

logo

NTDB id 1012276 ABU975 RS06260 WP 002387037.1
NTDB id 352 BSU 24240 NP 390304.2
consensus



80
80

logo

NTDB id 1012276 ABU975 RS06260 WP 002387037.1
NTDB id 352 BSU 24240 NP 390304.2
consensus



160
160

logo

NTDB id 1012276 ABU975 RS06260 WP 002387037.1
NTDB id 352 BSU 24240 NP 390304.2
consensus



240
240

logo

NTDB id 1012276 ABU975 RS06260 WP 002387037.1
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consensus



320
319

logo

NTDB id 1012276 ABU975 RS06260 WP 002387037.1
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consensus



400
399

logo

NTDB id 1012276 ABU975 RS06260 WP 002387037.1
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consensus



465
479

logo

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545
559

logo

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consensus

Sequence logo for the 10th position of the 12-mer motif. The logo shows the conservation of amino acids at each position. The top row shows the most conserved residues: R, H, A, Q, E, L, L, A, M, A, K, K. The bottom row shows the least conserved residues: R, H, A, K, E, L, L, K, Q, A, D, Q. The positions are numbered 1 to 12 below the sequences. The 10th position is highlighted with a red background and a red exclamation mark below it, indicating high conservation.

557
576

- non conserved
- similar
- $\geq 50\%$ conserved