

LGN0114 – Biologia Celular

Estrutura e Função dos Ácidos Nucleicos

No. 4356 April 25, 1953 NATURE 737

equipment, and to Dr. G. E. R. Deacon and the captain and officers of R.R.S. *Discovery II* for their part in making the observations.

¹ Young, T. R., G. E. R. Deacon, E. L., and J. J. W. V., *Phil. Mag.*, **46**, 149 (1950).

² Longuet-Higgins, H. S., *Nucl. Acids, Res., Suppl.*, **5**, 285 (1949).

³ Van der Meer, W. S., *Woods Hole Paper in Phys. Oceanog. Meteor.*, **11**, 151 (1960), 8.

⁴ Rhoads, V. W., *Arch. Met. Atmos. Phys. (Berkeley)*, **2** (11) (1955).

MOLECULAR STRUCTURE OF NUCLEIC ACIDS

A Structure for Deoxyribose Nucleic Acid
WE wish to suggest a structure for the salt of deoxyribose nucleic acid (D.N.A.). This structure has novel features which are of considerable biological interest.

A structure for nucleic acid has already been proposed by Pauling and Corey¹. They kindly made their manuscript available to us in advance of publication. Their model consists of three inter-twined chains, with the phosphates near the fibre axis, and the bases on the outside. In our opinion, this structure is unsatisfactory for two reasons: (1) We believe that the material which gives the X-ray diagrams is the salt, not the free acid. Without the acidic hydrogen atoms it is not clear what forces would hold the structure together, especially as the negatively charged phosphates near the axis will repel each other. (2) Some of the van der Waals distances appear to be too small.

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We wish to put forward a radically different structure for the salt of deoxyribose nucleic acid. This structure has two helical chains each coiled round the same axis (see diagram). We have made the usual chemical assumptions, namely, that each chain consists of phosphate di-ester groups joining 3',5'-deoxy-ribofuranose residues with 3',5' linkages. The two chains (but not their bases) are related by a dyad perpendicular to the fibre axis. Both chains follow right-handed helices, but owing to the dyad the sequences of the atoms in the two chains run in opposite directions. Each chain loosely resembles Furbert's model No. 1; that is, the bases are on the inside of the helix and the phosphates on the outside. The configuration of the sugar and the atoms near it is close to Furbert's 'standard configuration', the sugar being roughly perpendicular to the attached base. There

is a residue on each chain every 3.4 Å. in the z-direction. We have assumed an angle of 36° between adjacent residues in the same chain, so that the structure repeats after 10 residues on each chain, that is, after 34 Å. The distance of a phosphorus atom from the fibre axis is 10 Å. As the phosphates are on the outside, cations have easy access to them. The structure is an open one, and its water content is rather high. At lower water contents we would expect the bases to tilt so that the structure could become more compact.

The novel feature of the structure is the manner in which the two chains are held together by the purine and pyrimidine bases. The planes of the bases are perpendicular to the fibre axis. They are joined together in pairs, a single base from one chain being hydrogen-bonded to a single base from the other chain, so that the two lie side by side with identical z-co-ordinates. One of the pair must be a purine and the other a pyrimidine for bonding to occur. The hydrogen bonds are made as follows: purine position 1 to pyrimidine position 1; purine position 6 to pyrimidine position 6.

If it is assumed that the bases only occur in the structure in the most plausible tautomeric forms (that is, with the keto rather than the enol configurations) it is found that only specific pairs of bases can bond together. These pairs are: adenine (purine) with thymine (pyrimidine), and guanine (purine) with cytosine (pyrimidine).

In other words, if an adenine forms one member of a pair, on either chain, then on these assumptions the other member must be thymine; similarly for guanine and cytosine. The sequence of bases on a single chain does not appear to be restricted in any way. However, if only specific pairs of bases can be formed, it follows that if the sequence of bases on one chain is given, then the sequence on the other chain is automatically determined.

It has been found experimentally^{2,3} that the ratio of the amounts of adenine to thymine, and the ratio of guanine to cytosine, are always very close to unity for deoxyribose nucleic acid.

It is probably impossible to build this structure with a ribose sugar in place of the deoxyribose, as the extra oxygen atom would make too close a van der Waals contact.

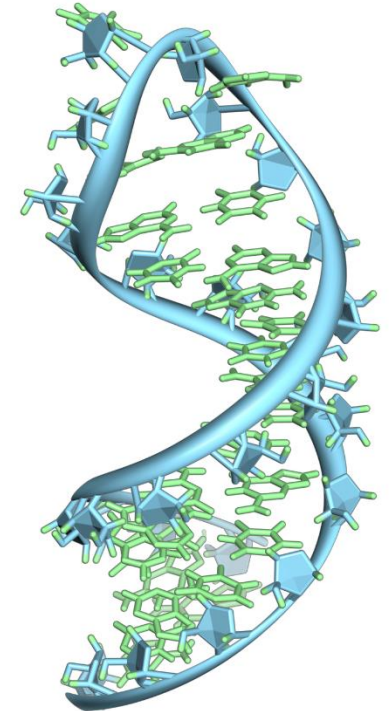
The previously published X-ray data^{4,5} on deoxyribose nucleic acid are insufficient for a rigorous test of our structure. So far as we can tell, it is roughly compatible with the experimental data, but it must be regarded as unproved until it has been checked against more exact results. Some of these are given in the following communications. We were not aware of the details of the results presented there when we devised our structure, which rests mainly though not entirely on published experimental data and stereochemical arguments.

It has not escaped our notice that the specific pairing we have postulated immediately suggests a possible copying mechanism for the genetic material. Full details of the structure, including the conditions assumed in building it, together with a set of co-ordinates for the atoms, will be published elsewhere.

We are much indebted to Dr. Jerry Donohue for constant advice and criticism, especially on inter-atomic distances. We have also been stimulated by a knowledge of the general nature of the unpublished experimental results and ideas of Dr. M. H. F. Wilkins, Dr. R. E. Franklin and their co-workers at



This figure is partly diagrammatic; the two chains are represented by two phosphate-sugar chains, and the helix is shown as a pair of lines. The vertical line marks the fibre axis.

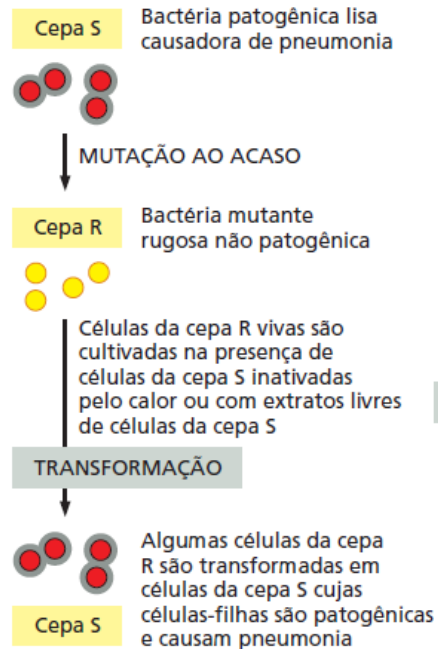


Aula 4

Antonio Figueira
figueira@cena.usp.br

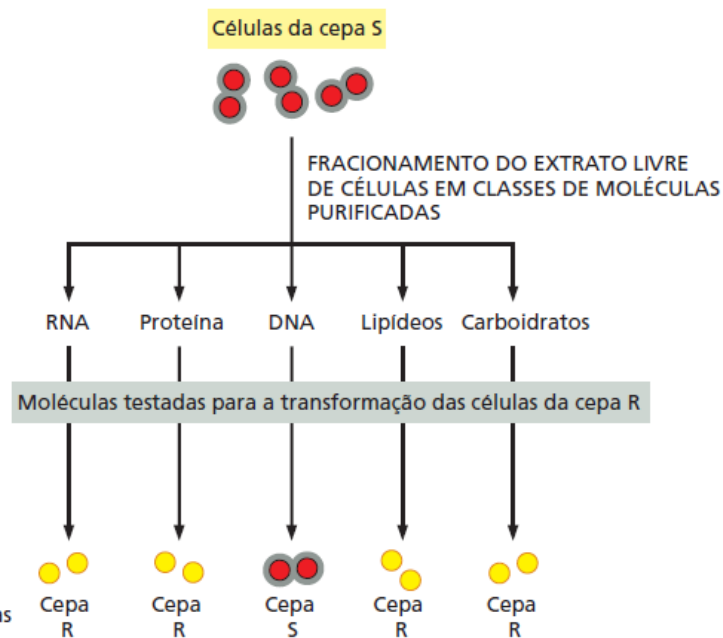
DNA como material genético

Fred Griffith 1929



CONCLUSÃO: as moléculas que podem transmitir as informações hereditárias estão presentes nas células da cepa S.

(A)



CONCLUSÃO: a molécula que carrega a informação hereditária é o DNA.

(B)

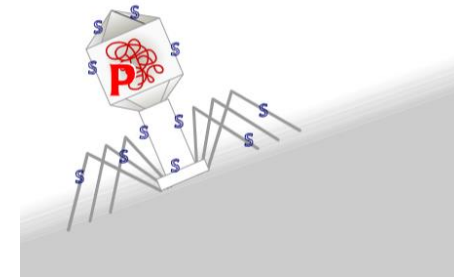


Courtesy of the Rockefeller Archive Center.
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Oswald Avery
Colin MacLeod
Maclyn McCarty
1944



