

Aula 3 – 10/06

Tópicos Especiais em Química Medicinal

**Tópicos Especiais
em Química Medicinal**
Código: BMF-777
Carga Horária: 45 horas
Créditos: 3 créditos

Editora UFRJ
35

Biorreceptores



O paradigma de Fischer



Emil Fischer

1852-1919

1902

E. Fischer, *Ber. Dtsch. Chem. Ges.* 1890, 23, 799



**LOCK & KEY
CONCEPT**



Estratégias da
Química Medicinal



**Novos
Fármacos
Século XX**

Biorreceptor
macrobiomolécula
baseado no sítio de
reconhecimento



Ligante
micromolécula

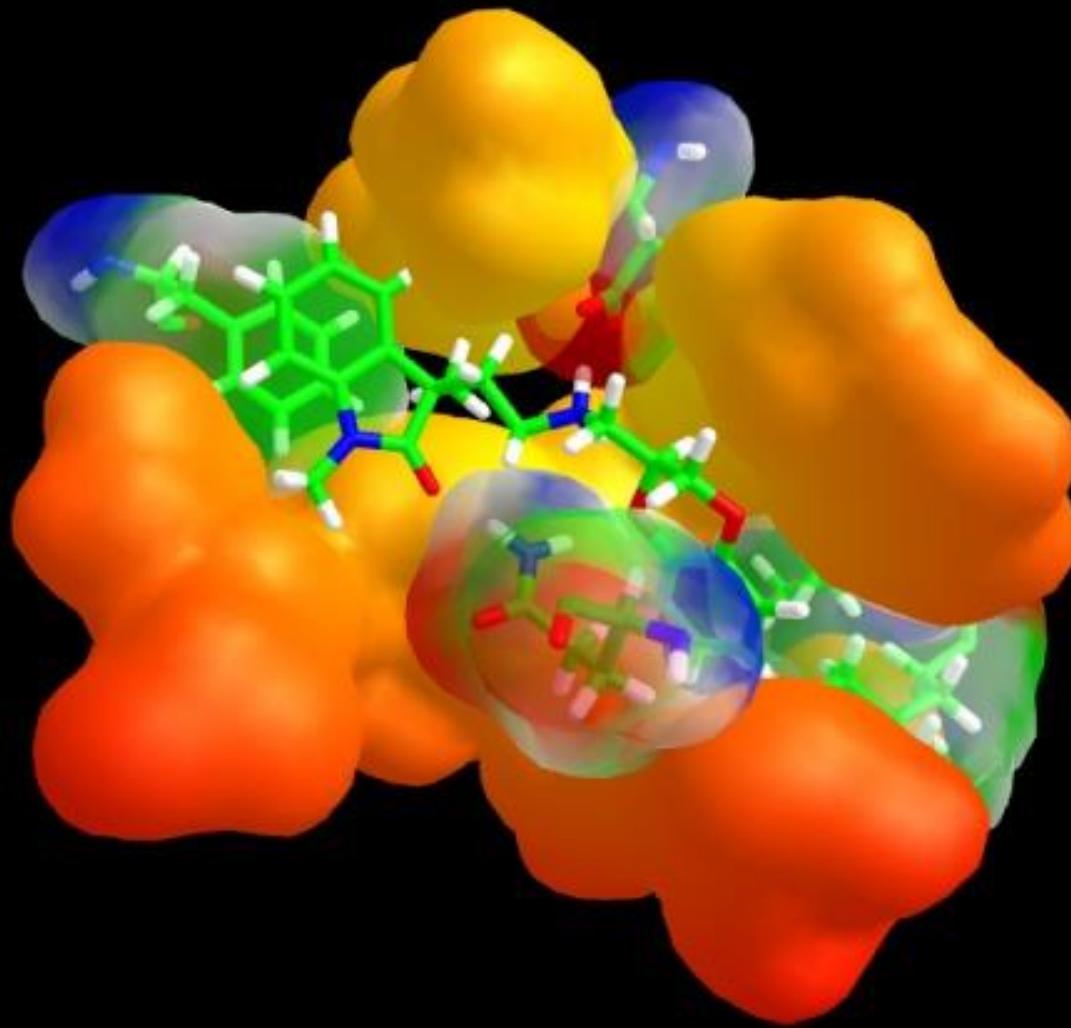
**Desenho
baseado no ligante
/ análogo-ativo**

chave



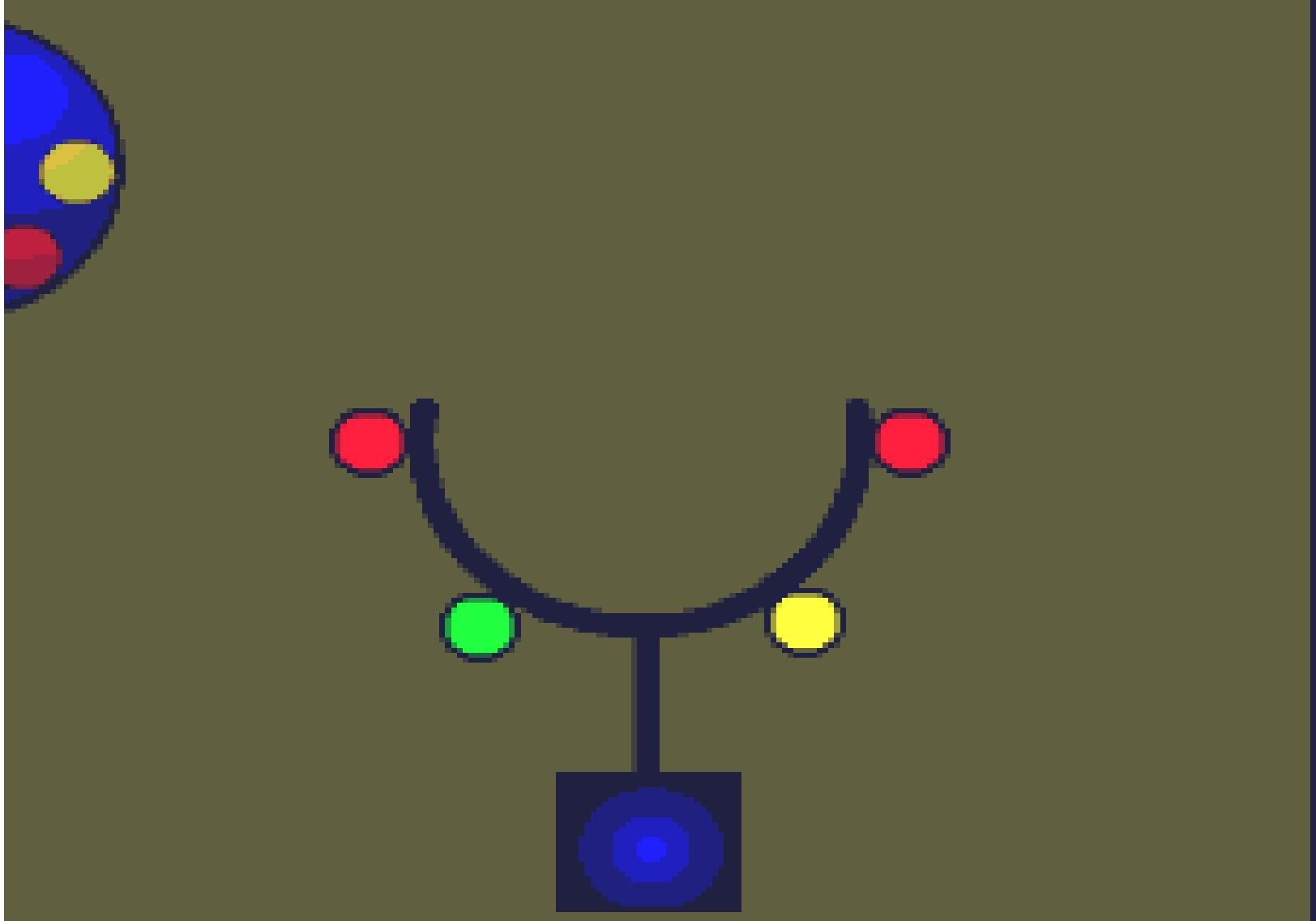
fechadura

As interações moleculares



Fármaco - Biorreceptor

Reconhecimento & complementaridade molecular



Potência & eficácia

Os fármacos atuam em alvos terapêuticos...

