

CSE8803/CX4803

Machine Learning in Computational Biology

Lecture 12: RNA Secondary Structure Prediction

Yunan Luo

Learning from structure

Date	Topic	Contents
1/10/2022	Introduction	Course intro & how to present papers
1/12/2022	Learning from sequence data	Dynamic programming & sequence alignment I
1/17/2022		No class (MLK Day)
1/19/2022		Sequence alignment II
1/24/2022		HMM & gene/motif finding
1/26/2022		HMM & Profile HMM
1/31/2022		Deep learning for DNA/protein sequence
2/2/2022	Learning from high-dim data	Learn from high-dim data: PCA, autoencoder & VAE
2/7/2022		Learn from high-dim data: MDS, tSNE, UMAP
2/9/2022		Clustering I
2/14/2022		Clustering II
2/16/2022	Phase 1 presentations	Clustering III
2/21/2022		Student presentation 1-3
2/23/2022		Student presentation 4-6
2/28/2022	Learning from structure data	RNA structure prediction
3/2/2022		Deep learning for structures (protein structure prediction)
3/7/2022	Phase 2 presentations	Student presentation 7-9
3/9/2022	Learning from network data	Network basics & traditional ML for graphs
3/14/2022		Network embeddings
3/16/2022	Phase 3 presentations	Student presentation 10-12
3/21/2022	Spring break	No class (Spring Break)
3/23/2022		No class (Spring Break)
3/28/2022	Learning from network data	Graphical Models
3/30/2022		Deep learning for networks (graph neural networks)

Central Dogma of Molecular Biology

Three fundamental molecules:

1. DNA

Information storage.

2. RNA

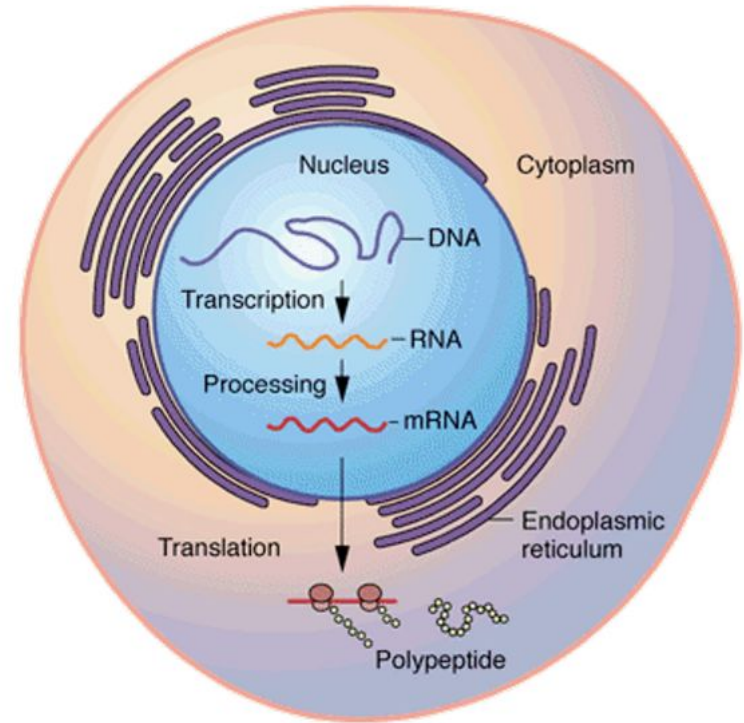
Old view: Mostly a “messenger”.

New view: Performs many important functions, through **3-D structure!**

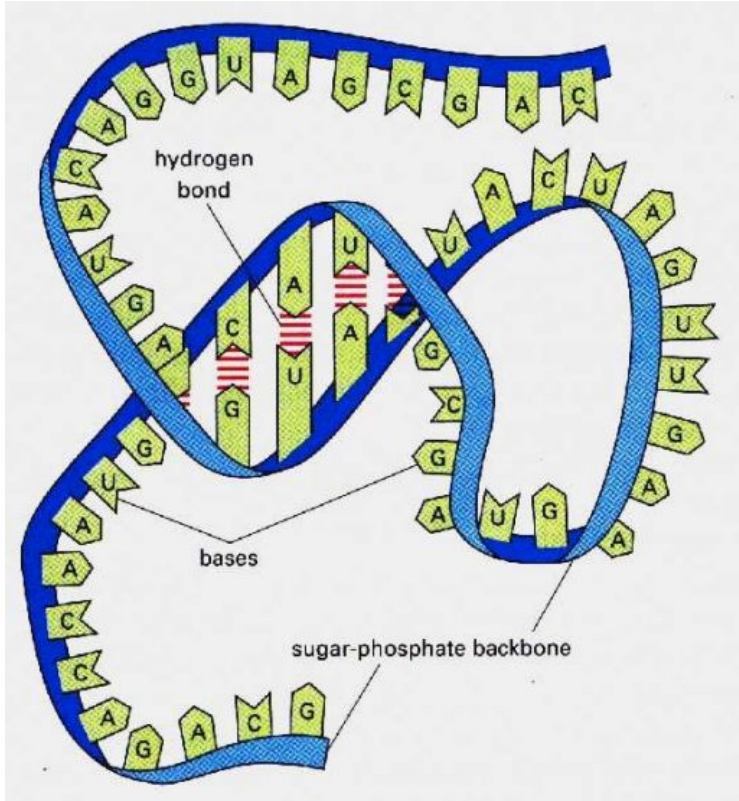
3. Protein

Perform most cellular functions
(biochemistry, signaling, control, etc.)

DNA → RNA → Protein



RNA



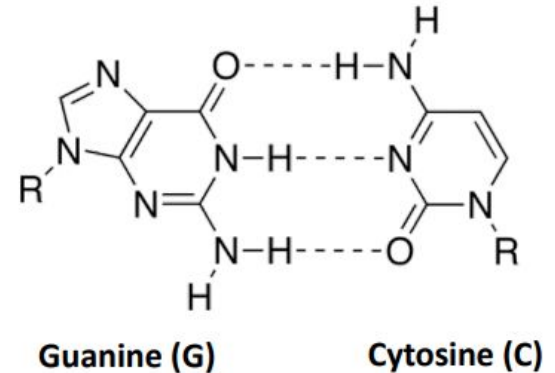
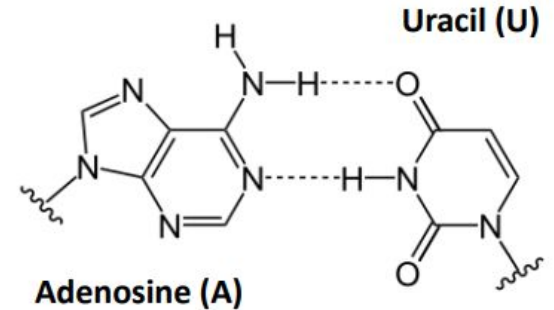
- **Single-stranded**
 - A (adenine)
 - C (cytosine)
 - U (uracil)
 - G (guanine)
- Can fold into **structures** due to nucleotide complementarity.
 $A \leftrightarrow U, C \leftrightarrow G$
- Comes in many flavors:
mRNA, rRNA, tRNA, tmRNA, snRNA, snoRNA, scaRNA, aRNA, asRNA, piwiRNA, etc.