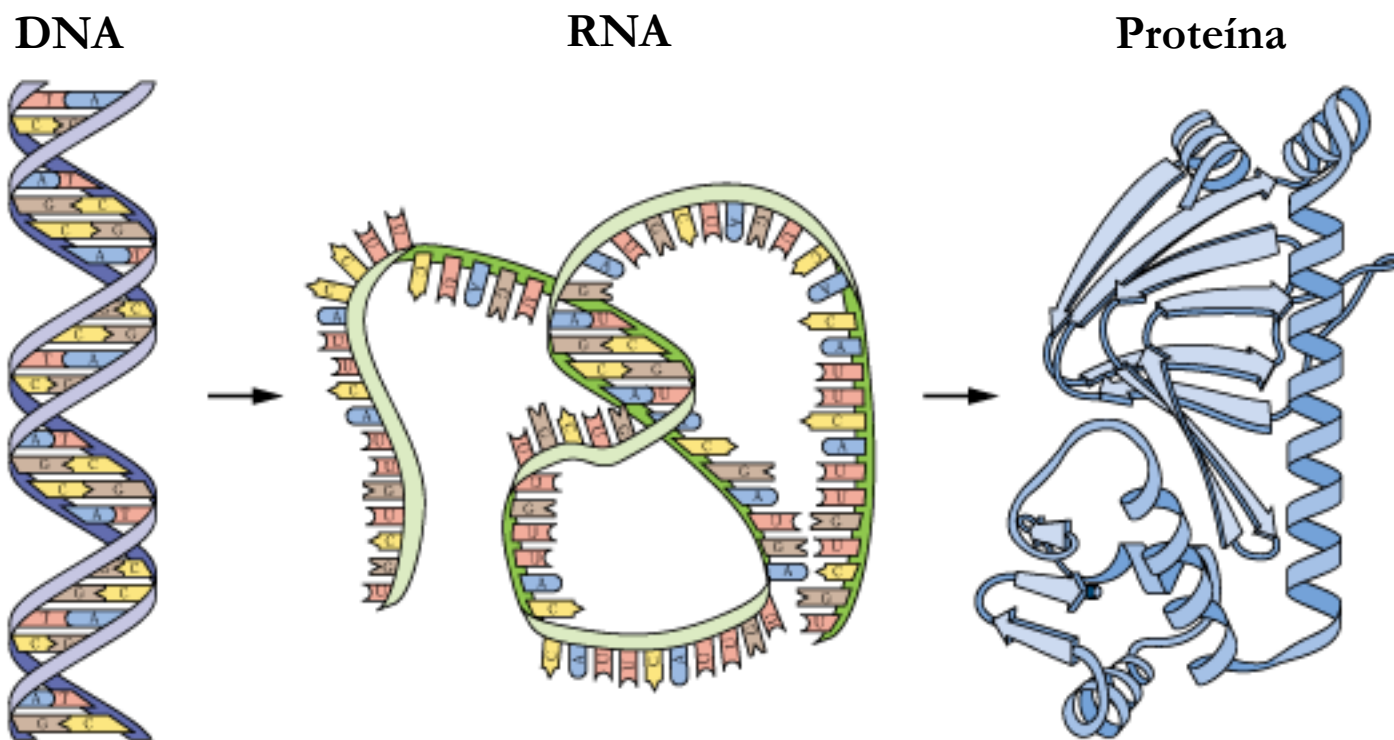
Alfred Hershey 1952



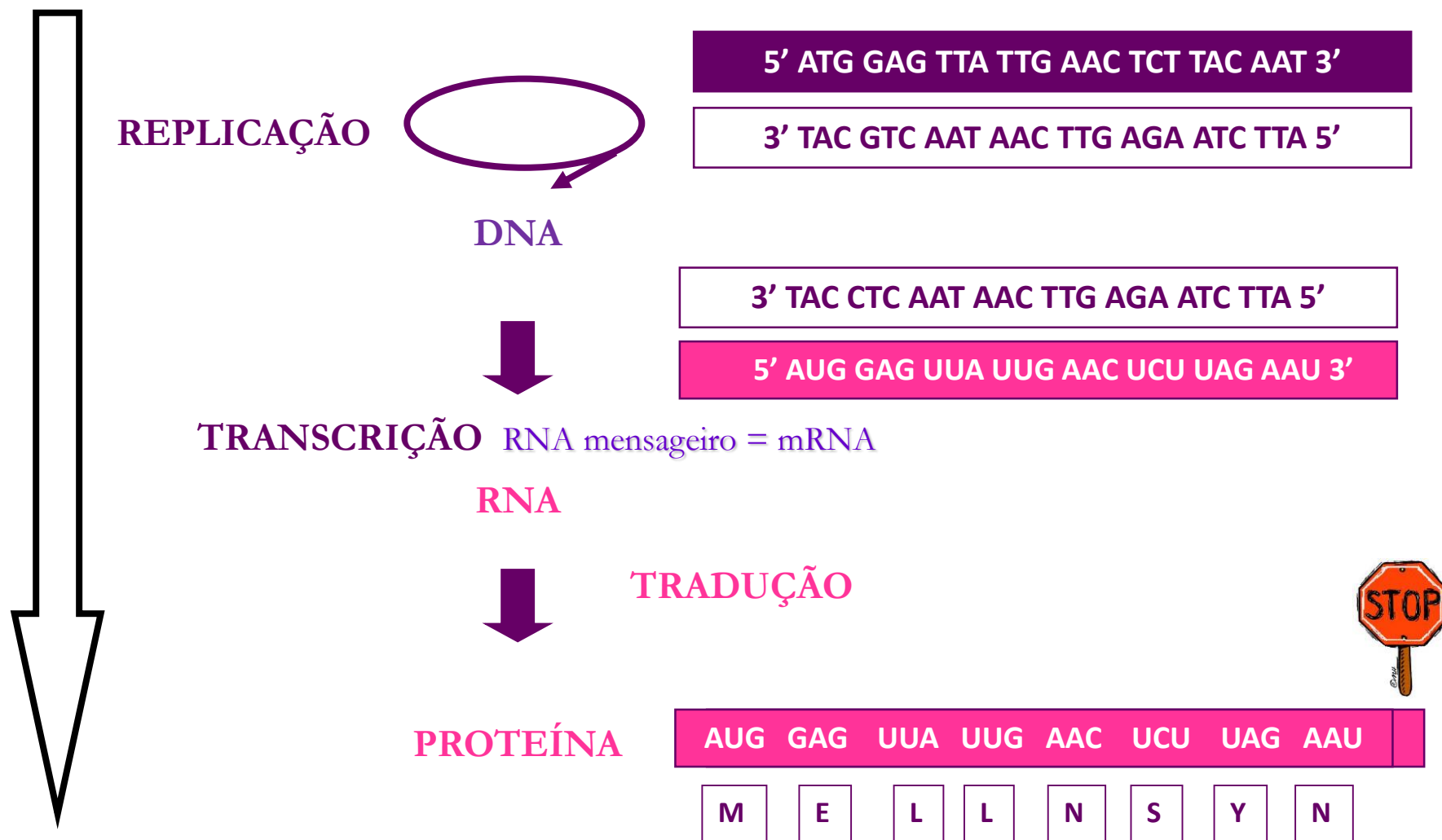
<http://www.dnaftb.org/17/animation.html>

DOGMA CENTRAL DA BIOLOGIA

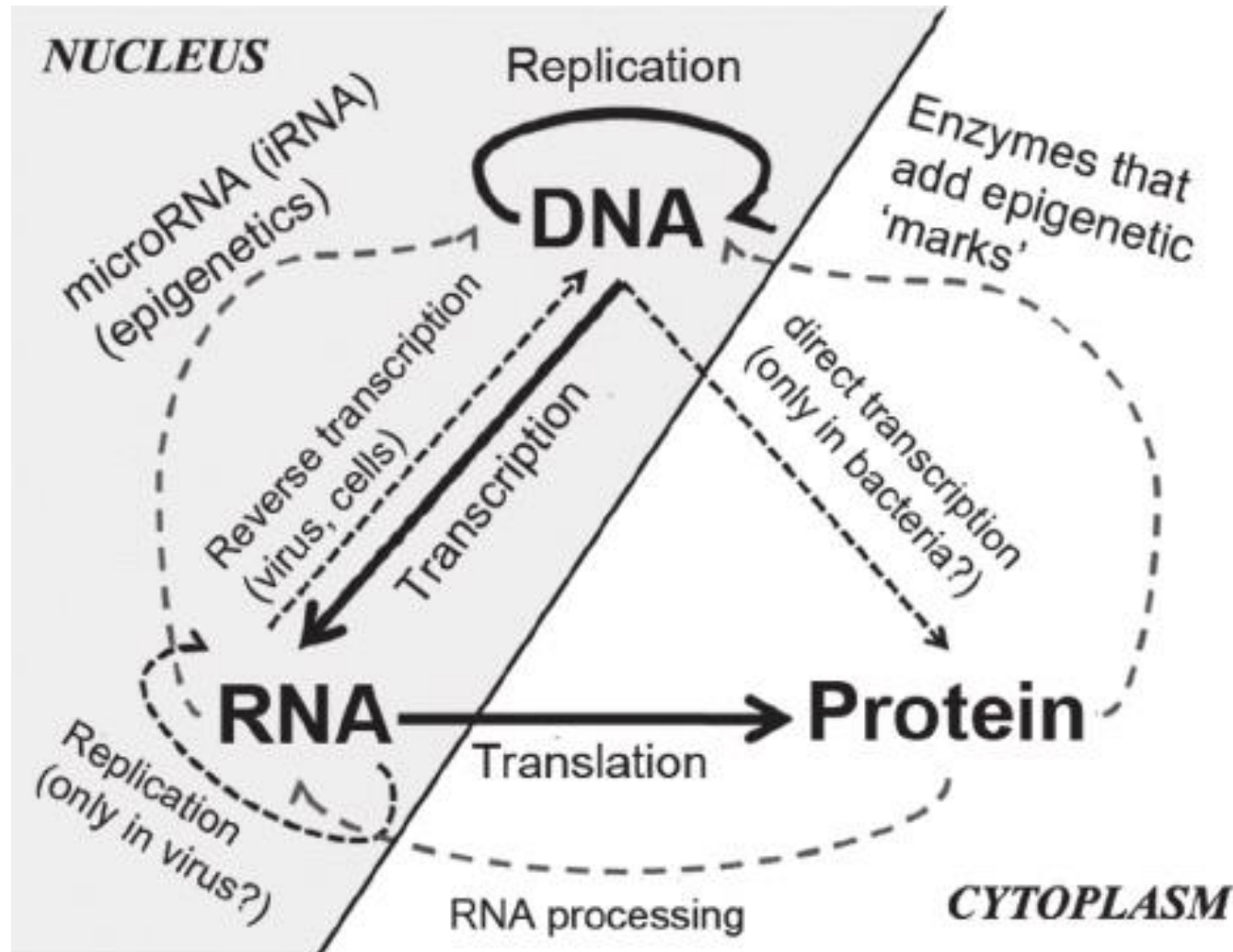
A informação genética, armazenada nos cromossomos, é transferida às células filhas através da **replicação do DNA**, sendo expressa através da **transcrição em mRNA** e traduzida subsequentemente em cadeias polipeptídicas (**proteínas**)



Fluxo da Informação Genética



Fluxo da Informação Genética



Dogma que não é Dogma!

Função do Material Genético

1. Função genotípica: replicação

2. Função fenotípica: expressão gênica

3. Função evolutiva: variabilidade
(mutações, transferência horizontal de genes...)

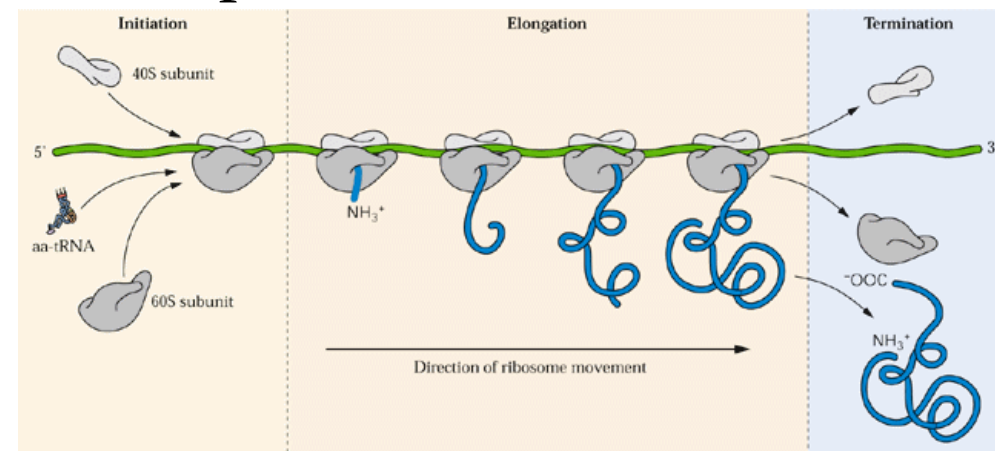
Ácidos Nucleicos

- **DNA:** Armazenamento da informação genética
 - **Estabilidade**
- **RNA:** síntese de macromoléculas - várias funções
 - **RNA ribossomal (rRNA)** - componentes estruturais de ribossomos
 - **RNA mensageiro (mRNA)** - contém a informação genética para a sequência de aminoácidos das proteínas
 - **RNA transferência (tRNA)** - identifica e transporta os aminoácidos até o ribossomo
 - **snRNA, snoRNA, microRNA, ncRNA**

