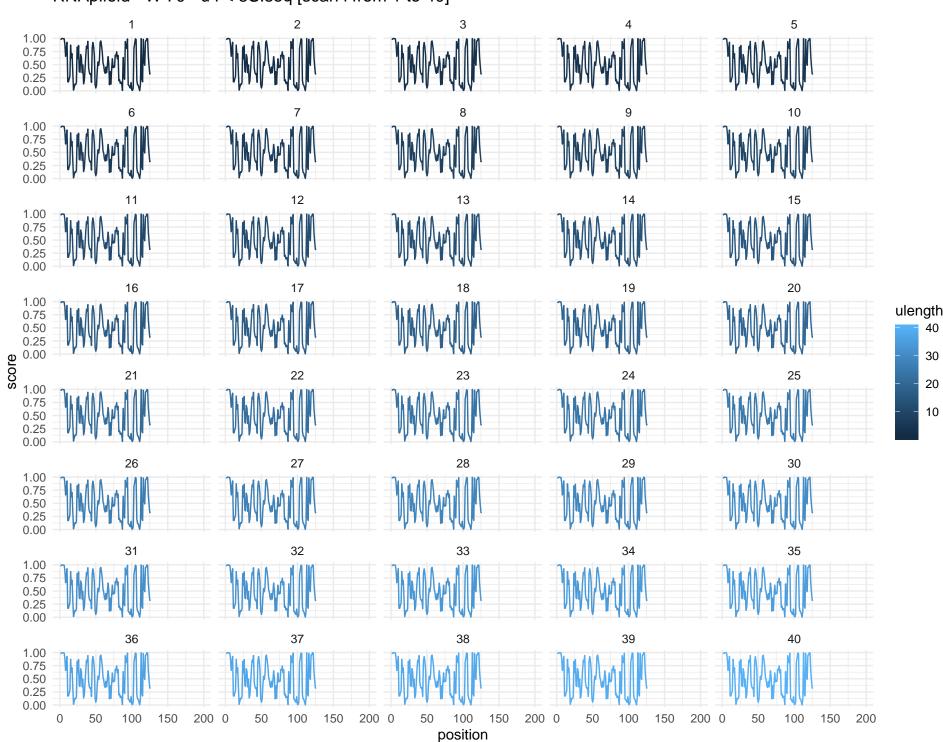
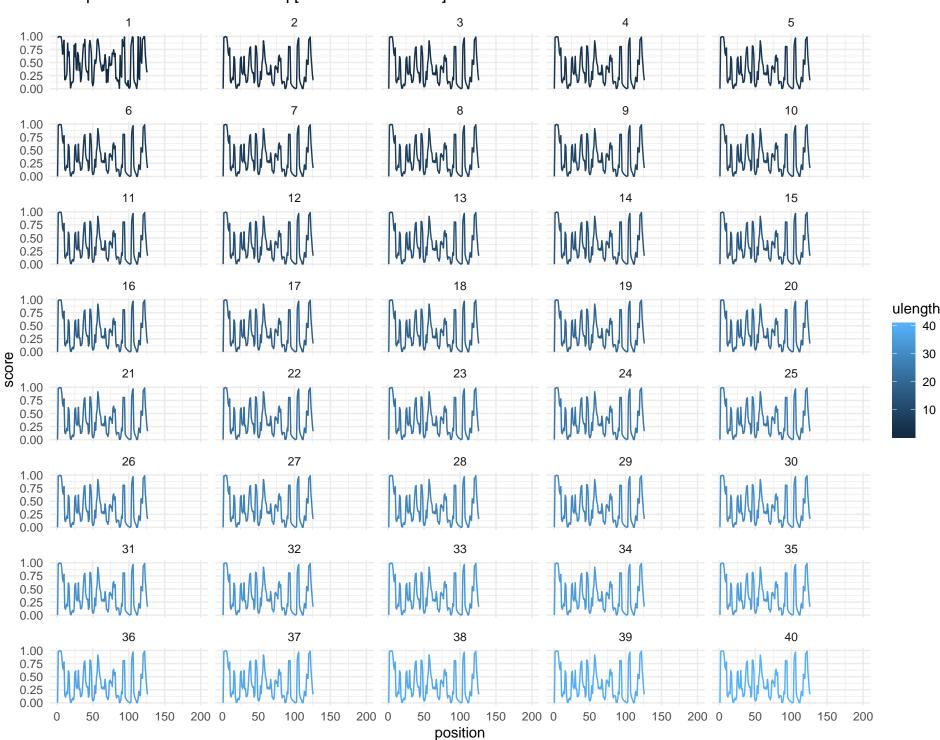
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



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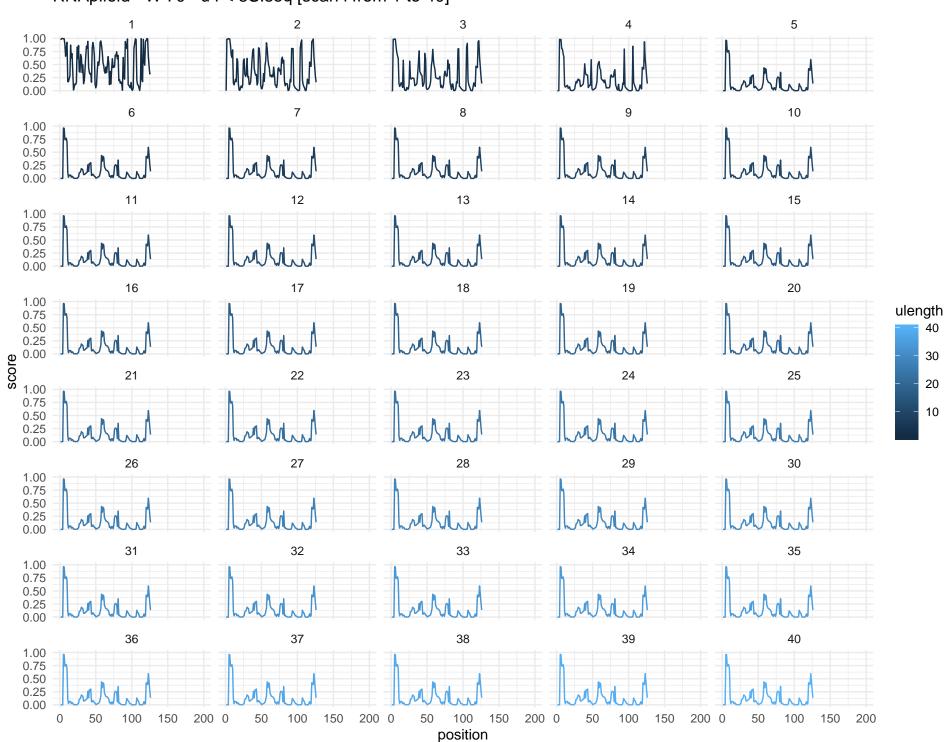
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



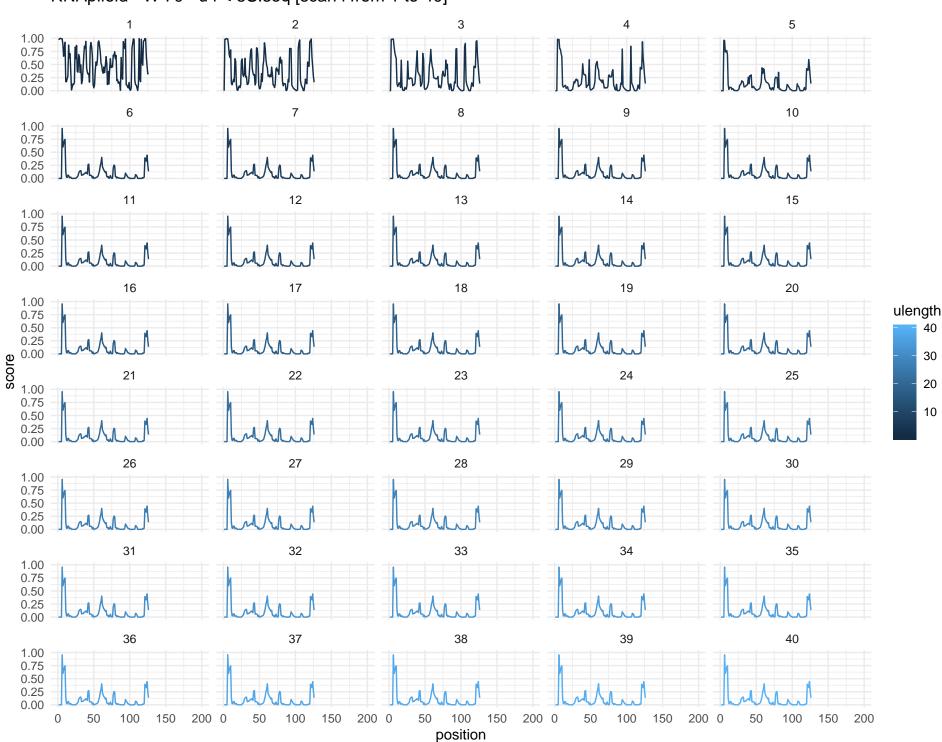
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