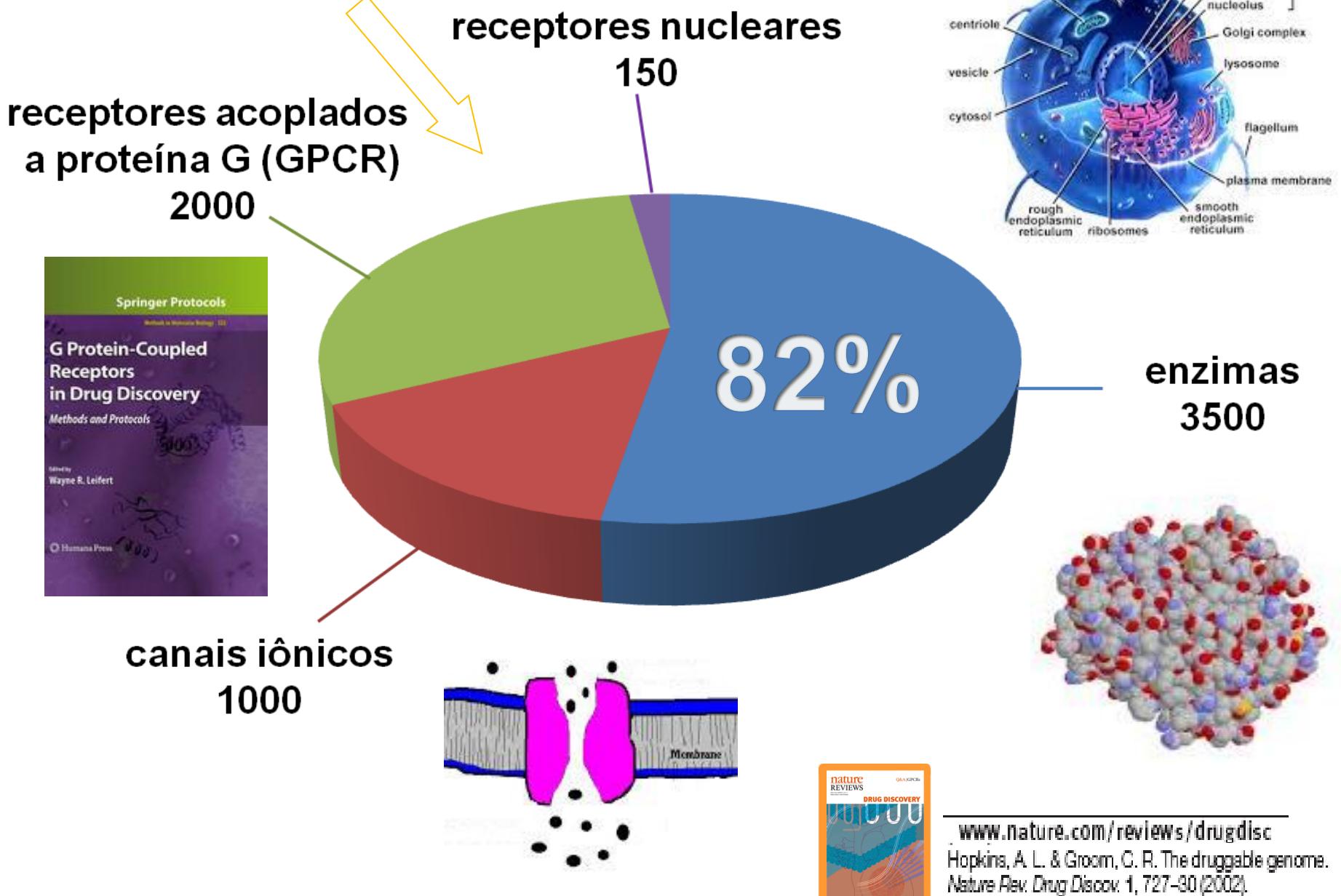
... os biorreceptores.

São 4 os tipos de alvos-
terapêuticos envolvidos na

resposta de todos os fármacos:

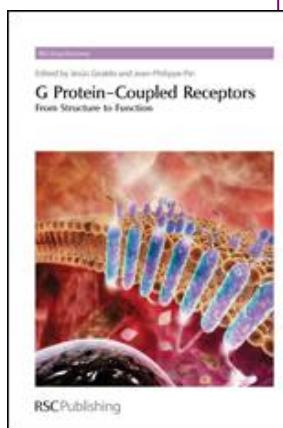
GPCR; Quinases; RH; Nucleares

A maioria dos biorreceptores dos fármacos contemporâneos são enzimas ...