RNA – Nucleotide Complementarity

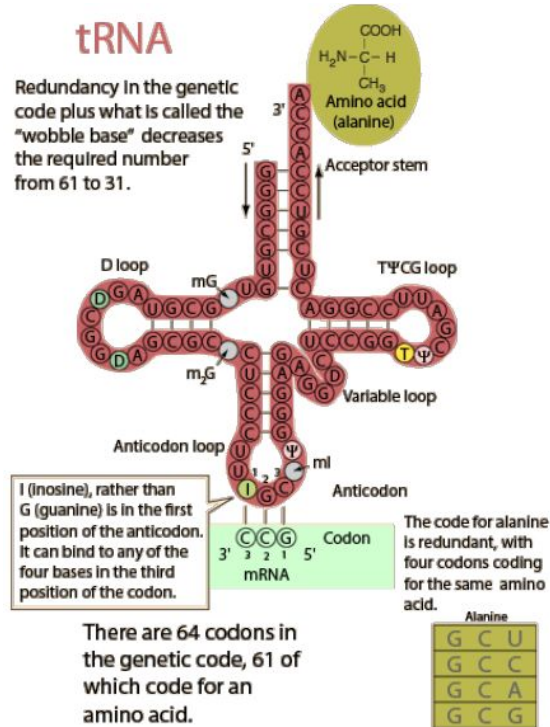
RNA can fold into structures due to
nucleotide complementarity:
 $A \leftrightarrow U$ and $G \leftrightarrow C$

$A \leftrightarrow U$ (2 hydrogen bonds)
is slightly weaker than
 $G \leftrightarrow C$ (3 hydrogen bonds)

$G \leftrightarrow C$ also observed but not as stable



Transfer RNA (tRNA) Secondary Structure



		Second base				
		U	C	A	G	
First base	U	UUU } Phenyl-alanine F UUC } UUA } Leucine L UUG }	UCU } Serine S UCC } UCA } UCG }	UAU } Tyrosine Y UAC } UAA } Stop codon UAG } Stop codon	UGU } Cysteine C UGC } UGA } Stop codon UGG } Tryptophan W	U C A G
	C	CUU } Leucine L CUC } CUA } CUG }	CCU } Proline P CCC } CCA } CCG }	CAU } Histidine H CAC } CAA } Glutamine Q CAG }	CGU } Arginine R CGC } CGA } CGG }	U C A G
	A	AUU } Isoleucine I AUC } AUA } AUG } Methionine start codon M	ACU } Threonine T ACC } ACA } ACG }	AAU } Asparagine N AAC } AAA } Lysine K AAG }	AGU } Serine S AGC } AGA } Arginine R AGG }	U C A G
	G	GUU } Valine V GUC } GUA } GUG }	GCU } Alanine A GCC } GCA } GCG }	GAU } Aspartic acid D GAC } GAA } Glutamic acid E GAG }	GGU } Glycine G GGC } GGA } GGG }	U C A G

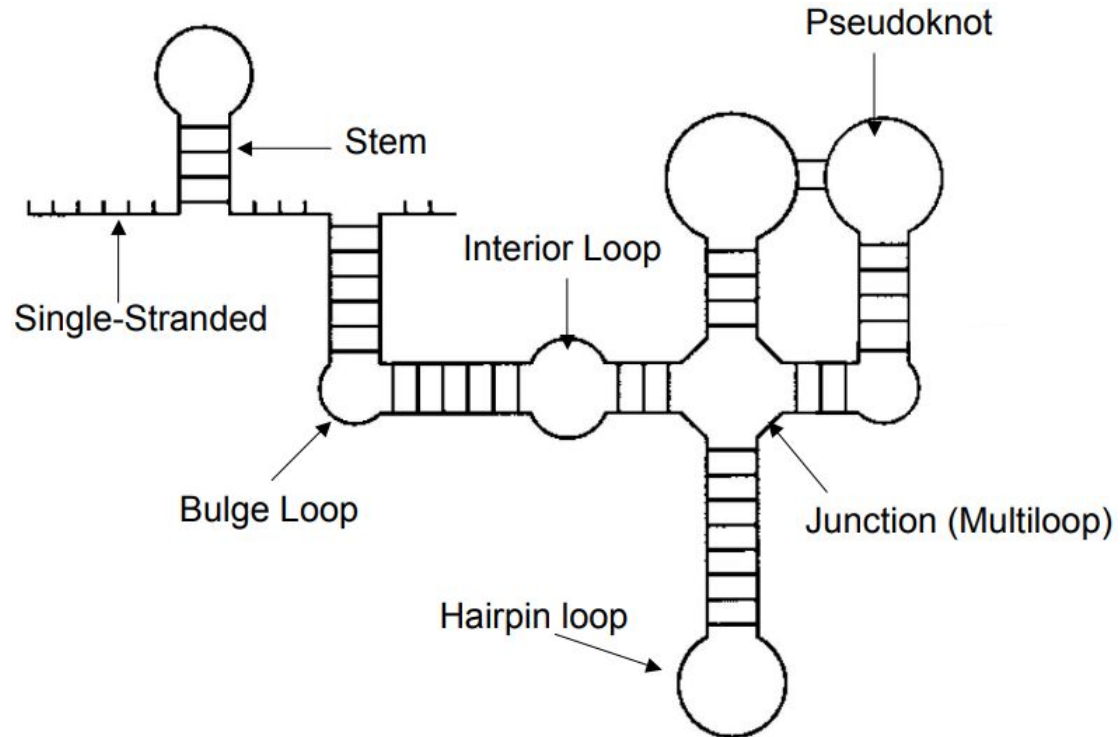
Third base

<http://bioinfo.bisr.res.in/project/crat/pictures/codon.jpg>

RNA Secondary Structure Elements

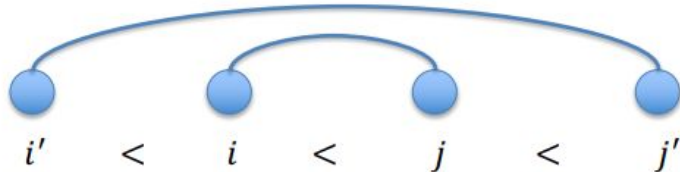
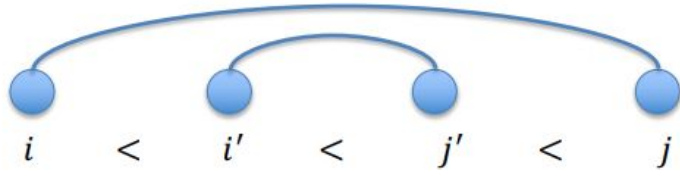
Each base/nucleotide participates in at most one pairing

Secondary structure is determined by a set of non-overlapping base/nucleotide pairs

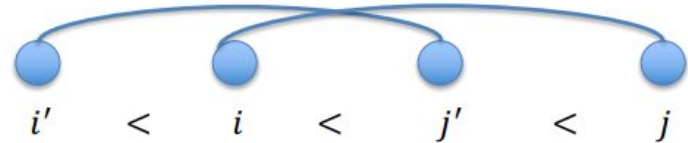
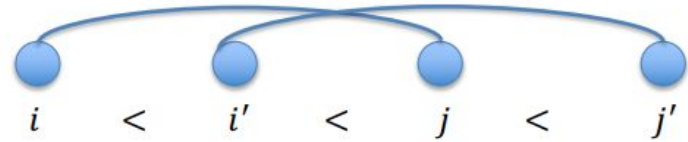


Nesting and Pseudoknot

Base pairs (i, j) and (i', j') are **nested**
provided
 $i < i' < j' < j$ or $i' < i < j < j'$



