

logo

NTDB id 56279 H175 RS05780 WP 000058585.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



85
85

logo

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consensus



170
169

logo

NTDB id 56279 H175 RS05780 WP 000058585.1
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consensus



255
254

logo

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consensus



339
339

logo

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424
424

logo

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509
508

logo

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consensus



594
593

[illegible]

764
760

[illegible][illegible]

Sequence logo showing amino acid conservation across two sequences. The top sequence is ENIGKDKDITEGVITDIAPYKMGNKAACCTFCNFKSVCQFDESLEDNQFRSLKDMKDSEANEKTRREEVVGGE... (1171) and the bottom sequence is QEAGEQITDGRVSIIEPYKMKNKTPTCTYCFAKFSVCQFDESLENEVRFLKABKDKTILEWIKKEADGNEHS (1166). The logo highlights conserved residues with color-coded bars above each position.

 non conserved
 similar
 $\geq 50\%$ conserved