

# Bioquímica Estrutural:

Estudo da estrutura das moléculas biológicas e dos princípios que a regem.

( ~ **Biologia Estrutural** )

# Macromoléculas biológicas

- **DNA:** repositório da informação genética na maioria dos organismos vivos
- **RNA:** transferência (e repositório) de informação genética, matriz para a síntese proteica, funções estruturais, etc...
- **Proteínas:** componentes estruturais (pele, ossos, músculo, cabelo, etc...), catálise de reacções bioquímicas (enzimas), transmissão de sinais, regulação, transdução de energia, etc.,etc., etc!..
- **Lípidos:** componentes essenciais das membranas biológicas, sinalização
- **Polissacáridos:** armazenamento de energia, função estrutural

Sequência



Estrutura



Função

# Fluxo de informação biológica

Gene ...TTAATAAGT...

↓ transcrição

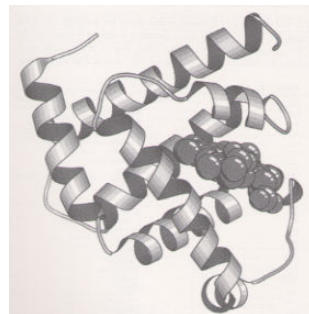
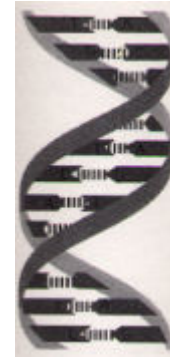
m-RNA ...UUAAUAAGU...

↓ splicing, tradução

cadeia  
polipeptídica ...LISVHDN...

↓ modificações pós-translacionais

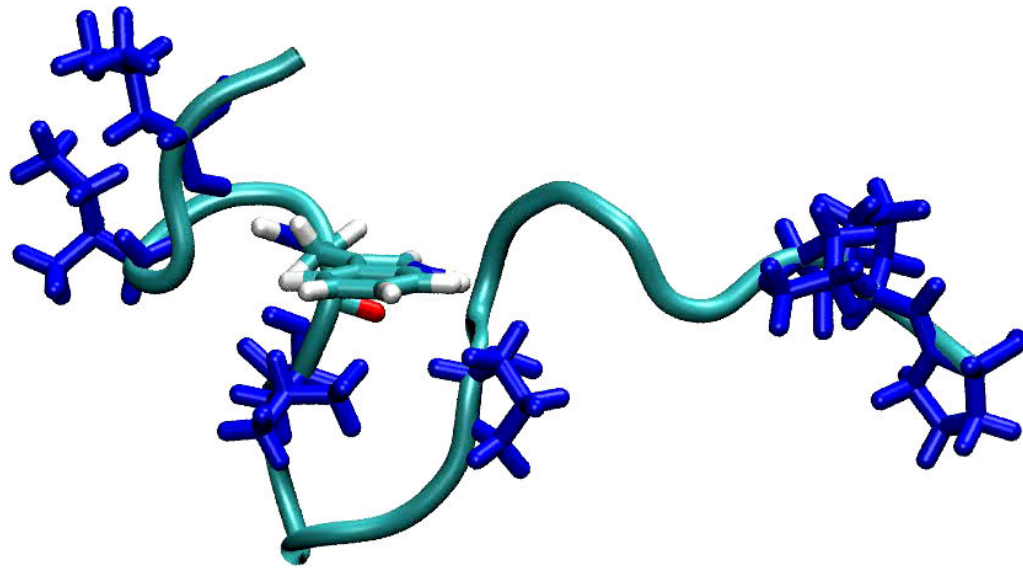
proteína



Dogma central da  
biologia molecular

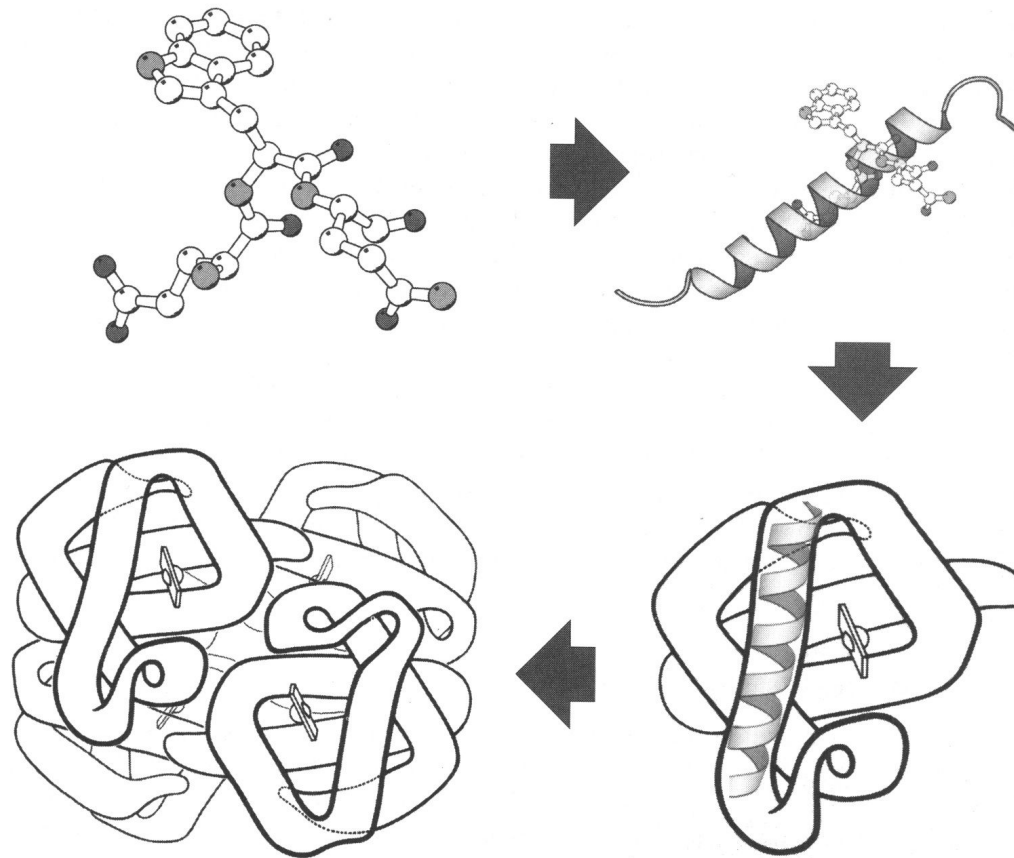
Excepções: vírus de RNA,  
priões, ribozimas (?)

# Sequência->Estrutura

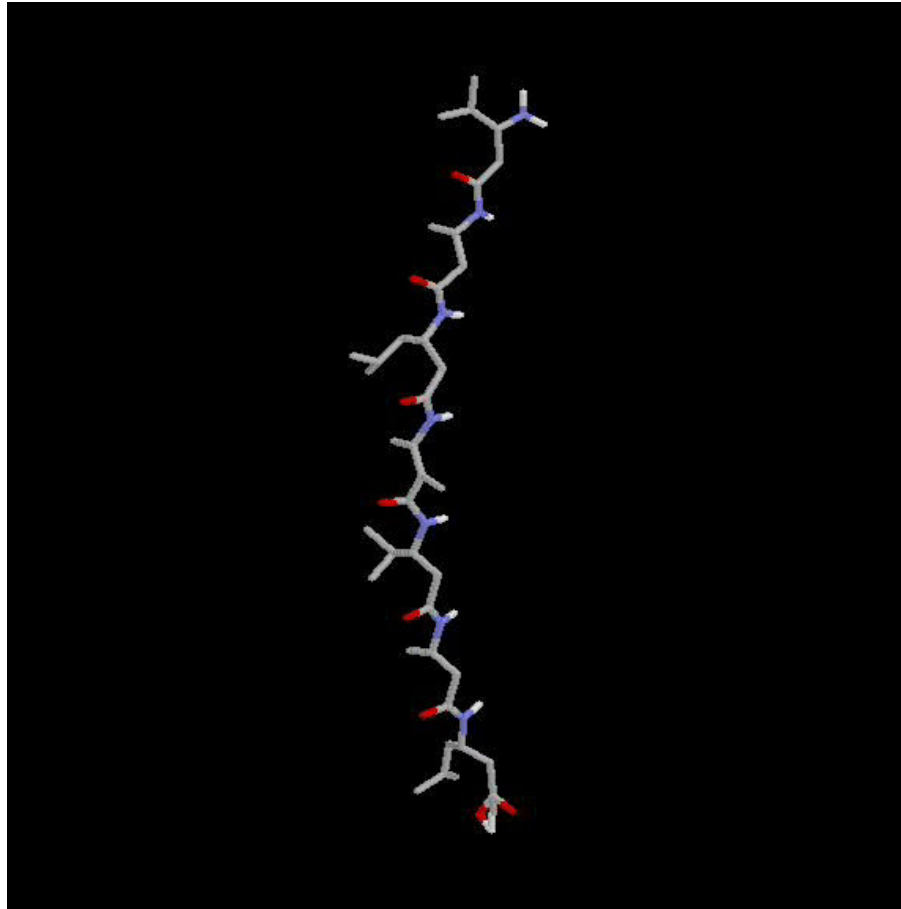


Muitas proteínas adquirem a sua estrutura tridimensional *espontaneamente (folding)*