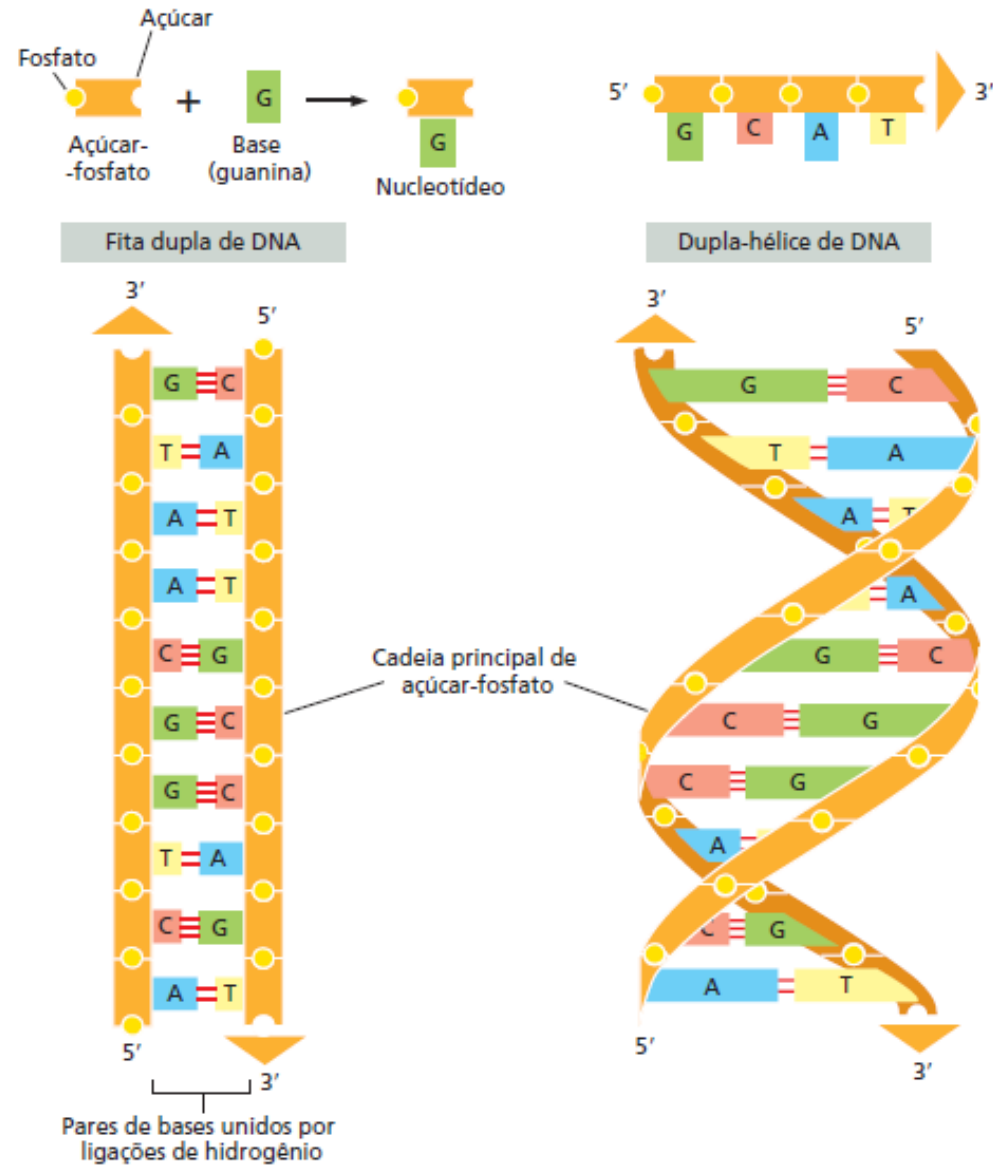
snRNA = *small nuclear RNA* - spliceossomos

snoRNA = *small nucleolar RNA* – montagem ribossomo

ncRNA = *non-coding RNA* – RNA regulatórios

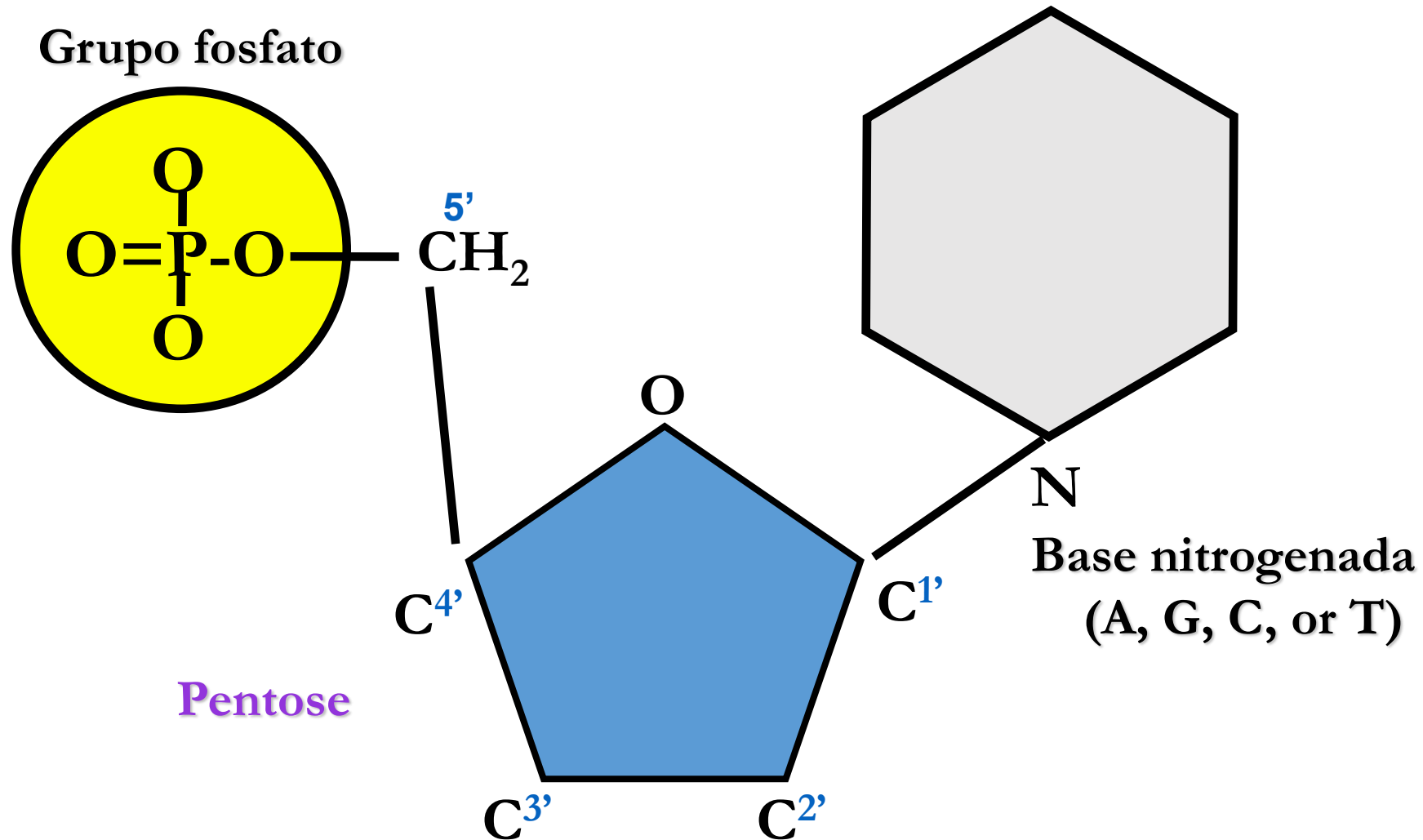


DNA



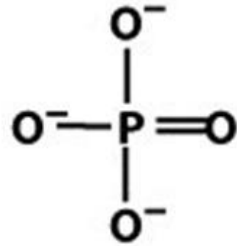
ÁCIDOS NUCLEICOS

São polímeros de nucleotídeos



Componentes dos Nucleotídeos

(1)
Um
grupo
fosfato:

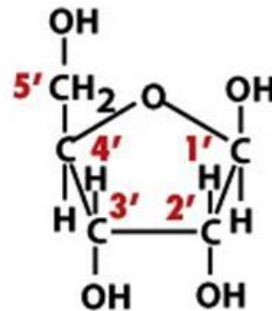


Carbono 5'

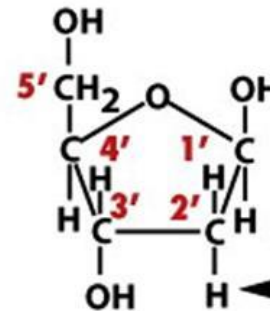
Nucleotídeo sem fosfato é nucleosídeo!!!

(2)
pentoses
(açúcares
de 5
carbonos)

(a) RNA:
Ribose



(b) DNA:
2-Desoxirribose

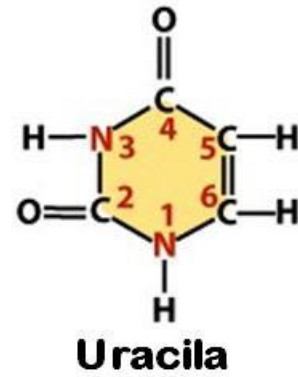


Carbono 2'

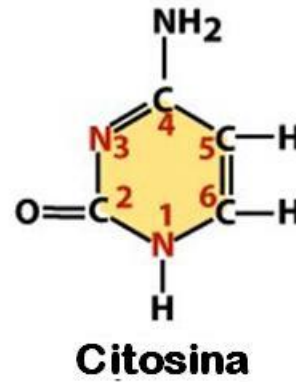
← Sem grupo hidroxila

(3)
Uma base
cíclica
contendo
Nitrogênio

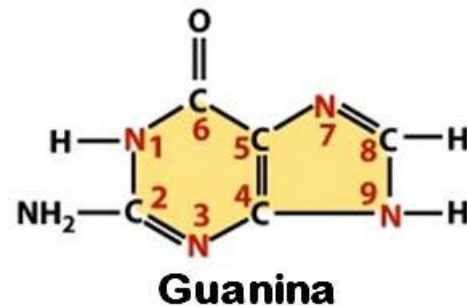
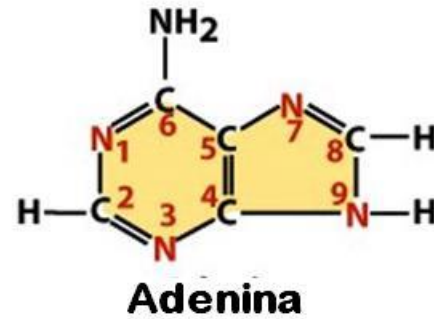
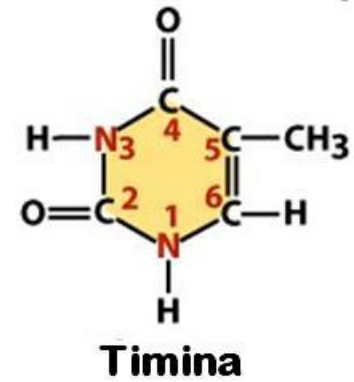
(a) RNA