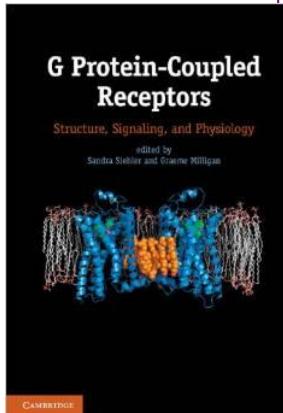




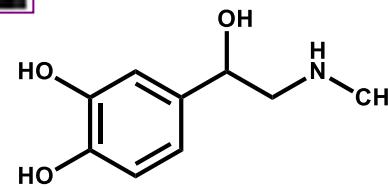
Receptores acoplados a proteína-G



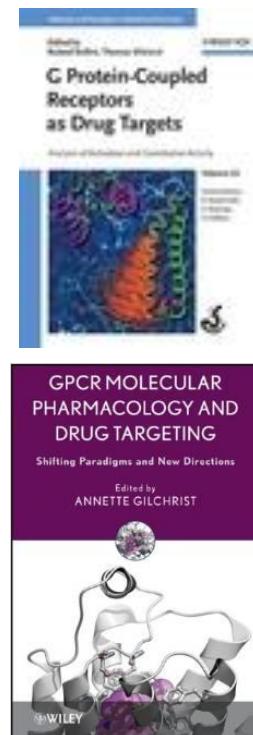
2012



Robert J. Lefkowitz
Duke University

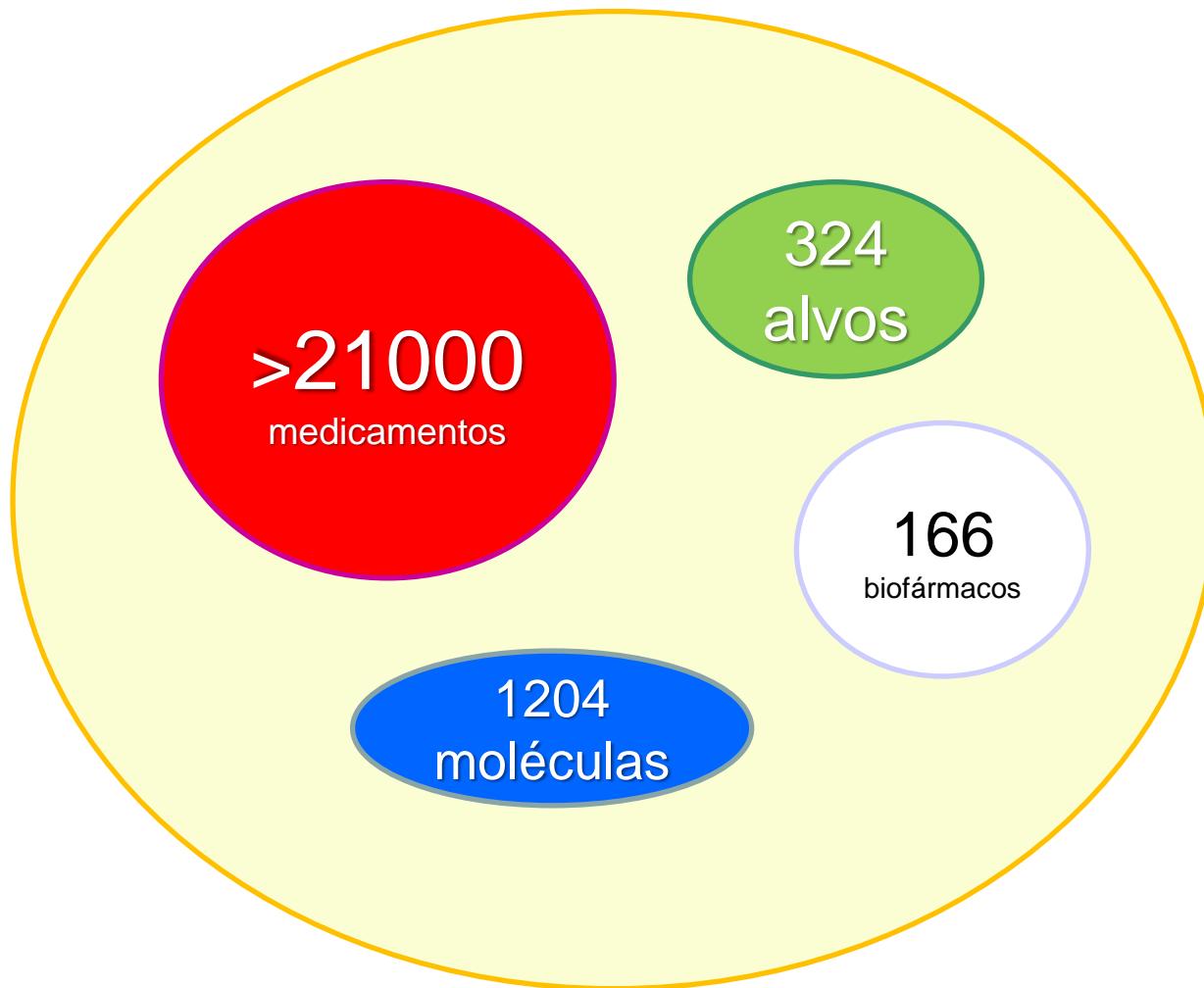


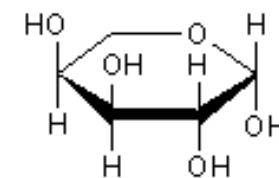
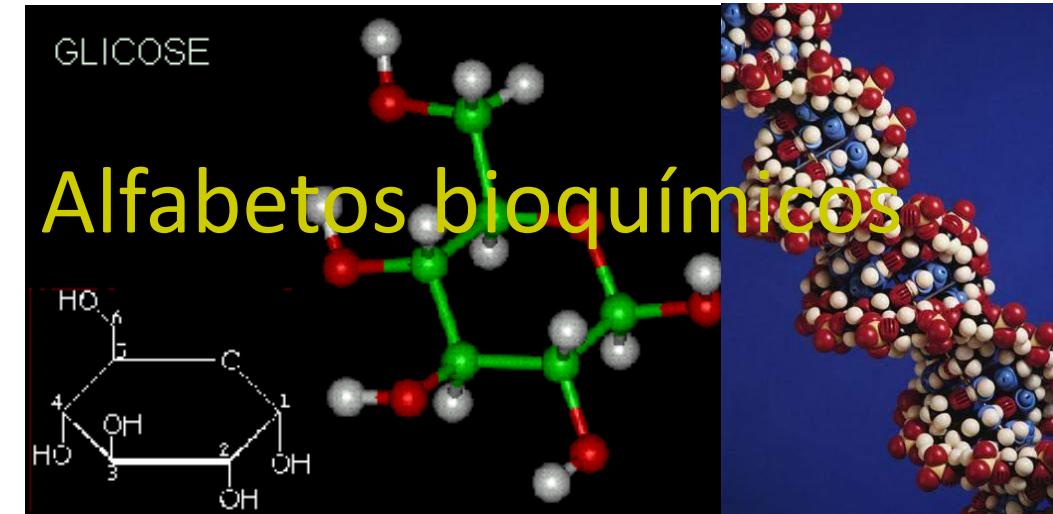
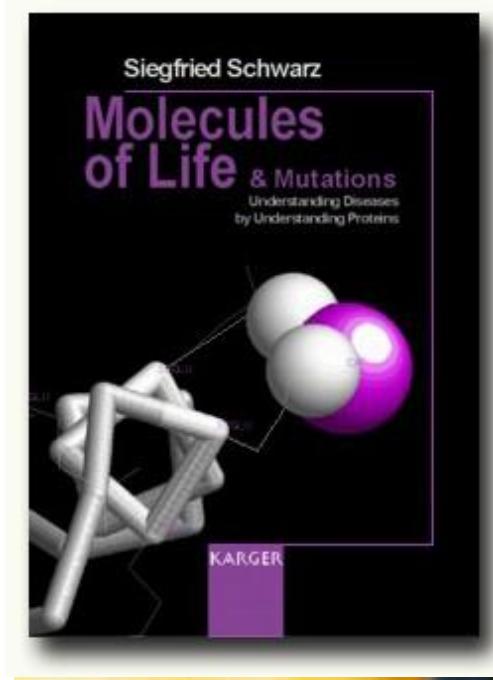
Brian K. Kobilka
Duke University



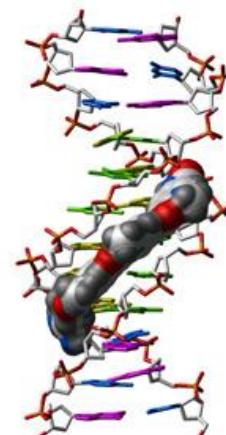
- a) Howard Hughes Medical Institute & Duke University Medical Center, Durham, NC, USA
b) Stanford University School of Medicine, Stanford, CA, USA

GPCR's





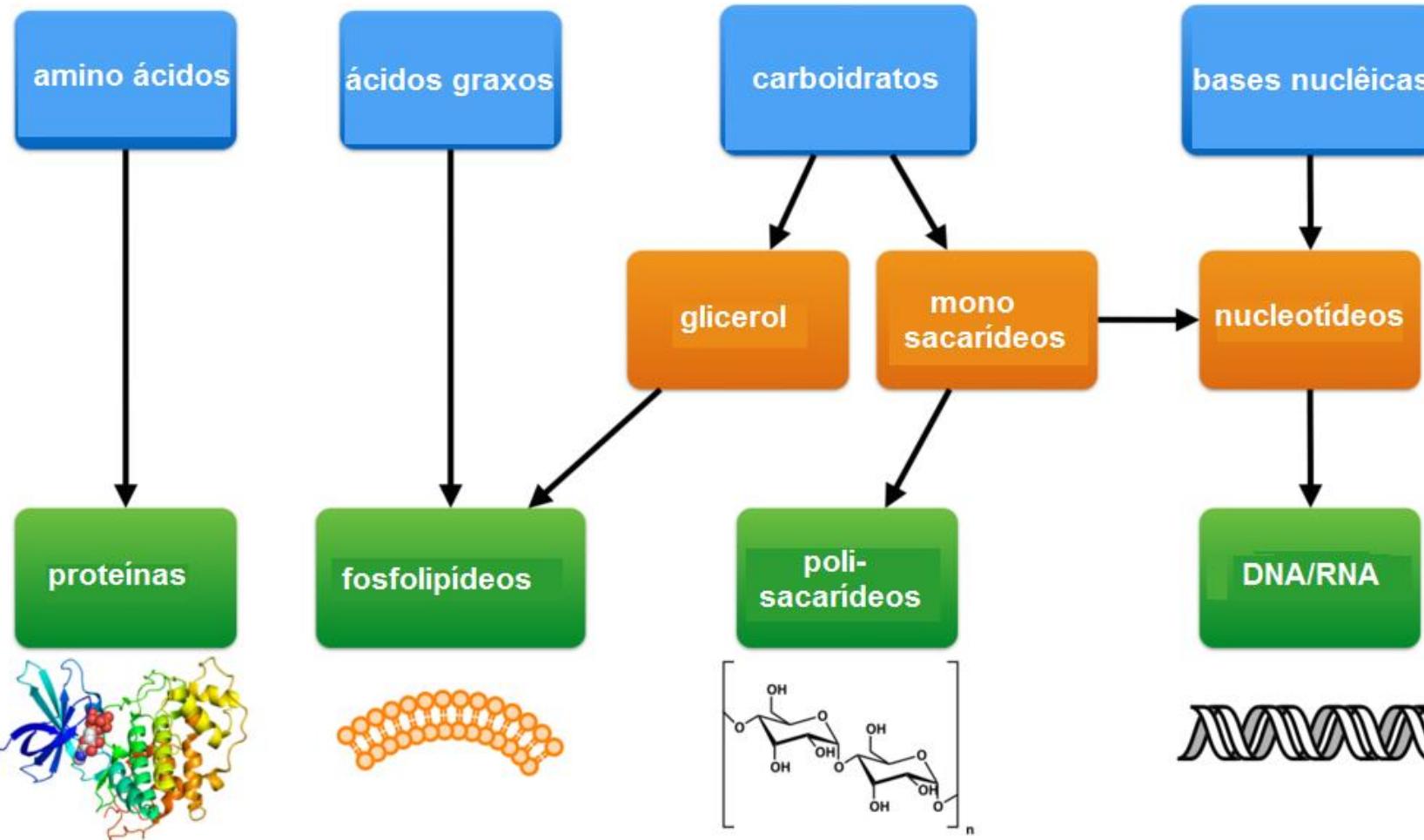
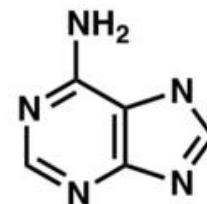
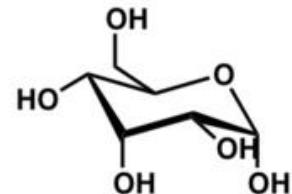
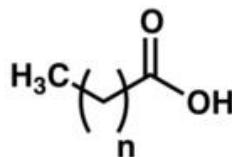
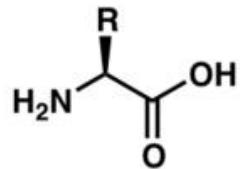
β -D-Arabinose