Base pairs (i, j) and (i', j') form a **pseudoknot** provided
 $i < i' < j < j'$ or $i' < i < j' < j$

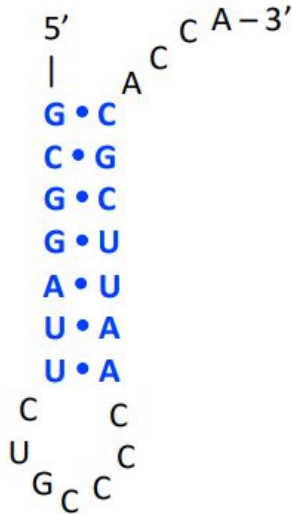


Most RNA molecules consist of nested base pairs

Nesting and Pseudoknot -- Examples

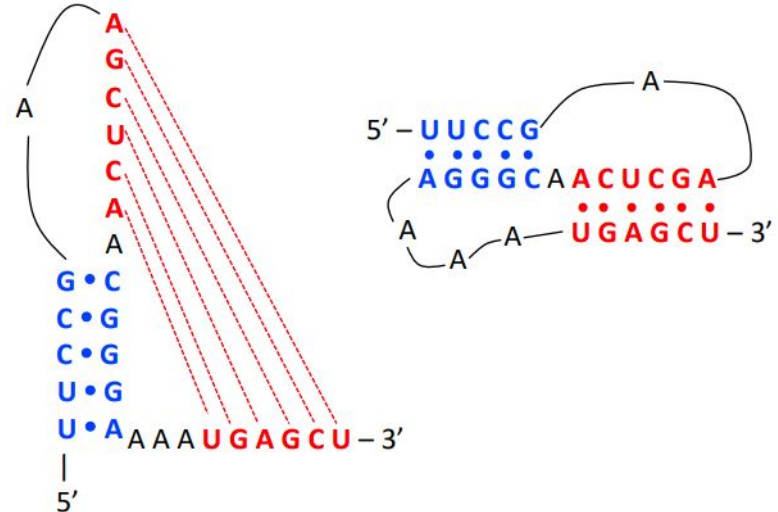
Nesting

5'-G C G G A U U C U G C C C C A A U U C G C A C C A-3'
 ((((((- - - - -)))))) - - - -



Pseudoknot

5'-U U C C G A A G C U C A A C G G G A A A A U G A G C U-3'
 (((((- (((((-))))) - - -)))))



Nussinov Algorithm

RNA can fold into structures due to nucleotide complementarity:

$A \leftrightarrow U$ and $G \leftrightarrow C$

Secondary structure is determined by a set of non-overlapping complementary base pairs

Question: How to find maximum number of such pairs?

5' – G C G G A U U C U G C C C C A A U U C G C A C C A – 3'

Nussinov Algorithm

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A \leftrightarrow U and G \leftrightarrow C

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Question: How to find maximum number of such pairs?

Need to constrain space of feasible solutions!

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ALGORITHMS FOR LOOP MATCHINGS*

RUTH NUSSINOV,[†] GEORGE PIECZENIK,[‡] JERROLD R. GRIGGS[¶]
AND DANIEL J. KLEITMAN[§]



Problem: Given RNA sequence $\mathbf{v} \in \{A, U, C, G\}^n$, find a *pseudoknot-free secondary structure* with the maximum number of complementary base pairings

Nussinov Algorithm – Dynamic Programming

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5' – G C G G A U U C U G C C C C A A U U C G C A C C A – 3'

Nussinov Algorithm – Dynamic Programming

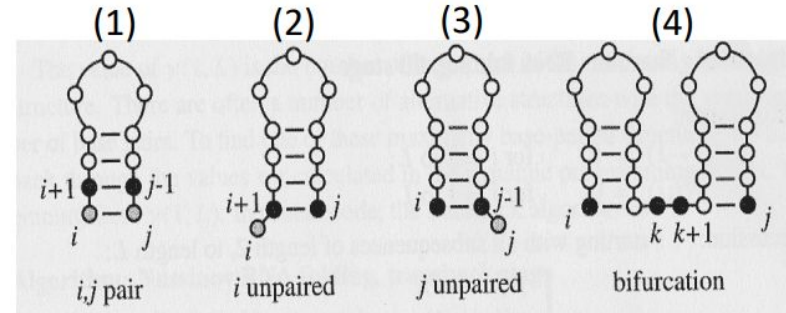
Problem: Given RNA sequence $\mathbf{v} \in \{A, U, C, G\}^n$, find a *pseudoknot-free secondary structure* with the maximum number of complementary base pairings

Let $s[i,j]$ denote the maximum number of pseudoknot-free complementary base pairings in subsequence v_i, \dots, v_j

Nussinov Algorithm – Dynamic Programming

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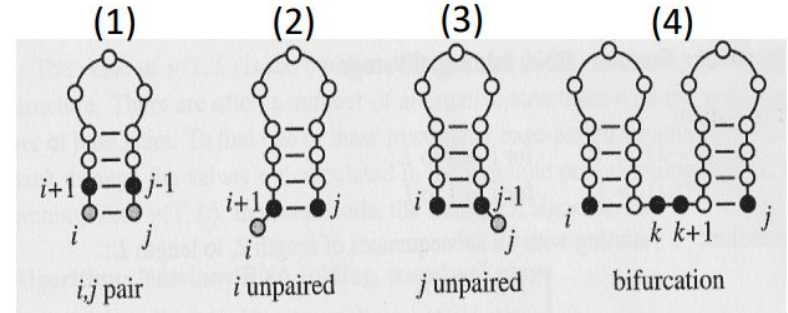
Let $s[i,j]$ denote the maximum number of pseudoknot-free complementary base pairings in subsequence $\mathbf{v}_i, \dots, \mathbf{v}_j$



Nussinov Algorithm – Dynamic Programming

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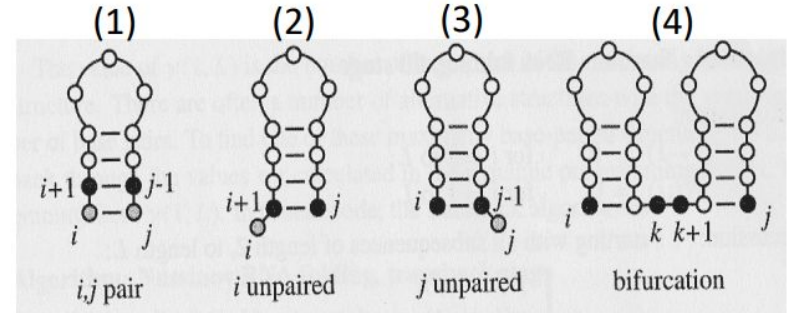
$$s[i, j] = \max \begin{cases} 0, & \text{if } i \geq j, \\ s[i + 1, j - 1] + 1, & \text{if } i < j \text{ and } (v_i, v_j) \in \Gamma, \quad (1) \\ s[i + 1, j - 1], & \text{if } i < j \text{ and } (v_i, v_j) \notin \Gamma, \quad (1^*) \\ s[i + 1, j], & \text{if } i < j, \quad (2) \\ s[i, j - 1], & \text{if } i < j, \quad (3) \\ \max_{i < k < j} \{s[i, k] + s[k + 1, j]\}, & \text{if } i < j, \quad (4) \end{cases}$$

$$\Gamma = \{(G, C), (C, G), (A, U), (U, A)\}$$

Nussinov Algorithm – Dynamic Programming

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Let $s[i, j]$ denote the maximum number of pseudoknot-free complementary base pairings in subsequence $\mathbf{v}_i, \dots, \mathbf{v}_j$



$$s[i, j] = \max \begin{cases} 0, & \text{if } i \geq j, \\ s[i + 1, j - 1] + 1, & \text{if } i < j \text{ and } (v_i, v_j) \in \Gamma, \text{ (1)} \\ s[i + 1, j - 1], & \text{if } i < j \text{ and } (v_i, v_j) \notin \Gamma, \text{ (1*)} \\ s[i + 1, j], & \text{if } i < j, \text{ (2)} \\ s[i, j - 1], & \text{if } i < j, \text{ (3)} \\ \max_{i < k < j} \{s[i, k] + s[k + 1, j]\}, & \text{if } i < j, \text{ (4)} \end{cases}$$

Question:
Which case is redundant?