(b) DNA e RNA

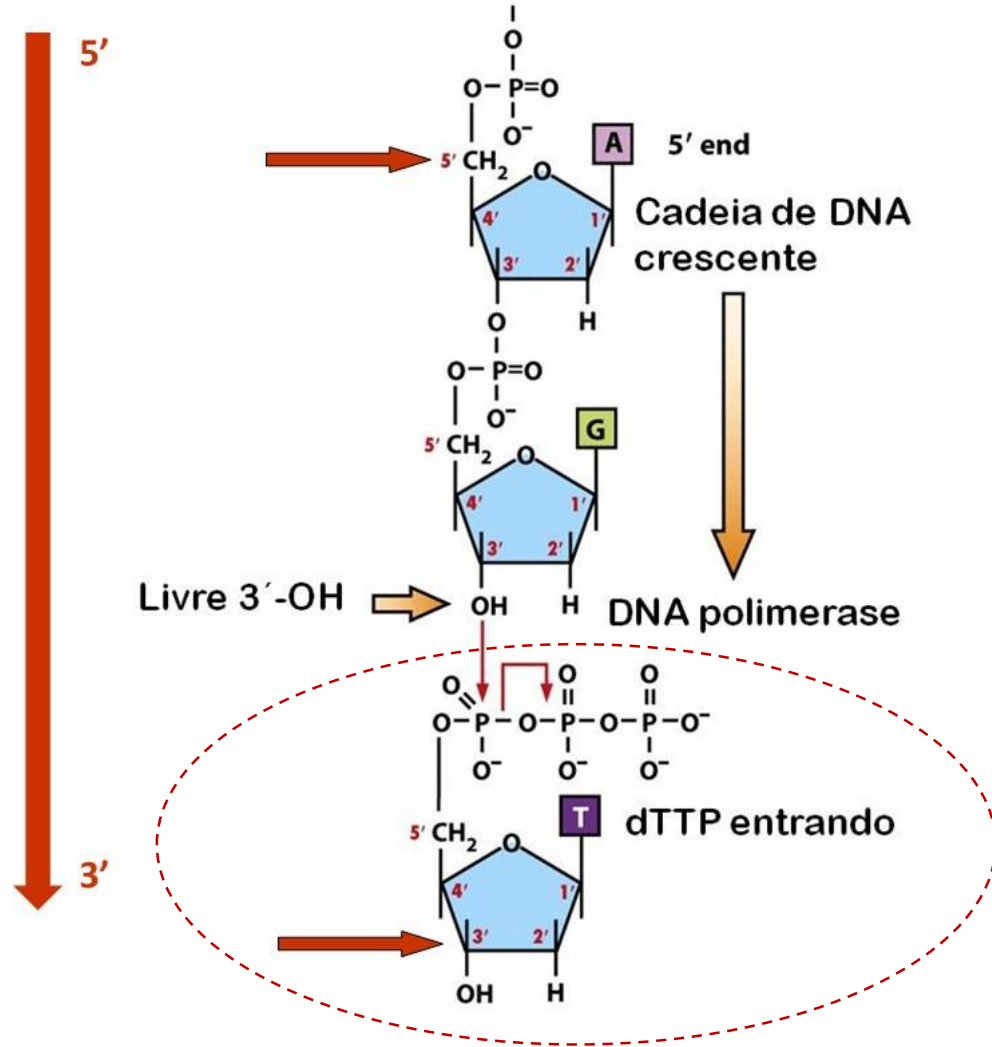


(c) DNA



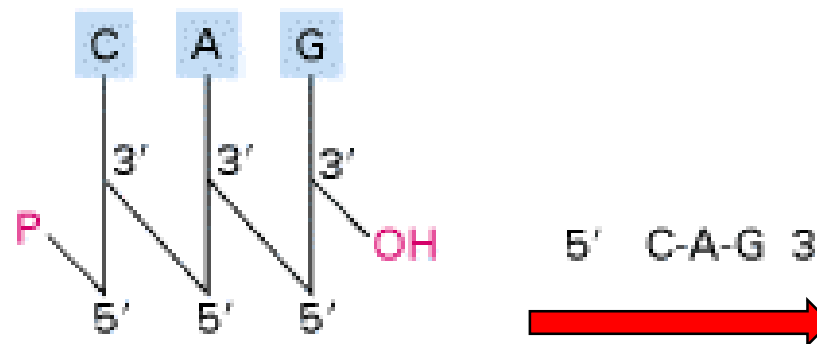
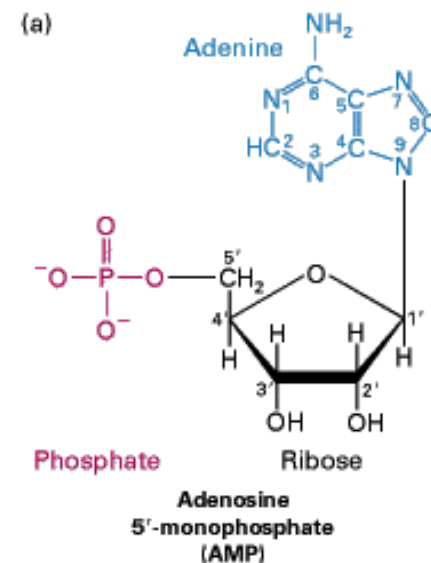
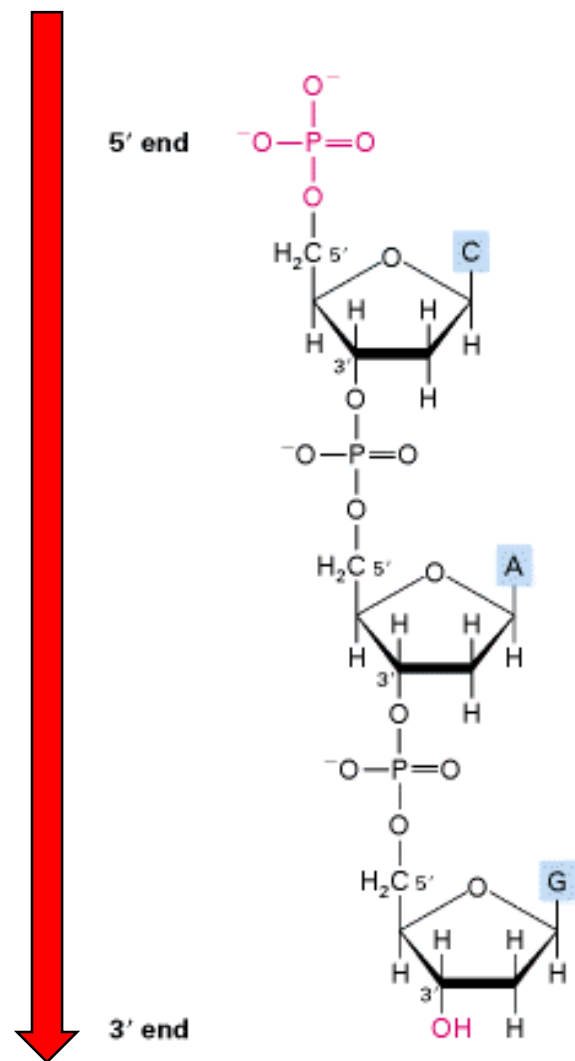
Purinas: A, G
Pirimidinas: U, T, C

Ácidos Nucleicos são formados por ligações fosfodiester



Nucleotídeo livre está na forma de **trifosfato**

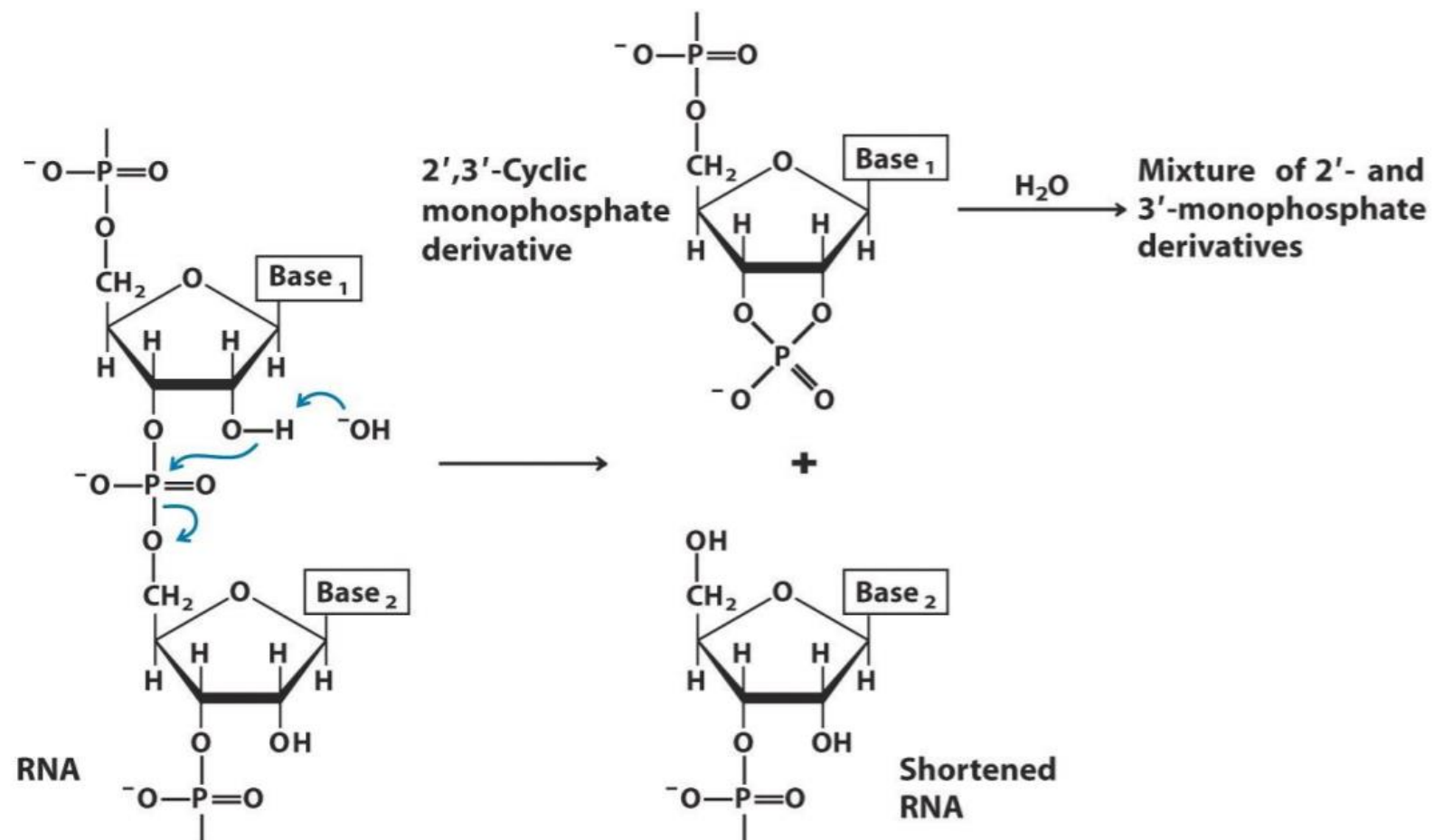
Ligações fosfodiéster - polarização 5' – 3'



- entre o carbono 3' do nucleotídeo de “cima” e o carbono 5' do nucleotídeo de “baixo”

Porque o RNA é mais lábil?

Efeito de pH alcalino



Porque o Timina (DNA) x Uracil (RNA)?

Deaminação de Citosina a Uracil

Ocorre naturalmente

Afeta o significado da sequência

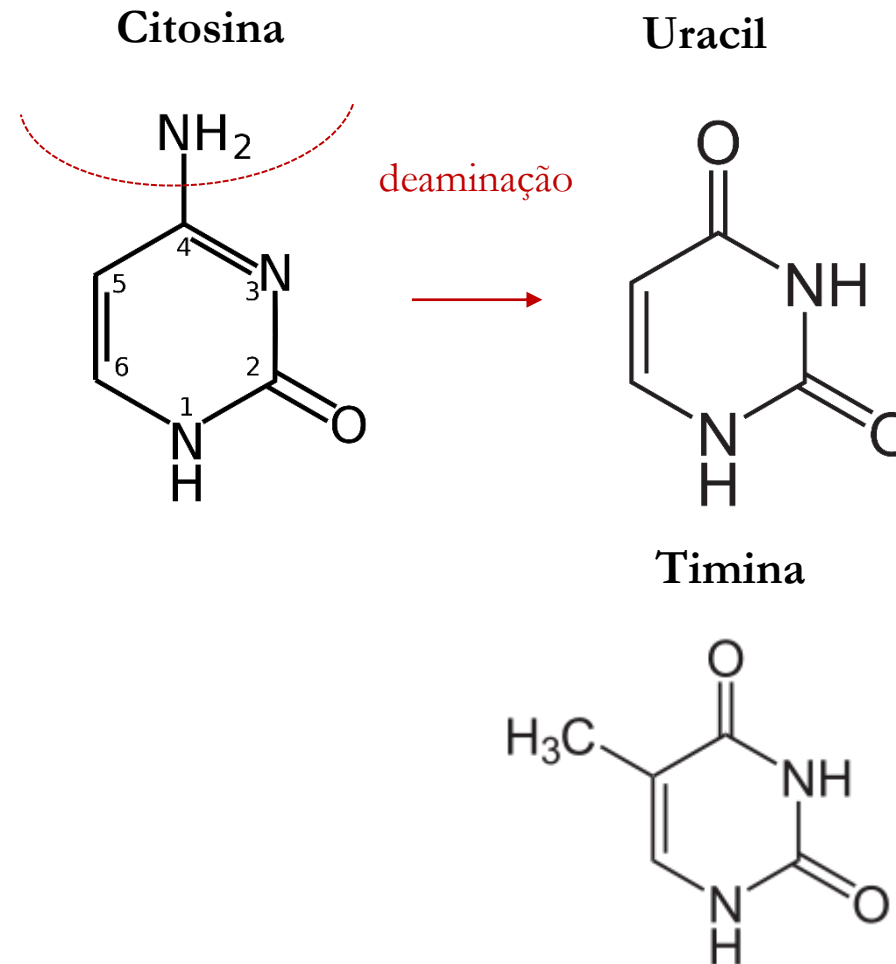
Perda de amina em C $\rightarrow$ U

Reconhecido em DNA

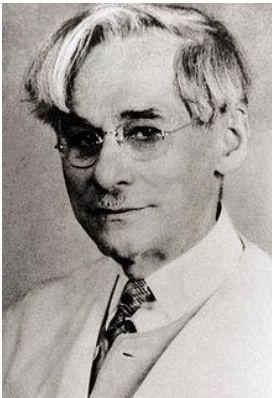
Pareamento em DNA

C – G passaria a ser U-G

Na replicação incorreria em erro



Hipótese dos Tetranucleotídeos



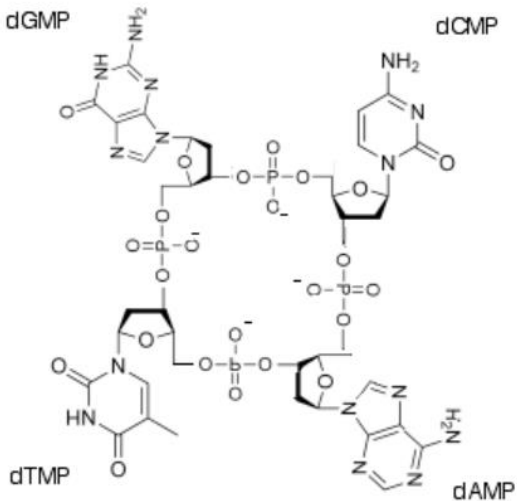
Phoebus Levene



Erwin Chargaff

Regra de Chargaff

DNA SOURCE	ADENINE	THYMINE	GUANINE	CYTOSINE
Calf Thymus	1.7	1.6	1.2	1.0
Beef Spleen	1.6	1.5	1.3	1.0
Yeast	1.8	1.9	1.0	1.0
Tubercle Bacillus	1.1	1.0	2.6	2.4



equipment, and to Dr. G. E. R. Deacon and the captain and officers of R.R.S. *Discovery II* for their part in making the observations.