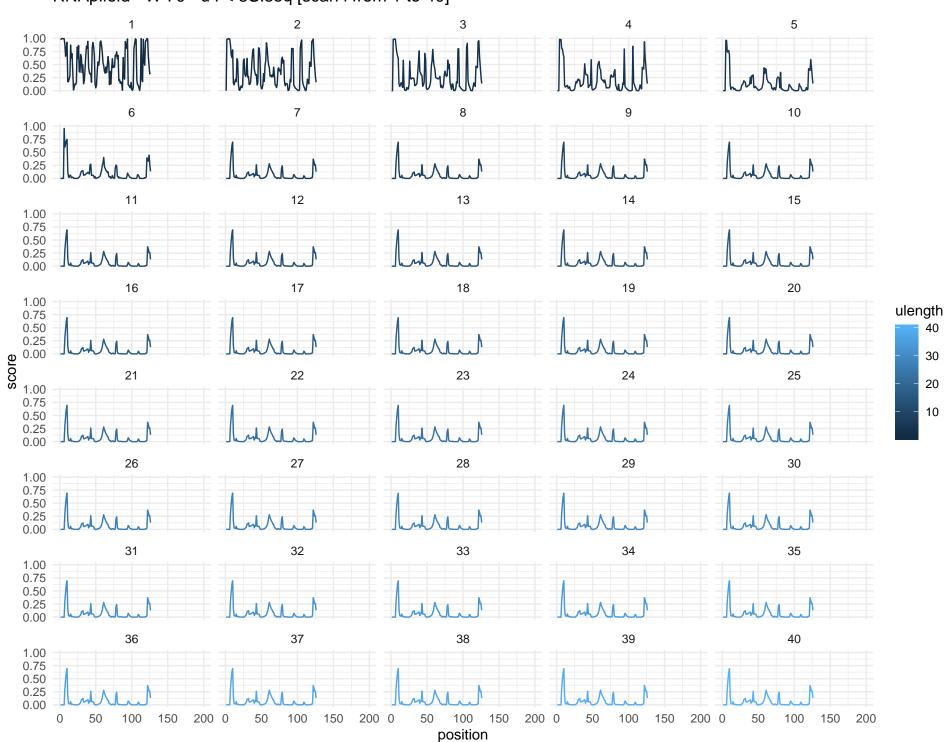
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



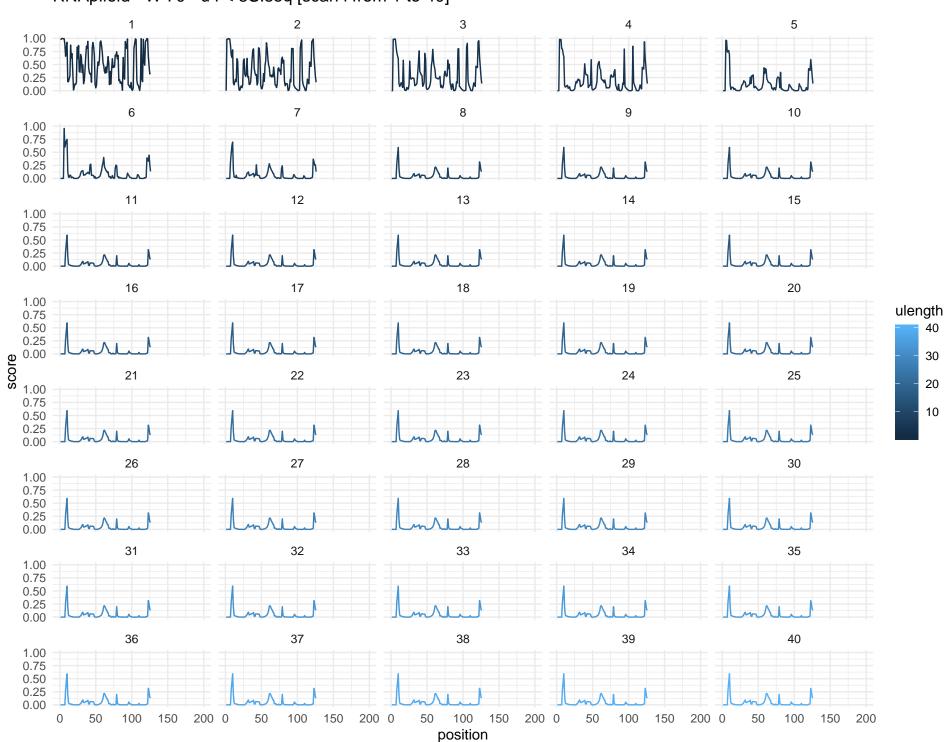
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



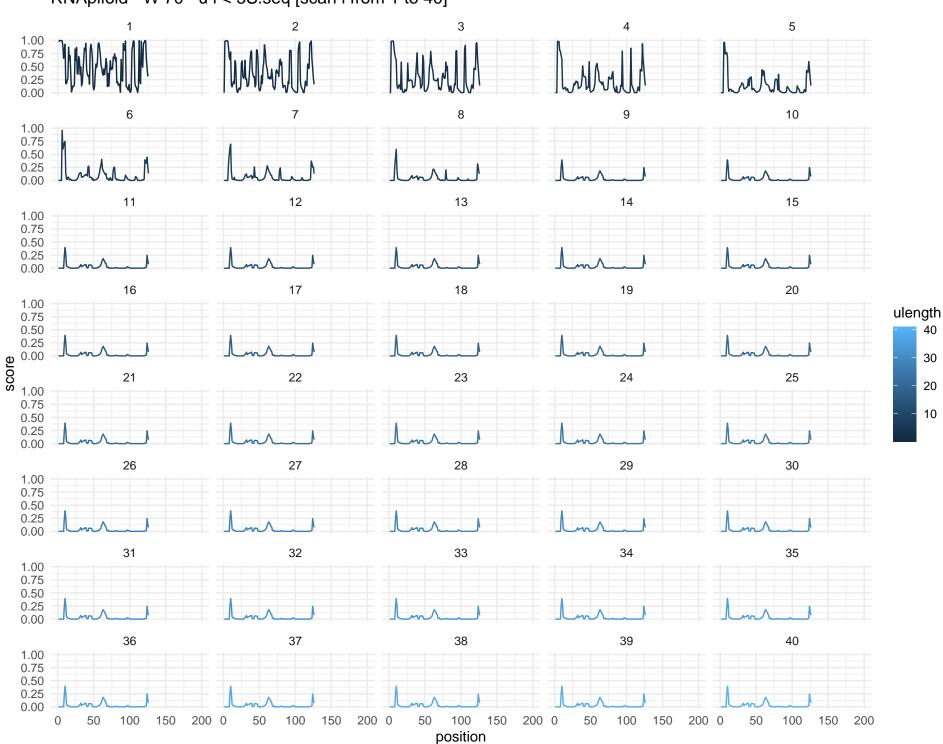
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



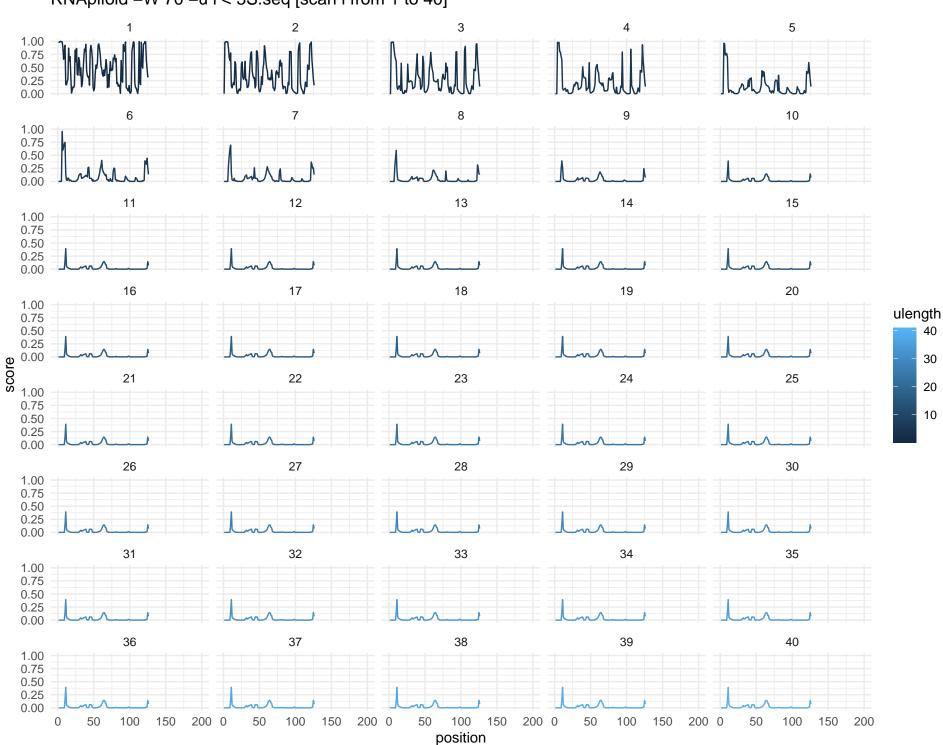
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



