

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...



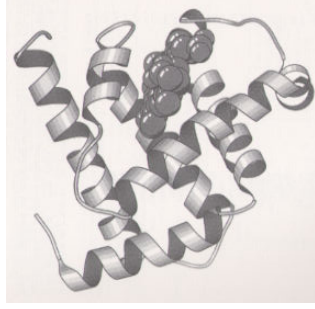
m-RNA ...UUAUAAGU...



cadeia ...LISVHDN...
polipeptídica



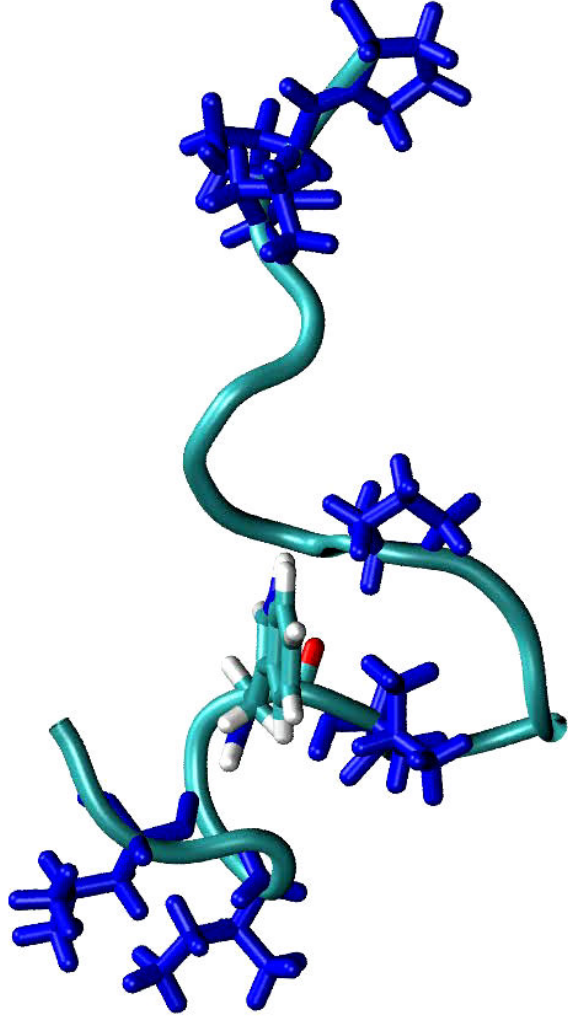
proteína



Dogma central da
biologia molecular

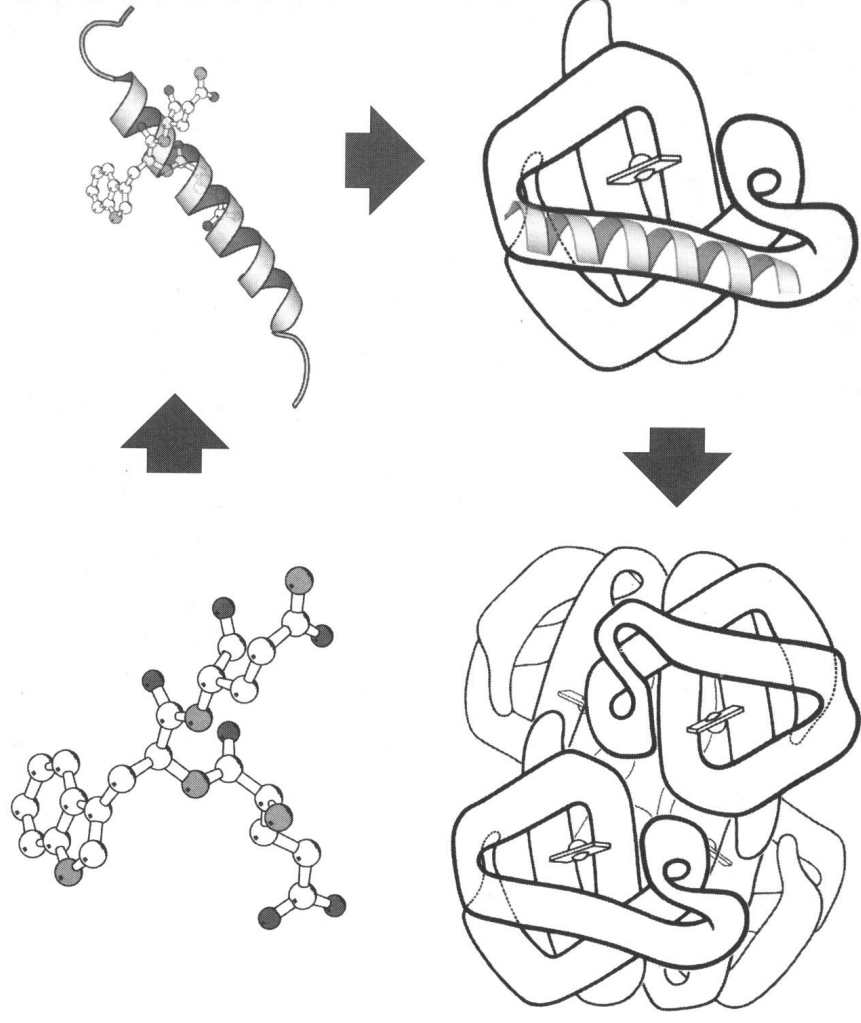
Exceções: vírus de RNA,
príões, ribozimas (?)

Sequência->Estrutura

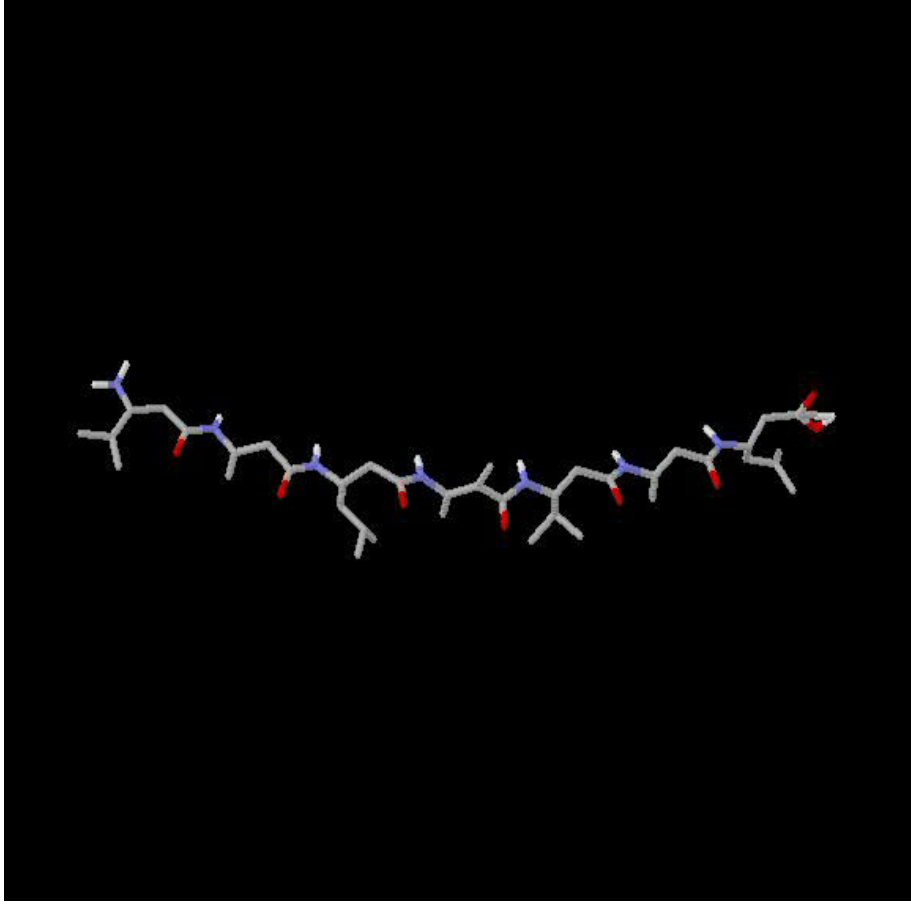


Muitas proteínas adquirem a sua estrutura tridimensional *espontaneamente (folding)*