Develop intuition

$$s[i, j] = \max \begin{cases} 0, & \text{if } i \geq j, \\ s[i+1, j-1] + 1, & \text{if } i < j \text{ and } (v_i, v_j) \in \Gamma, \\ s[i+1, j-1], & \text{if } i < j \text{ and } (v_i, v_j) \notin \Gamma, \\ s[i+1, j], & \text{if } i < j, \\ s[i, j-1], & \text{if } i < j, \\ \max_{i < k < j} \{s[i, k] + s[k+1, j]\}, & \text{if } i < j, \end{cases}$$

(((((- - - - -) - -))))																								
1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23		
G	C	U	C	G	G	G	U	U	C	C	C	U	A	U	U	C	A	A	G	A	G	C		
0																							G	1
0	0																						C	2
0	0	0																					U	3
0	0	0	0																				C	4
0	0	0	0	0																			G	5
0	0	0	0	0	0																		G	6
0	0	0	0	0	0	0																	G	7
0	0	0	0	0	0	0	0																U	8
0	0	0	0	0	0	0	0	0															U	9
0	0	0	0	0	0	0	0	0	0														C	10
0	0	0	0	0	0	0	0	0	0	0													C	11
0	0	0	0	0	0	0	0	0	0	0	0												C	12
0	0	0	0	0	0	0	0	0	0	0	0	0											U	13
0	0	0	0	0	0	0	0	0	0	0	0	0	0										A	14
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0									U	15
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0								U	16
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0							C	17
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0						A	18
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0					A	19
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0				G	20
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0			A	21
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		G	22
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	C	23

Develop intuition

$$s[i, j] = \max \begin{cases} 0, & \text{if } i \geq j, \\ s[i+1, j-1] + 1, & \text{if } i < j \text{ and } (v_i, v_j) \in \Gamma, \\ s[i+1, j-1], & \text{if } i < j \text{ and } (v_i, v_j) \notin \Gamma, \\ s[i+1, j], & \text{if } i < j, \\ s[i, j-1], & \text{if } i < j, \\ \max_{i < k < j} \{s[i, k] + s[k+1, j]\}, & \text{if } i < j, \end{cases}$$

[illegible]

Develop intuition

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(((((((-	-	-))	-	(-)	-	-))))			
1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23		
G	C	U	C	G	G	G	U	U	C	C	C	U	A	U	U	C	A	A	G	A	G	C		
0																							G	1
0	0																						C	2
0	0	0																					U	3
0	0	0	0																				C	4
0	0	0	0	0																			G	5
0	0	0	0	0	0																		G	6
0	0	0	0	0	0	0																	G	7
0	0	0	0	0	0	0	0																U	8
0	0	0	0	0	0	0	0	0															U	9
0	0	0	0	0	0	0	0	0	0														C	10
0	0	0	0	0	0	0	0	0	0	0													C	11
0	0	0	0	0	0	0	0	0	0	0	0												C	12
0	0	0	0	0	0	0	0	0	0	0	0	0											U	13
0	0	0	0	0	0	0	0	0	0	0	0	0	0										A	14
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0									U	15
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0								U	16
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0							C	17
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0						A	18
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0					A	19
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0				G	20
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0			A	21
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		G	22
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	C	23

Develop intuition

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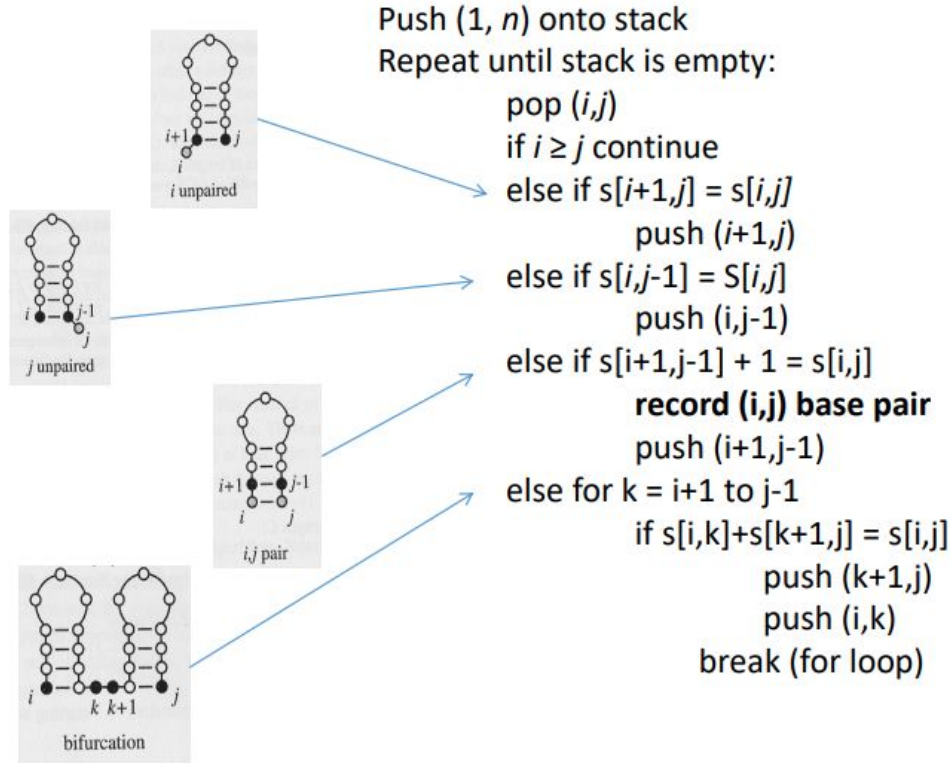
[illegible]

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[illegible]

Nussinov Algorithm – Traceback Step



Filling in the matrix

$$s[i, j] = \max \begin{cases} 0, & \text{if } i \geq j, \\ s[i + 1, j - 1] + 1, & \text{if } i < j \text{ and } (v_i, v_j) \in \Gamma, \\ s[i + 1, j - 1], & \text{if } i < j \text{ and } (v_i, v_j) \notin \Gamma, \\ s[i + 1, j], & \text{if } i < j, \\ s[i, j - 1], & \text{if } i < j, \\ \max_{i < k < j} \{s[i, k] + s[k + 1, j]\}, & \text{if } i < j, \end{cases}$$

		1		<i>j</i>						<i>n</i>
		G	G	G	A	A	A	U	C	C
<i>i</i>	1	G								
	G									
	G									
	A									
	A									
	A									
	U									
	C									
<i>n</i>	C									