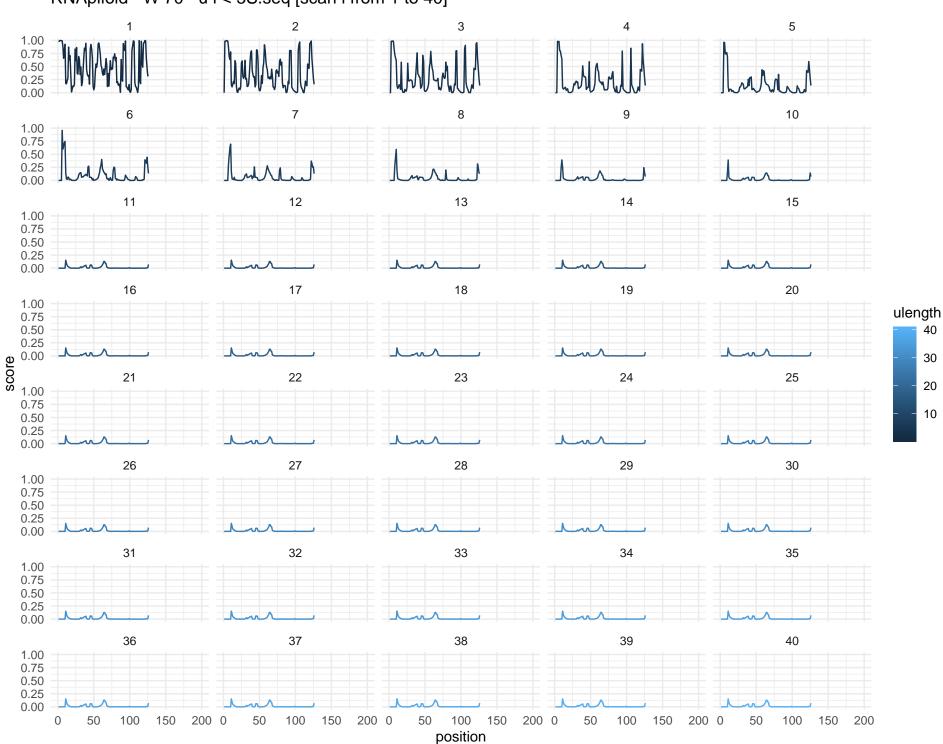
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



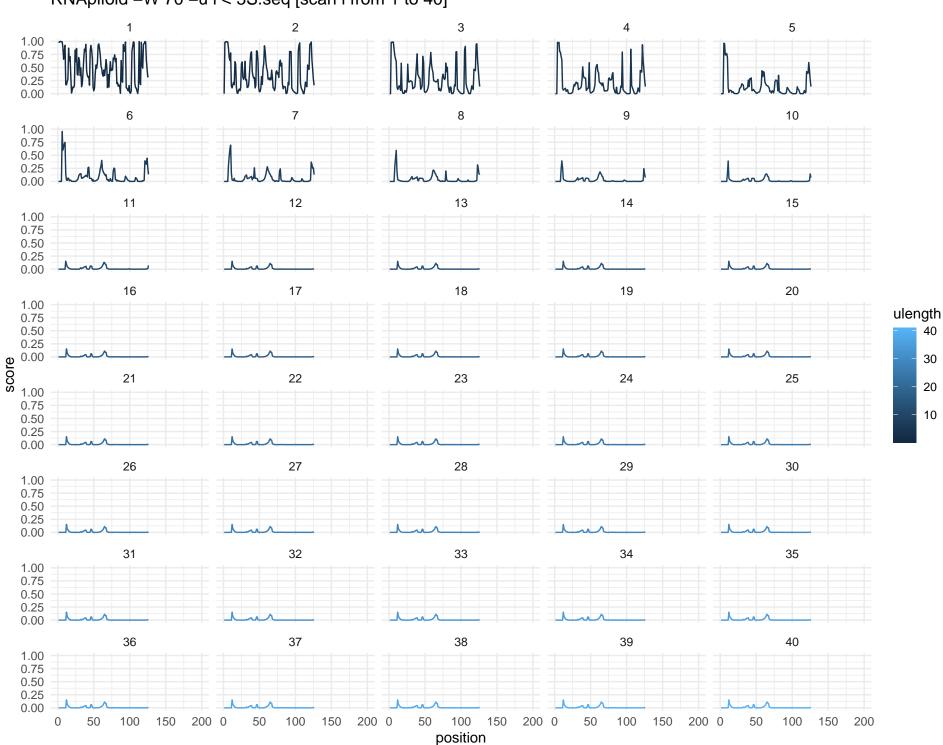
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



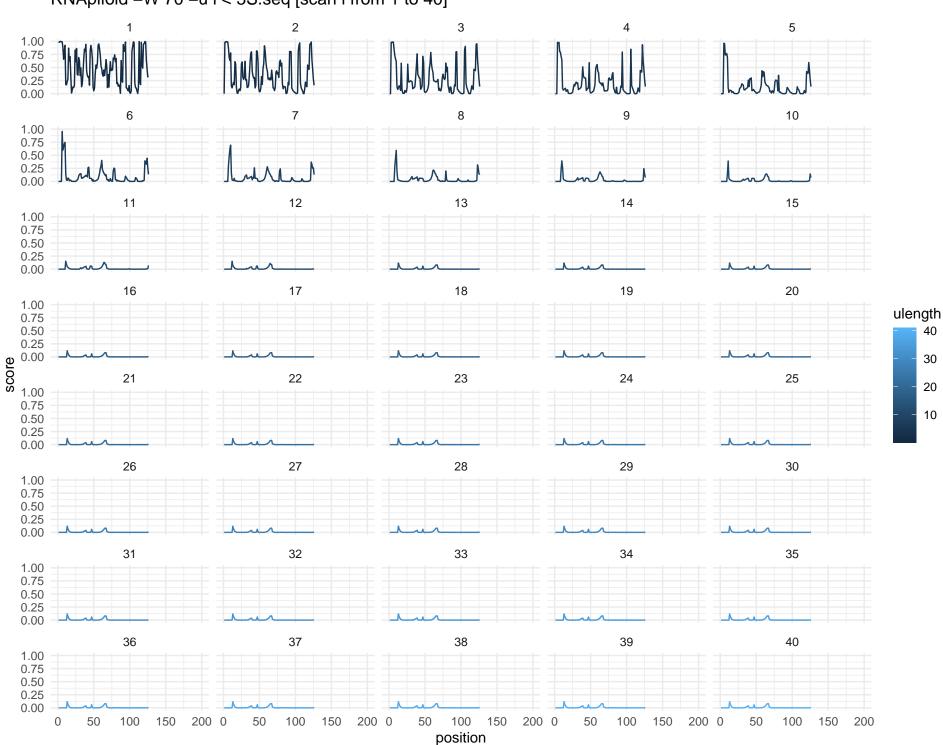
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



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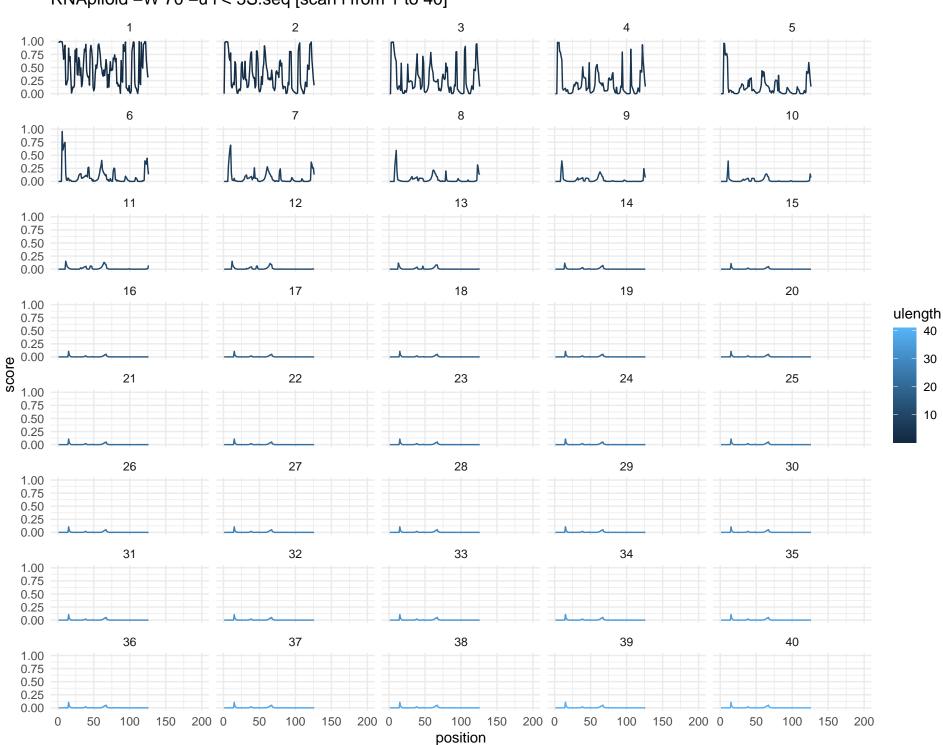
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