Model Compound Bound to the Minor
Groove of a DNA Molecule

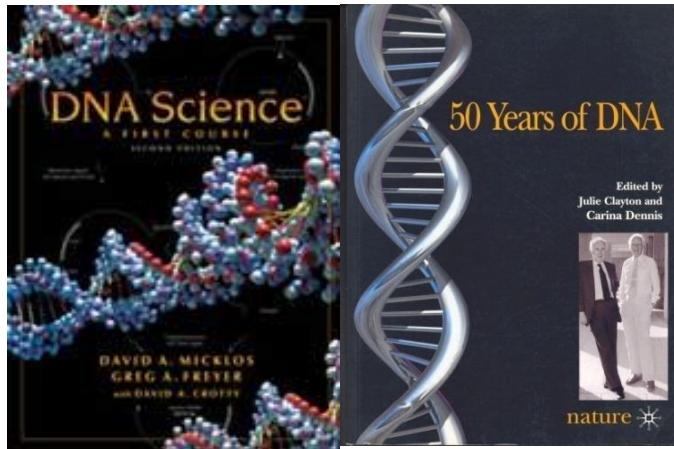
*Carbohidratos
Lipídeos
ácidos nucléicos
proteínas*

Alfabetos bioquímicos

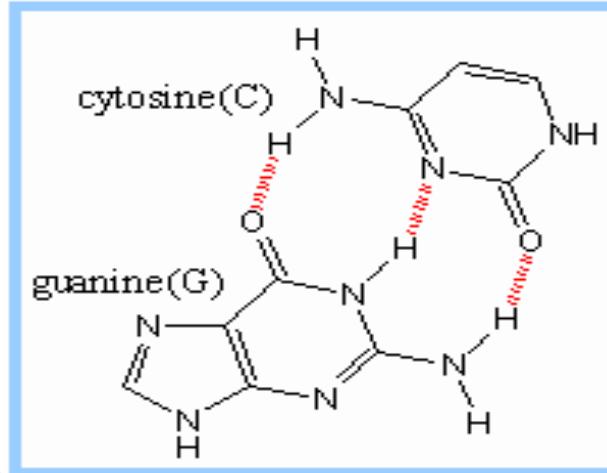




“for their discoveries concerning the molecular structure of nucleic acids and its significance for information transfer in living material”
Prêmio Nobel de Medicina e Fisiologia 1962



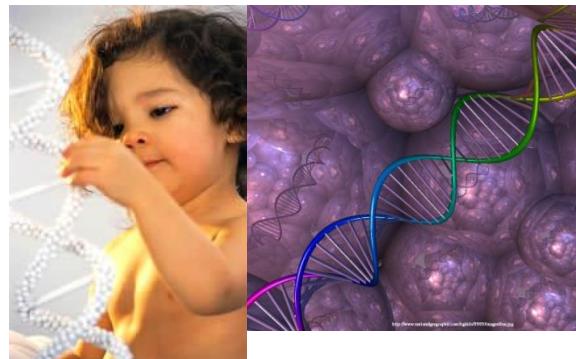
J. Clayton & C. Dennis, Eds., “50 Years of DNA”,
Nature Pub. 2003.



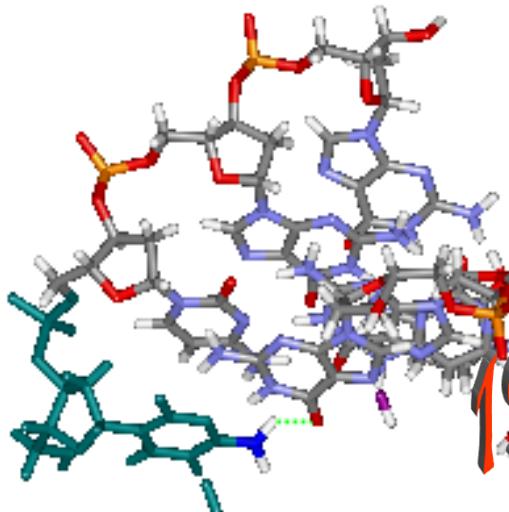
65 anos
65 anos



Ligações de Hidrogênio entre
Guanina (G) / Citosina (C) e
Adenina (A) / Timidina (T)



Moléculas da vida...



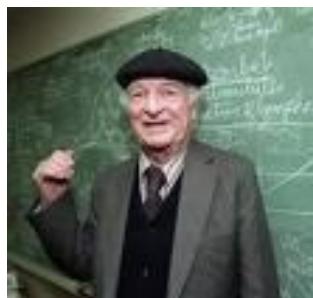
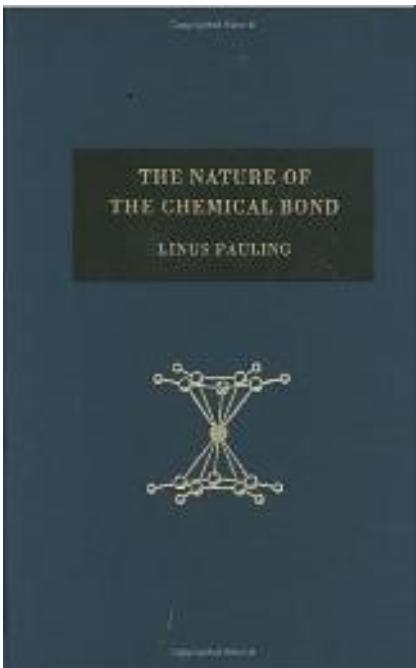
65 anos