Filling in the matrix

$$s[i, j] = \max \begin{cases} 0, & \text{if } i \geq j, \\ s[i + 1, j - 1] + 1, & \text{if } i < j \text{ and } (v_i, v_j) \in \Gamma, \\ s[i + 1, j - 1], & \text{if } i < j \text{ and } (v_i, v_j) \notin \Gamma, \\ s[i + 1, j], & \text{if } i < j, \\ s[i, j - 1], & \text{if } i < j, \\ \max_{i < k < j} \{s[i, k] + s[k + 1, j]\}, & \text{if } i < j, \end{cases}$$

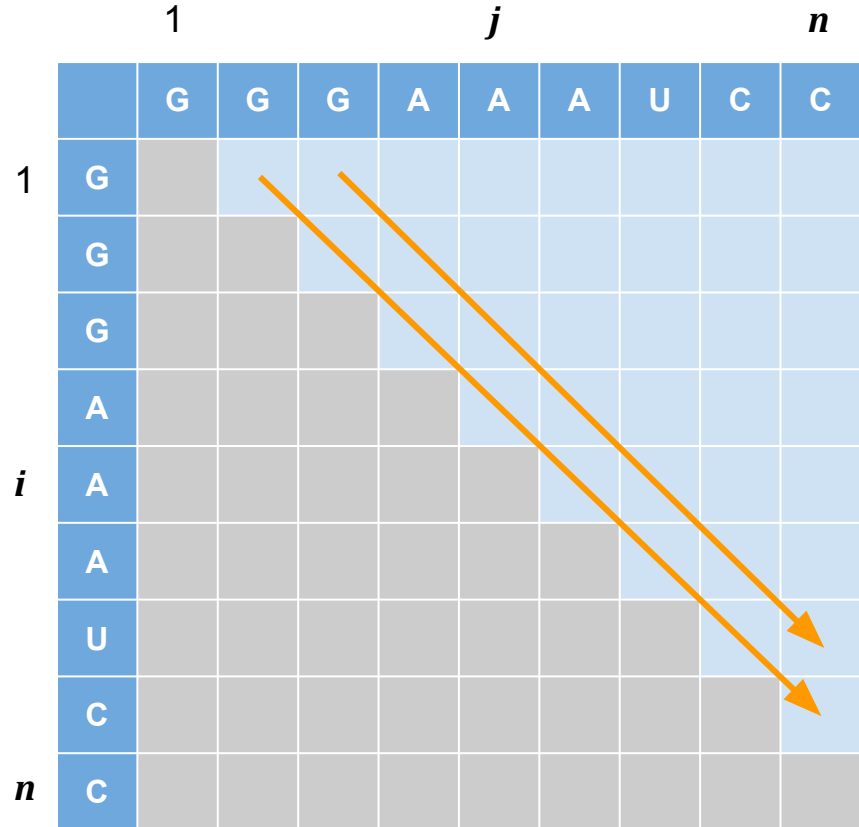
In order of increasing $j - i$

		1	<i>j</i>						<i>n</i>	
		G	G	G	A	A	A	U	C	C
1	G									
	G									
	G									
	A									
<i>i</i>	A									
	A									
	U									
	C									
<i>n</i>	C									

Filling in the matrix

$$s[i, j] = \max \begin{cases} 0, & \text{if } i \geq j, \\ s[i + 1, j - 1] + 1, & \text{if } i < j \text{ and } (v_i, v_j) \in \Gamma, \\ s[i + 1, j - 1], & \text{if } i < j \text{ and } (v_i, v_j) \notin \Gamma, \\ s[i + 1, j], & \text{if } i < j, \\ s[i, j - 1], & \text{if } i < j, \\ \max_{i < k < j} \{s[i, k] + s[k + 1, j]\}, & \text{if } i < j, \end{cases}$$

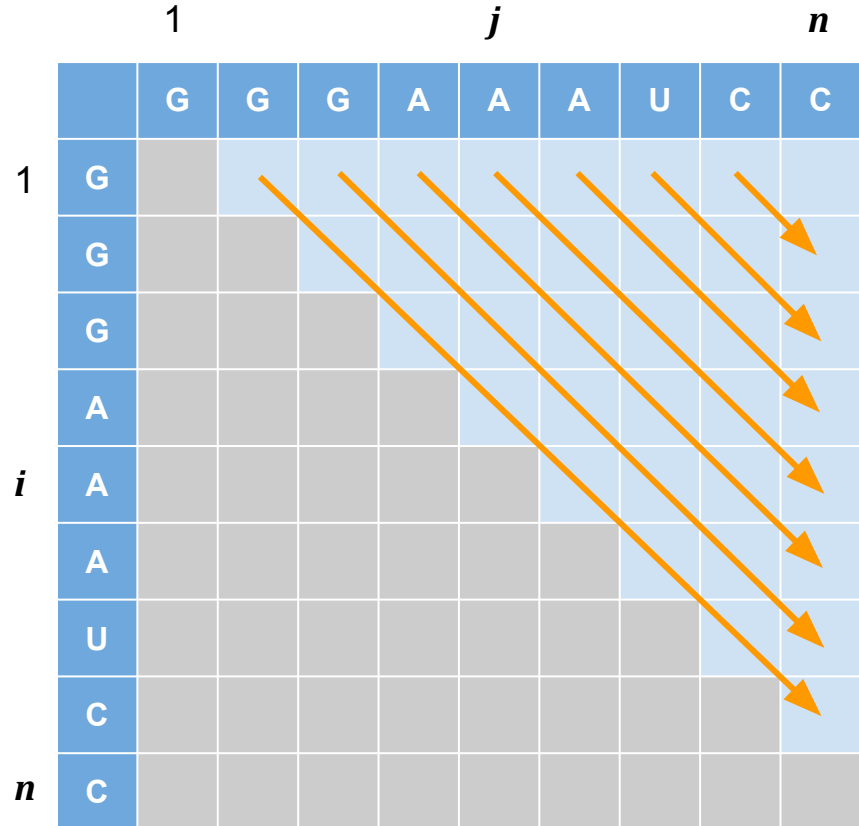
In order of increasing $j - i$



Filling in the matrix

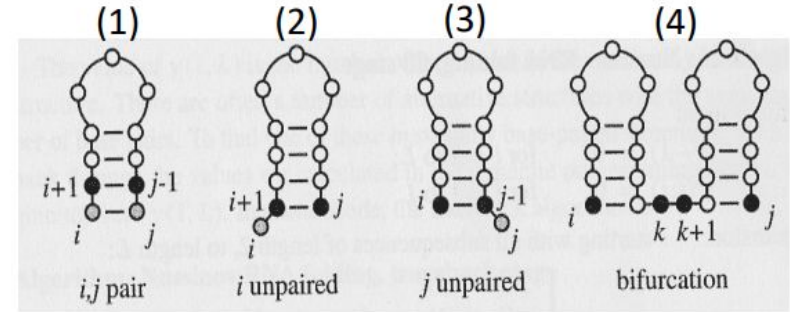
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In order of increasing $j - i$