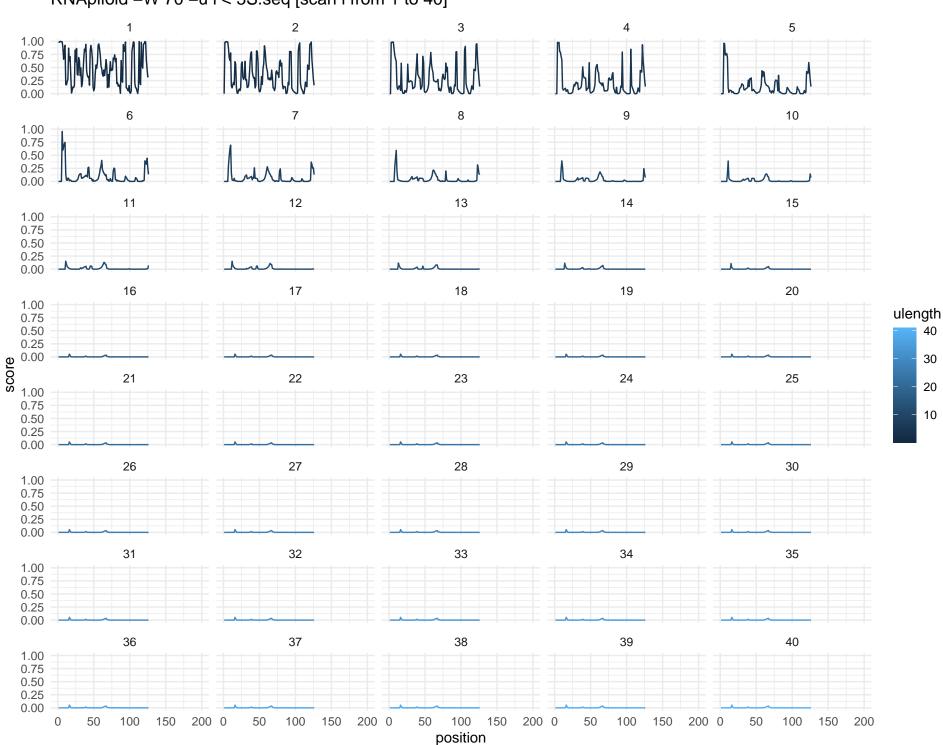
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



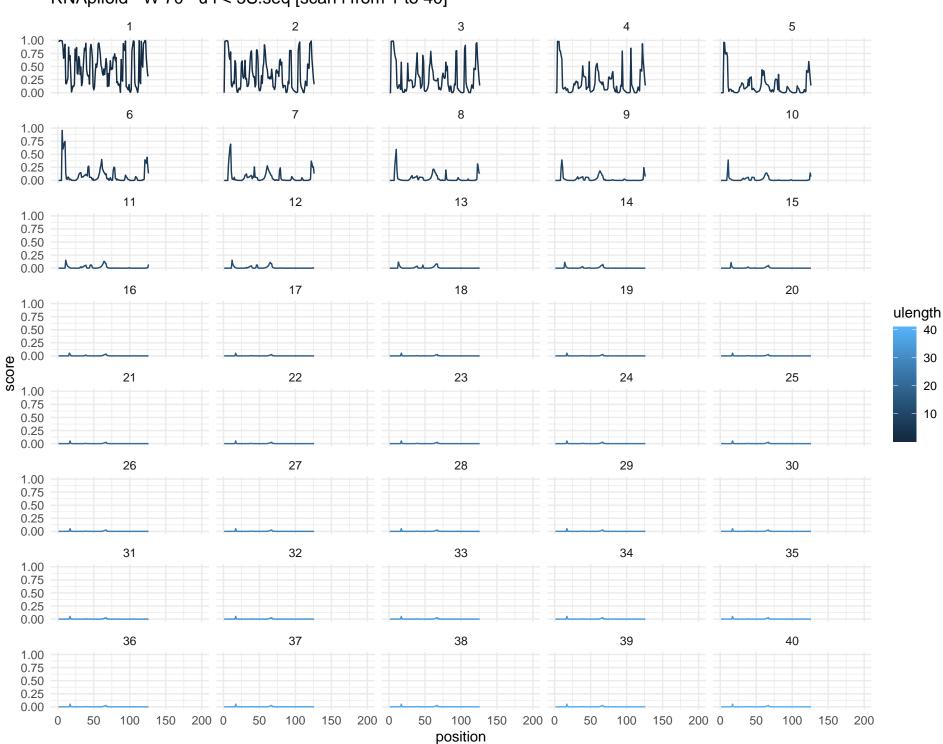
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



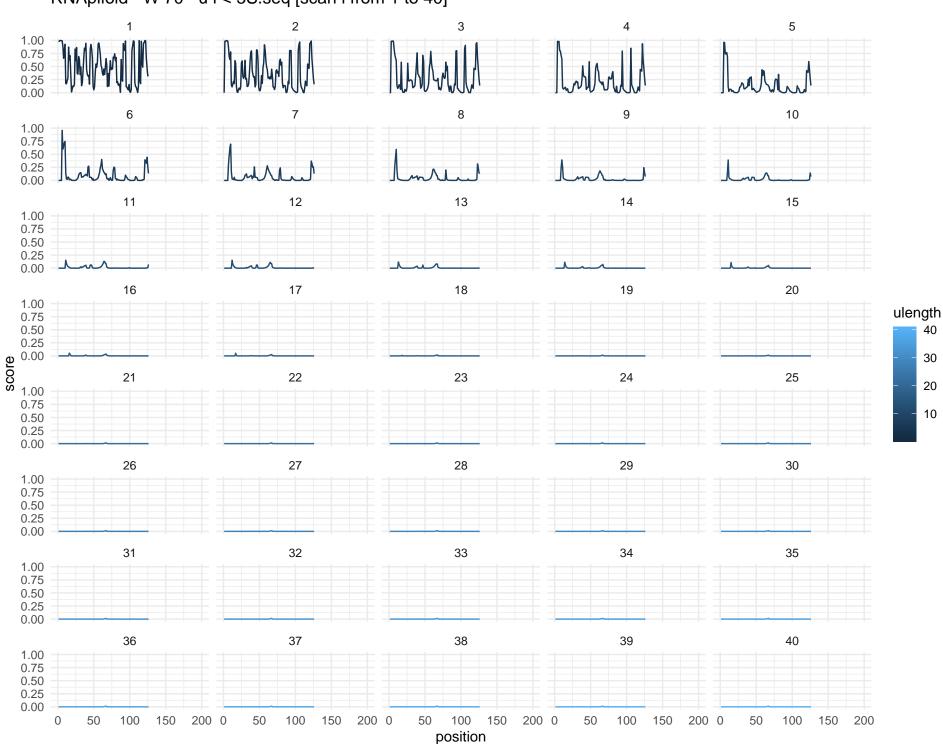
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



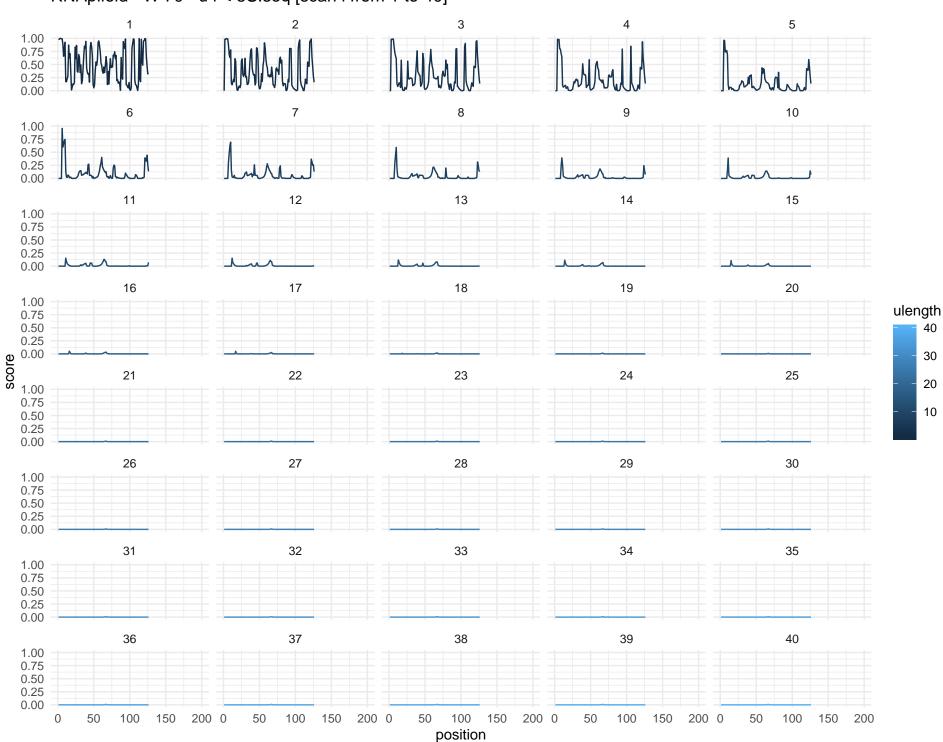
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