Níveis de organização da estrutura

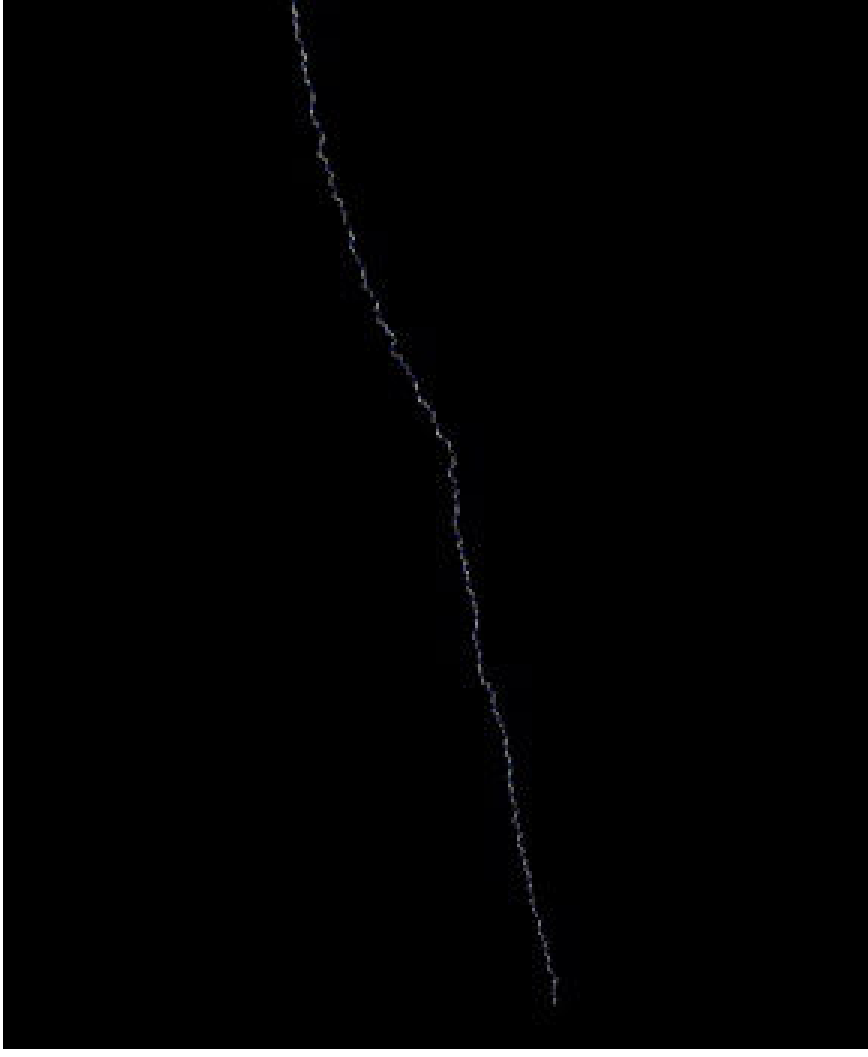


Formação hierárquica da estrutura



Formação espontânea de uma hélice α
(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 nucleico-proteína
- 1671 ácidos nucleicos
- 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

The image displays a vast collection of protein structures from the Protein Data Bank, categorized by their function and structure. Key features include:

- Top Section:** Large-scale molecular machines and complexes, including the ribosome (1ubw), DNA polymerase (1aa), RNA polymerase (1hs), and myosin (1m7).
- Middle Section:** Numerous individual enzymes and structural proteins, each labeled with its PDB ID and name, such as lysozyme (1lze), insulin (2ins), hemoglobin (1hhb), and many others.
- Bottom Section:** Smaller proteins and peptides, including cytochrome c (1j7c) and various kinases.

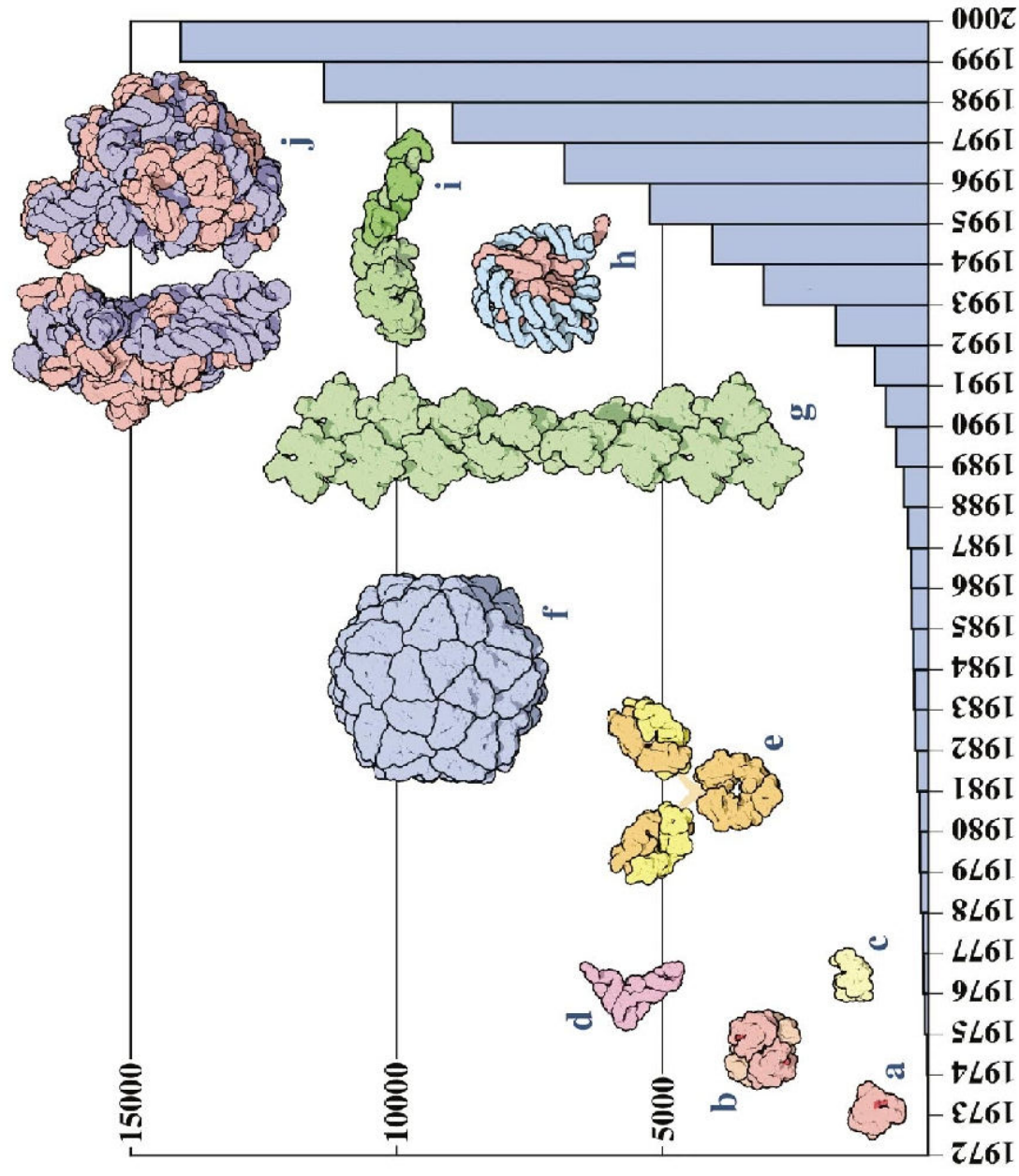
The structures are color-coded and arranged to show the diversity and complexity of biological macromolecules.