- ¹Young, F. B., Gerrard, H., and Jevons, W., *Phil. Mag.*, **40**, 149 (1920).
- ²Longuet-Higgins, M. S., *Mon. Not. Roy. Astr. Soc., Geophys. Supp.*, **5**, 255 (1949).
- ³Von Arx, W. S., *Woods Hole Papers in Phys. Oceanogr. Meteor.*, **11** (5) (1950).
- ⁴Elmann, V. W., *Arkiv. Mat. Astron. Fysik. (Stockholm)*, **2** (11) (1955).

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This figure is purely diagrammatic. The two ribbons symbolize the two phosphate-sugar chains, and the horizontal rods the pairs of bases holding the chains together. The vertical line marks the fibre axis.

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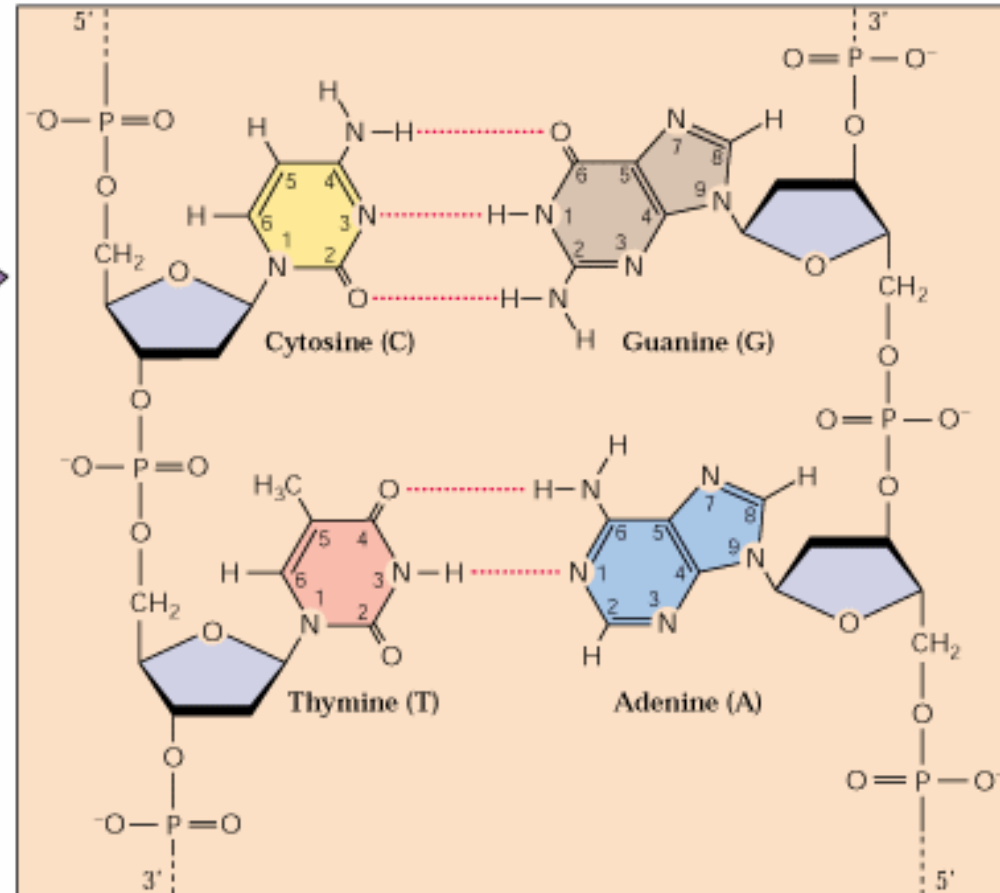
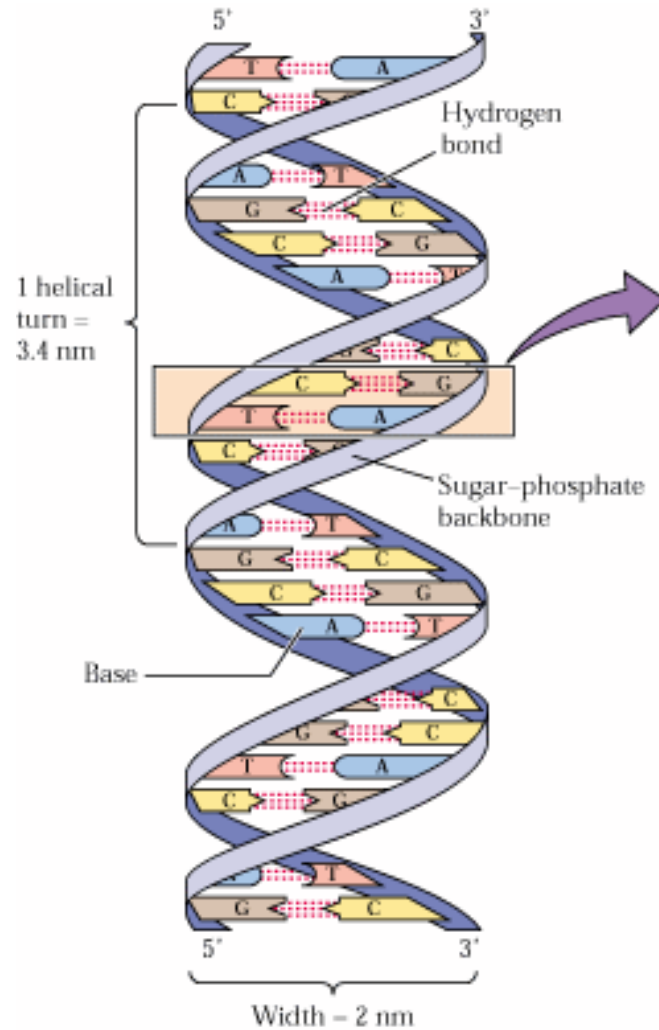
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DNA - Ácido Desoxirribonucleico



DNA – Fita Dupla!

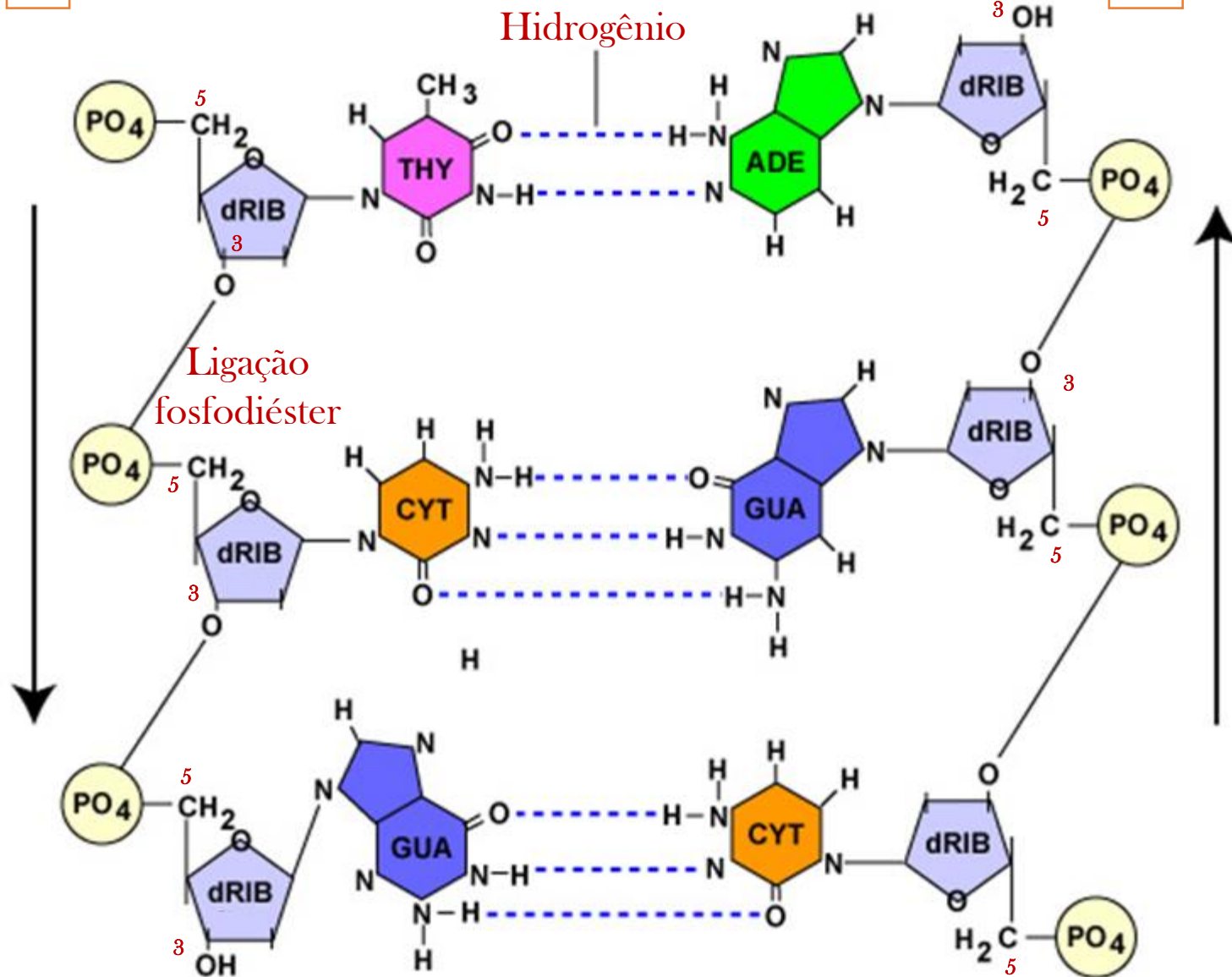
- Duas cadeias independentes
- Dupla hélice, sentido direito
- Hélices **antiparalelas**
- Complementariedade das bases
 - Eixo externo **hidrofílico** - deoxiribose + fosfato
 - Bases hidrofóbicas (planas) no interior
 - Bases ligadas por pontes de H

5'

3'