# Níveis de organização da estrutura



# Formação hierárquica da estrutura



Formação espontânea de uma hélice  $\alpha$   
(simulação)

# Formação hierárquica da estrutura



"Simulação" do *folding* ubiquitina



# De onde provêm a informação estrutural ?

Combinação de vários tipos de conhecimento:

- Teoria da ligação química
- Geometria de moléculas pequenas
- Métodos experimentais para a determinação da estrutura de biomoléculas:
  - Cristalografia de raios X
  - Ressonância Mag. Nuclear
  - Outros métodos

# Que informação temos disponível ?

**Número de estruturas tridimensionais** (coordenadas atómicas):

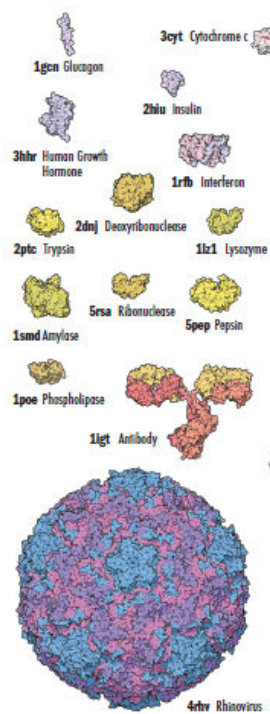
- 35767 proteínas
- 1579 complexos ácidos nucleíco-proteína
- 1671 ácidos nucleícos
- 18 glícidos

**Total:** 39051 estruturas

**Métodos experimentais de determinação da estrutura:**

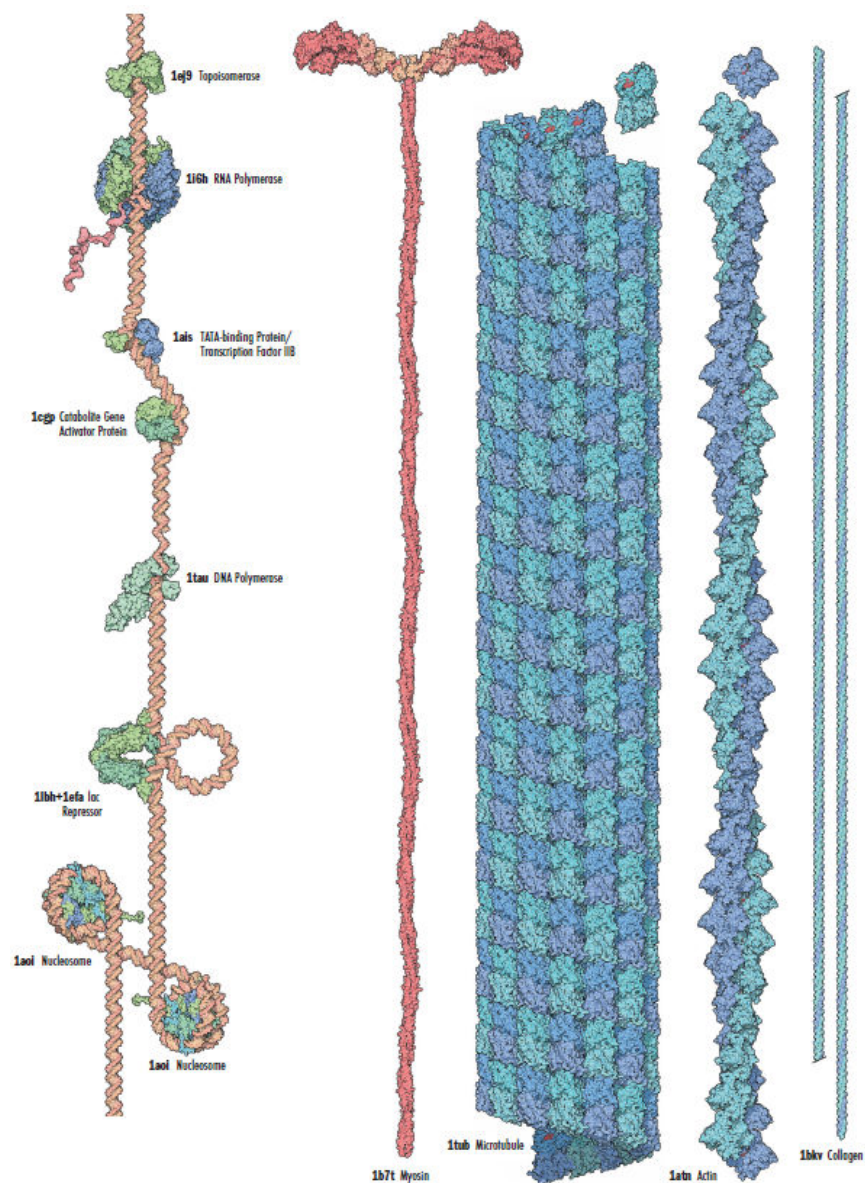
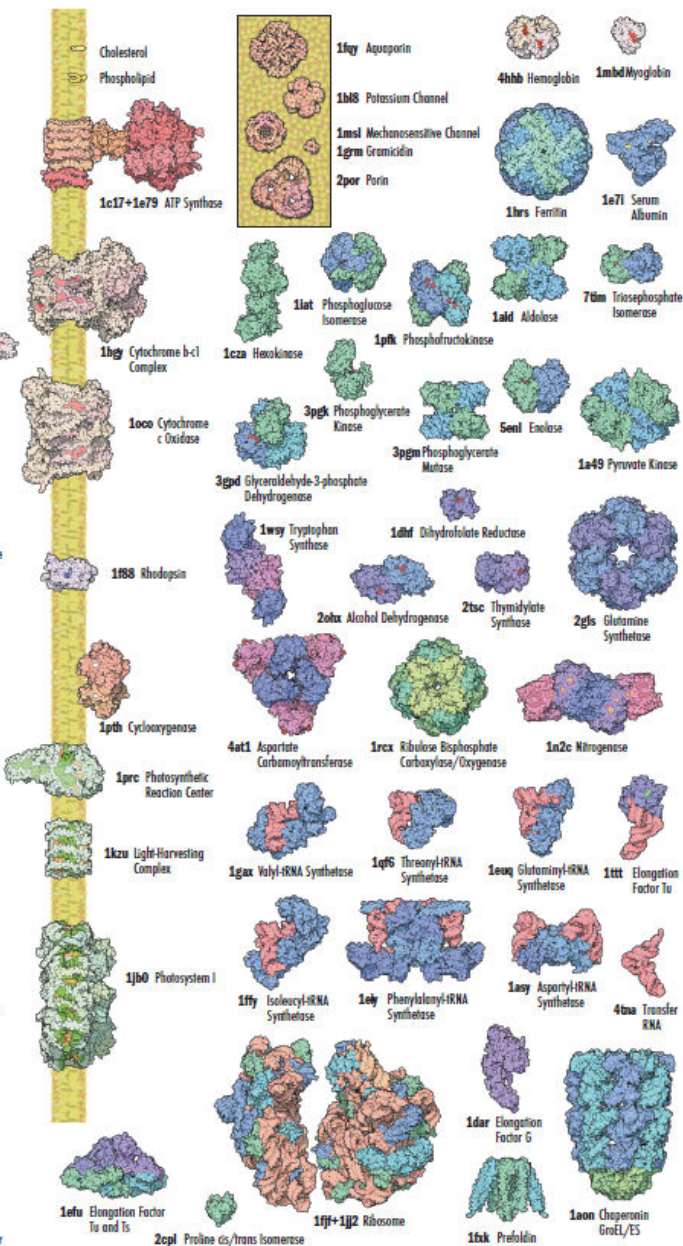
- X-ray: 33126
- NMR: 5707
- Microscopia: 134
- Outros: 84