<http://www.pdb.org/> • 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

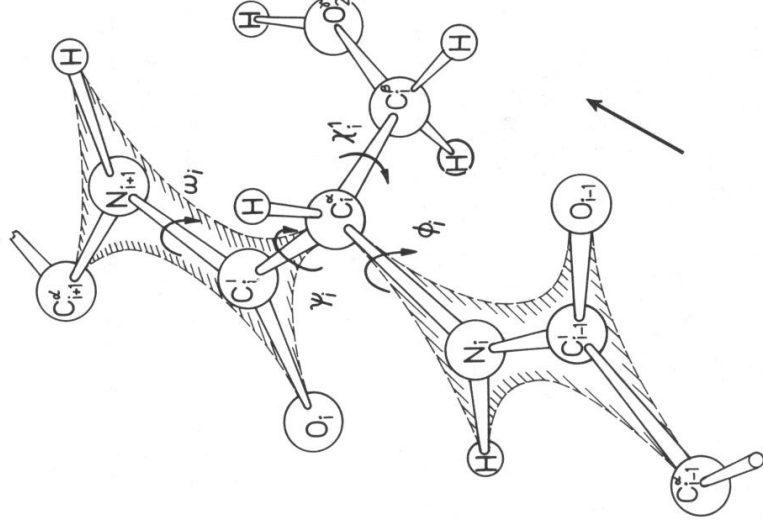
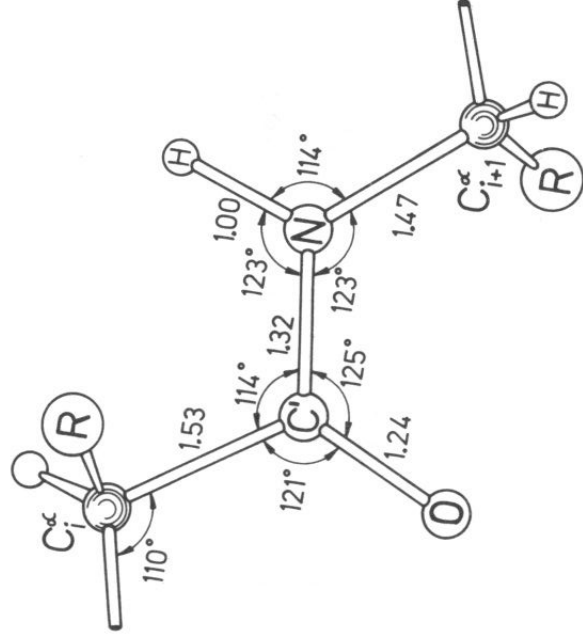
“Perhaps the most remarkable features of the molecule are its *complexity* and *lack of symmetry*. The arrangement seems to be almost totally lacking the kind of regularities which one instinctively anticipates and it is more complicated than has been predicted by any theory of protein structure”

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



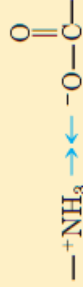

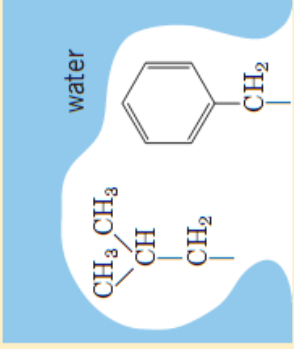


- As macromoléculas biológicas parece, numa primeira análise, distanciar-se dos princípios simples de geometria e simetria que sabemos reger a estrutura das moléculas pequenas.

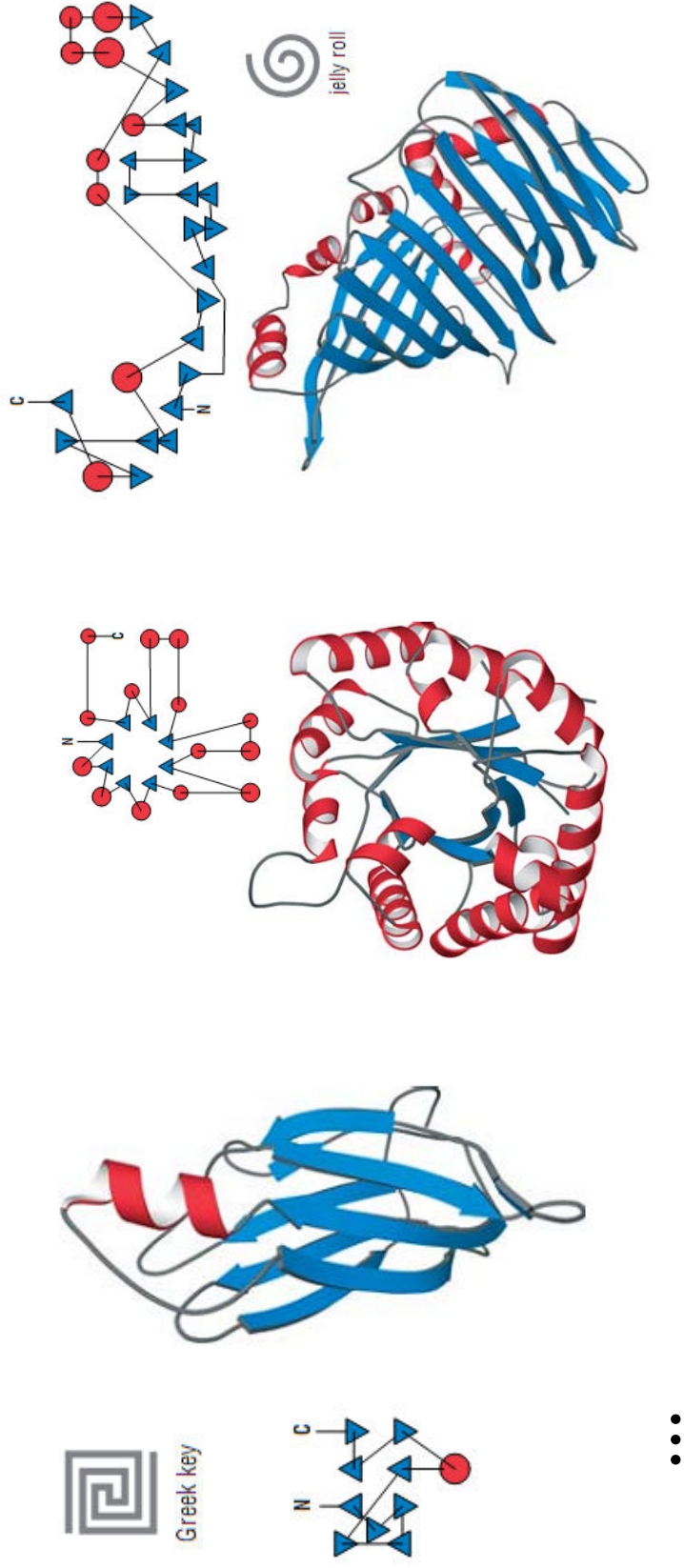
- Ligações covalentes, geometria molecular:



• Interacções não-covalentes:

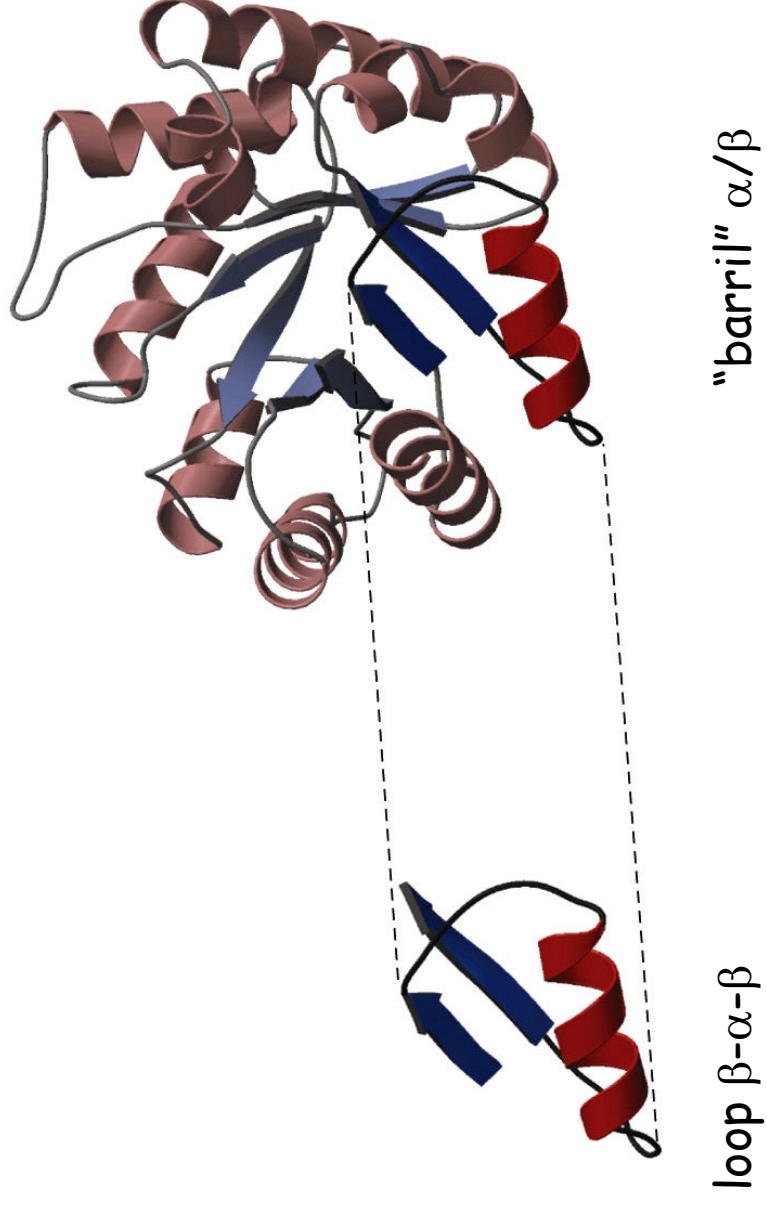
Hydrogen bonds Between neutral groups	
Between peptide bonds	
Ionic interactions	
Attraction	
Repulsion	
Hydrophobic interactions	
van der Waals interactions	Any two atoms in close proximity