Ponte de
Hidrogênio

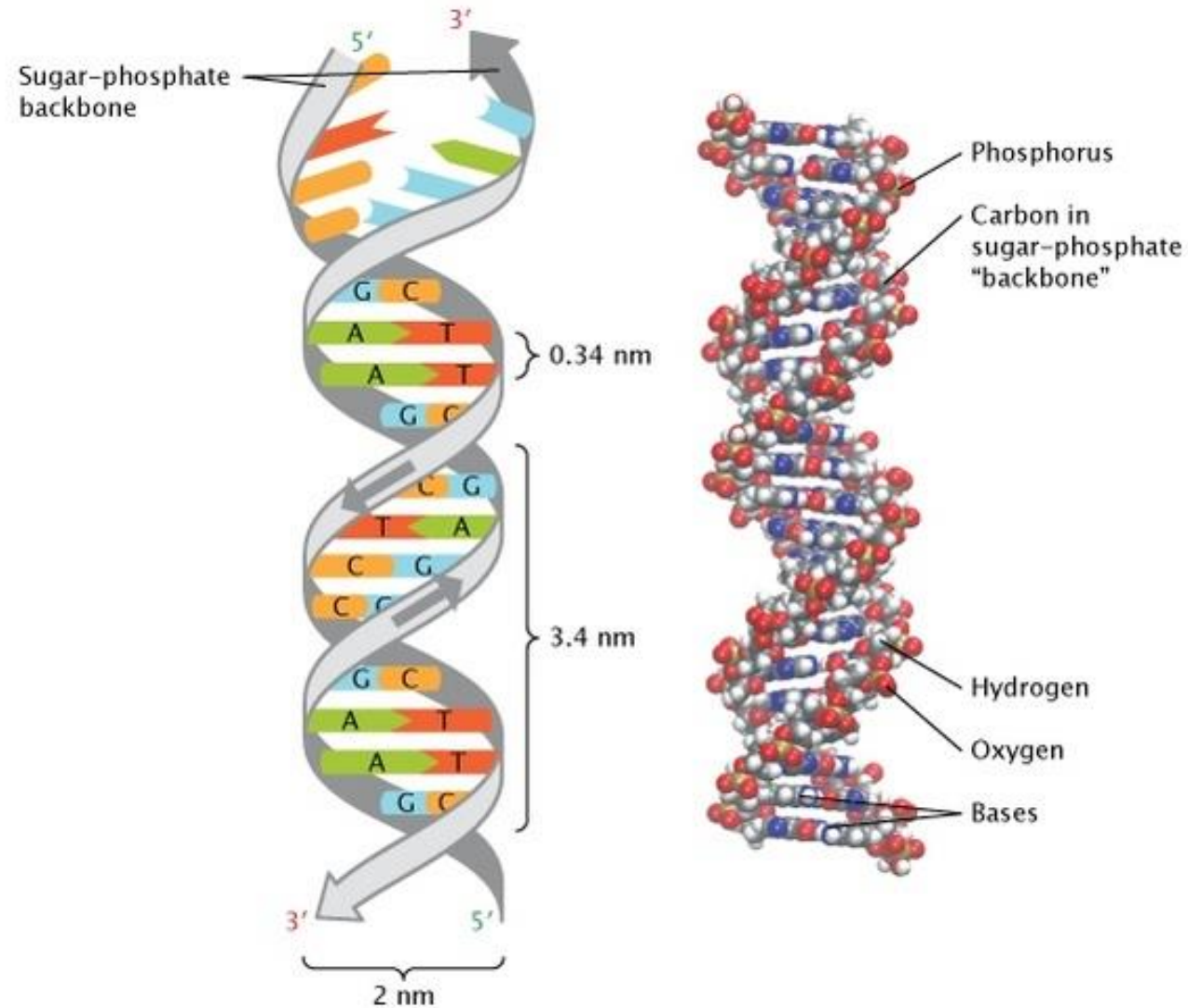
Ligação
fosfodiéster



3'

5

DNA – Fita Dupla!



Características da Dupla Hélice

- Contém duas fitas de polinucleotídeos (“corrimão”) **antiparalelas**
- O esqueleto de cada fita é formado por desoxirribose e fosfato
- O grupo fosfato ligado ao carbono 5’ de uma desoxirribose se liga **covalentemente** ao terminal hidroxila do carbono 3’ da próxima unidade
- As purinas e pirimidinas estão voltadas para dentro da hélice
- Cada base forma **pontes de H** com uma base oposta a ela, formando um par de bases
- **3,4 Å** separam os planos (“degraus”), onde bases adjacentes estão localizadas
- A dupla hélice faz uma volta completa com **10 nucleotídeos (34 Å)**
- Existem em média 25 pontes de H dentro de cada volta completa da hélice, promovendo uma **estabilidade de ligação tão forte** como uma ligação covalente
- O diâmetro da hélice é cerca de 20 Å

Principais Tipos de RNA

RNAs ocorrem no núcleo e citoplasma

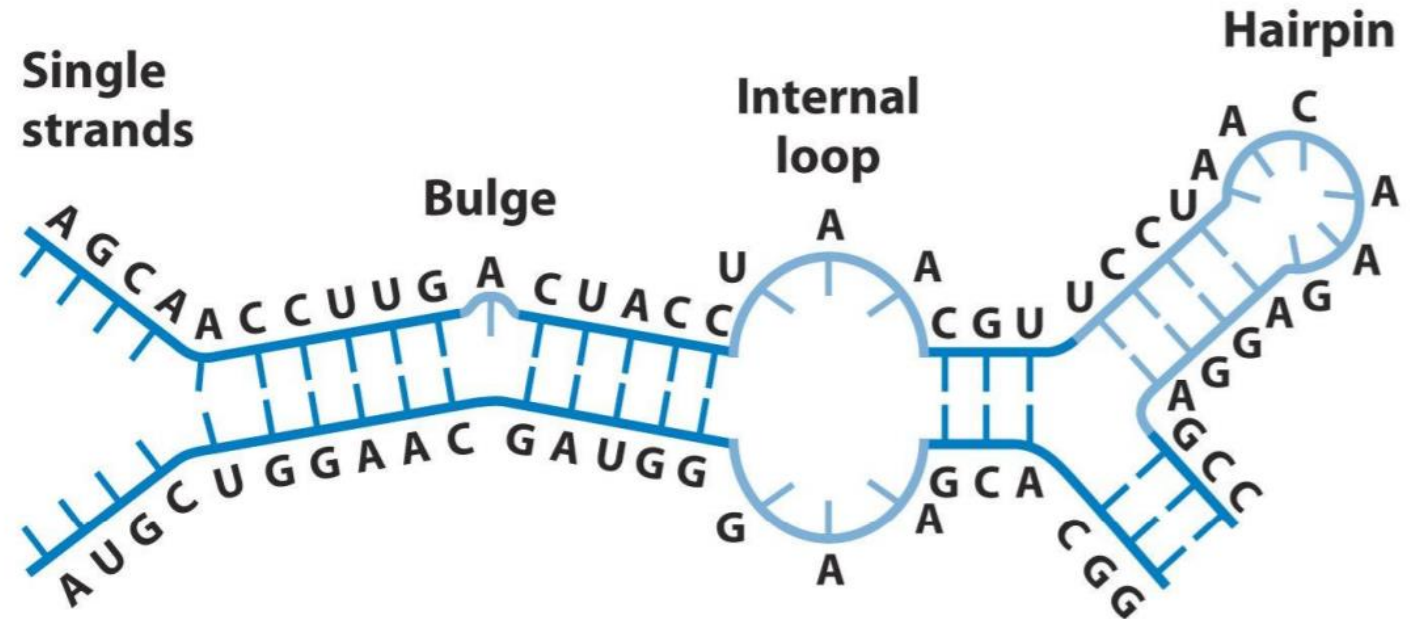
RNA mensageiro (mRNA): contém a informação genética para a sequência de aminoácidos das proteínas

RNA transferência (tRNA): identifica e transporta os aminoácidos até o ribossomo

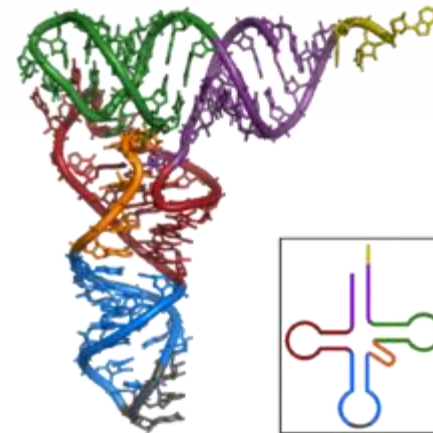
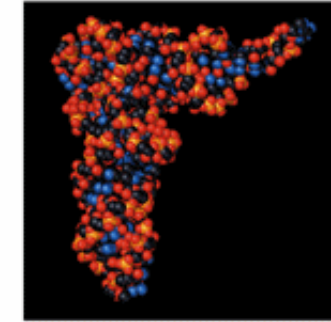
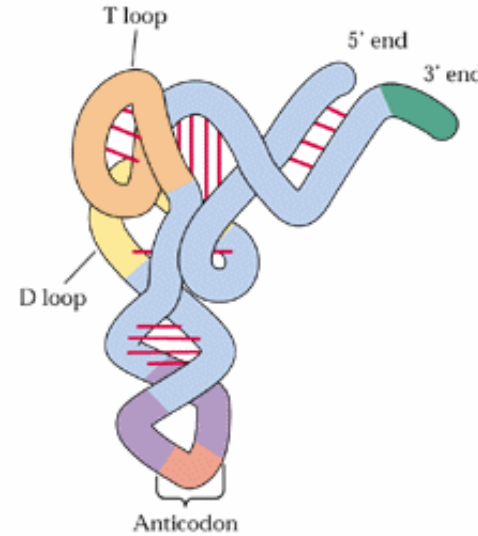
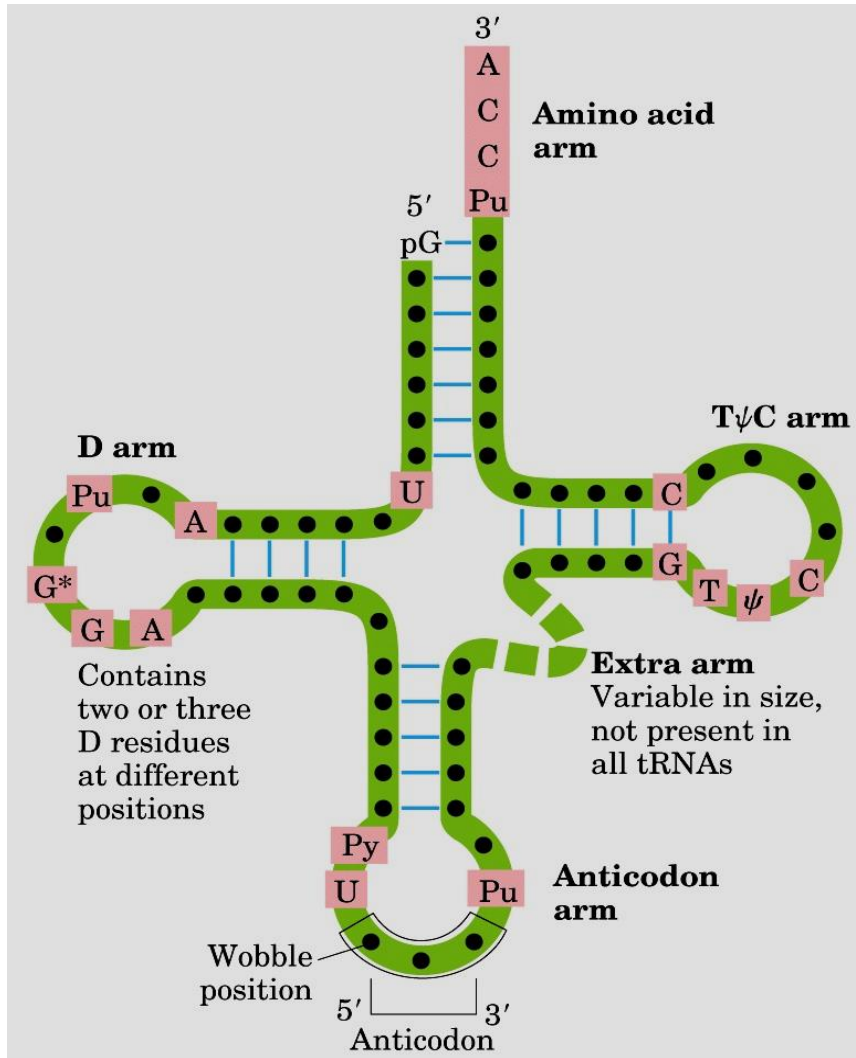
RNA ribossômico (rRNA): constituinte dos ribossomos