DNA

1953-2018

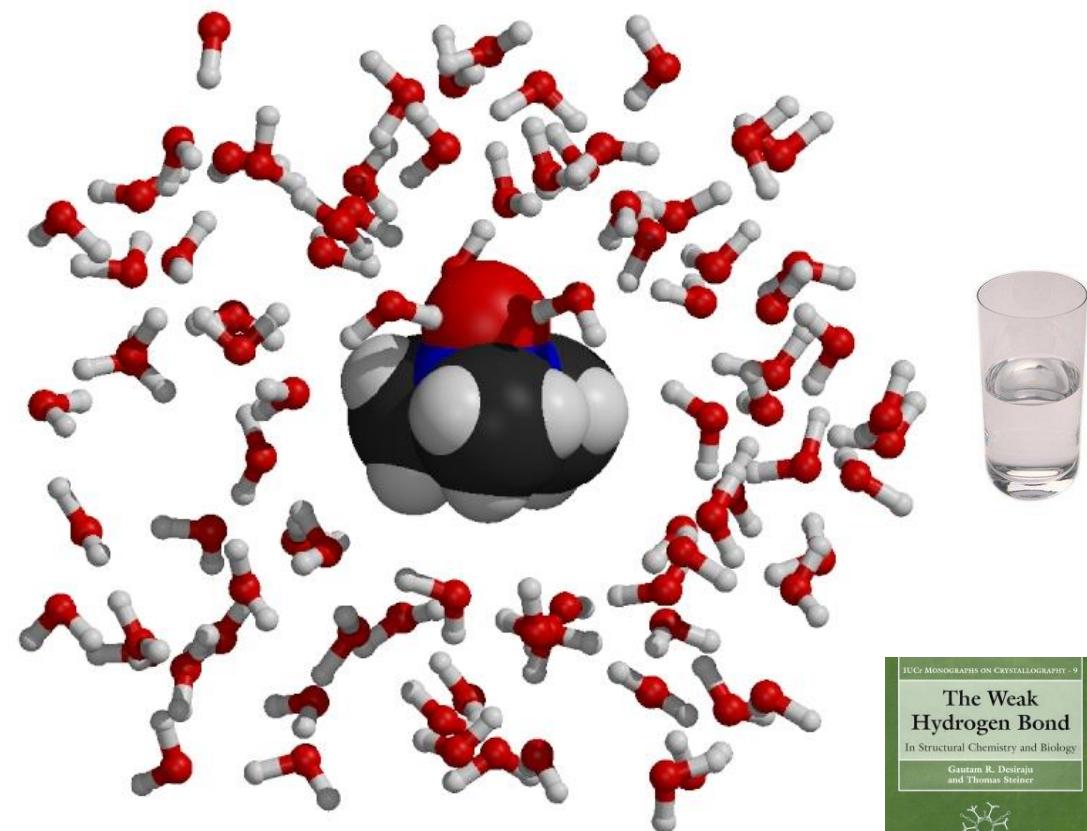


A importância das “ligações” frágeis...



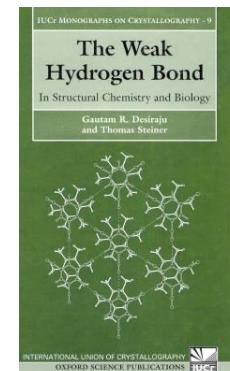
Linus C. Pauling
1901-1994

1954, 1962



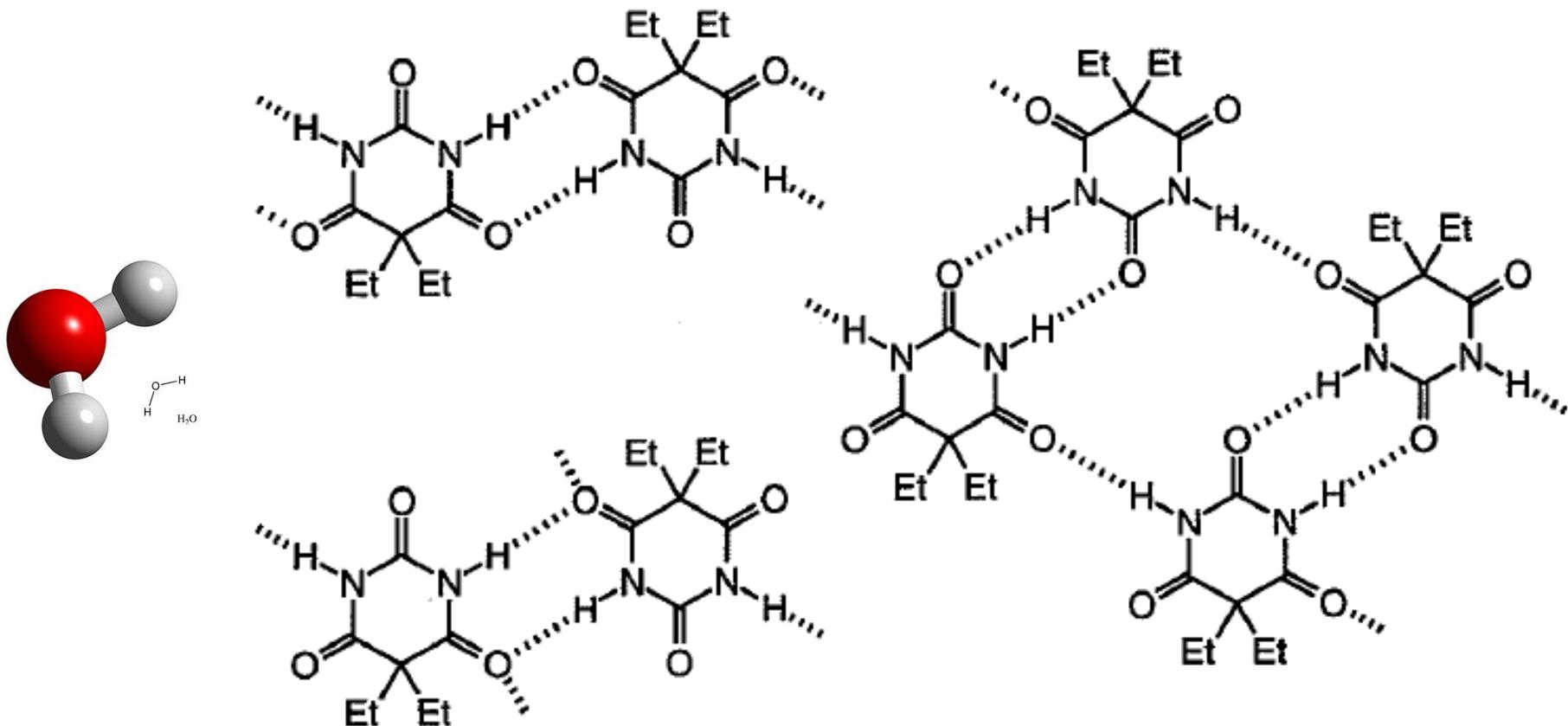
“ligações”
de hidrogênio ...