# MOLECULAR MACHINERY: A Tour of the Protein Data Bank

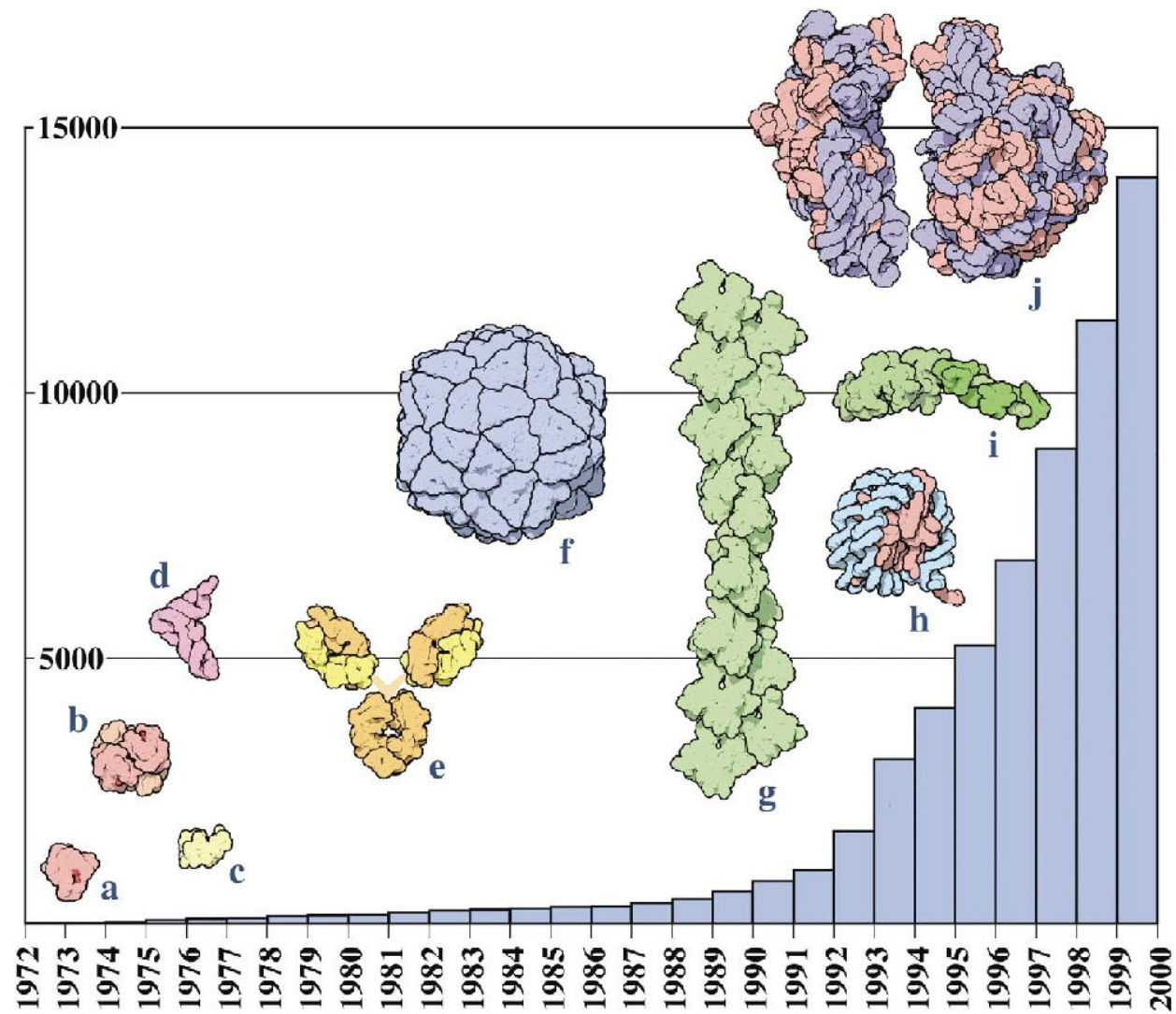


**PDB**  
PROTEIN DATA BANK

<http://www.pdb.org/> • [info@rcsb.org](mailto:info@rcsb.org)  
RESEARCH COLLABORATORY FOR  
STRUCTURAL BIOINFORMATICS  
RUTGERS, THE STATE UNIVERSITY OF NEW JERSEY  
SAN DIEGO SUPERCOMPUTER CENTER  
NATIONAL INSTITUTE OF STANDARDS AND TECHNOLOGY



# Crescimento do Protein Data Bank



# Princípios que regem a estrutura das biomoléculas

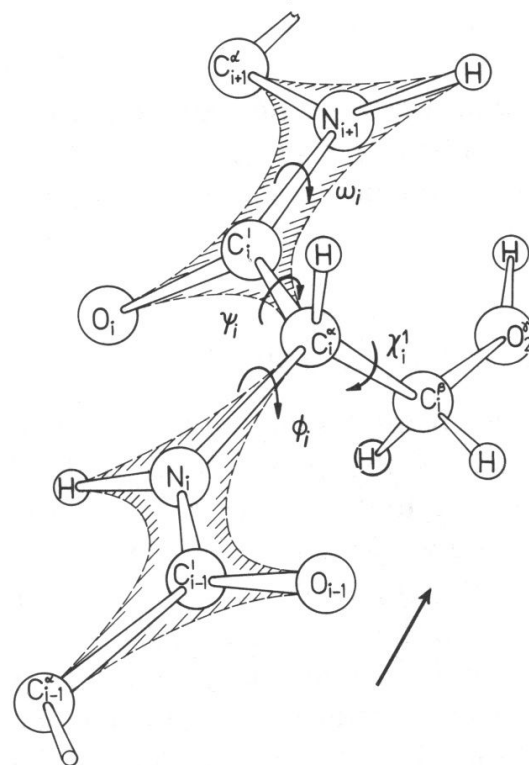
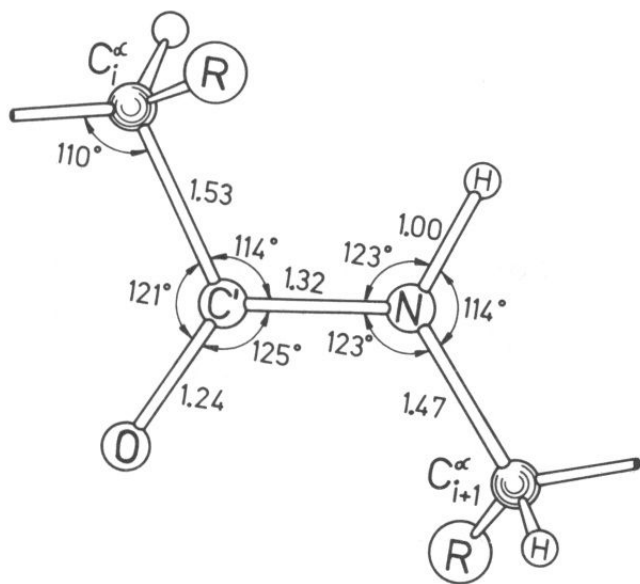
"Perhaps the most remarkable features of the molecule are its *complexity* and *lack of symmetry*. The arrangement seems to be almost totally lacking the kind of regularities which one instinctively anticipates and it is more complicated than has been predicted by any theory of protein structure"

J.C. Kendrew *et al.*, 1958



- As macromoléculas biológicas parece, numa primeira análise, distanciar-se dos princípios simples de geometria e simetria que sabemos reger a estrutura das moléculas pequenas.

- Ligações covalentes, geometria molecular:

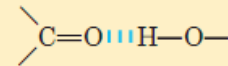




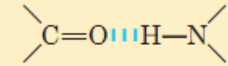
## •Interacções não-covalentes:

Hydrogen bonds

Between neutral groups

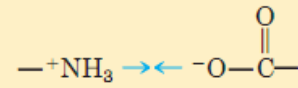


Between peptide bonds



Ionic interactions

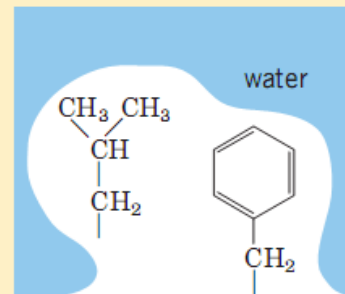
Attraction



Repulsion



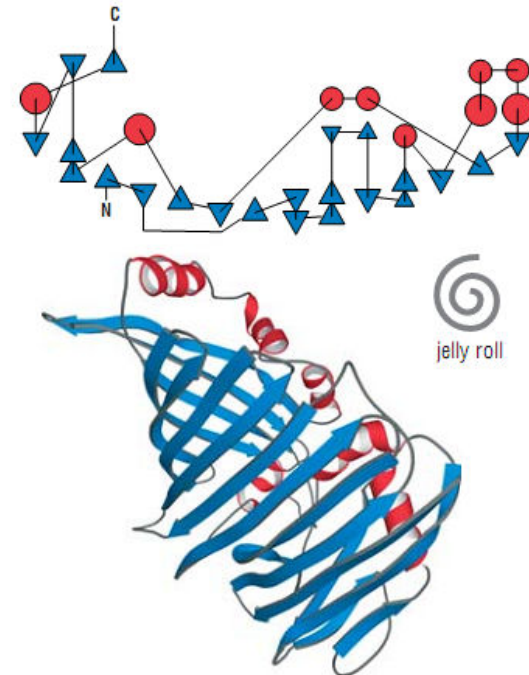
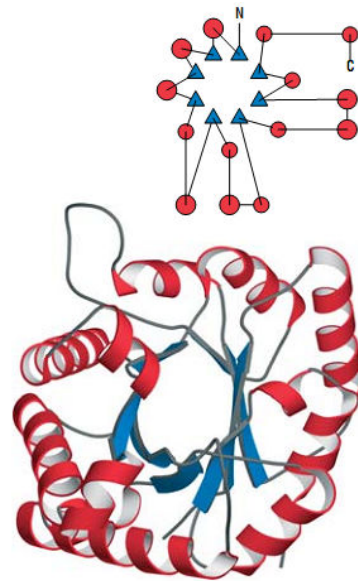
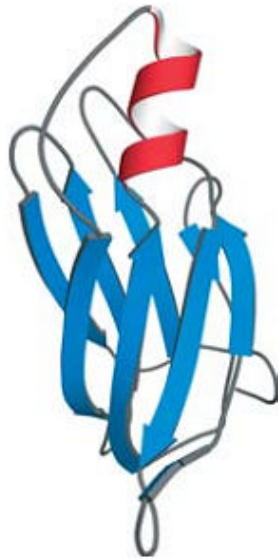
Hydrophobic interactions