Estrutura de RNA

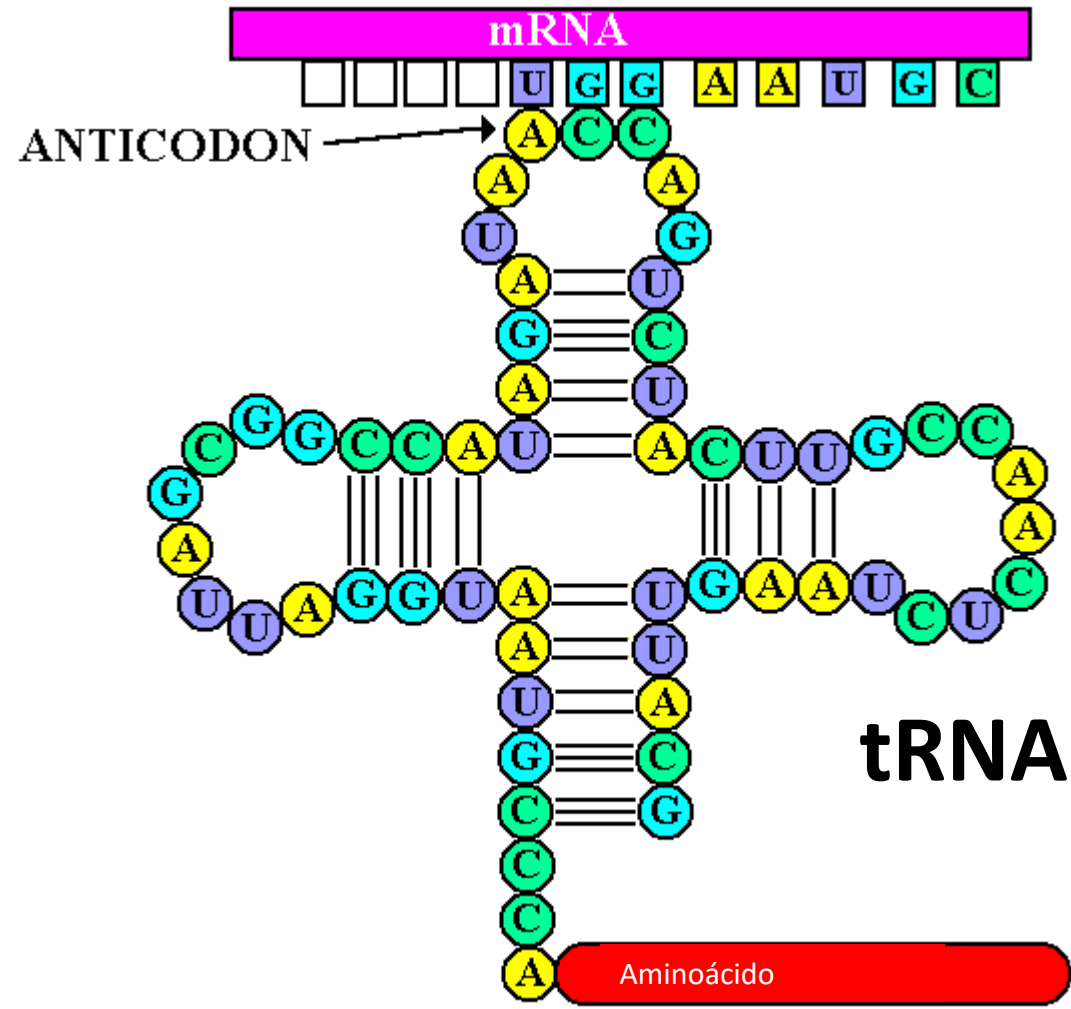
- RNA não possuem estrutura regular como DNA
- As estruturas de RNA são complexas e únicas
- Pareamento de bases similar a DNA forma estruturas



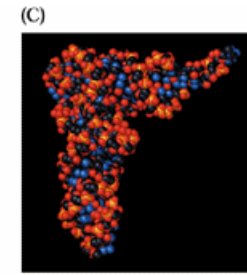
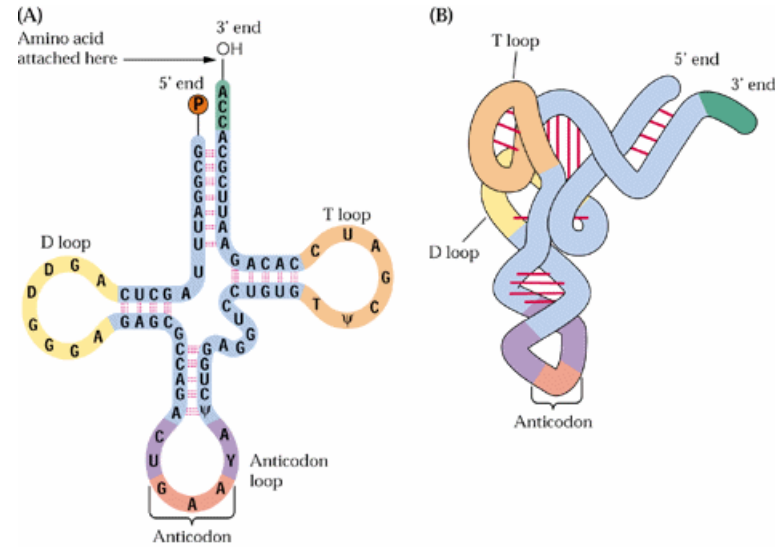
RNA transportador - tRNA



Reconhece códons em mRNA - anticódon

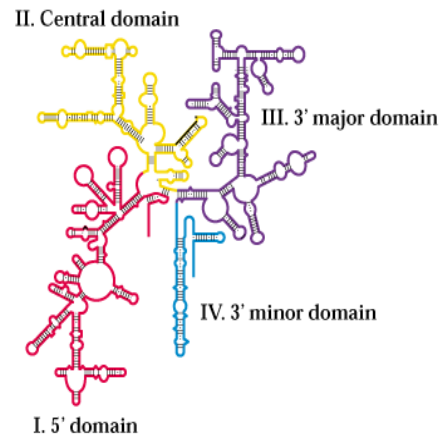


tRNA

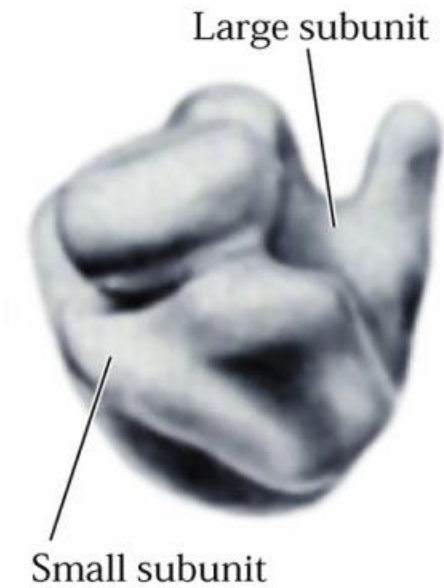
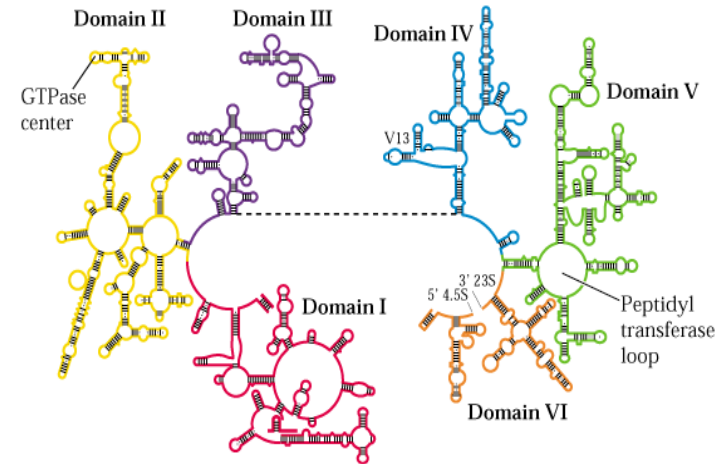


rRNA

(A) Tobacco plastid 16S rRNA

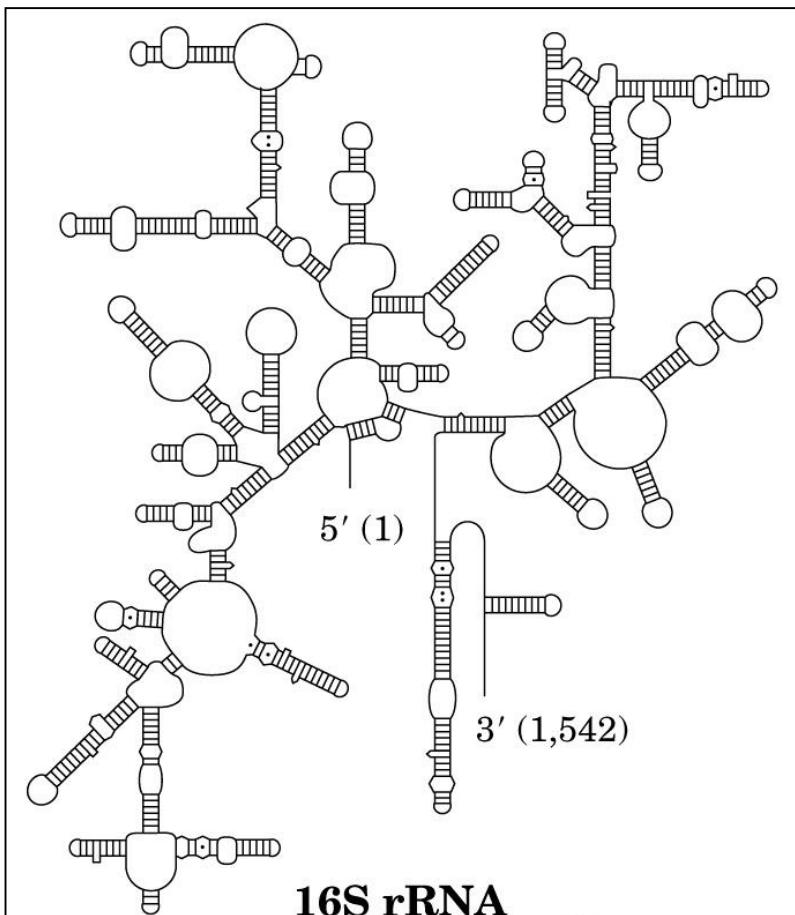


(B) Tobacco plastid 23S and 4.5S rRNAs



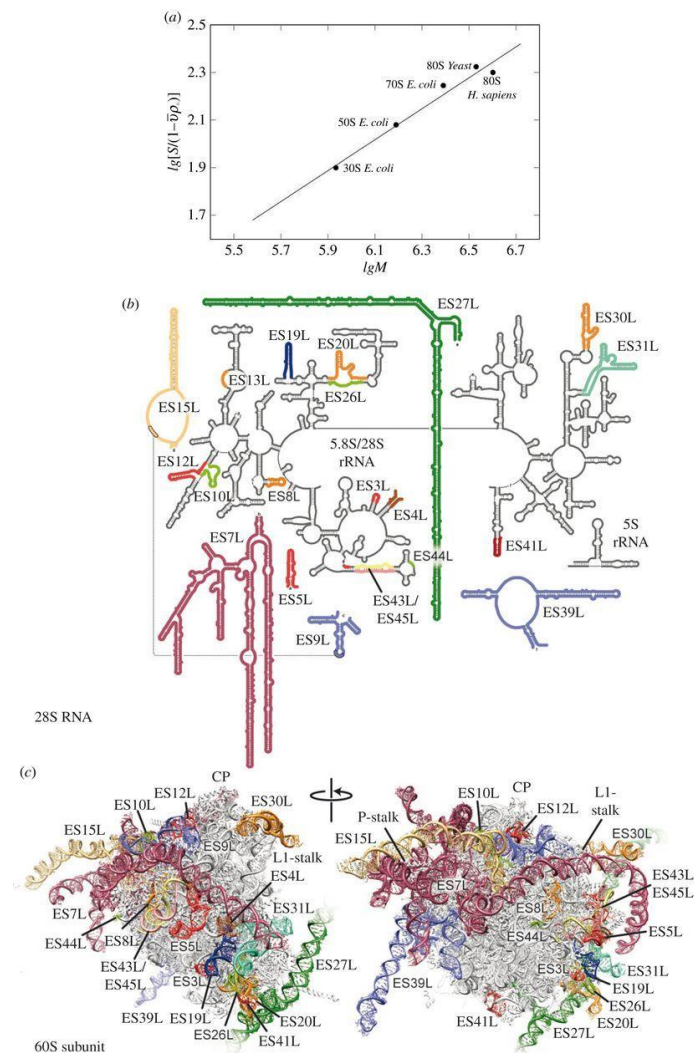
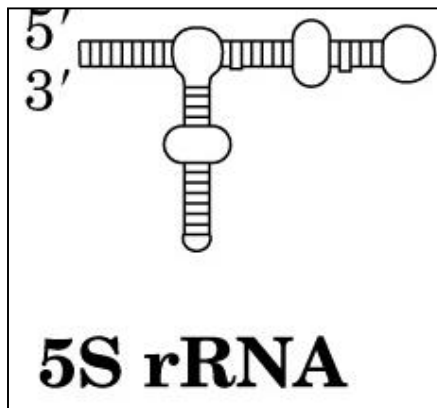
RNA ribossômico - rRNA