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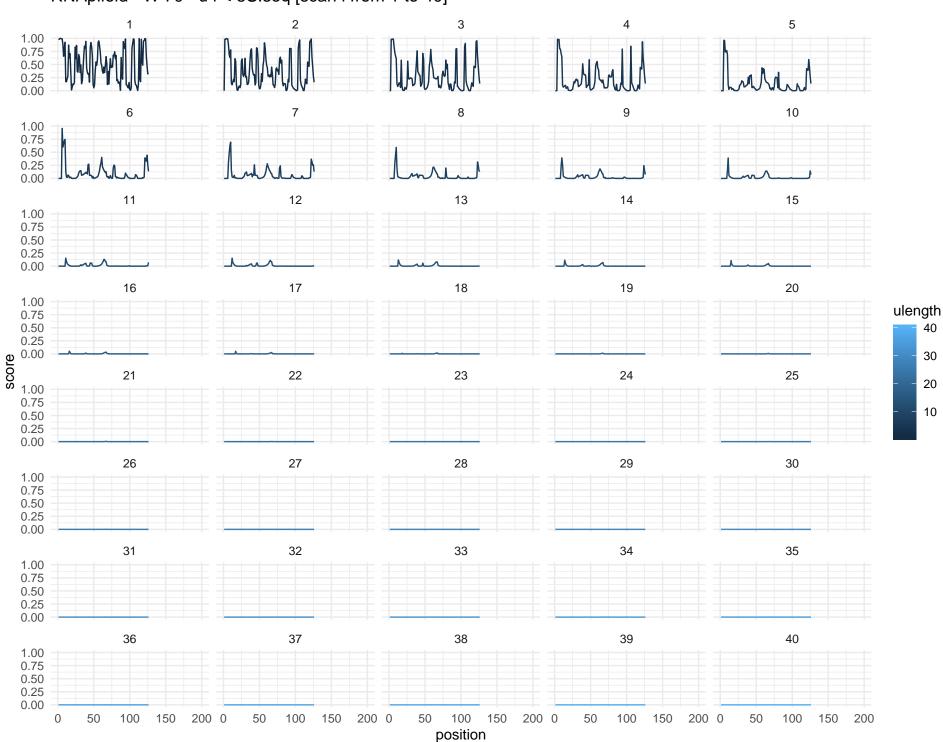
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



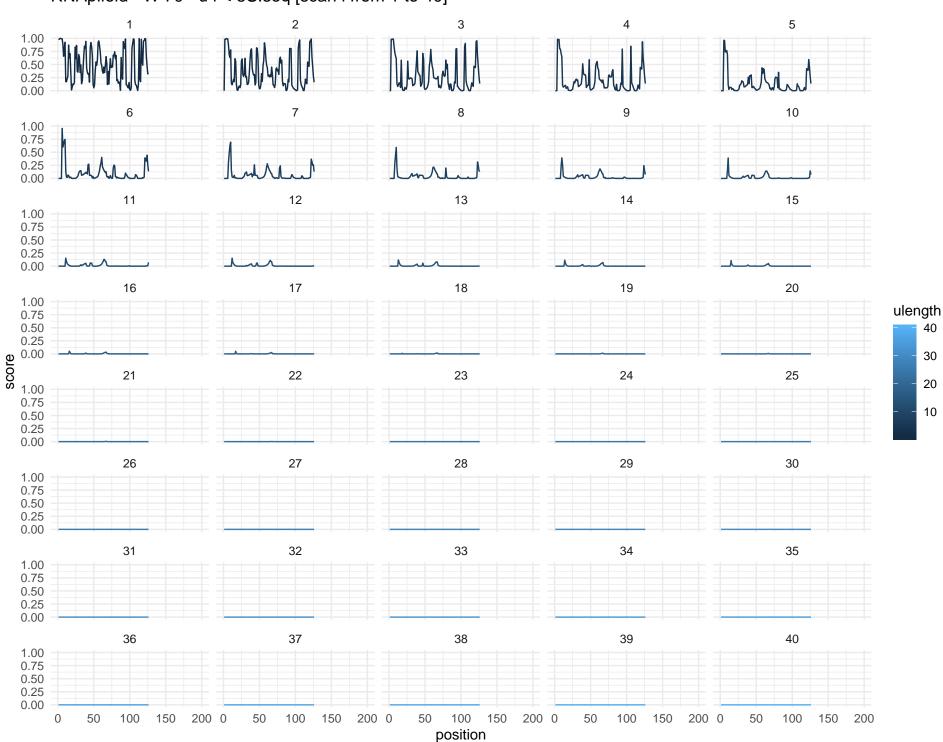
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



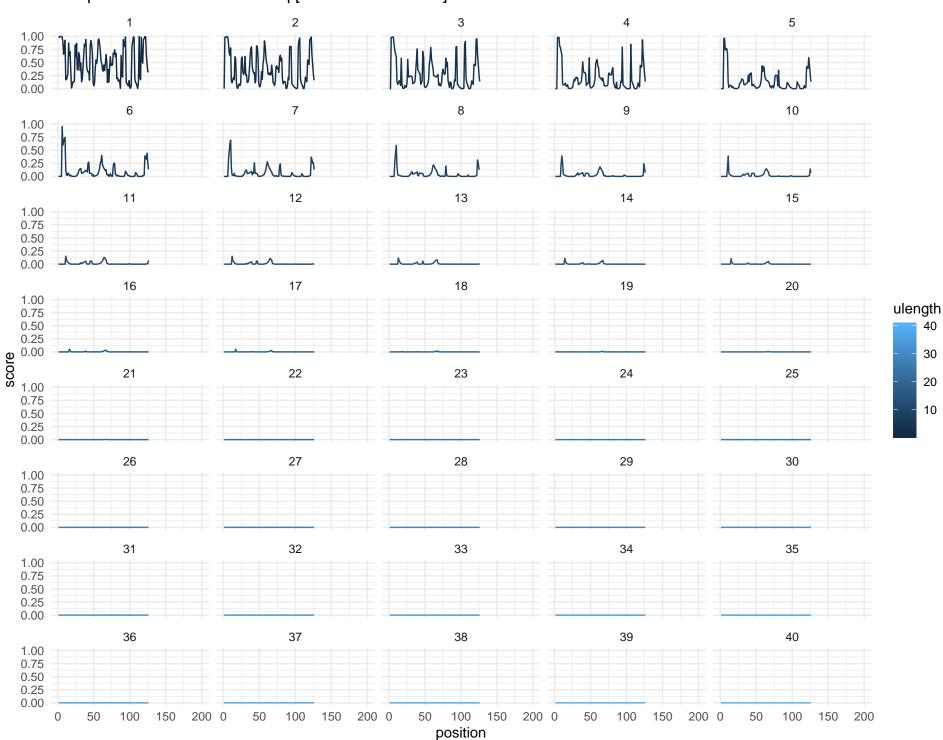
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