van der Waals interactions

Any two atoms in  
close proximity

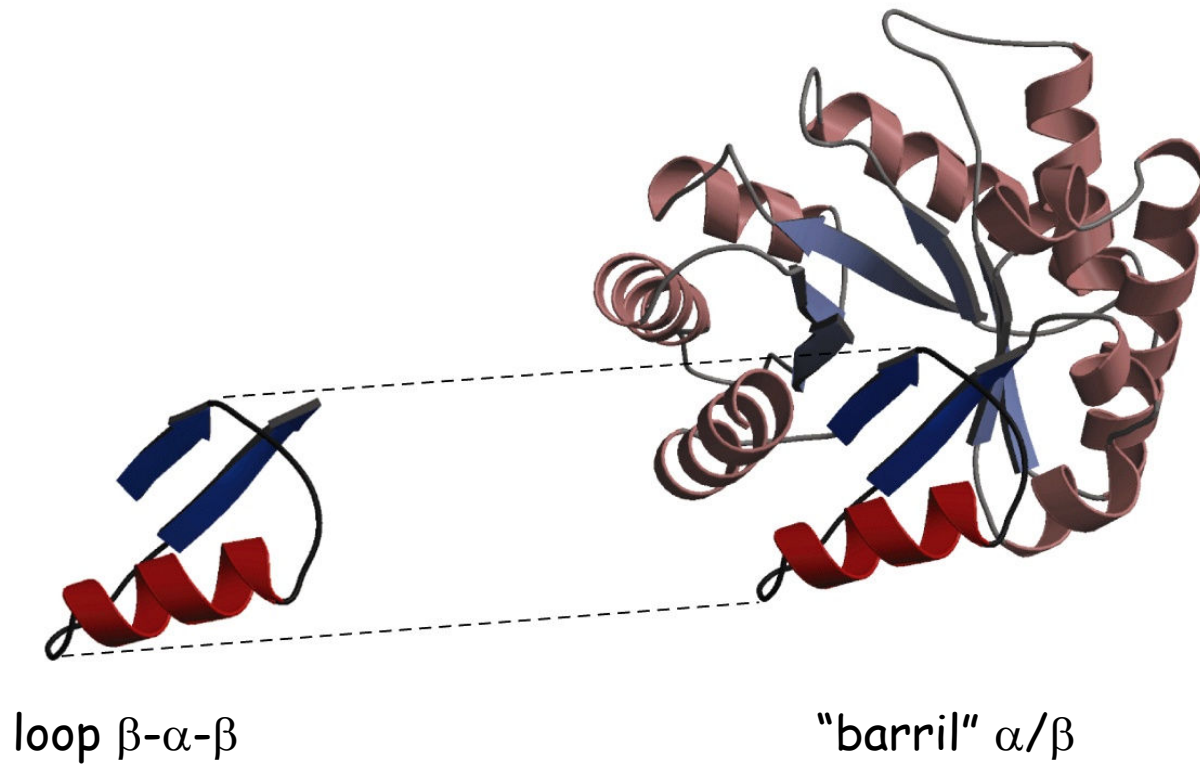
- Princípios arquitetônicos



...

Recorrência de padrões estruturais na arquitectura das biomoléculas.

- Princípios arquitetônicos (cont.):



Formação de estruturas a partir da associação de unidades estruturais

- Oligomerização:

(a) dimer



(b) trimer



(c) planar tetramer



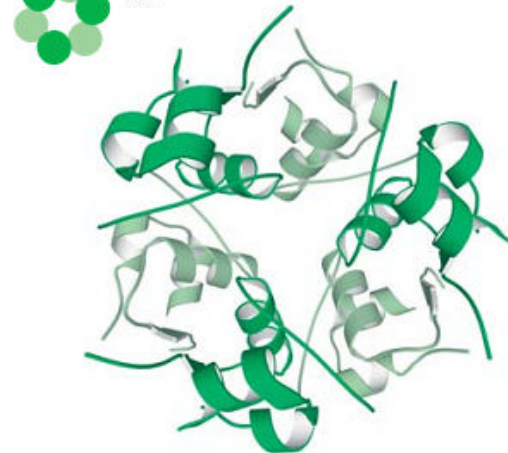
(d) tetramer



(e) pentamer



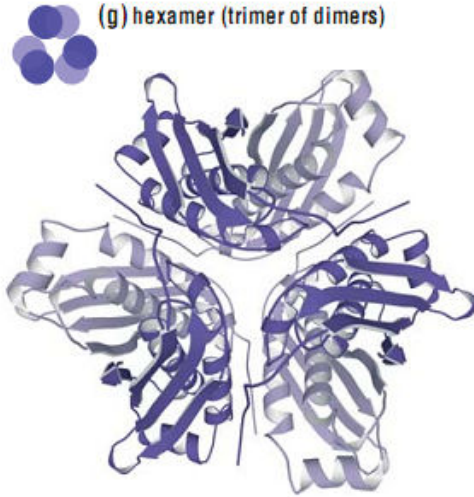
(f) planar hexamer



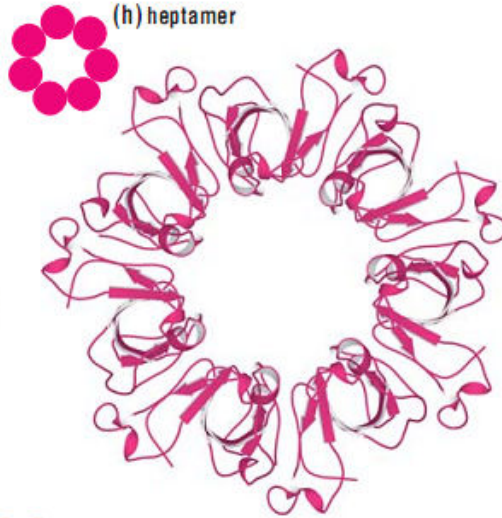


- Oligomerização(cont.):

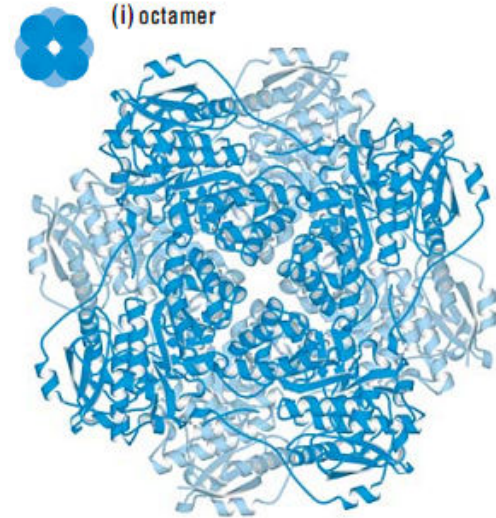
(g) hexamer (trimer of dimers)