Possuem estrutura tridimensional específica visando promover a estabilidade e atividade catalítico nos ribossomos



Exemplos de rRNAs:

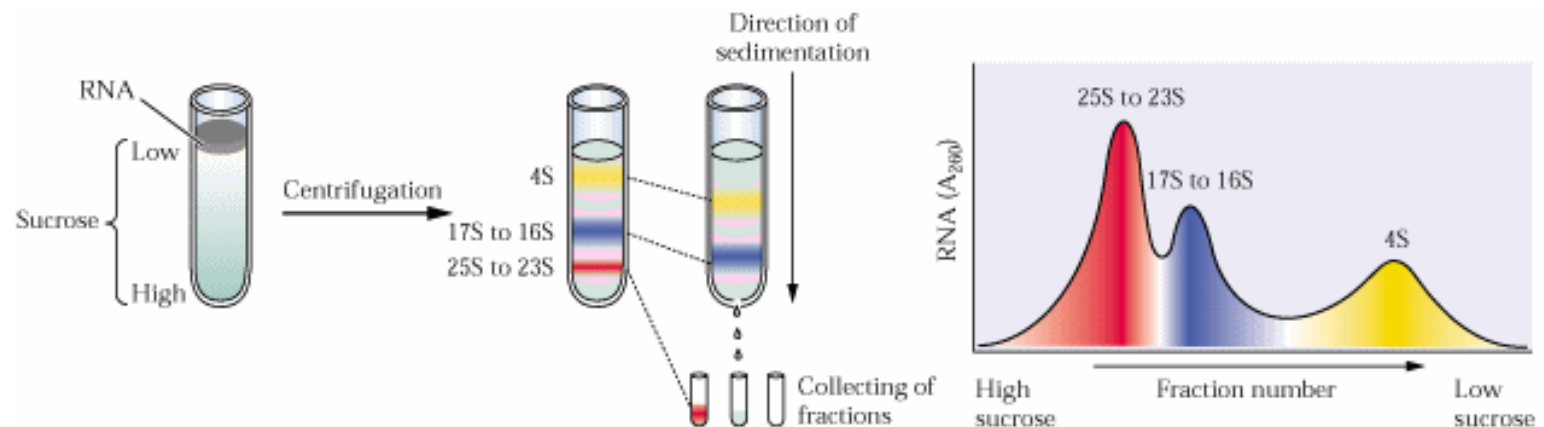
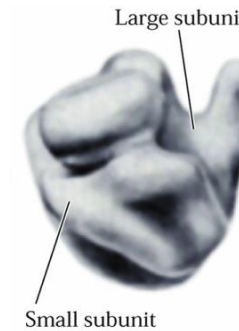
- Estrutura secundária
com grampos e alças



Ácidos Nucleicos - RNA

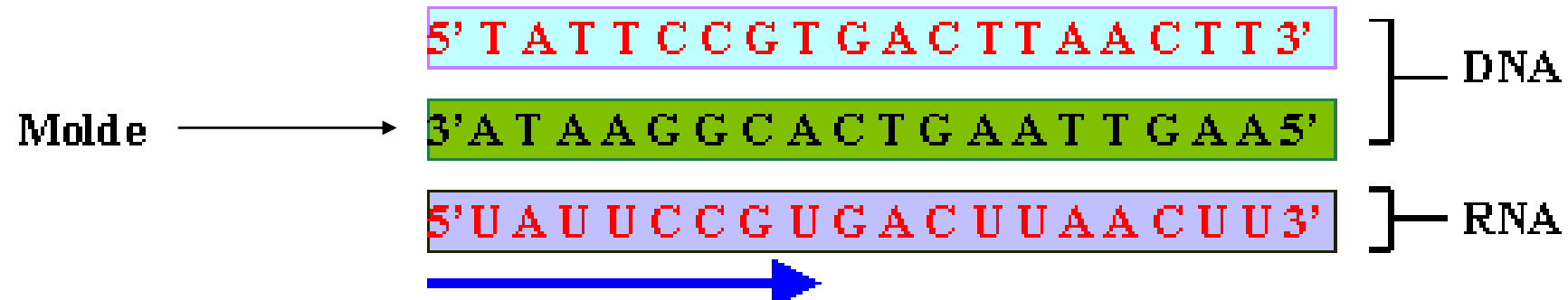
RNA ribossomal - rRNA

- mais abundante na célula ($>80\%$)
- síntese nuclear
- combinado com proteínas - ribossomo
- procarioto 70S - (23S + 5S) (16S)
- eucarioto 80S - (28S+5.8S+5S) (18S)



RNA mensageiro - mRNA

1 trinca de bases nitrogenadas = 1 códon



Sentido da transcrição

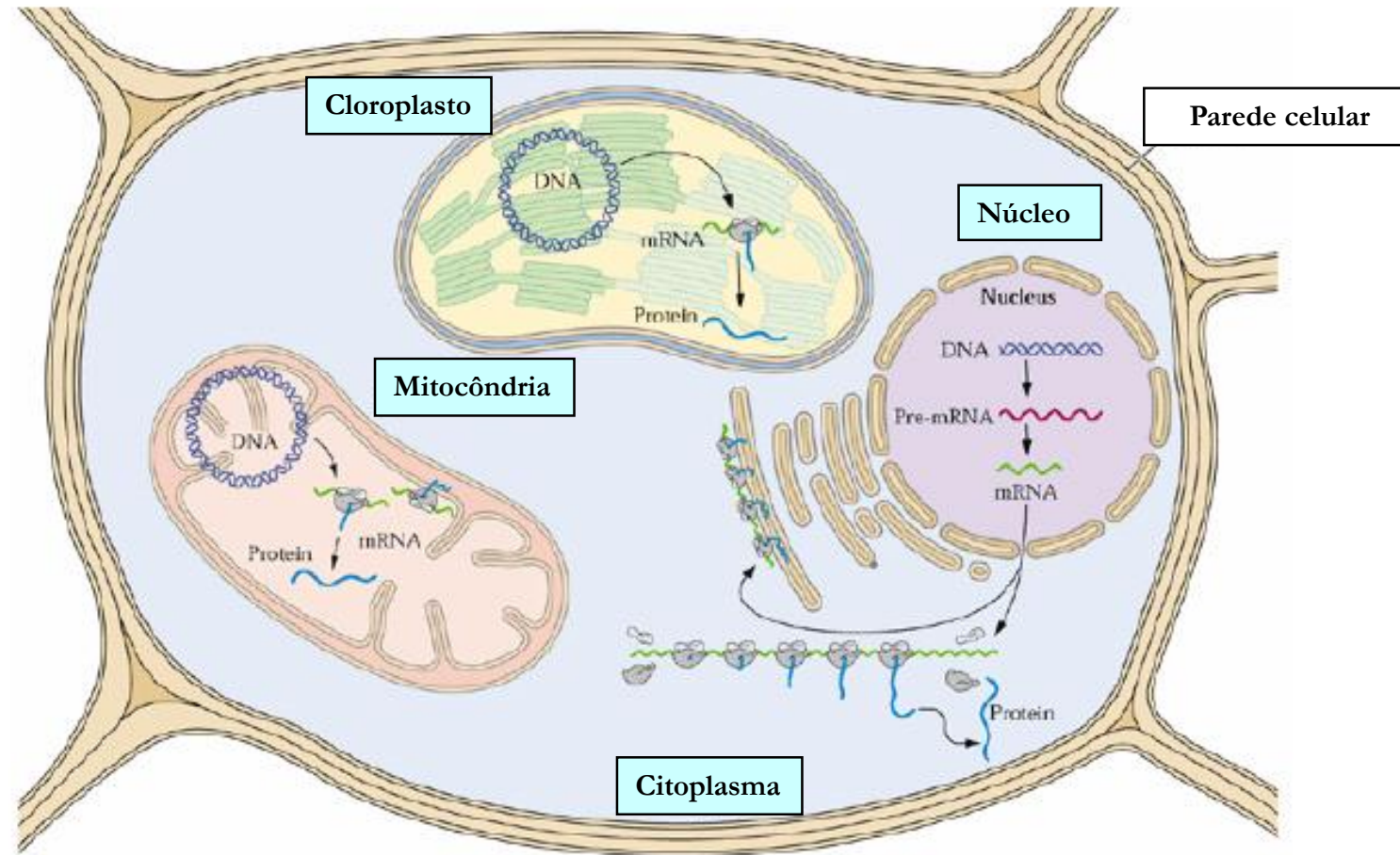
		Segunda Base				
		U	C	A	G	
Primeira Base	U	UUU } Fenil-alanina UUC } UUA } Leucina UUG }	UCU } UCC } Serina UCA } UCG }	UAU } Tirosina UAC } UAA } Stop codon UAG } Stop codon	UGU } Cysteine UGC } UGA } Stop codon UGG } Tryptophan	Terceira Base
	C	CUU } CUC } Leucina CUA } CUG }	CCU } CCC } Prolina CCA } CCG }	CAU } Histidina CAC } CAA } Glutamina CAG }	CGU } CGC } Arginina CGA } CGG }	
	A	AUU } Isoleucina AUC } AUA } AUG } Metionina start codon	ACU } ACC } Treonina ACA } ACG }	AAU } Asparagina AAC } AAA } Lisina AAG }	AGU } Serina AGC } AGA } Arginina AGG }	
	G	GUU } Valina GUC } GUA } GUG }	GCU } GCC } Alanina GCA } GCG }	GAU } Ácido Aspártico GAC } GAA } Ácido Glutâmico GAG }	GGU } GGC } Glicina GGA } GGG }	



Código Genético Universal

Amino acid	3-Letter code	1-Letter code	Codons
Alanine	Ala	A	GCC, GCU, GCG, GCA
Arginine	Arg	R	CGC, CGG, CGU, CGA, AGA, AGG
Asparagine	Asn	N	AAU, AAC
Aspartic acid	Asp	D	GAU, GAC
Cysteine	Cys	C	UGU, UGC
Glutamic acid	Glu	E	GAA, GAG
Glutamine	Gln	Q	CAA, CAG
Glycine	Gly	G	GGU, GGC, GGA, GGG
Histidine	His	H	CAU, CAC
Isoleucine	Ile	I	AUU, AUC, AUA
Leucine	Leu	L	UUA, UUG, CUA, CUG, CUU, CUC
Lysine	Lys	K	AAA, AAG
Methionine	Met	M	AUG <i>start codon -iniciador</i>
Phenylalanine	Phe	F	UUC, UUU
Proline	Pro	P	CCU, CCC, CCA, CCG
Serine	Ser	S	UCU, UCC, UCA, UCG, AGU, AGC
Threonine	Thr	T	ACU, ACC, ACA, ACG
Tyrosine	Tyr	Y	UAU, UAC
Tryptophan	Trp	W	UGG
Valine	Val	V	GUU, GUC, GUA, GUG
"Stop"	—	—	UAA, UAG, UGA <i>stop codon -terminador</i>

Três genomas em plantas: cromossomial, plastidial e mitocondrial



Estudo Dirigido

1. Diferenças fundamentais entre DNA e RNA;
2. Estrutura e função do DNA;
3. Principais características da dupla hélice do DNA;
4. Principais tipos e funções dos RNAs.

Capítulo 2 – Componentes químicos das Células

Capítulo 5 – DNA e Cromossomos

Alberts, B.; Bray, D.; Hopkin, K.; Johnson, A.; Lewis, J.; Raff, M.; Roberts, K.; Walter, P.
2011. *Fundamentos da Biologia Celular*. 3ª Edição brasileira. Artmed, Porto Alegre