- Princípios arquitectónicos



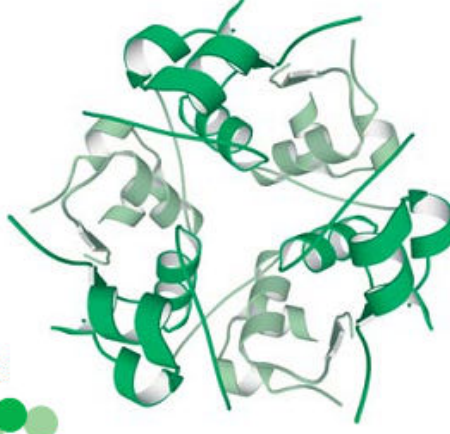
Recorrência de padrões estruturais na arquitectura das biomoléculas.

- Princípios arquitectónicos (cont.):

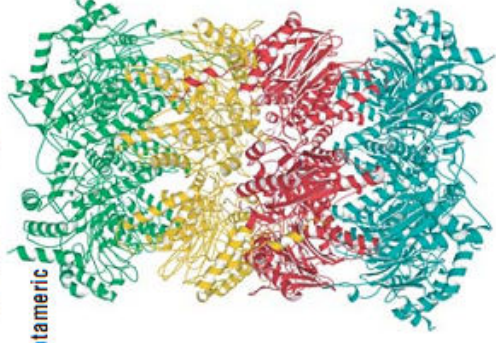
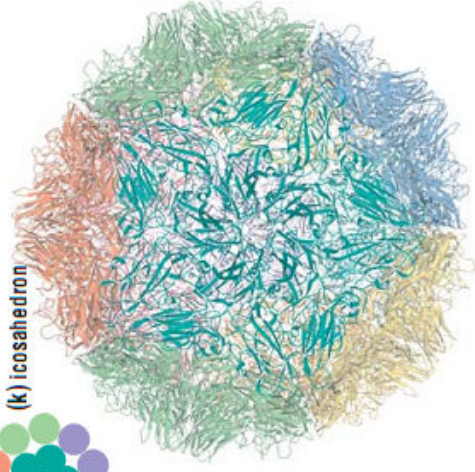
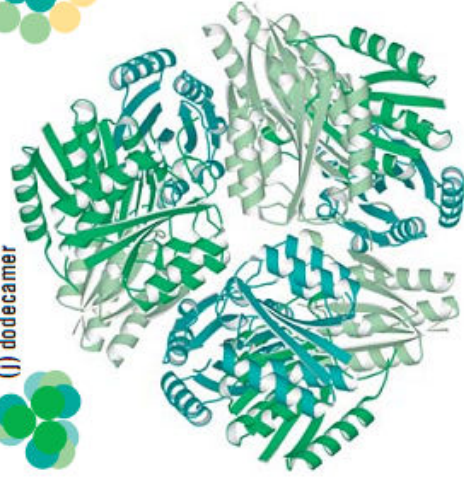
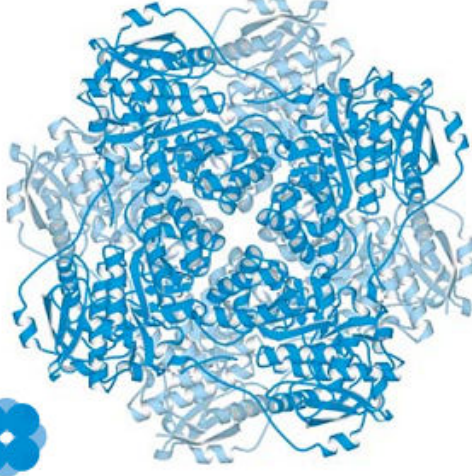
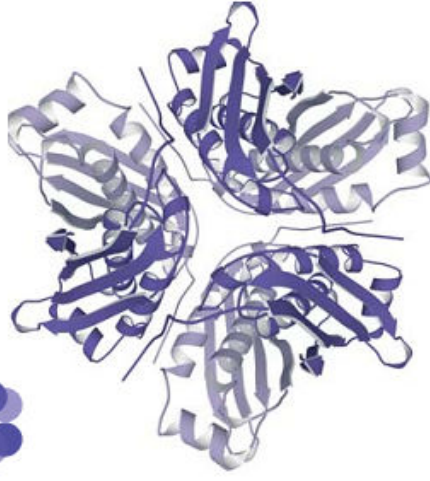


Formação de estruturas a partir da associação de unidades estruturais

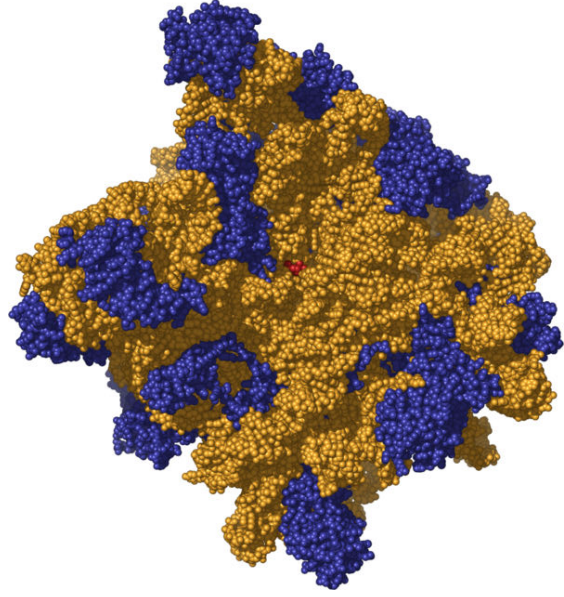
- Oligomerização:



- Oligomerização(cont.):