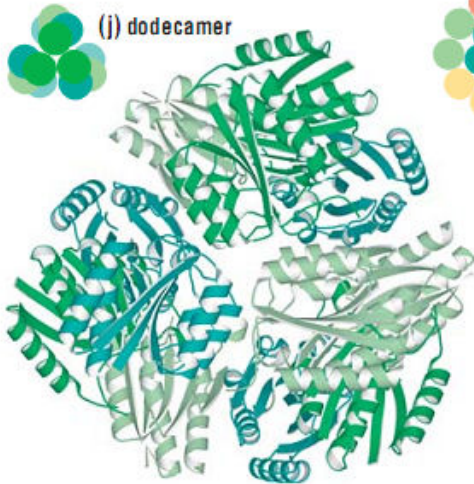
(h) heptamer



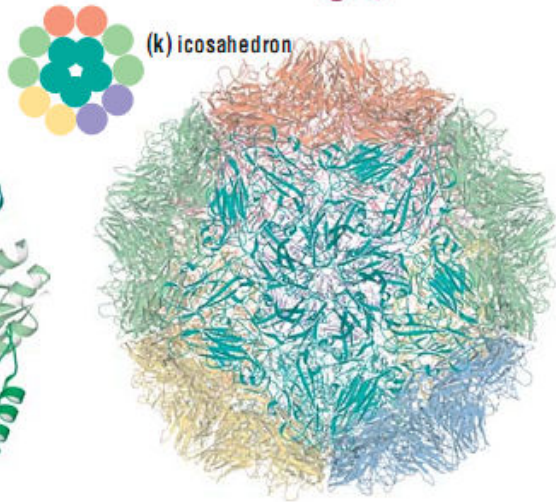
(i) octamer



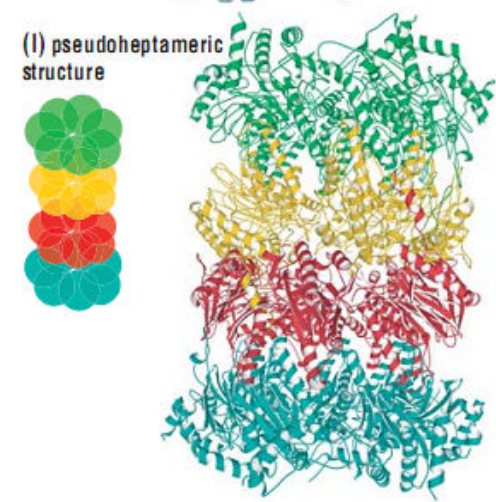
(j) dodecamer



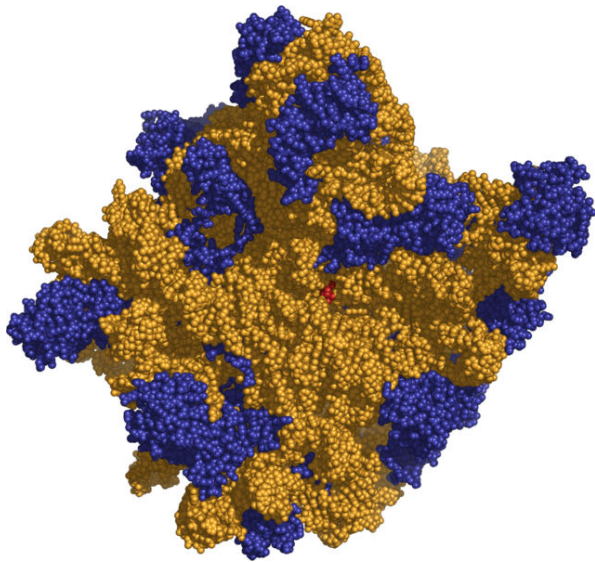
(k) icosahedron



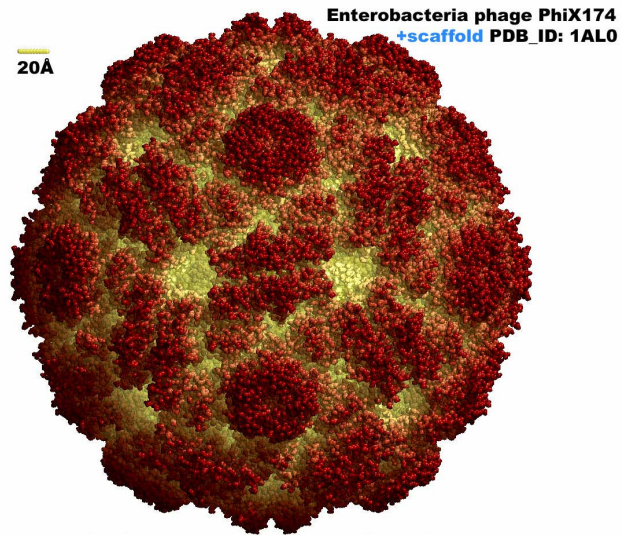
(l) pseudoheptameric structure



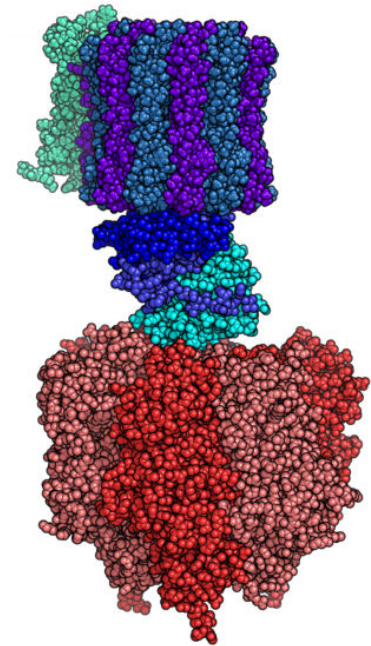
- Formação de estruturas supramacromoleculares



Ribossoma



Vírus

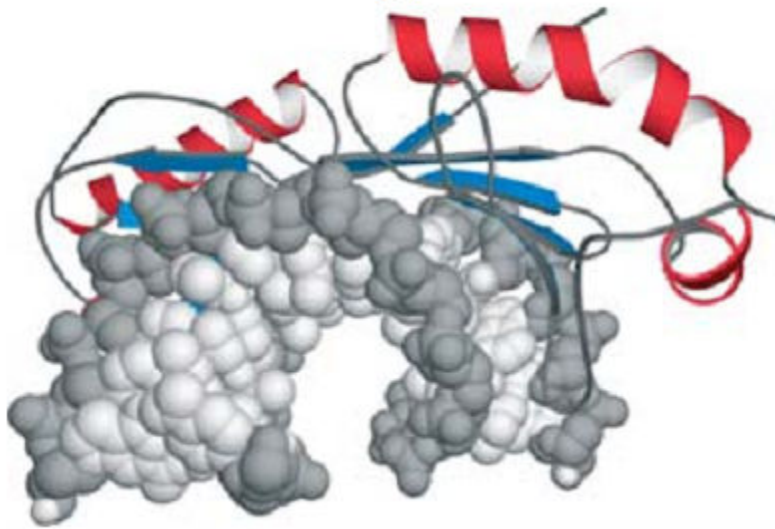


ATPase

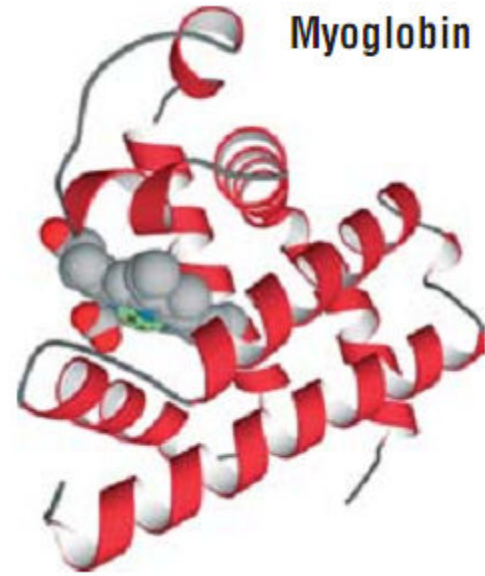
Função

# "Binding"

TATA binding protein

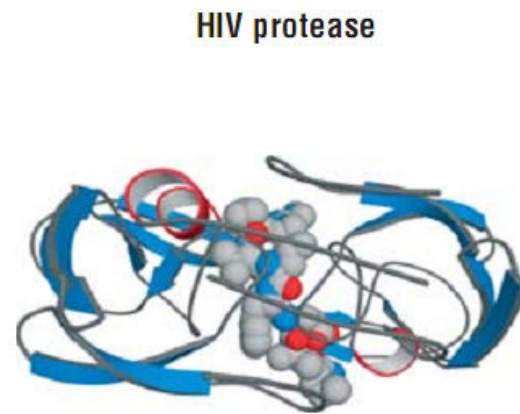
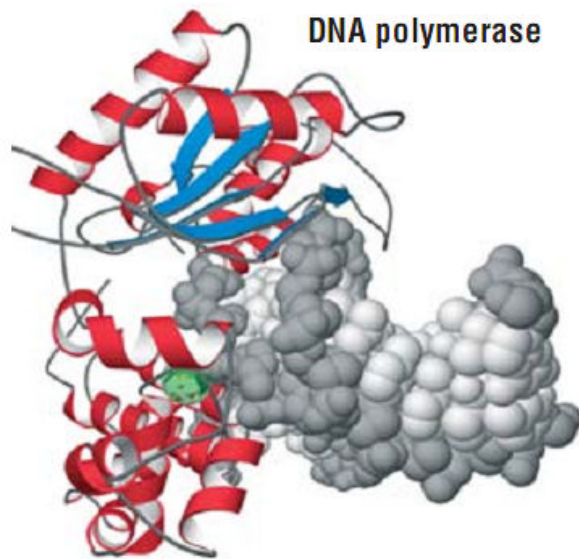


