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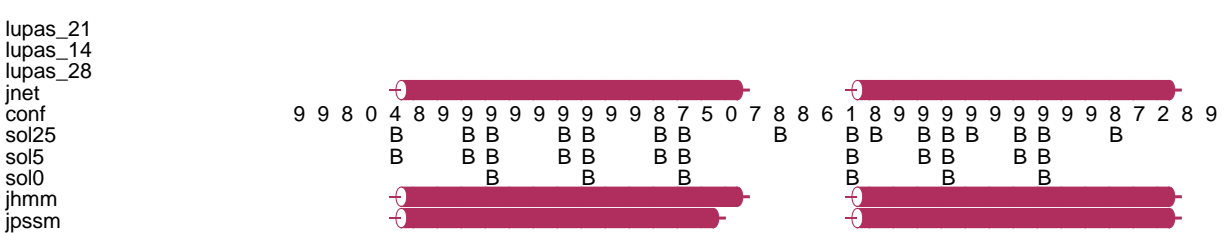
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Multiple sequence alignment

Position markers: 1 ... 10 ... 20 ... 30

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