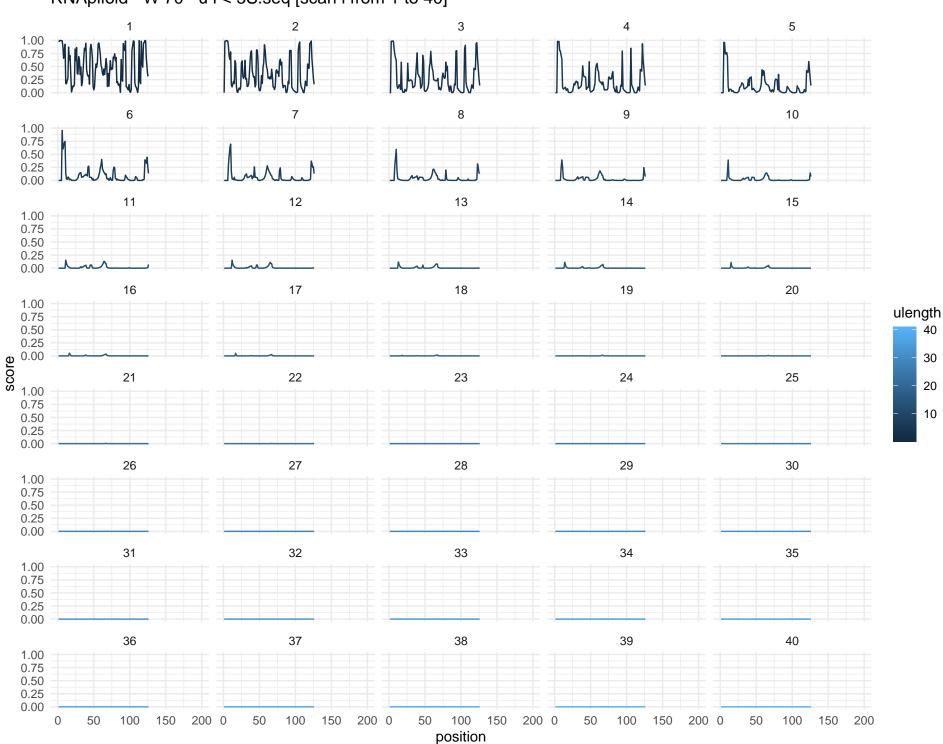
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



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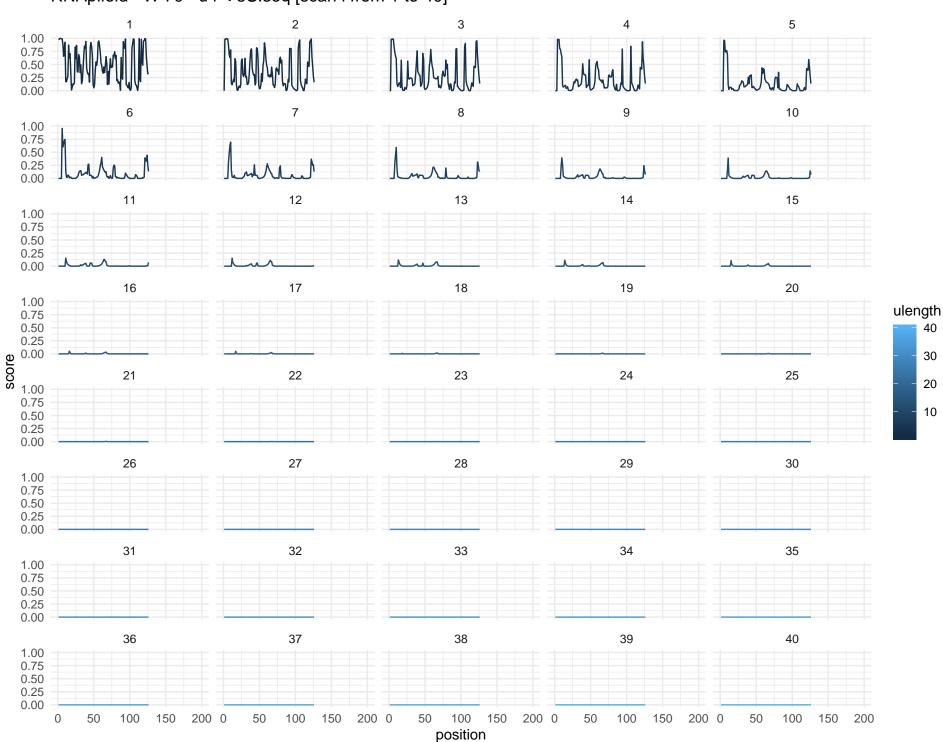
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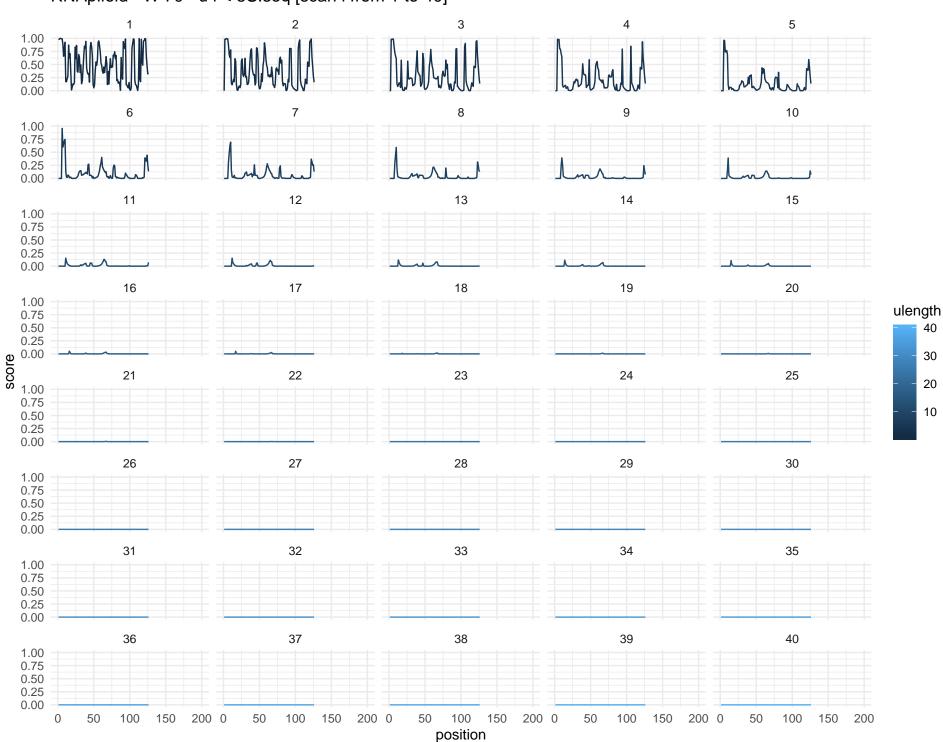
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



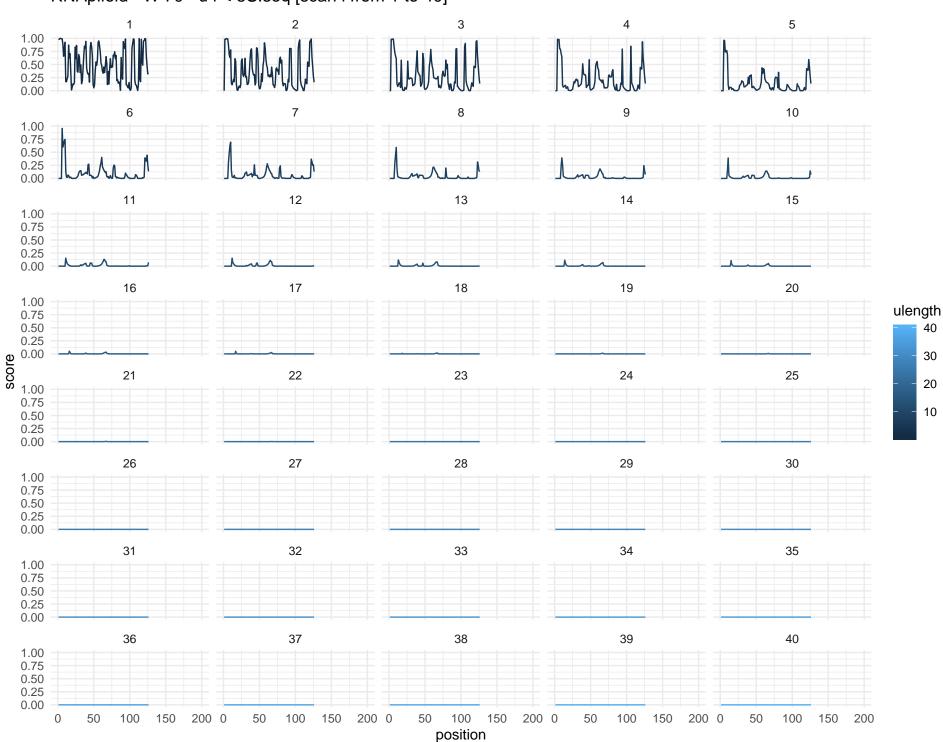
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