Myoglobin

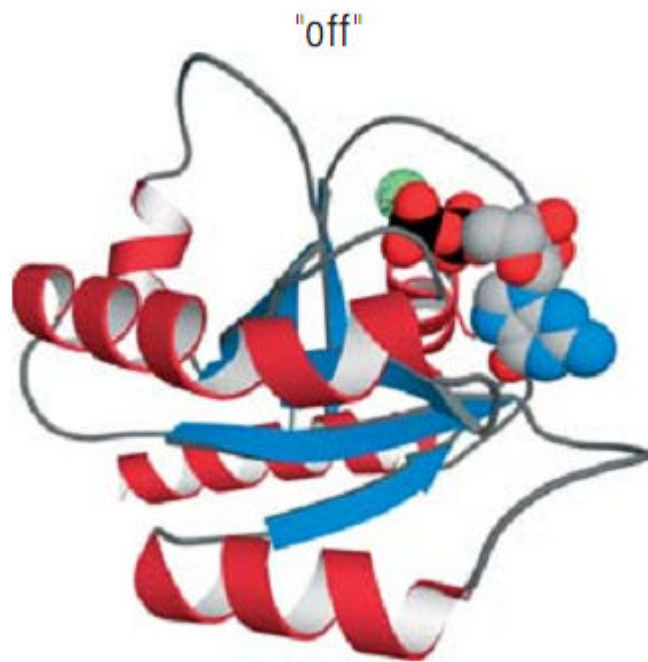




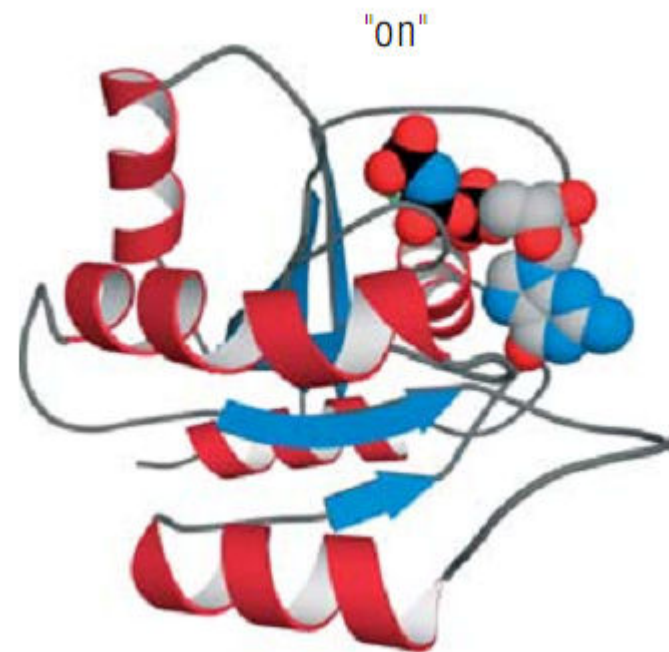
# Catálise



# "Switching"

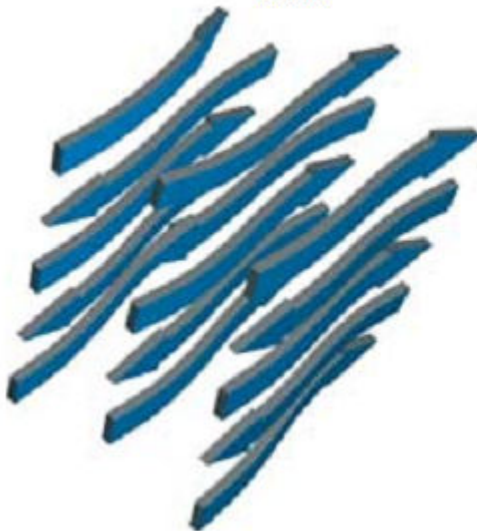


Ras

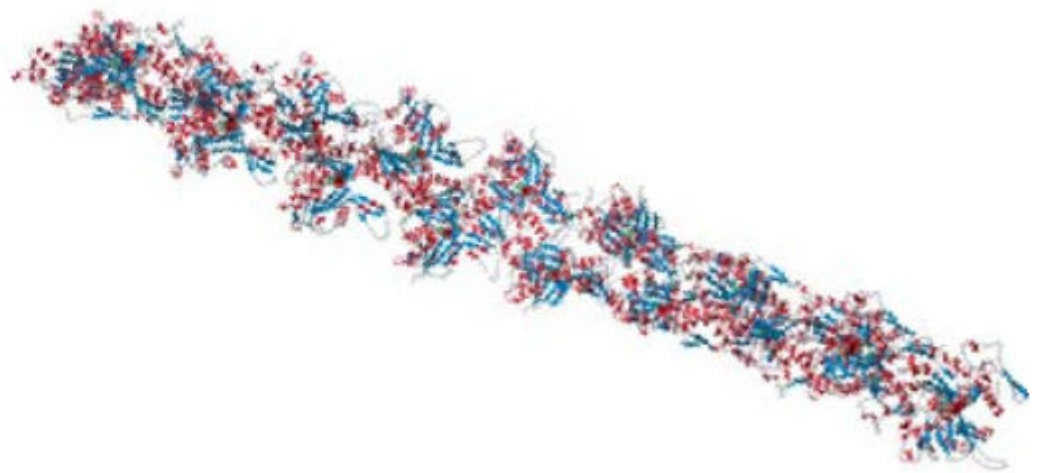


# Estrutura

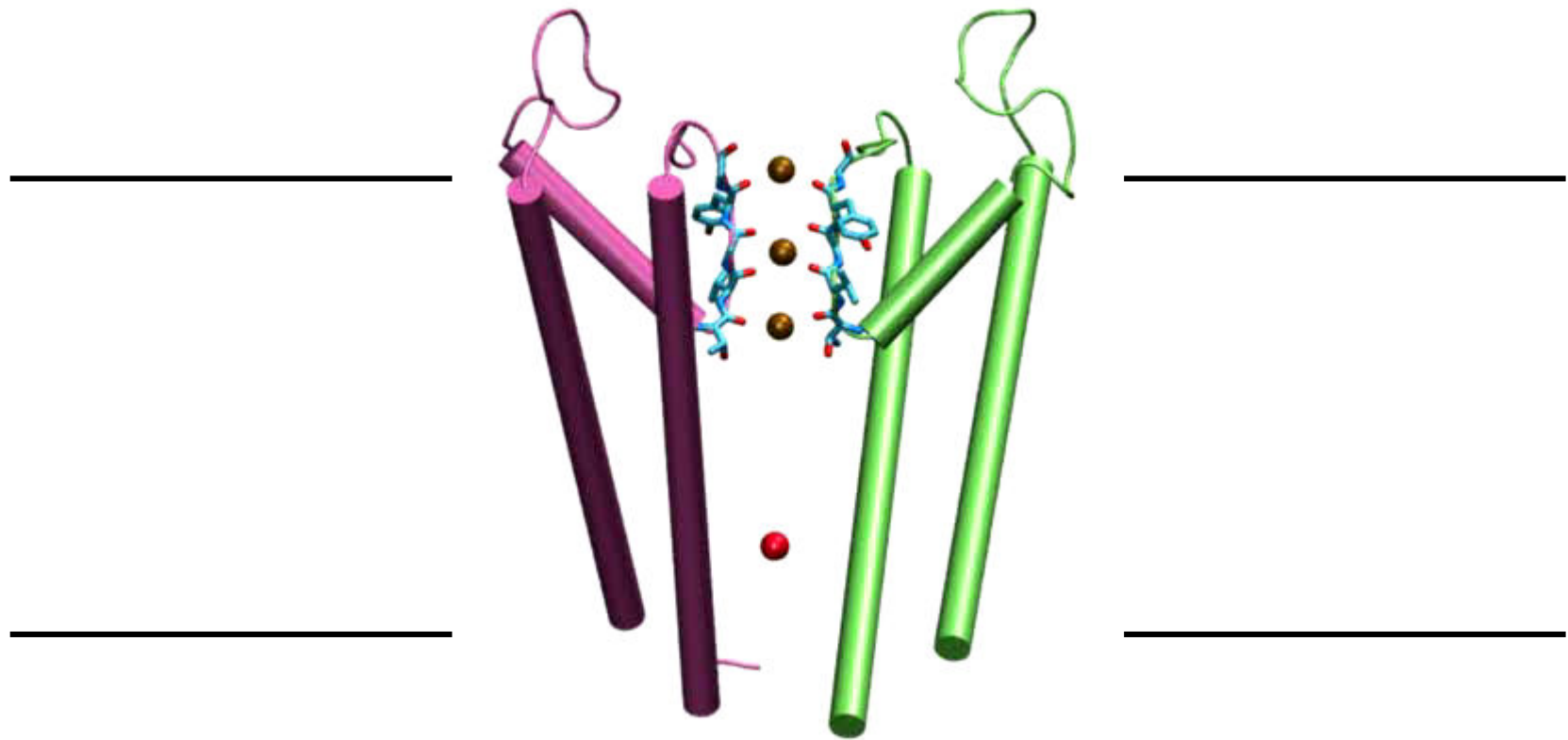
Silk



F-actin



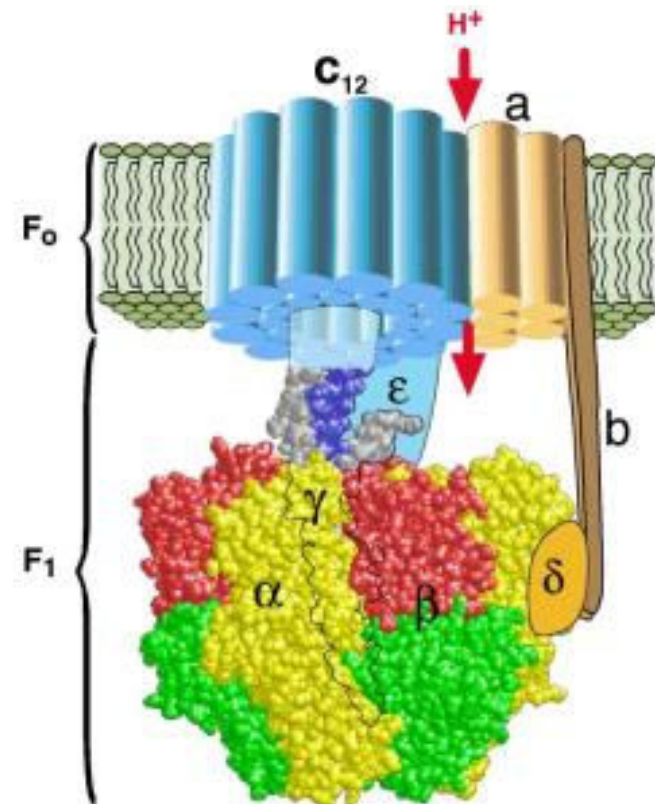
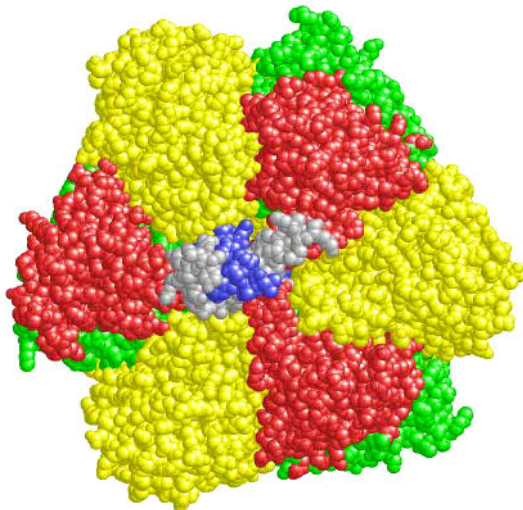
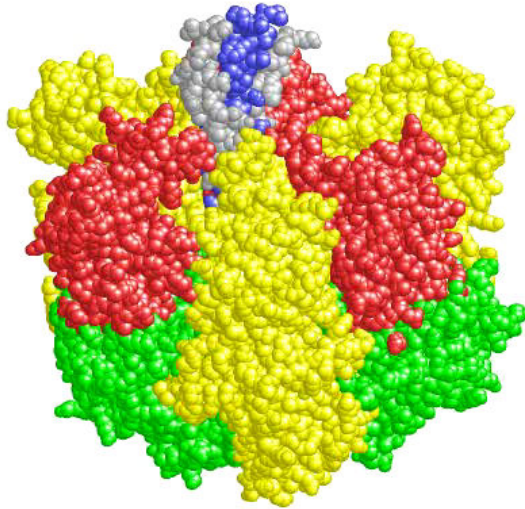
# Permeabilidade