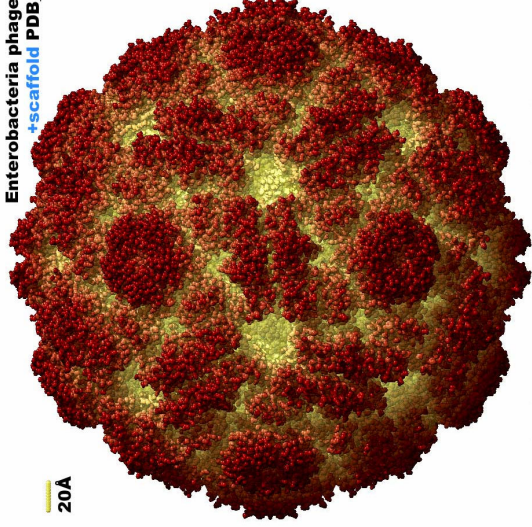


- Formação de estruturas supramacromoleculares

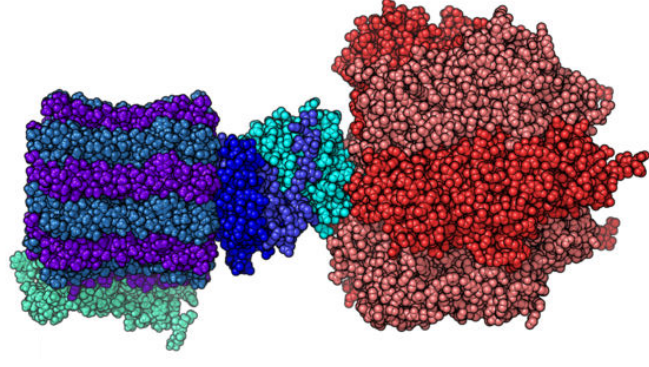


Ribossoma

Enterobacteria phage PhiX174
+scaffold PDB ID: 1AL0



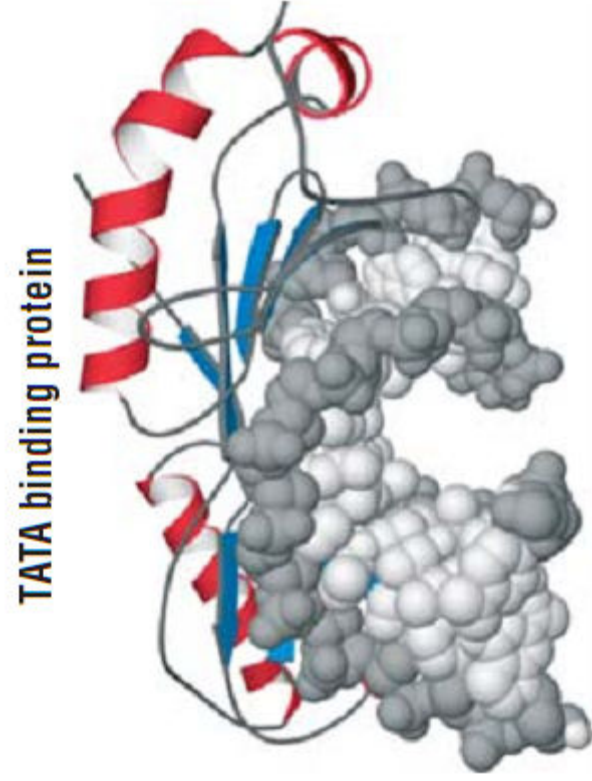
Vírus



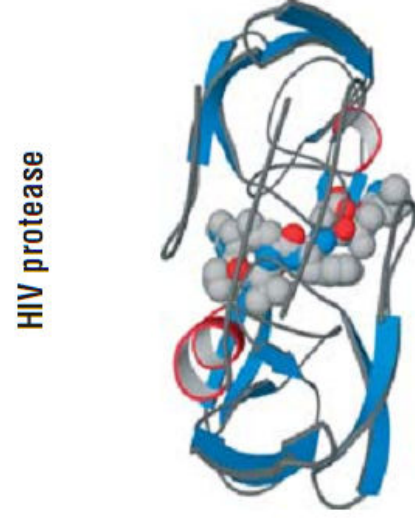
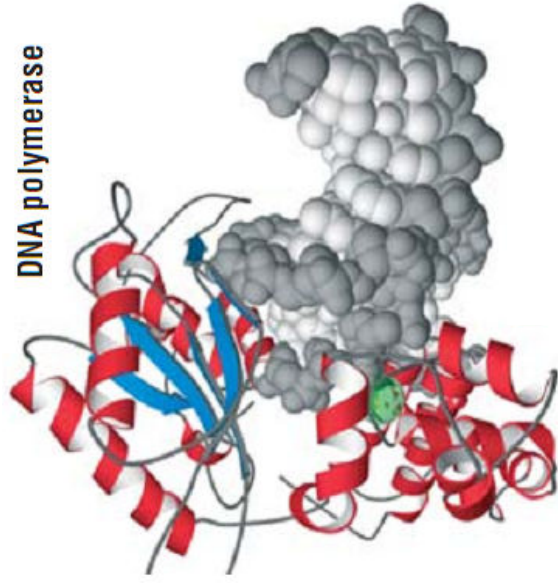
ATPase

Função

"Binding"

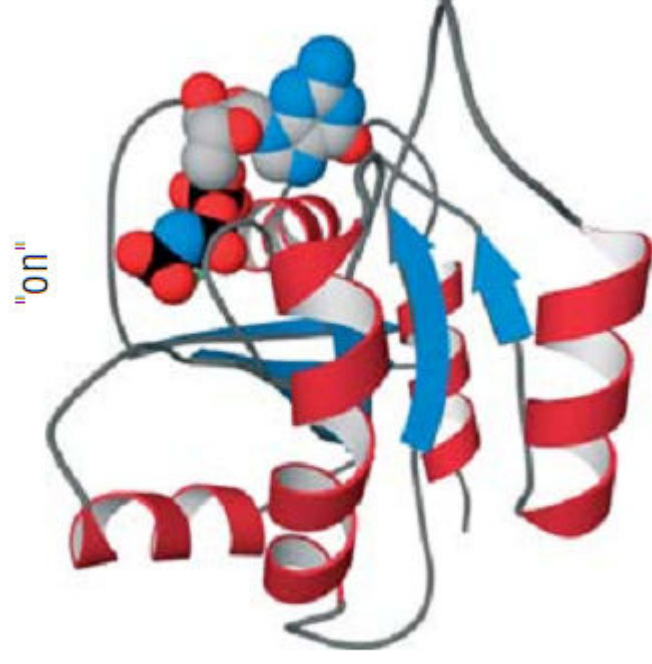
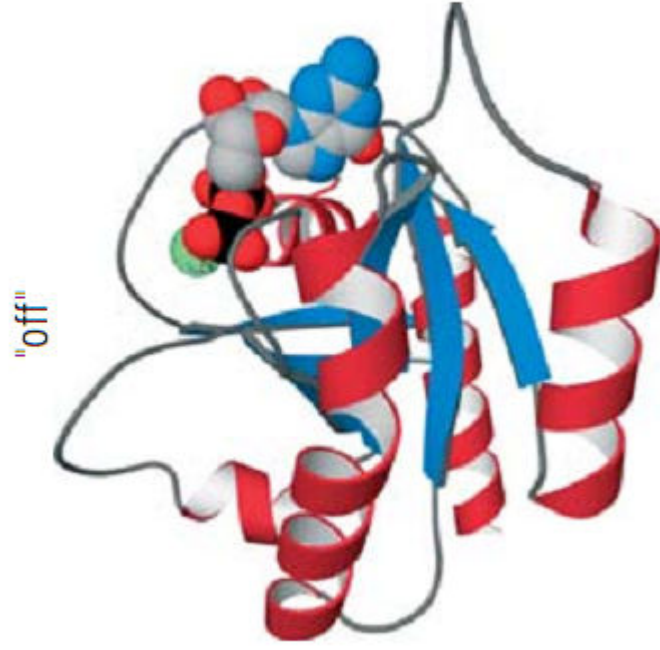


Catálise



"Switching"