Nussinov Algorithm – Example

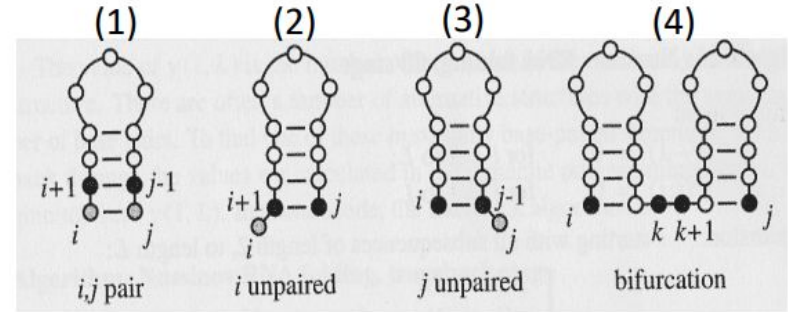
		<i>1</i>			<i>j</i>			<i>n</i>		
		G	G	G	A	A	A	U	C	C
<i>1</i>	G	0								
	G	0	0							
	G	0	0	0						
	A	0	0	0	0					
<i>i</i>	A	0	0	0	0	0				
	A	0	0	0	0	0	0			
	U	0	0	0	0	0	0	0		
	C	0	0	0	0	0	0	0	0	
<i>n</i>	C	0	0	0	0	0	0	0	0	0



$$s[i, j] = \max \begin{cases} 0, & \text{if } i \geq j, \\ s[i+1, j-1] + 1, & \text{if } i < j \text{ and } (v_i, v_j) \in \Gamma, \quad (1) \\ s[i+1, j], & \text{if } i < j \text{ and } (v_i, v_j) \notin \Gamma, \quad (1^*) \\ s[i+1, j], & \text{if } i < j, \quad (2) \\ s[i, j-1], & \text{if } i < j, \quad (3) \\ \max_{i < k < j} \{s[i, k] + s[k+1, j]\}, & \text{if } i < j, \quad (4) \end{cases}$$

Nussinov Algorithm – Example

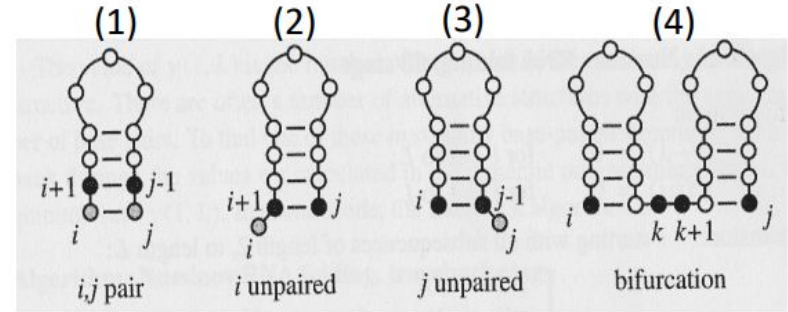
		<i>1</i>			<i>j</i>			<i>n</i>		
		G	G	G	A	A	A	U	C	C
<i>1</i>	G	0	0							
	G	0	0	0						
	G	0	0	0	0					
	A	0	0	0	0	0				
<i>i</i>	A	0	0	0	0	0	0			
	A	0	0	0	0	0	0	1		
	U	0	0	0	0	0	0	0	0	
	C	0	0	0	0	0	0	0	0	0
<i>n</i>	C	0	0	0	0	0	0	0	0	0



$$s[i, j] = \max \begin{cases} 0, & \text{if } i \geq j, \\ s[i+1, j-1] + 1, & \text{if } i < j \text{ and } (v_i, v_j) \in \Gamma, \text{ (1)} \\ s[i+1, j], & \text{if } i < j \text{ and } (v_i, v_j) \notin \Gamma, \text{ (1*)} \\ s[i, j-1], & \text{if } i < j, \text{ (2)} \\ \max_{i < k < j} \{s[i, k] + s[k+1, j]\}, & \text{if } i < j, \text{ (3) (4)} \end{cases}$$

Nussinov Algorithm – Example

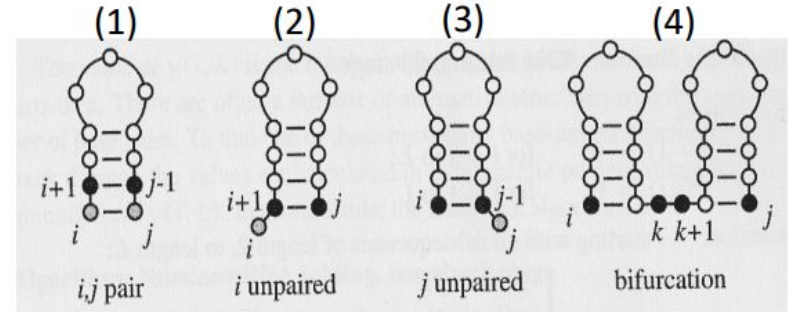
		<i>1</i>			<i>j</i>			<i>n</i>		
		G	G	G	A	A	A	U	C	C
<i>1</i>	G	0	0	0						
	G	0	0	0	0					
	G	0	0	0	0	0				
	A	0	0	0	0	0	0			
<i>i</i>	A	0	0	0	0	0	0	1		
	A	0	0	0	0	0	0	1	1	
	U	0	0	0	0	0	0	0	0	0
	C	0	0	0	0	0	0	0	0	0
<i>n</i>	C	0	0	0	0	0	0	0	0	0



$$s[i, j] = \max \begin{cases} 0, & \text{if } i \geq j, \\ s[i + 1, j - 1] + 1, & \text{if } i < j \text{ and } (v_i, v_j) \in \Gamma, \text{ (1)} \\ s[i + 1, j], & \text{if } i < j \text{ and } (v_i, v_j) \notin \Gamma, \text{ (1*)} \\ s[i, j - 1], & \text{if } i < j, \text{ (2)} \\ \max_{i < k < j} \{s[i, k] + s[k + 1, j]\}, & \text{if } i < j, \text{ (3) (4)} \end{cases}$$

Nussinov Algorithm – Example

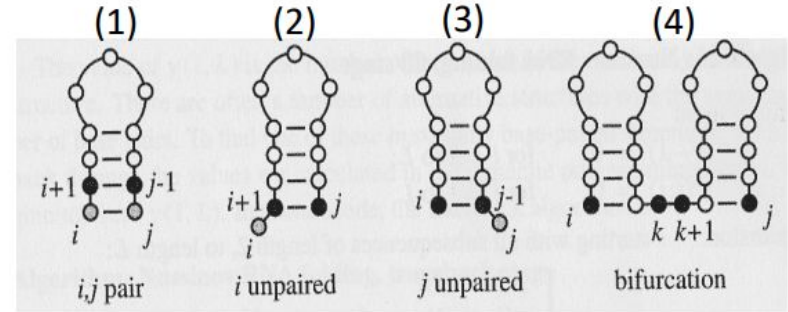
		<i>1</i>			<i>j</i>			<i>n</i>		
		G	G	G	A	A	A	U	C	C
<i>1</i>	G	0	0	0	0					
	G	0	0	0	0	0				
	G	0	0	0	0	0	0			
	A	0	0	0	0	0	0	1		
<i>i</i>	A	0	0	0	0	0	0	1	1	
	A	0	0	0	0	0	0	1	1	1
	U	0	0	0	0	0	0	0	0	0
	C	0	0	0	0	0	0	0	0	0
<i>n</i>	C	0	0	0	0	0	0	0	0	0



$$s[i, j] = \max \begin{cases} 0, & \text{if } i \geq j, \\ s[i + 1, j - 1] + 1, & \text{if } i < j \text{ and } (v_i, v_j) \in \Gamma, \text{ (1)} \\ s[i + 1, j - 1], & \text{if } i < j \text{ and } (v_i, v_j) \notin \Gamma, \text{ (1*)} \\ s[i + 1, j], & \text{if } i < j, \text{ (2)} \\ s[i, j - 1], & \text{if } i < j, \text{ (3)} \\ \max_{i < k < j} \{s[i, k] + s[k + 1, j]\}, & \text{if } i < j, \text{ (4)} \end{cases}$$