As macromoléculas biológicas funcionam como máquinas moleculares

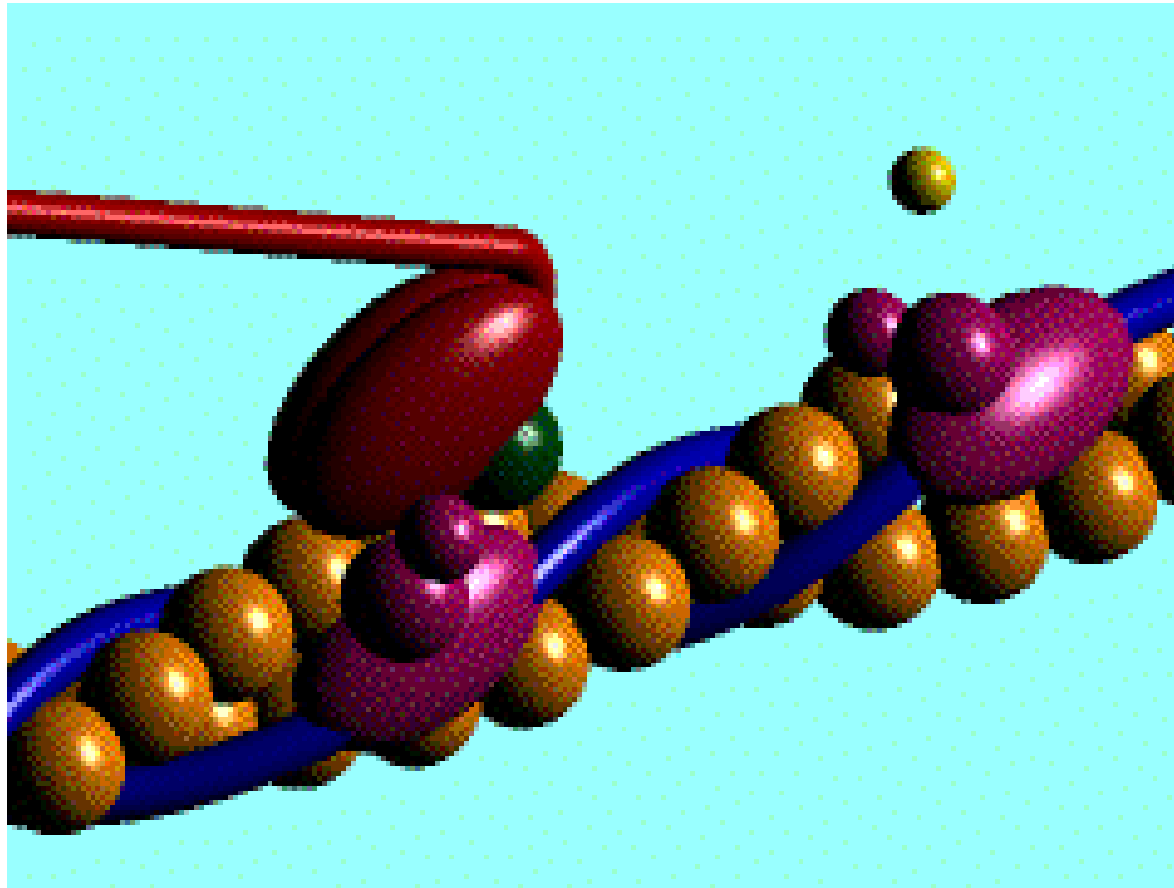
# Conversão de energia

## ATP sintase



H. Wang and G. Oster (1998). Nature 396:279-282.

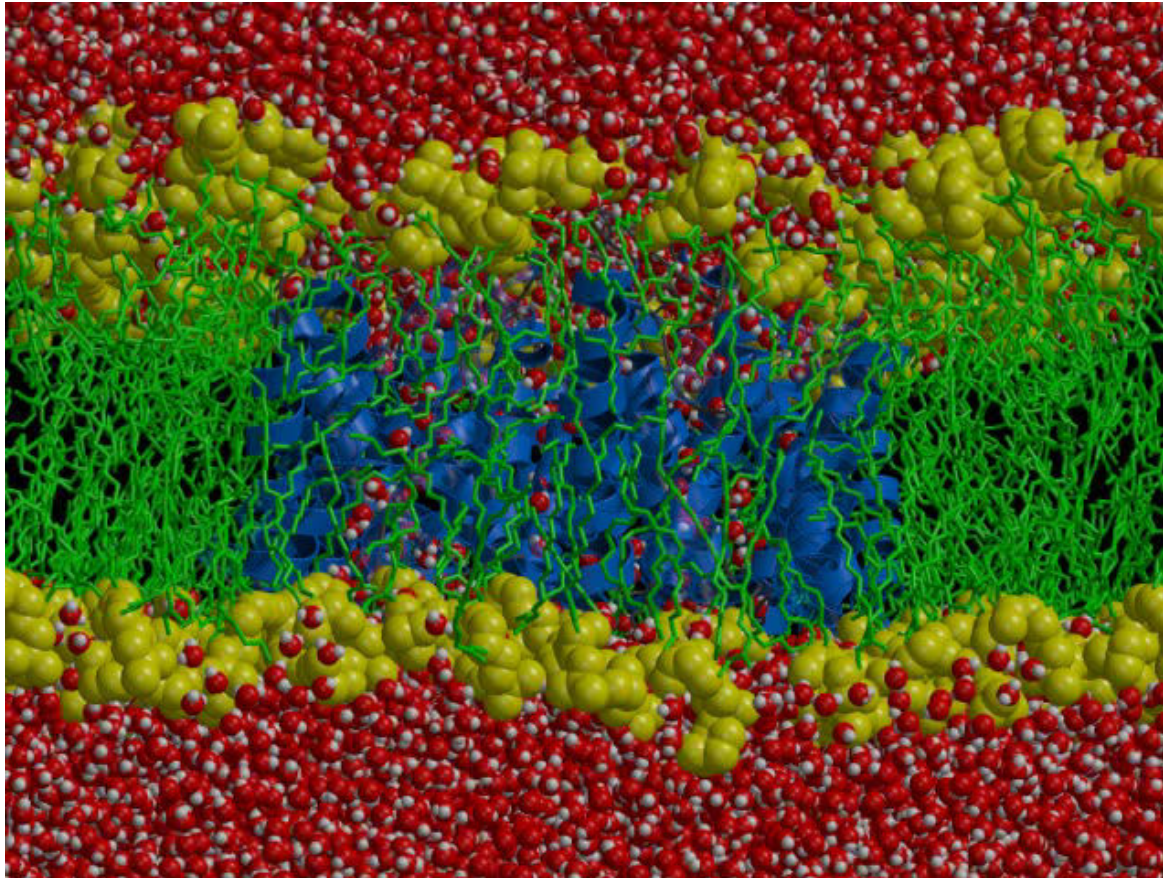
# Motilidade



Actina+Miosina



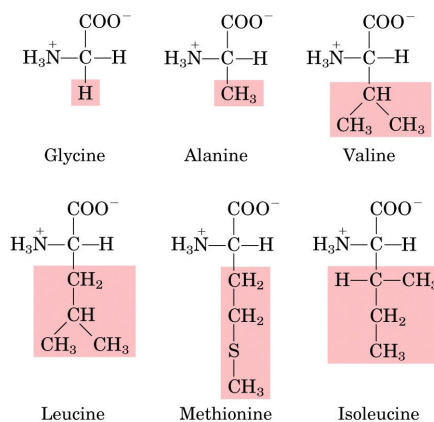
# Compartimentação



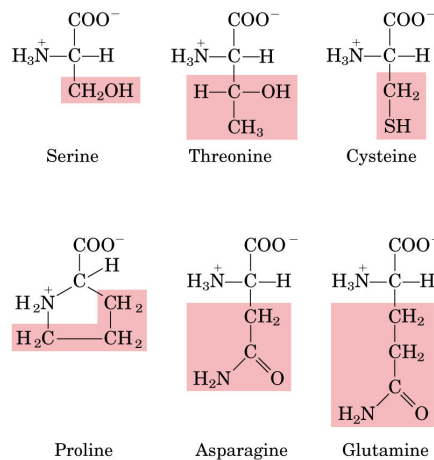
Membrana+aquaporina

# Proteínas

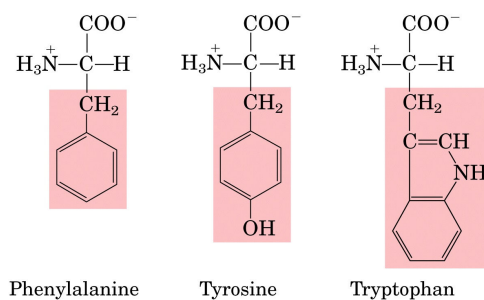
## Nonpolar, aliphatic R groups



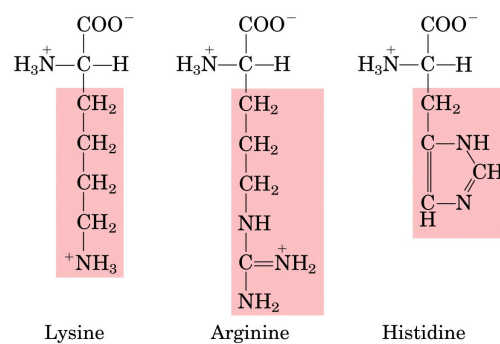
## Polar, uncharged R groups



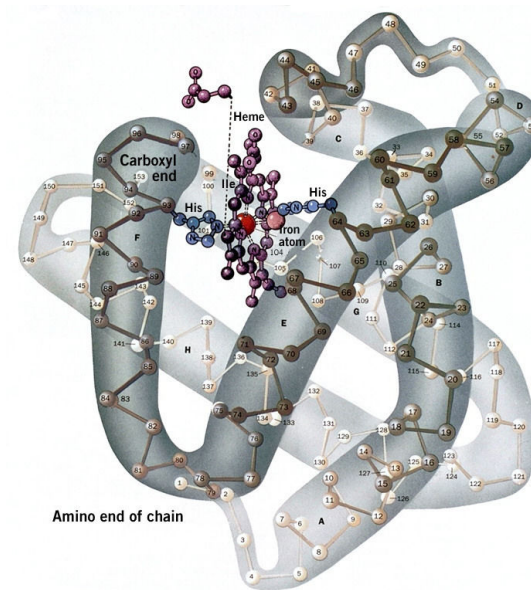
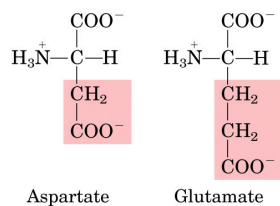
