

logo

NTDB id 624 LCA RS00040 WP 011373726.1

NTDB id 113 BSU 40900 NP 391970.1

NTDB id 597 KW2 RS01935 WP 021036727.1

NTDB id 498 SM12261 RS01980 WP 000282464.1

NTDB id 526 SMSK321 RS08825 WP 000282440.1

NTDB id 59034 QMM27 RS08470 WP 000282467.1

NTDB id 268 KZH43 RS08520 WP 000282467.1

NTDB id 227 SPD RS09110 WP 000282467.1

NTDB id 193 SPR RS08685 WP 000282467.1

NTDB id 158 SP RS09595 WP 000282442.1

NTDB id 409 SMU RS08940 WP 002263454.1

NTDB id 583 DLJ51 RS09560 WP 002962590.1

consensus

Sequence logo for the first block showing amino acid conservation across 12 sequences. The x-axis represents positions from 1 to 100. The y-axis represents the log-odds of an amino acid being at that position. The logo shows high conservation at the N-terminus and C-terminus, with a more variable middle section.

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NTDB id 583 DLJ51 RS09560 WP 002962590.1

consensus

Sequence logo for the second block showing amino acid conservation across 12 sequences. The x-axis represents positions from 1 to 100. The y-axis represents the log-odds of an amino acid being at that position. The logo shows high conservation at the N-terminus and C-terminus, with a more variable middle section.

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165

122

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NTDB id 158 SP RS09595 WP 000282442.1

NTDB id 409 SMU RS08940 WP 002263454.1

NTDB id 583 DLJ51 RS09560 WP 002962590.1

consensus

Sequence logo for the third block showing amino acid conservation across 12 sequences. The x-axis represents positions from 1 to 100. The y-axis represents the log-odds of an amino acid being at that position. The logo shows high conservation at the N-terminus and C-terminus, with a more variable middle section.

170

172

129

131

131

131

131

131

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131

131

non conserved

similar

≥ 50% conserved