Ras

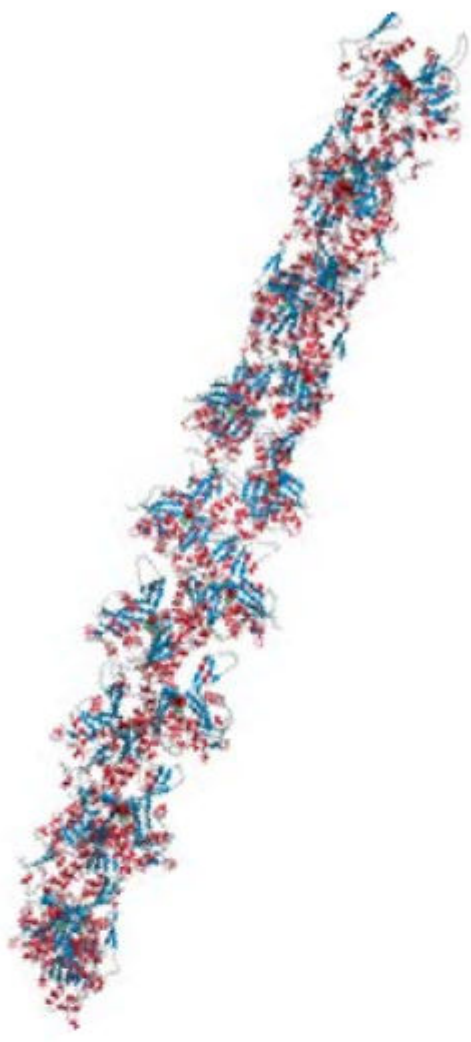


Estrutura

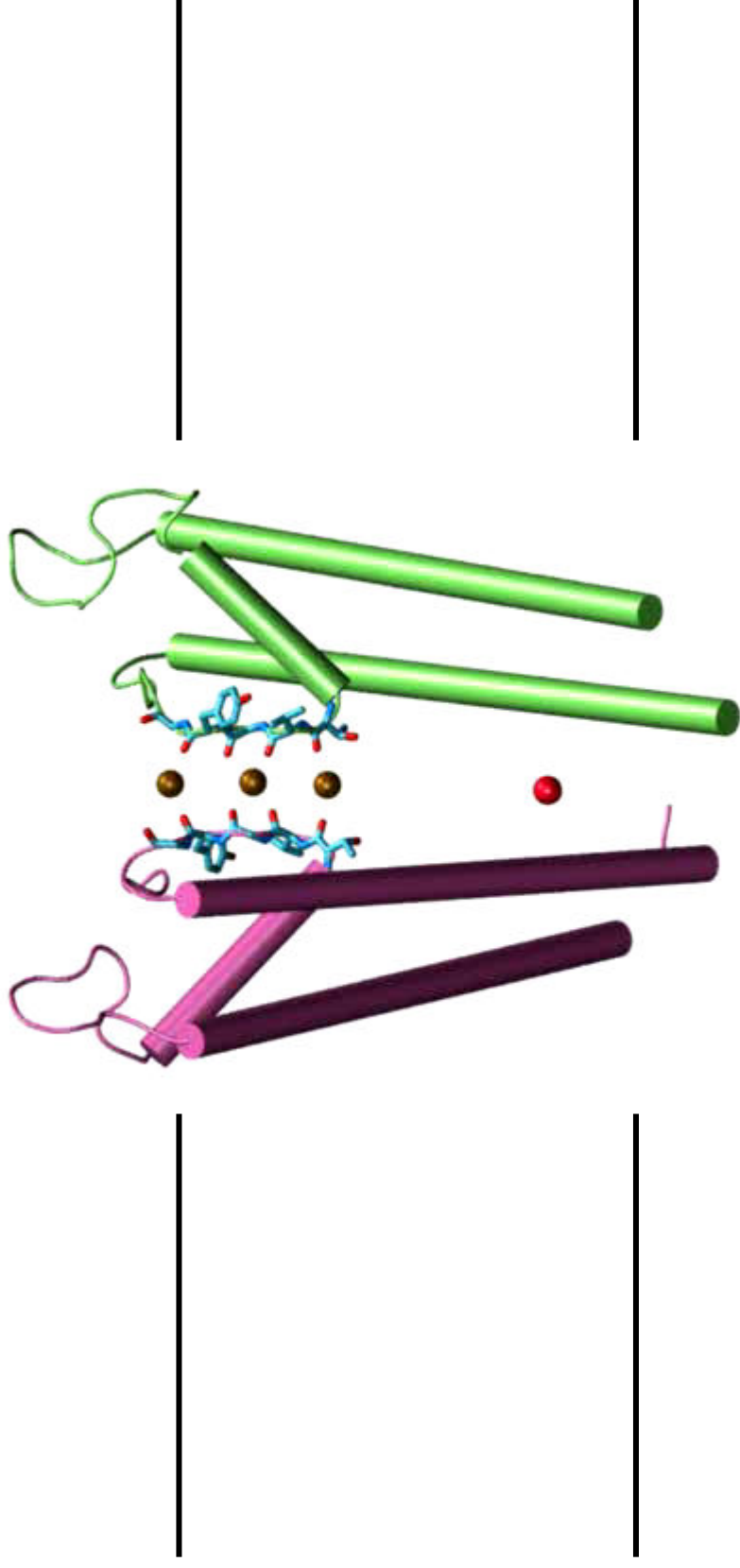
Silk



F-actin



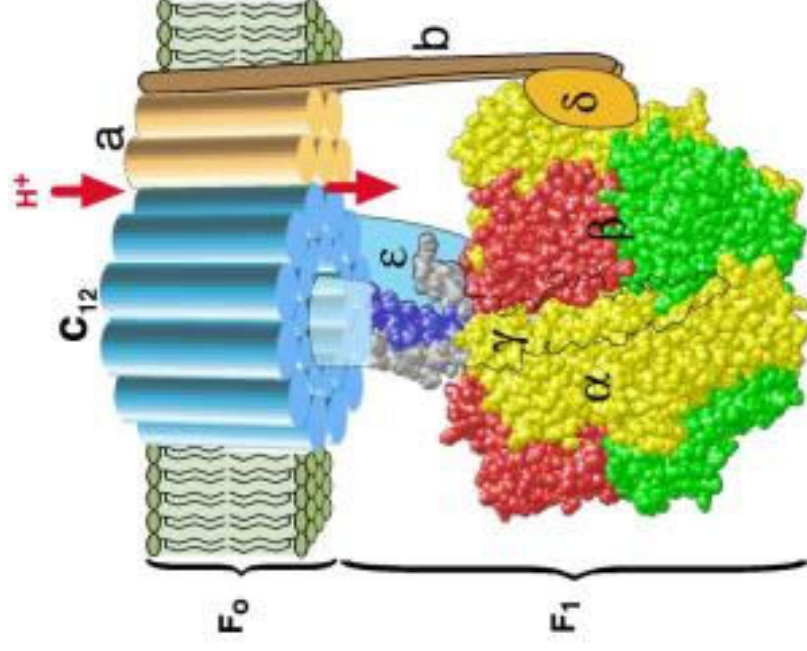
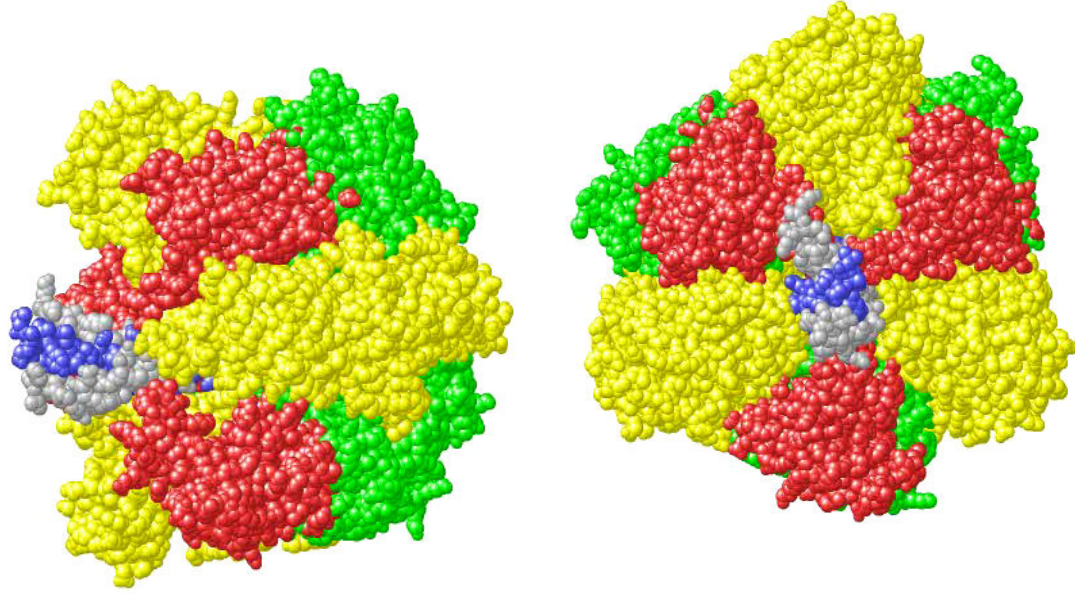
Permeabilidade



