

Combinatorial approaches to RNA folding

Part I: Basics

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Math 4500, Fall 2016

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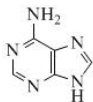
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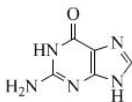
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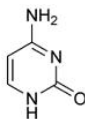
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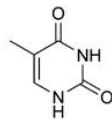
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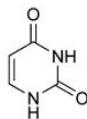
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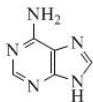
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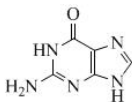
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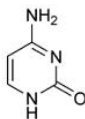
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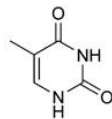
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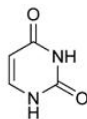
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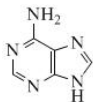
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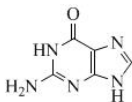
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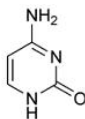
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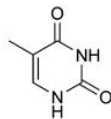
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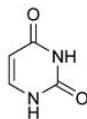
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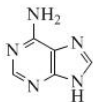
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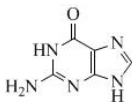
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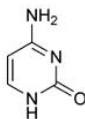
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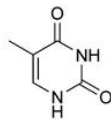
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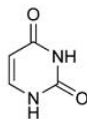
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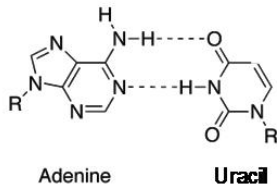
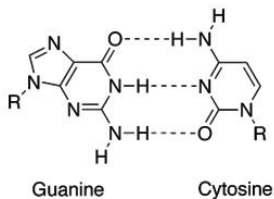
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Bases can **bond**: A with U, and C with G. (*Watson–Crick* base pairs.)



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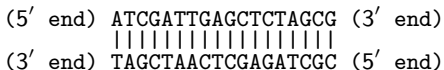
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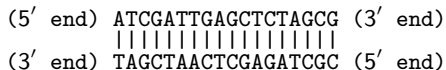
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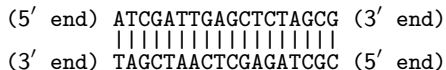
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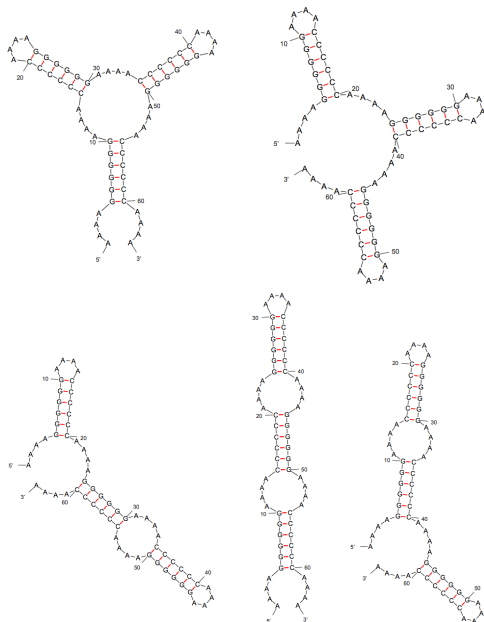
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RNA consists of a single strand. It can fold and bond to itself. It is much less structurally constrained than DNA!

How does RNA fold? [image from C. Heitsch; Georgia Tech]



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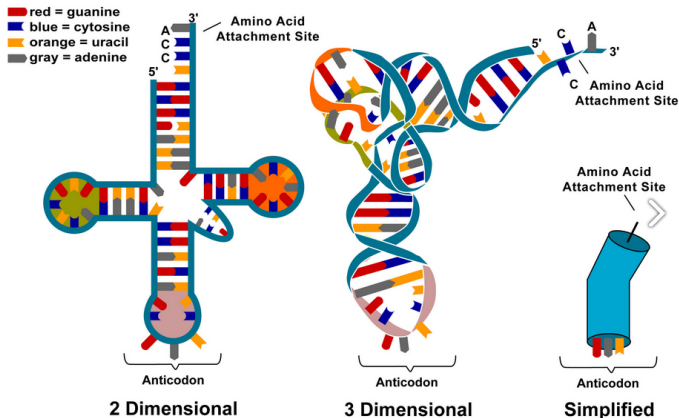
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- *Primary structure*: The raw sequence of nucleotides.
- *Secondary structure*: The bonding between nucleotides on a single strand.
- *Tertiary structure*: Embedding (e.g., twisting, knotting, etc.) of the strand in 3-dimensional space.



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Before we proceed, we will need to establish a combinatorial framework for describing RNA strands.