Nussinov Algorithm – Example

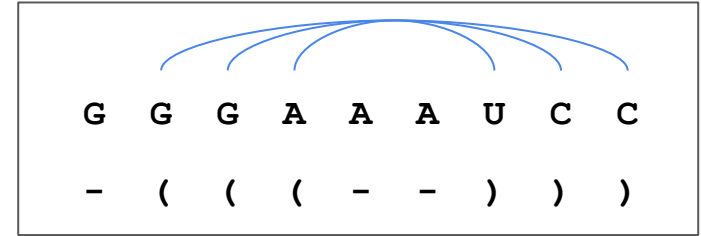
		<i>1</i>			<i>j</i>			<i>n</i>		
<i>1</i>		G	G	G	A	A	A	U	C	C
	G	0	0	0	0	0	0	1	2	3
	G	0	0	0	0	0	0	1	2	3
	G	0	0	0	0	0	0	1	2	2
<i>i</i>	A	0	0	0	0	0	0	1	1	1
	A	0	0	0	0	0	0	1	1	1
	A	0	0	0	0	0	0	1	1	1
	U	0	0	0	0	0	0	0	0	0
<i>n</i>	C	0	0	0	0	0	0	0	0	0
	C	0	0	0	0	0	0	0	0	0



$$s[i, j] = \max \begin{cases} 0, & \text{if } i \geq j, \\ s[i + 1, j - 1] + 1, & \text{if } i < j \text{ and } (v_i, v_j) \in \Gamma, \text{ (1)} \\ s[i + 1, j - 1], & \text{if } i < j \text{ and } (v_i, v_j) \notin \Gamma, \text{ (1*)} \\ s[i + 1, j], & \text{if } i < j, \text{ (2)} \\ s[i, j - 1], & \text{if } i < j, \text{ (3)} \\ \max_{i < k < j} \{s[i, k] + s[k + 1, j]\}, & \text{if } i < j, \text{ (4)} \end{cases}$$

Nussinov Algorithm – Example

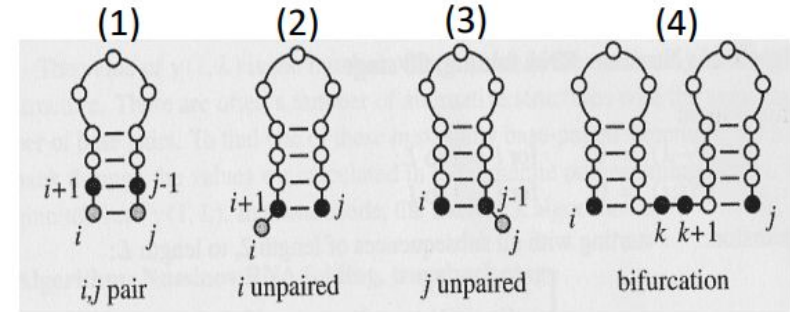
		<i>1</i>			<i>j</i>			<i>n</i>		
<i>1</i>		G	G	G	A	A	A	U	C	C
	G	0	0	0	0	0	0	1	2	3
	G	0	0	0	0	0	0	1	2	3
	G	0	0	0	0	0	0	1	2	2
	A	0	0	0	0	0	0	1	1	1
<i>i</i>	A	0	0	0	0	0	0	1	1	1
	A	0	0	0	0	0	0	1	1	1
	U	0	0	0	0	0	0	0	0	0
	C	0	0	0	0	0	0	0	0	0
<i>n</i>	C	0	0	0	0	0	0	0	0	0



Nussinov Algorithm – Example With Bifurcation

		<i>1</i>				<i>j</i>				<i>n</i>
		G	C	A	C	G	A	C	G	
<i>1</i>	G	0	1	1	1	2	2	2	3	
	C	0	0	0	0	1	1	1	2	
	A	0	0	0	0	1	1	1	2	
	C	0	0	0	0	1	1	1	2	
<i>i</i>	G	0	0	0	0	0	0	1	1	
	A	0	0	0	0	0	0	0	1	
	G	0	0	0	0	0	0	0	1	
	C	0	0	0	0	0	0	0	0	
<i>n</i>		0	0	0	0	0	0	0	0	

Where did we come from to get here?



$$s[i, j] = \max \begin{cases} 0, & \text{if } i \geq j, \\ s[i + 1, j - 1] + 1, & \text{if } i < j \text{ and } (v_i, v_j) \in \Gamma, \text{ (1)} \\ s[i + 1, j - 1], & \text{if } i < j \text{ and } (v_i, v_j) \notin \Gamma, \text{ (1*)} \\ s[i + 1, j], & \text{if } i < j, \text{ (2)} \\ s[i, j - 1], & \text{if } i < j, \text{ (3)} \\ \max_{i < k < j} \{s[i, k] + s[k + 1, j]\}, & \text{if } i < j, \text{ (4)} \end{cases}$$

GCACGACG

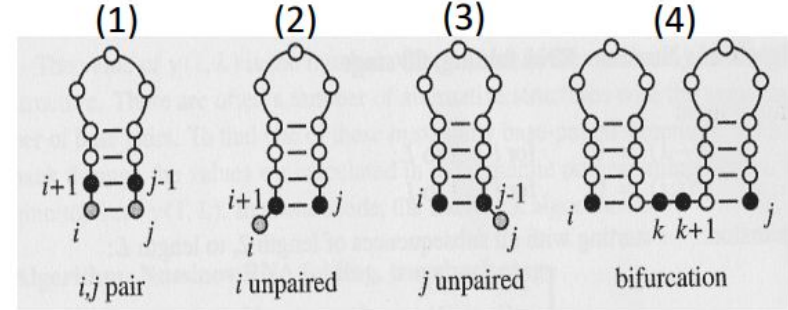
() . ((.))

Nussinov Algorithm – Example With Bifurcation

		<i>1</i>				<i>j</i>				<i>n</i>
		G	C	A	C	G	A	C	G	
<i>1</i>	G	0	1	1	1	2	2	2	3	
	C	0	0	0	0	1	1	1	2	
	A	0	0	0	0	1	1	1	2	
	C	0	0	0	0	1	1	1	2	
<i>i</i>	G	0	0	0	0	0	0	1	1	
	A	0	0	0	0	0	0	0	1	
	G	0	0	0	0	0	0	0	1	
	C	0	0	0	0	0	0	0	0	
<i>n</i>										

GCACGACG

() . ((.))



$$s[i, j] = \max \begin{cases} 0, & \text{if } i \geq j, \\ s[i+1, j-1] + 1, & \text{if } i < j \text{ and } (v_i, v_j) \in \Gamma, \text{ (1)} \\ s[i+1, j-1], & \text{if } i < j \text{ and } (v_i, v_j) \notin \Gamma, \text{ (1*)} \\ s[i+1, j], & \text{if } i < j, \text{ (2)} \\ s[i, j-1], & \text{if } i < j, \text{ (3)} \\ \max_{i < k < j} \{s[i, k] + s[k+1, j]\}, & \text{if } i < j, \text{ (4)} \end{cases}$$