As macromoléculas biológicas funcionam como máquinas moleculares

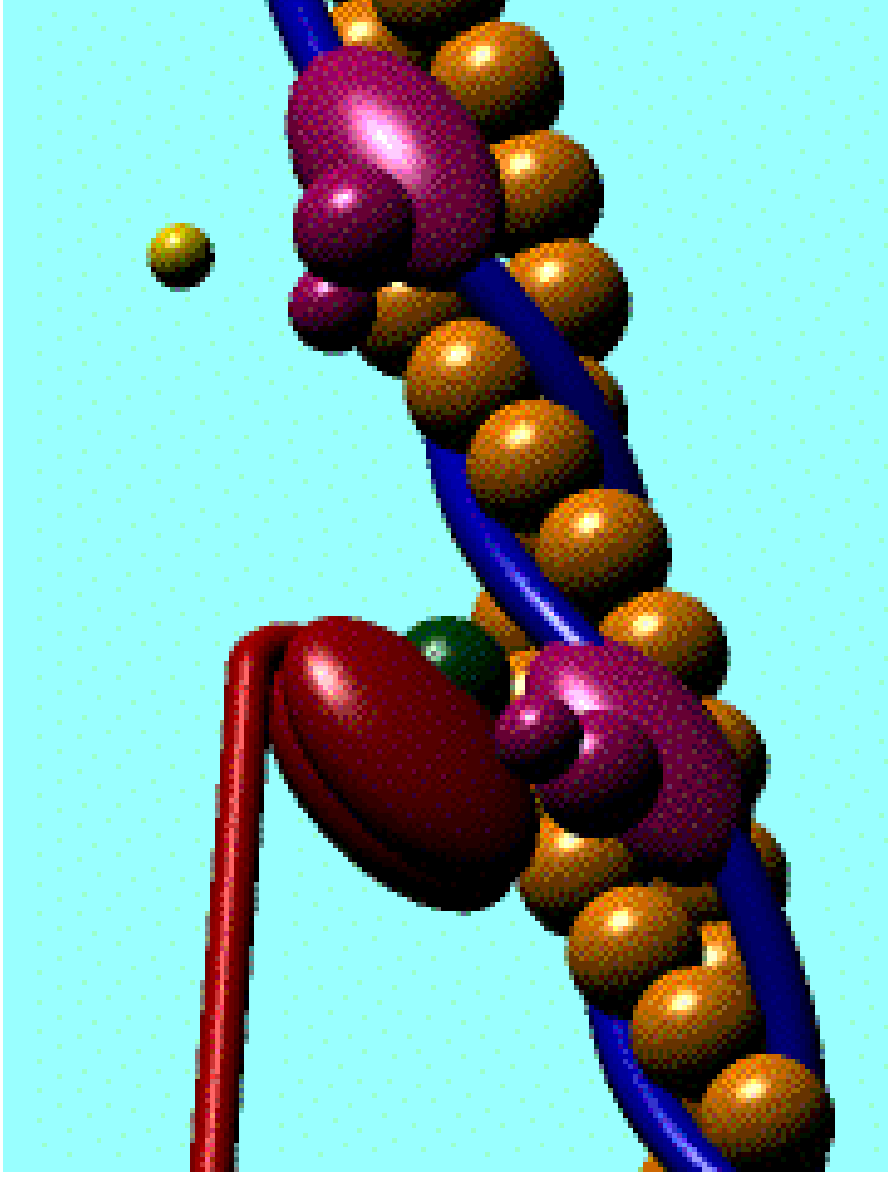
Conversão de energia

ATP sintase



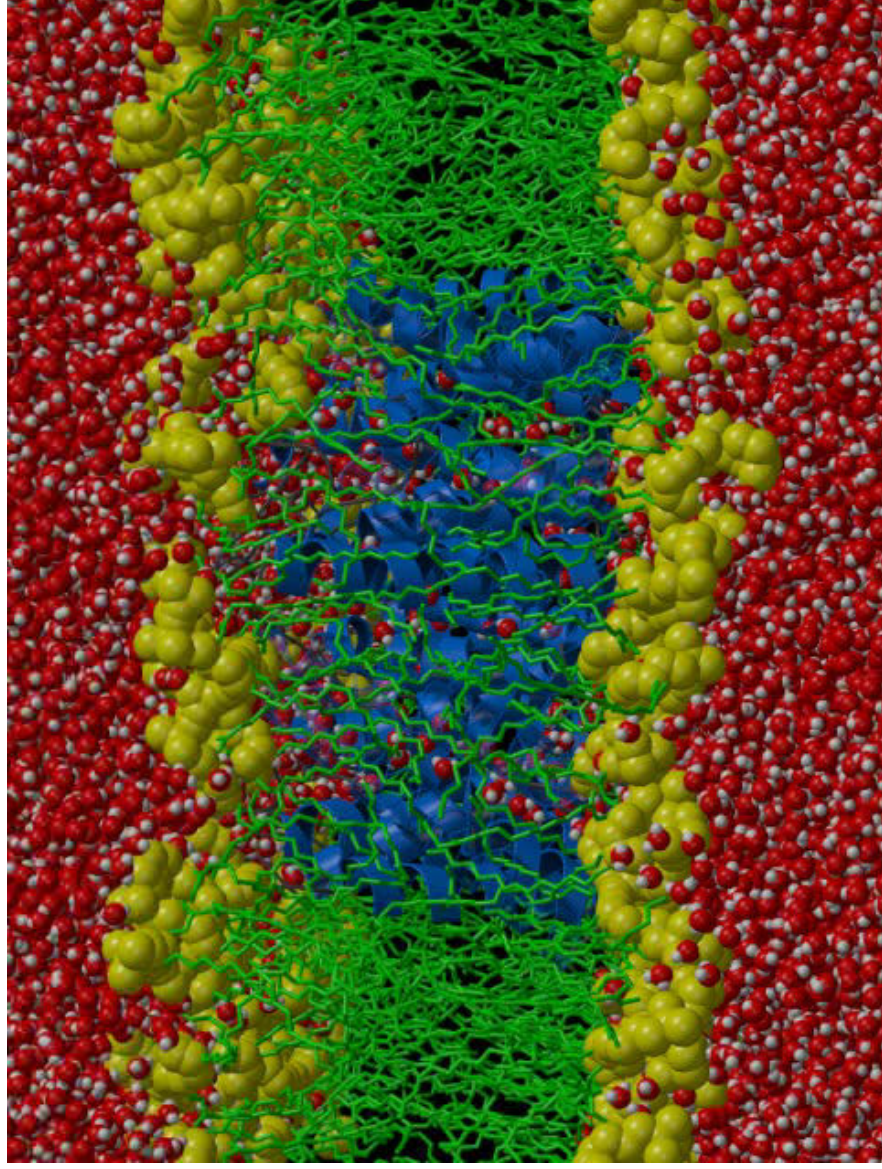
H. Wang and G. Oster (1998). Nature 396:279-282.