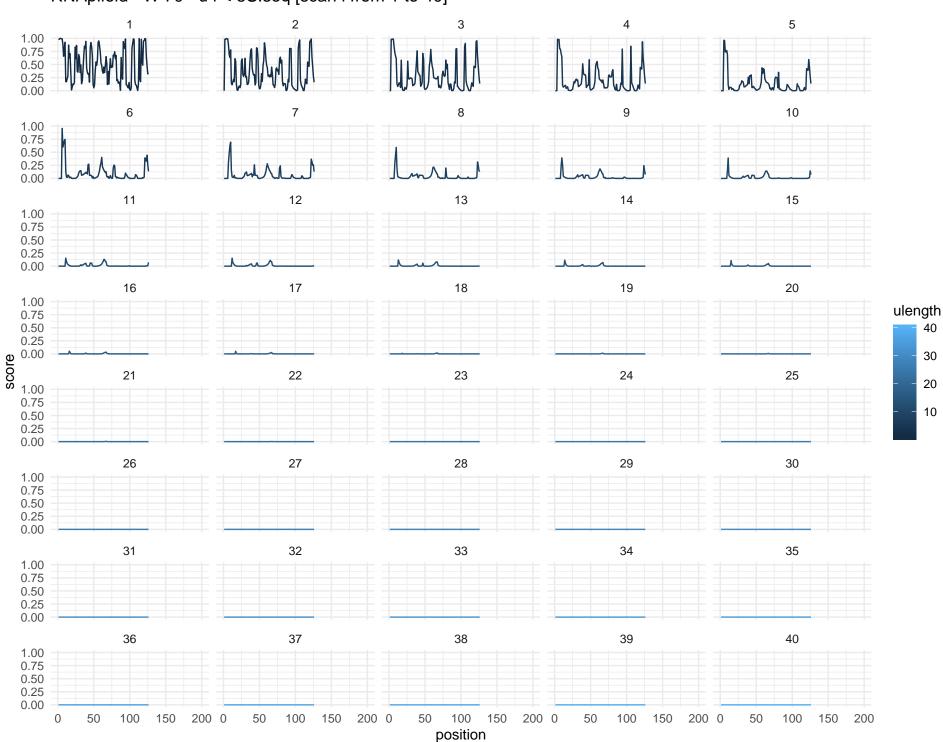
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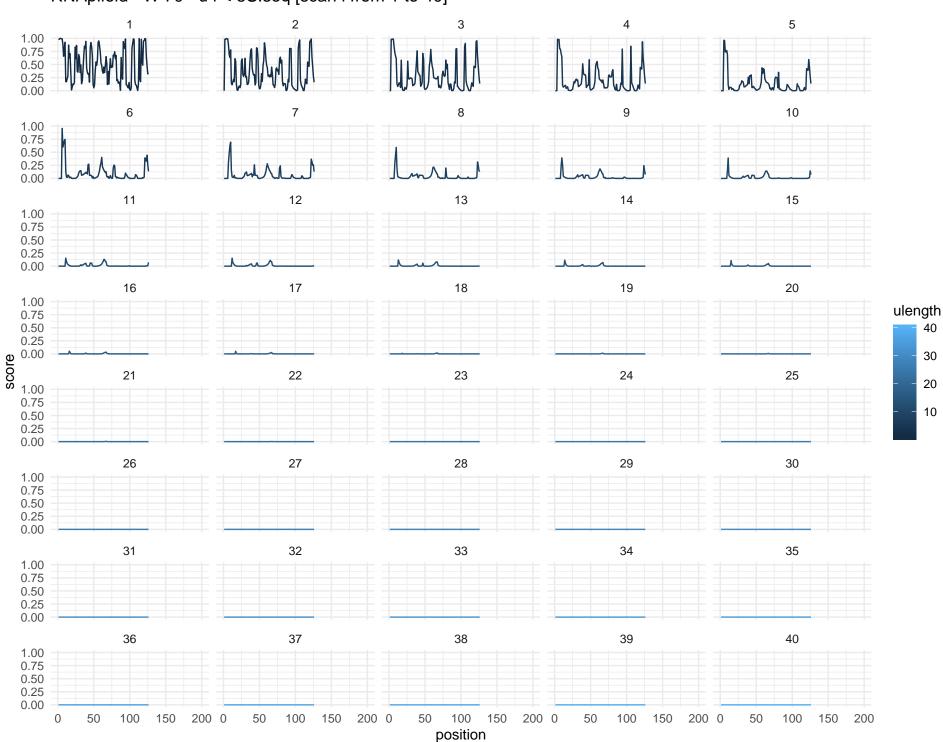
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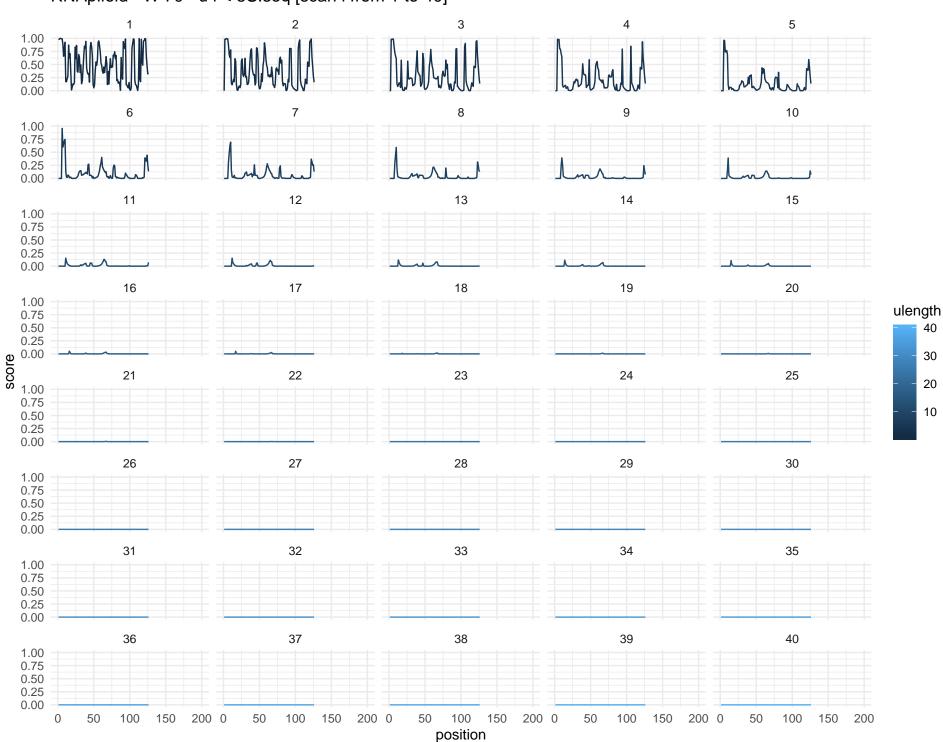
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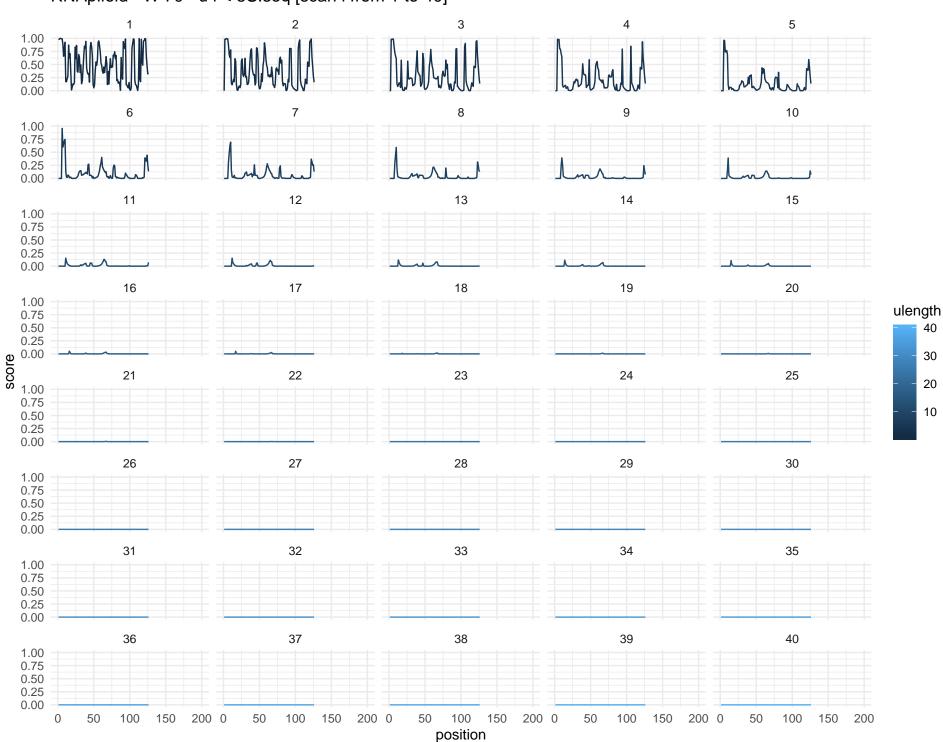
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



