
NTDB id 324 STU RS16140 WP 011226306.1
NTDB id 292 STER RS06940 WP 011681419.1
consensus
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NTDB id 336951 EQH41 RS01710 WP 000842584.1
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
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NTDB id 336951 EQH41 RS01710 WP 000842584.1
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
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-  non conserved
-  similar
-  > 50% conserved