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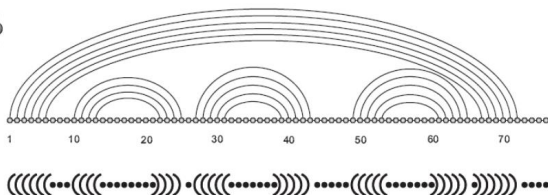
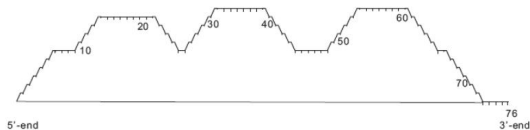
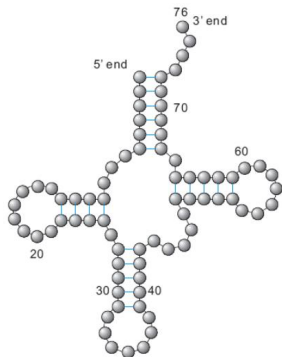
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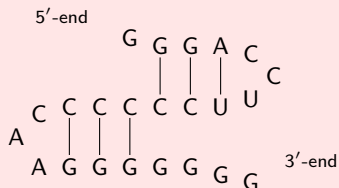
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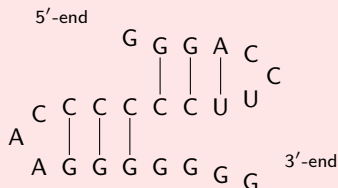


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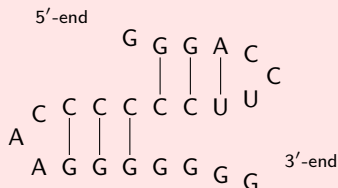
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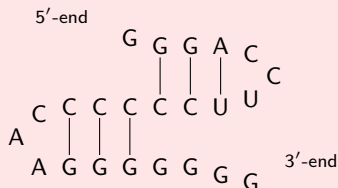
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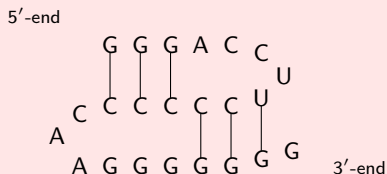
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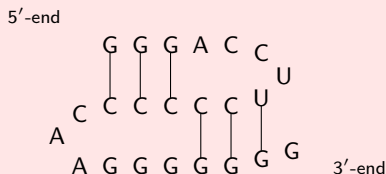
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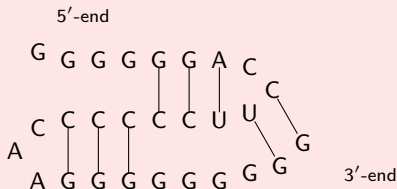
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An RNA structure is a **pseudoknot** if its arc diagram has crossings.

An arc diagram is **k -noncrossing** if there is no set of k mutually crossing arcs.

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- (i) Draw the corresponding **arc diagram**. What is the smallest k for which this is k -noncrossing .
- (ii) What if the first G bonds with the C “directly below” it (vertex 17). Does this change the k from the previous part?
- (iii) Draw a picture of a folded RNA strand (like the one above) that is 4-noncrossing but not 3-noncrossing.

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- How can we *uniformly generate* an RNA structure?

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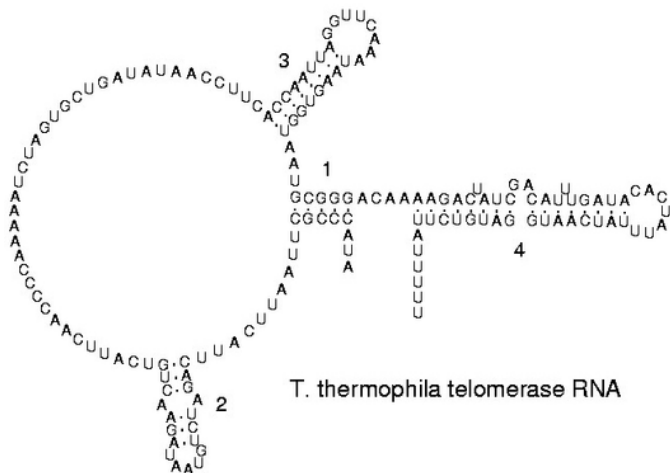
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Loop types

0. The vertices not accessible from any arcs form the unique 0-loop, or **null loop** L_0 .
1. A 1-loop is called a **hairpin loop**
2. There are three types of 2-loops: **bulge loops**, **interior loops**, and **stacked pairs**.
3. A k -loop for $k \geq 3$ is called a **multiloop**.

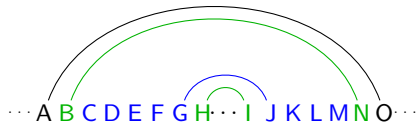
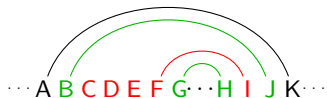
Loop decomposition

2-loops

Suppose (i', j') is the unique accessible base pair from (i, j) . Then the resulting 2-loop is:

- 2a. a **stacked pair** if $i - i' = j' - j = 1$;
- 2b. a **bulge loop** if exactly one of $i - i'$ and $j' - j$ is > 1 ;
- 2c. an **interior loop** if both $i - i'$ and $j' - j$ are > 1 ;

Two 2-loops: a bulge loop (left) and an interior loop (right). Each secondary structure also contains two 2-loops that are stacked pairs.



Loop decomposition with pseudoknotting

Things get a little more complicated when the diagram contains a pseudoknot, but there is still a well-defined decomposition. (We won't go into details.)