Motilidade



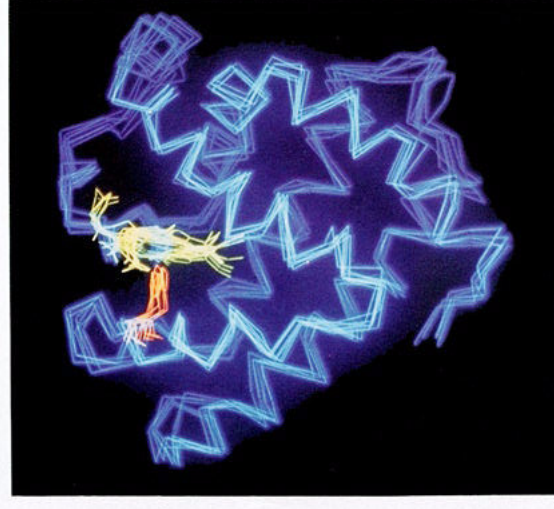
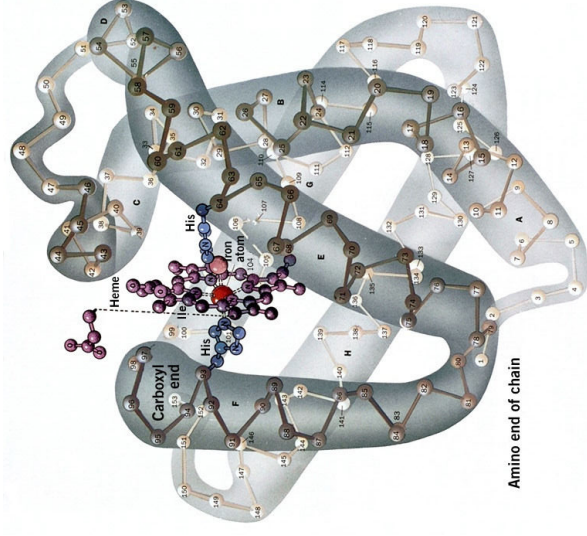
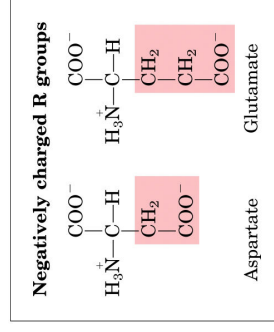
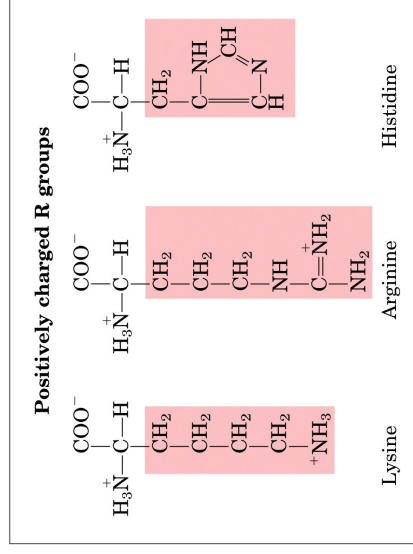
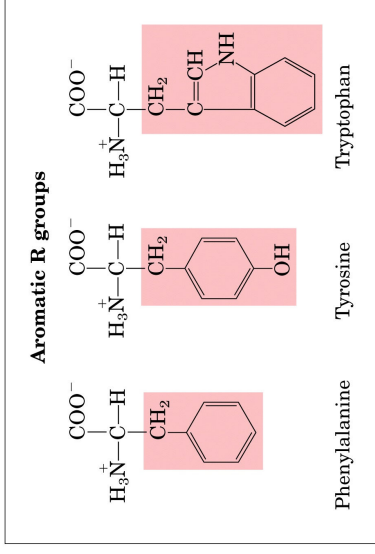
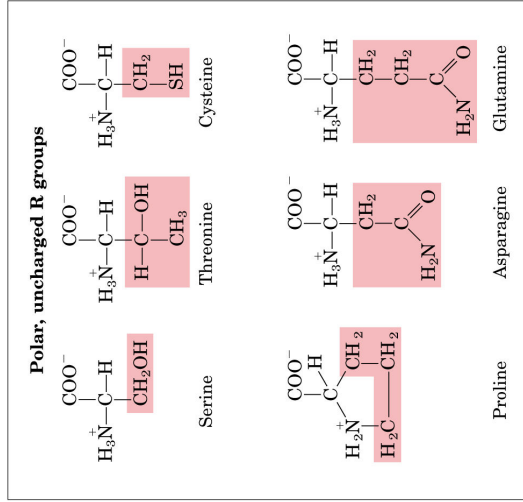
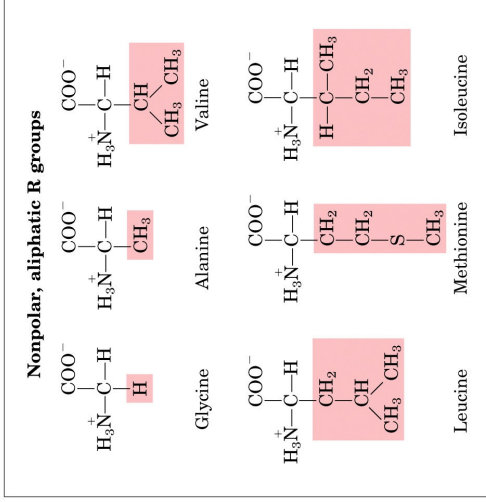
Actina+Miosina

Compartimentação



Membrana+aquaporina

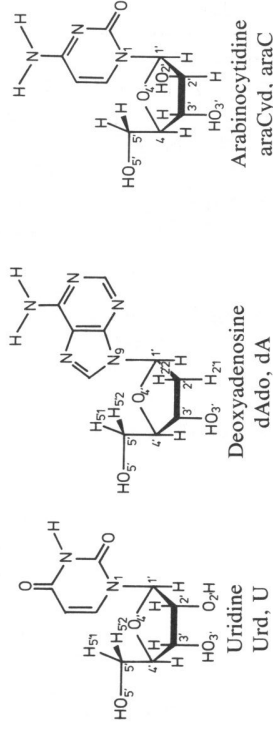
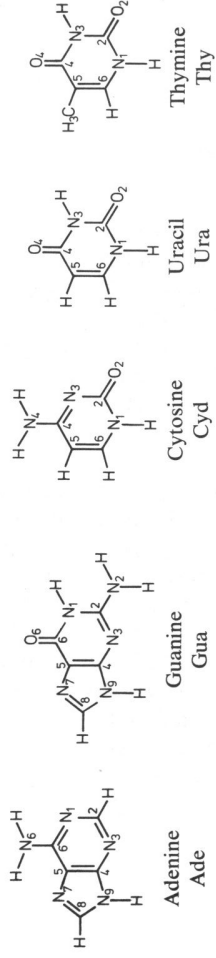
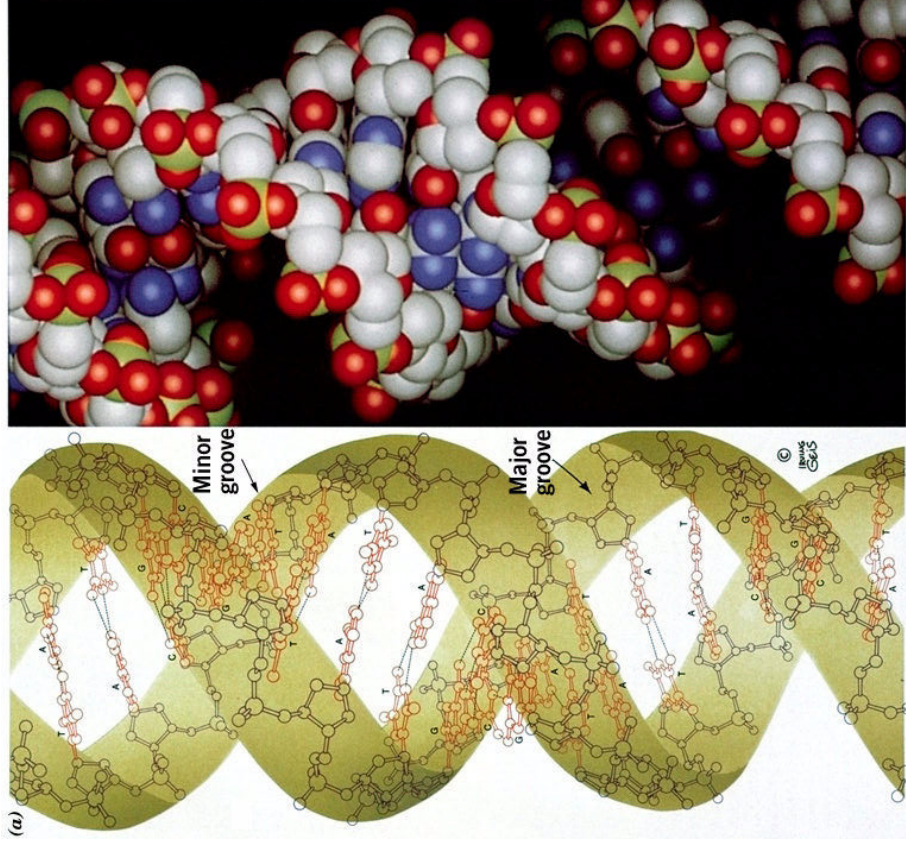
Proteínas



Mioglobina

(a)

Ácidos nucleicos



B-Dna