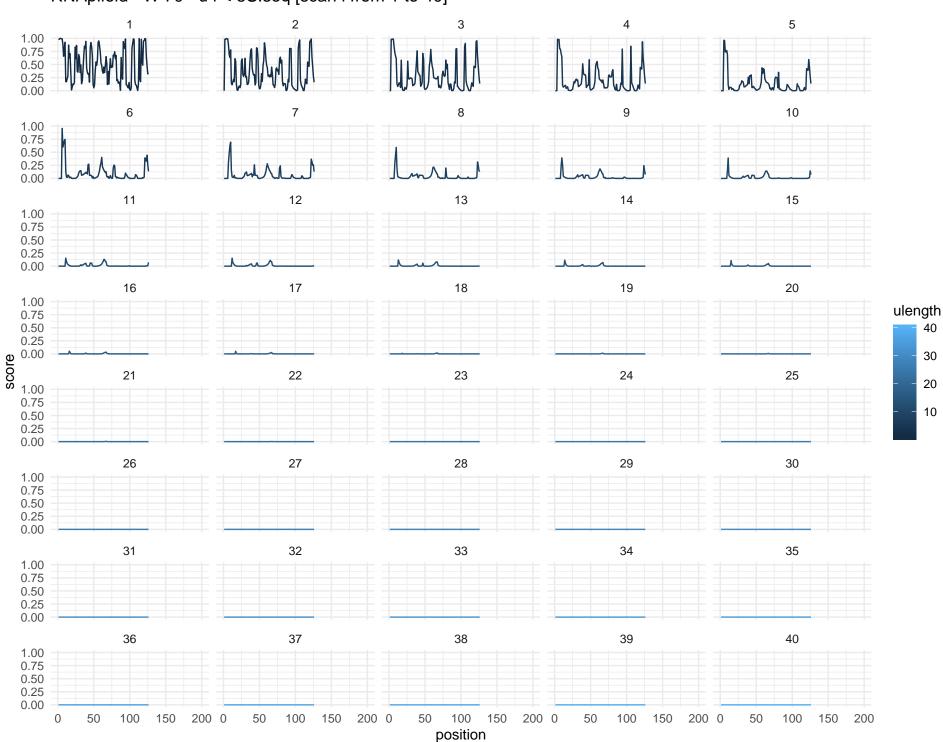
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



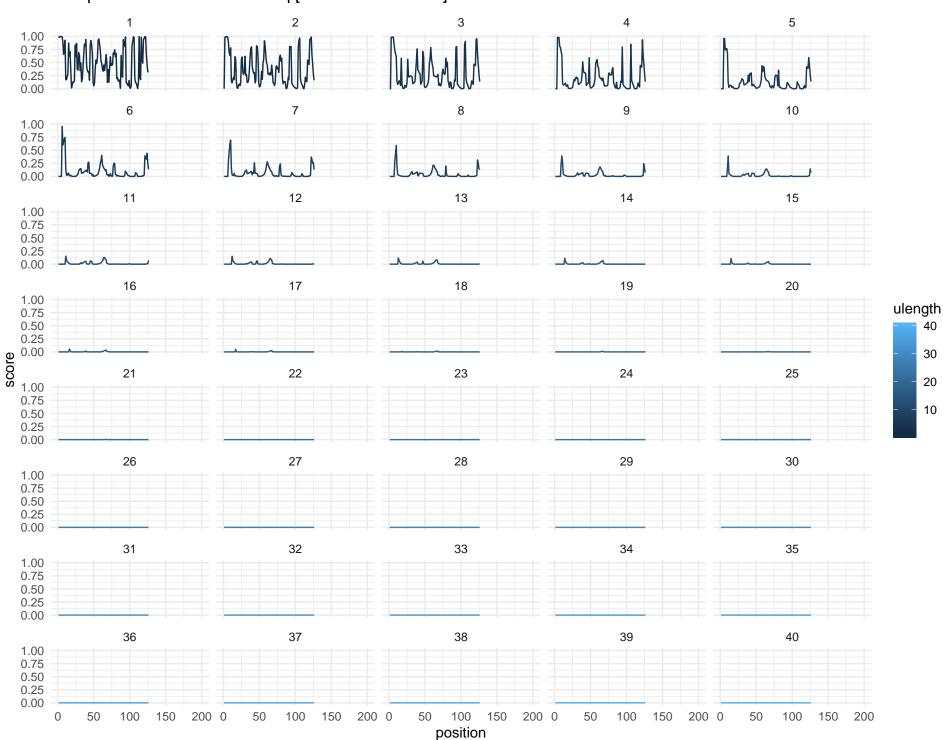
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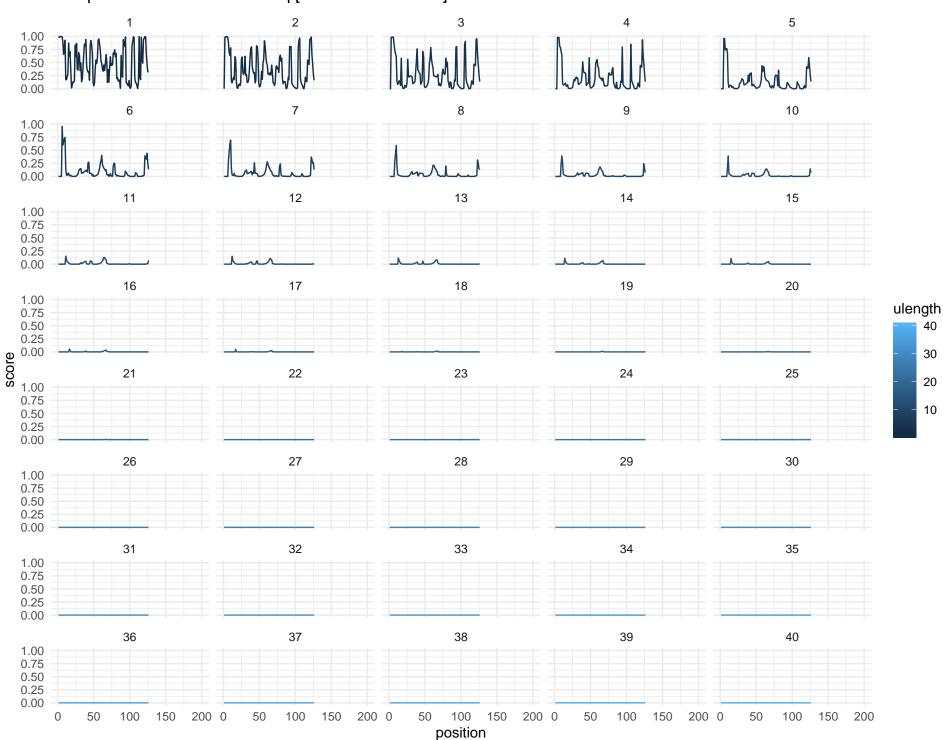
RNAplfold –W 70 –u i < 5S.seq [scan i from 1 to 40]



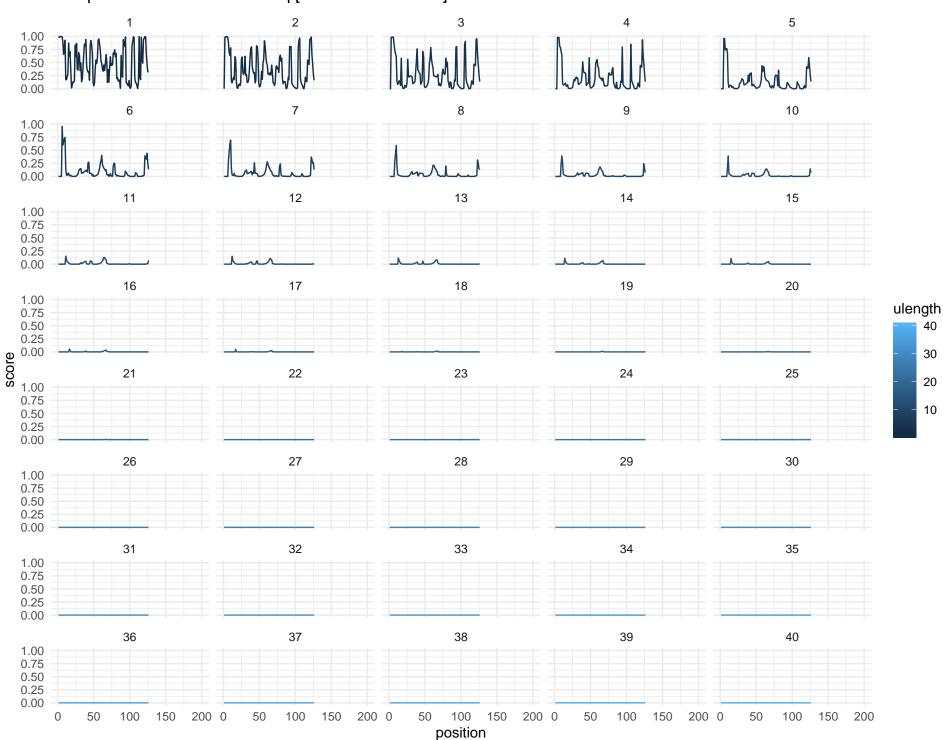
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