Nussinov Algorithm – Alternative Solutions

	G	G	G	A	A	A	U	C	C
G	0	0	0	0	0	0	1	2	3
G	0	0	0	0	0	0	1	2	3
G	0	0	0	0	0	0	1	2	2
A	0	0	0	0	0	0	1	1	1
A	0	0	0	0	0	0	1	1	1
A	0	0	0	0	0	0	1	1	1
U	0	0	0	0	0	0	0	0	0
C	0	0	0	0	0	0	0	0	0
C	0	0	0	0	0	0	0	0	0



	G	G	G	A	A	A	U	C	C
G	0	0	0	0	0	0	1	2	3
G	0	0	0	0	0	0	1	2	3
G	0	0	0	0	0	0	1	2	2
A	0	0	0	0	0	0	1	1	1
A	0	0	0	0	0	0	1	1	1
A	0	0	0	0	0	0	1	1	1
U	0	0	0	0	0	0	0	0	0
C	0	0	0	0	0	0	0	0	0
C	0	0	0	0	0	0	0	0	0

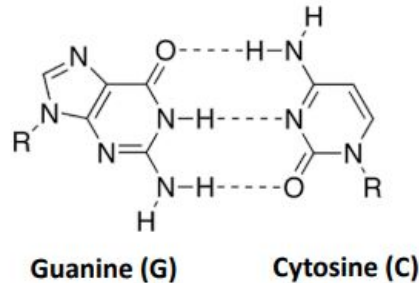
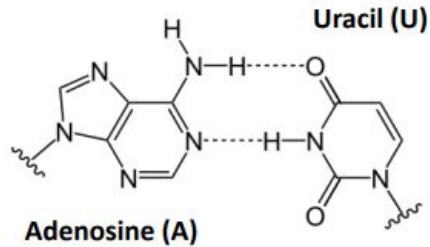
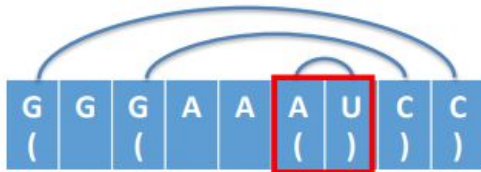


	G	G	G	A	A	A	U	C	C
G	0	0	0	0	0	0	1	2	3
G	0	0	0	0	0	0	1	2	3
G	0	0	0	0	0	0	1	2	2
A	0	0	0	0	0	0	1	1	1
A	0	0	0	0	0	0	1	1	1
A	0	0	0	0	0	0	1	1	1
U	0	0	0	0	0	0	0	0	0
C	0	0	0	0	0	0	0	0	0
C	0	0	0	0	0	0	0	0	0



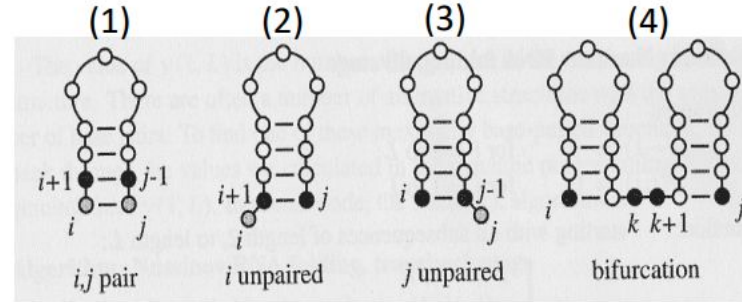
Does this make sense?

	G	G	G	A	A	A	U	C	C
G	0	0	0	0	0	0	1	2	3
G	0	0	0	0	0	0	1	2	3
G	0	0	0	0	0	0	1	2	2
A	0	0	0	0	0	0	1	1	1
A	0	0	0	0	0	0	1	1	1
A	0	0	0	0	0	0	1	1	1
U	0	0	0	0	0	0	0	0	0
C	0	0	0	0	0	0	0	0	0
C	0	0	0	0	0	0	0	0	0



Sharp loops are not preferred

Hairpin Loops with Minimum Length



$$s[i, j] = \max \begin{cases} 0, & \text{if } i + \ell \geq j, \\ s[i + 1, j - 1] + 1, & \text{if } i + \ell < j \text{ and } (v_i, v_j) \in \Gamma, \quad (1) \\ s[i + 1, j], & \text{if } i + \ell < j \text{ and } (v_i, v_j) \notin \Gamma, \quad (1^*) \\ s[i + 1, j], & \text{if } i + \ell < j, \quad (2) \\ s[i, j - 1], & \text{if } i + \ell < j, \quad (3) \\ \max_{i+\ell < k < j} \{s[i, k] + s[k + 1, j]\}, & \text{if } i + \ell < j, \quad (4) \end{cases}$$

A typical value of minimum loop length is 4

Time and space complexity

$$s[i, j] = \max \begin{cases} 0, & \text{if } i \geq j, \\ s[i + 1, j - 1] + 1, & \text{if } i < j \text{ and } (v_i, v_j) \in \Gamma, \\ s[i + 1, j - 1], & \text{if } i < j \text{ and } (v_i, v_j) \notin \Gamma, \\ s[i + 1, j], & \text{if } i < j, \\ s[i, j - 1], & \text{if } i < j, \\ \max_{i < k < j} \{s[i, k] + s[k + 1, j]\}, & \text{if } i < j, \end{cases}$$

- We have a subproblem for every interval (i, j)
- How many subproblems are there?

$$\binom{n}{2} = O(n^2)$$

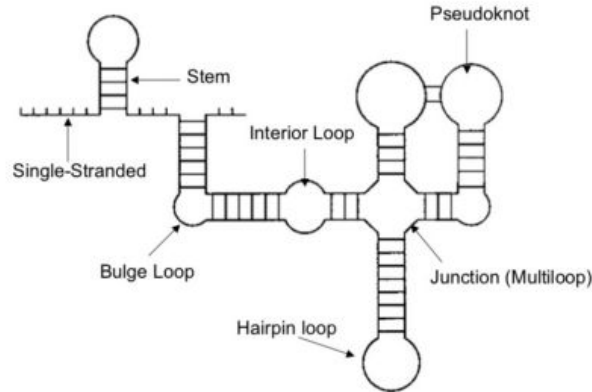
- Each takes $O(n)$ time to solve
 - have to search over all possible choices of k
- Total running time is $O(n^3)$
- Space complexity $O(n^2)$

RNA Secondary Structure Prediction in Practice

Rather than maximize number of compl. base pairs, minimize free energy (FE)

Zuker's algorithm: Dynamic programming w/ three matrices similar to affine gap penalties

- $V(i,j)$: FE of optimal structure of $s[i..j]$ assuming i,j form a base pair
- $VBI(i,j)$: FE of optimal structure of $s[i..j]$ assuming i,j closes a bulge or internal loop
- $VM(i,j)$: FE of optimal structure of $s[i..j]$ assuming i,j closes a multibranch loop



FE minimization with pseudoknots is NP-hard
[Lyngso and Pedersen, RECOMB 2000]

Summary

- RNA is a sequence of four bases/nucleotides {A, U, C, G}
- RNA folds into structures due to base/nucleotide complementarity
 - $A \leftrightarrow U$ and $C \leftrightarrow G$
- RNA secondary structure is defined by a set of non-overlapping complementary nucleotide pairs
- RNA folding rules:
 - If two bases are closer than 4 bases apart, they cannot pair (no sharp turns)
 - Each base is matched to at most one other base
 - The allowable pairs are {U, A} and {C, G}
 - Pairs cannot “cross.”
- Nussinov Algorithm: Dynamic programming to find pseudoknot-free structure with maximum number of complementary nucleotide pairs