ligação-H



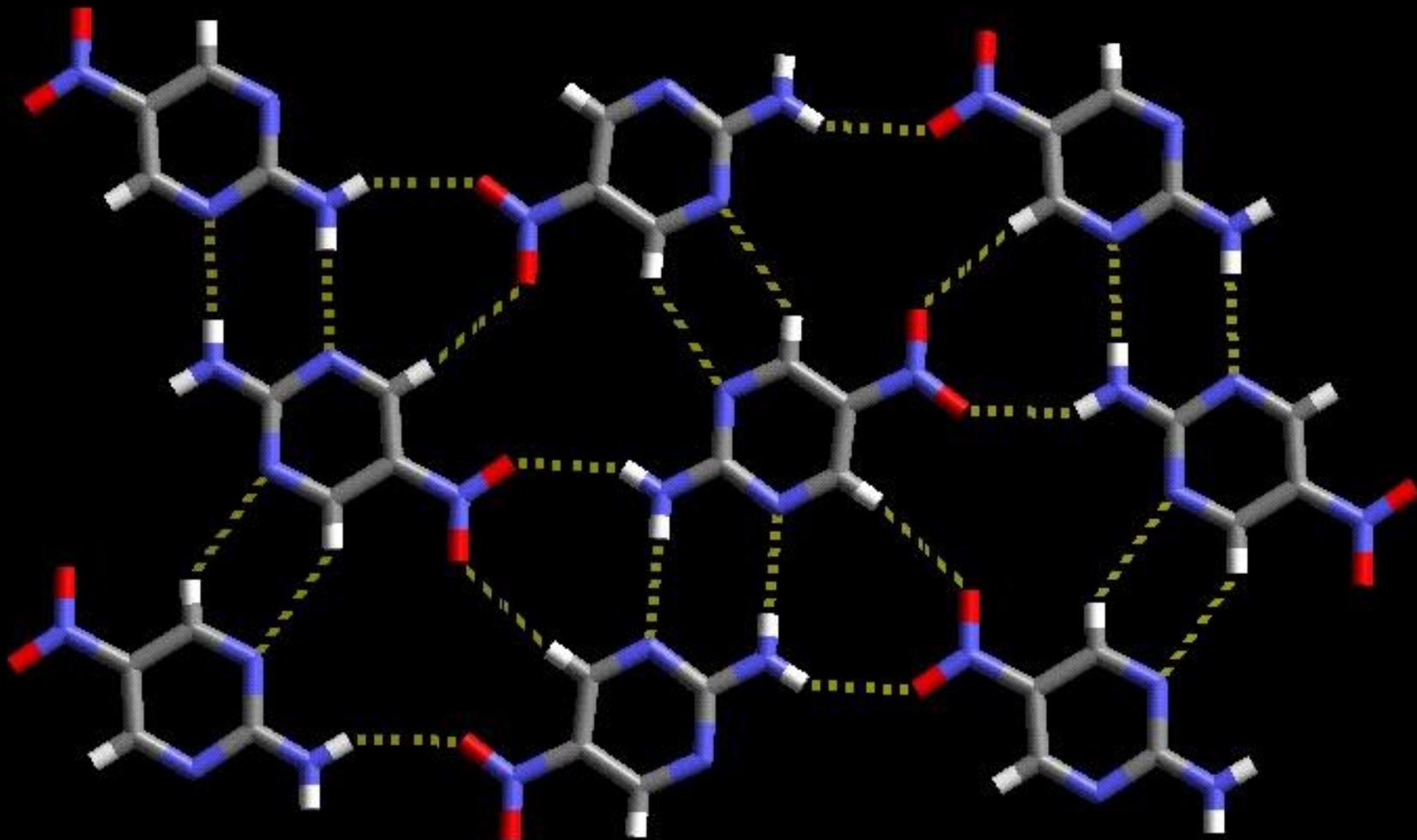


Consequências interação-H

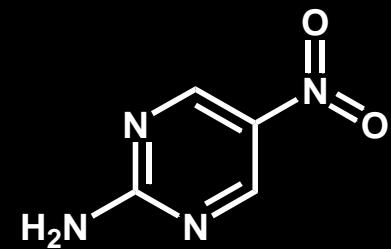


polymorphs of 5,5-diethylbarbituric acid.

ligação-H

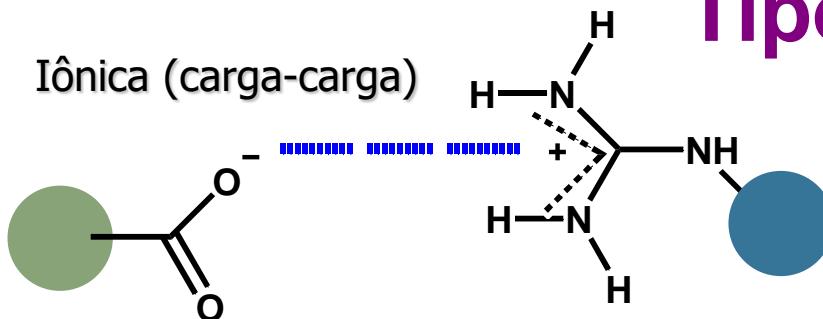


5-nitro-2-aminopirimidina



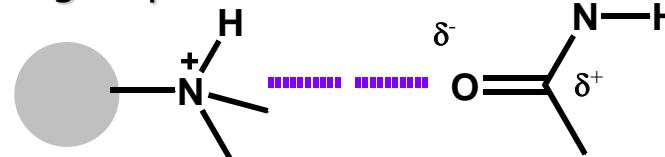
Tipos de interações F-R

Iônica (carga-carga)



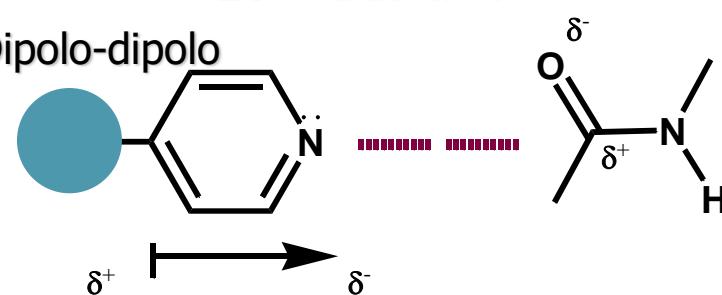
$\Delta G = 20-40 \text{ kJ/mol}$

Carga-dipolo



$\Delta G = 12-20 \text{ kJ/mol}$

Dipolo-dipolo

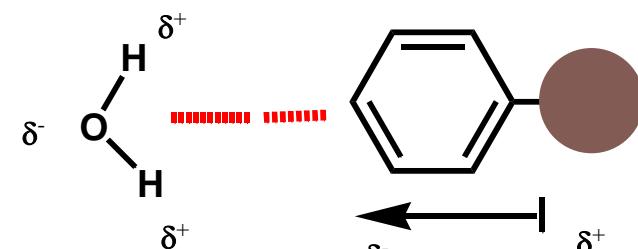


Carga-dipolo
induzido



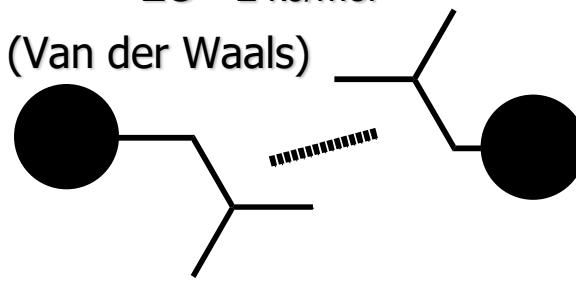
$\Delta G = 2-10 \text{ kJ/mol}$

Dipolo induzido-dipolo



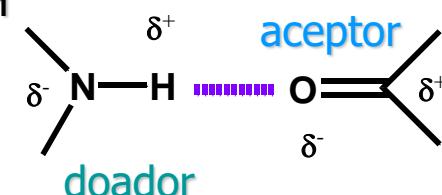
$\Delta G = 2 \text{ kJ/mol}$

Dispersão (Van der Waals)



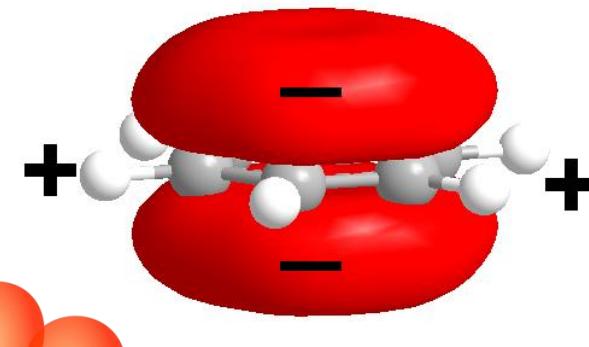
$\Delta G = 2-4 \text{ kJ/mol}$

Ligaçāo-H



$\Delta G = 4-30 \text{ kJ/mol}$

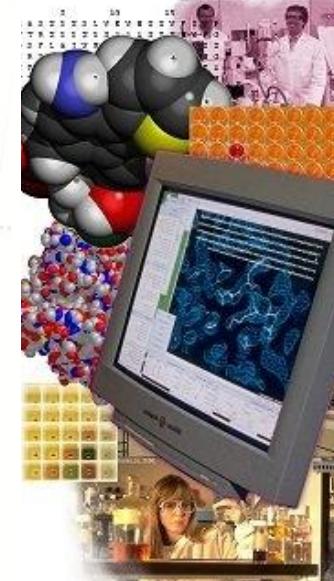
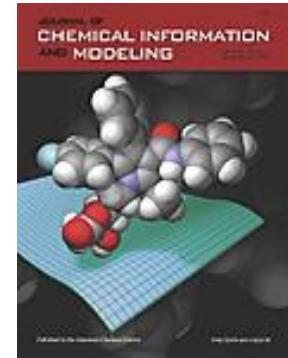
Interações π



cation-π		cation: Lys, Arg π donor: Phe, Trp, Tyr	0.2–2.5
$\pi-\pi$	<p>sandwich T-shaped parallel displaced</p>	Phe, Tyr, Trp, His	0.5–1.0
van der Waals	<p>drug side chains</p>	Ala, Val, Leu,Ile	0.5–1.0