## Aromatic R groups



## Positively charged R groups



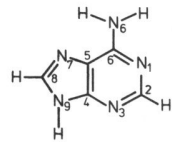
## Negatively charged R groups



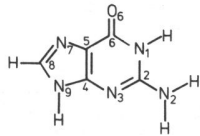
(a) Mioglobina



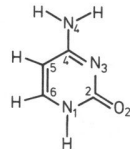
# Ácidos nucleicos



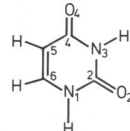
Adenine  
Ade



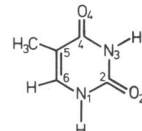
Guanine  
Gua



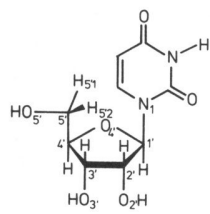
Cytosine  
Cyd



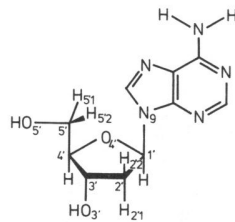
Uracil  
Ura



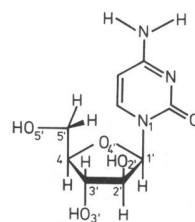
Thymine  
Thy



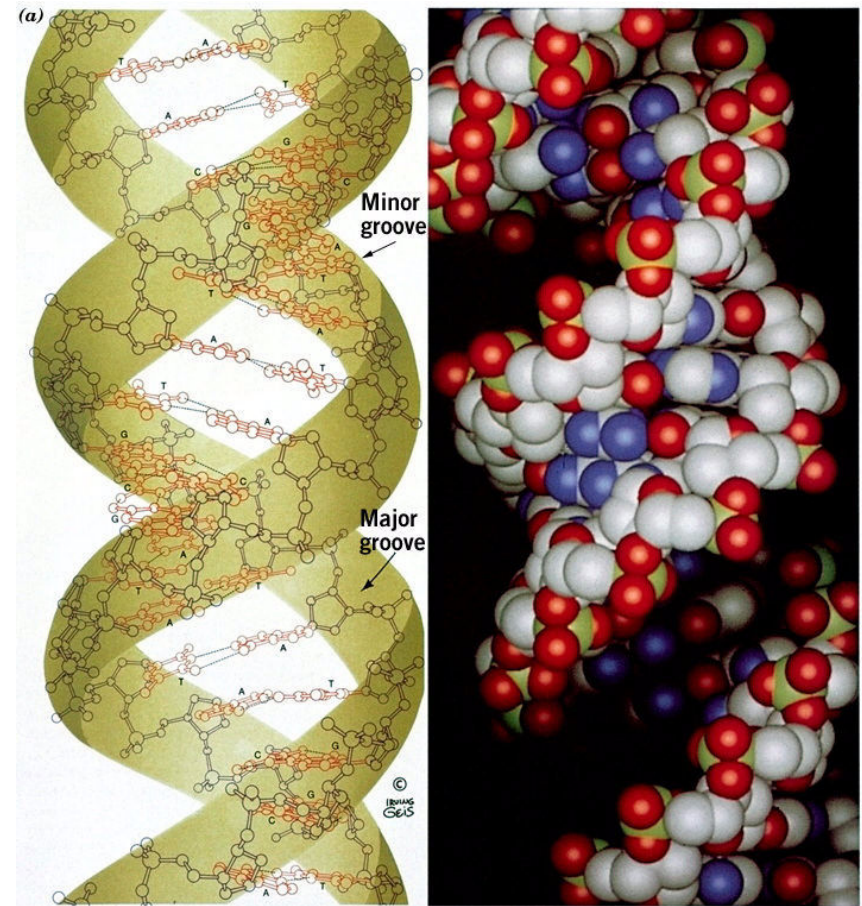
Uridine  
Urd, U



Deoxyadenosine  
dAdo, dA



Arabinocytidine  
araCyd, araC



B-Dna