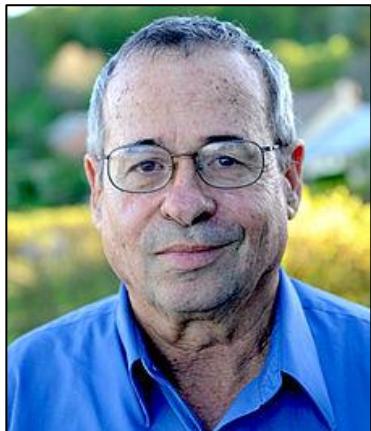
Química Computacional



Modelagem molecular



Química Computacional



Ariel Warshel
(1940 -)



2013

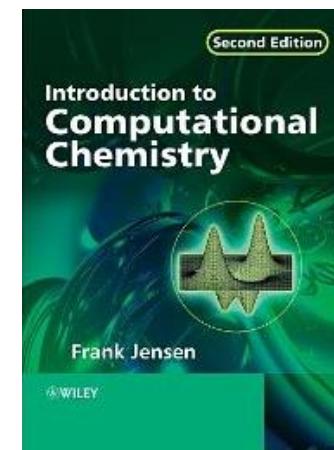


Michael Levitt
(1947 -)



Martin Karplus
(1930 -)

Modelagem & dinâmica molecular



"for the development of multiscale models for complex chemical systems"



Access the PDB FTP:

[RCSB PDB](#) | [PDBe](#) | [PDBj](#)

[Archive Download](#)

[Chemical Component Dictionary](#)

Deposit Data to the PDB:

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[PDBj](#) | [BMRB](#)

Search for Structures:

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Instructions to Journals

Documentation

[Format](#)

[Annotation and Policies](#)

Workshops and Task Forces

[X-ray Validation](#)

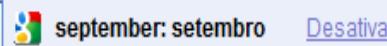
[NMR Validation](#)

wwPDBAC

Data Download Details

The wwPDB ftp site is quite large. As of September 21, 2008, it contains more than 443,000 files and requires over 94 GBbytes of storage. It will continue to grow with each weekly update. Fresh [downloads will require substantial amount](#) of time.

A number of download locations and options are provided.



Access is available from: [the RCSB PDB \(USA\)](#) | [PDBe \(UK\)](#) | [PDBj \(Japan\)](#)

Please note: <ftp://ftp.rcsb.org> is no longer updated. Please access the PDB archive using one of the FTP sites listed in the left menu.

RCSB PDB:

USING RSYNC PROTOCOL:

```
rsync --port=33444 ftp.wwpdb.org::
```

`ftp` Top level of ftp tree approximately 66 GB (/pub/pdb)

`ftp_data` Data directory within ftp archive Aproximately 65.7 GB (/pub/pdb/data)

`ftp_derived` Derived data directory within ftp archive Aproximately 134 MB (/pub/pdb/derived_data)

`ftp_doc` Doc directory within ftp archive Aproximately 234 MB (/pub/pdb/doc)

Downloading coordinate files in PDB Exchange Format (mmCIF): To download the entry files in PDB exchange format the following rsync command may be used:

```
rsync -rlpt -v -z --delete --port=33444 \
rsync.wwpdb.org::ftp_data/structures/divided/mmCIF/ ./mmCIF
```

Downloading coordinate files in PDBML Format (xml): To download the entry files in PDBML format the following rsync command may be used:



Docking Molecular

SELEÇÃO DOS LIGANTES

Quimioteca (FBDD)

[Quimioteca da LASSBio](#)

ANÁLISE CONFORMACIONAL (método semi-empírico AM1)

CAPÍTULO 5

UMA INTRODUÇÃO À MODELAGEM MOLECULAR APLICADA À QUÍMICA MEDICINAL 231



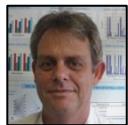
ESTRUTURA do BIORRECEPTOR

■ PDB*

■ Construção do modelo

FlexX / Gold

*PROTEIN DATA BANK



ANCORAMENTO DOS LIGANTES (hit)

Virtual Screening

SELEÇÃO DA ESTRUTURA DE REFERÊNCIA

IDENTIFICAÇÃO DO SÍTIO DE RECONHECIMENTO

Mapa farmacofórico 3D/SAR

DETERMINAÇÃO DA ENERGIA LIVRE DE INTERAÇÃO ($\Delta G_{\text{interação}}$)

Sybyl, Version 8.0, Tripos Associates

Spartan Pro 14.0, Wavefunction, Inc.

KNIME; Biovia Discovery

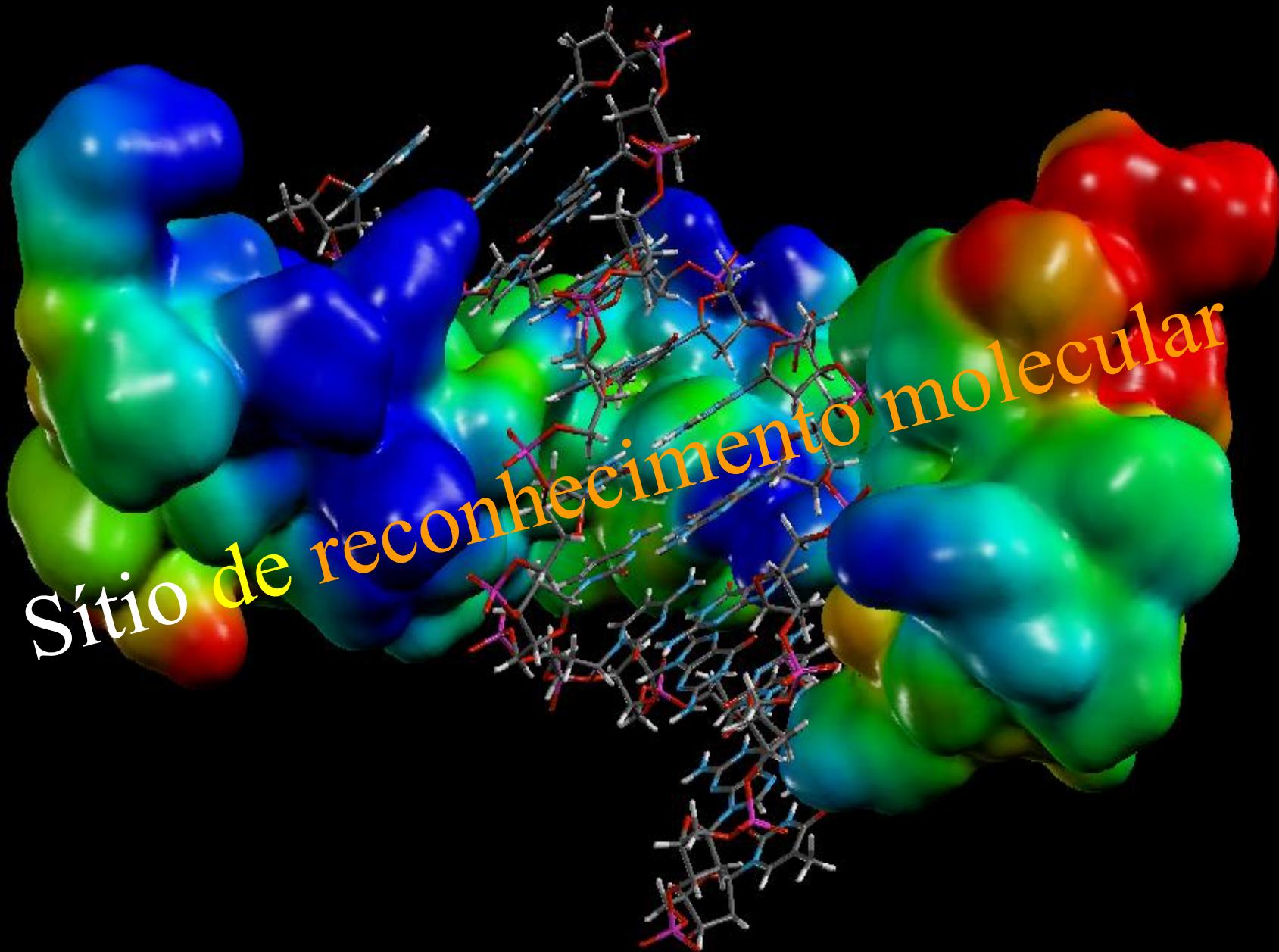
GOLD (Jones et al. 1997; Portal da CAPES); Glide (Schrödinger)

**Hit
($10 \mu\text{M}$; $>30 \mu\text{M}$)**

**Validação
(h_2I)**



Sítio de reconhecimento molecular



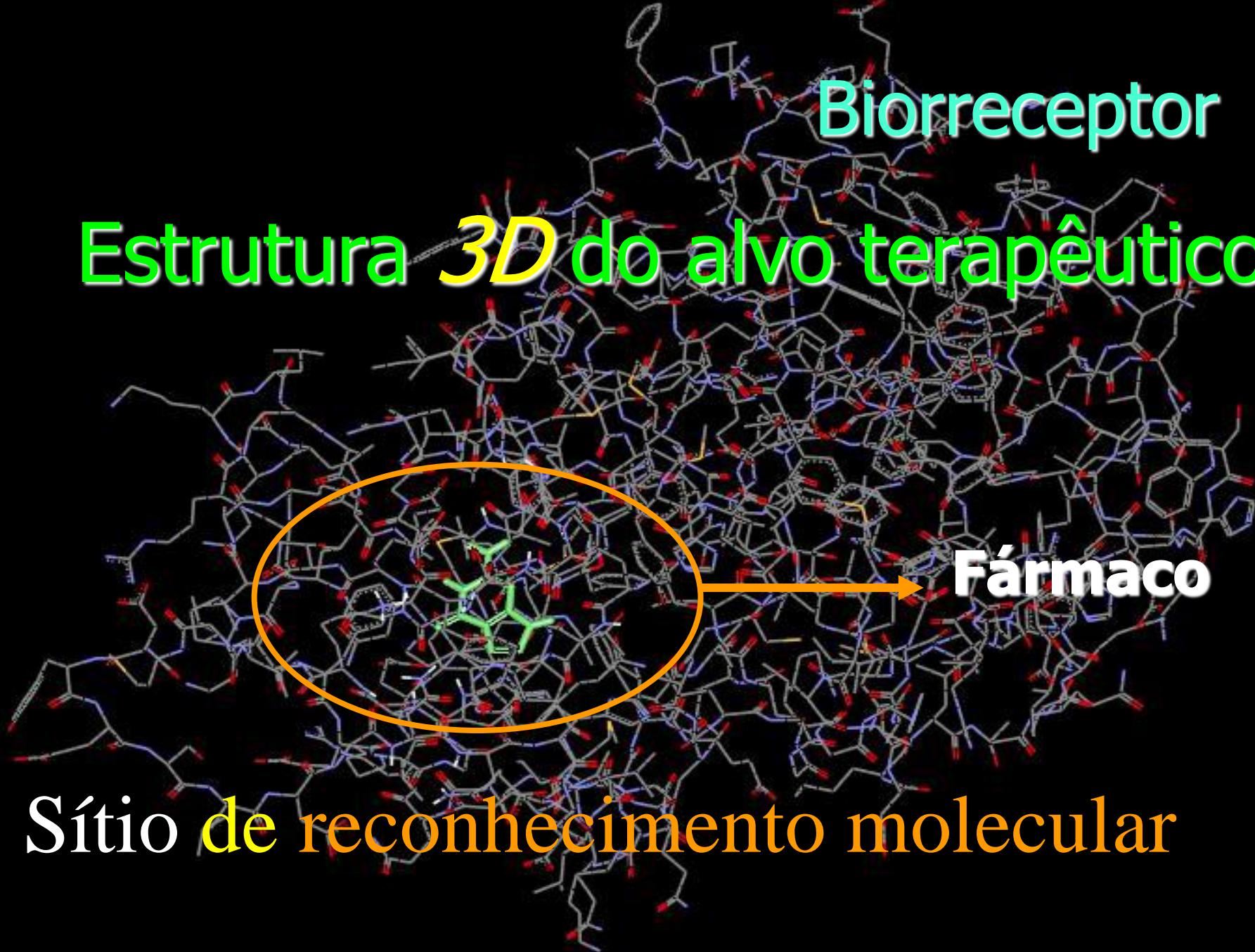


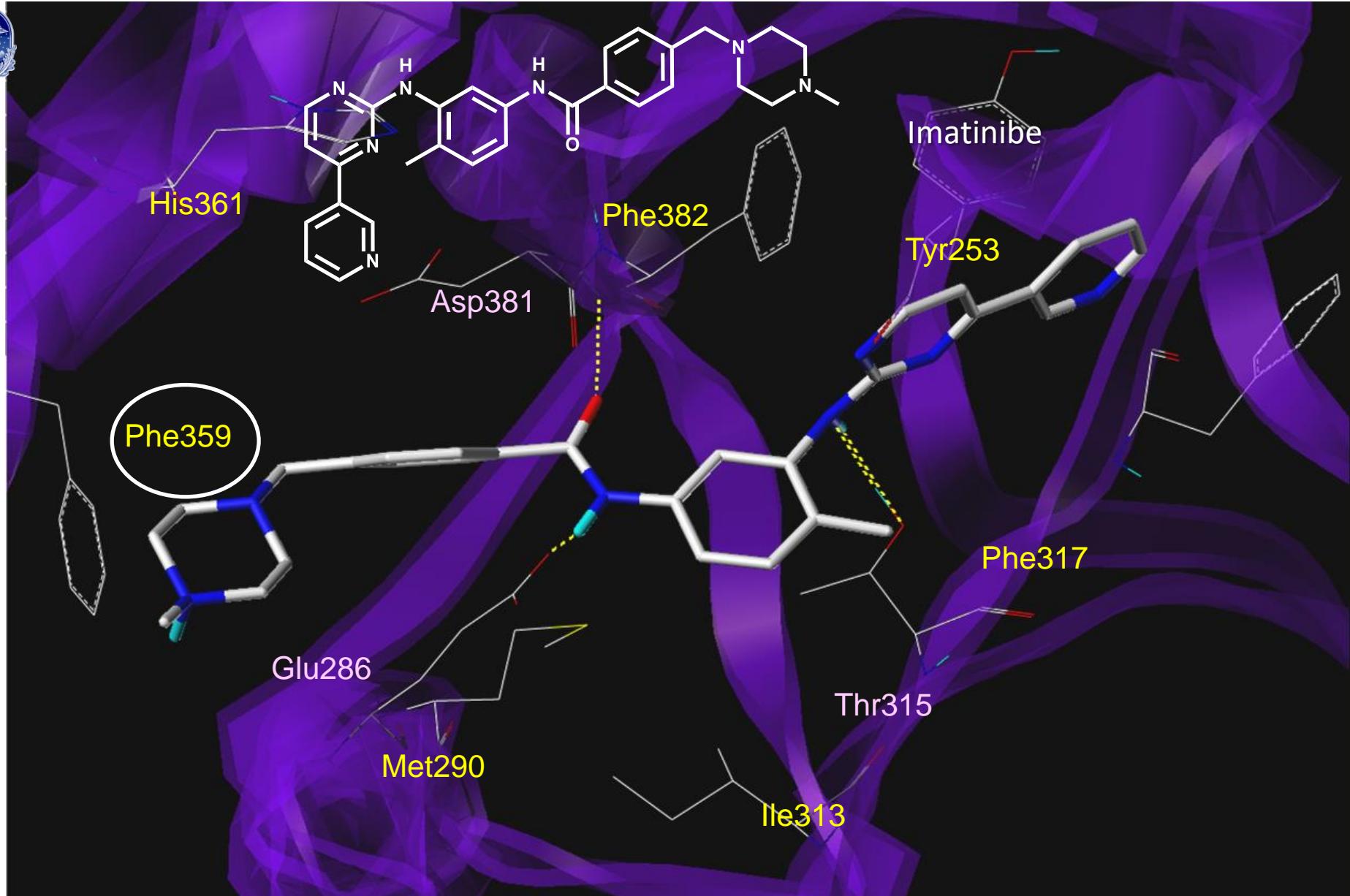
Biorreceptor

Estrutura 3D do alvo terapêutico

Fármaco

Sítio de reconhecimento molecular

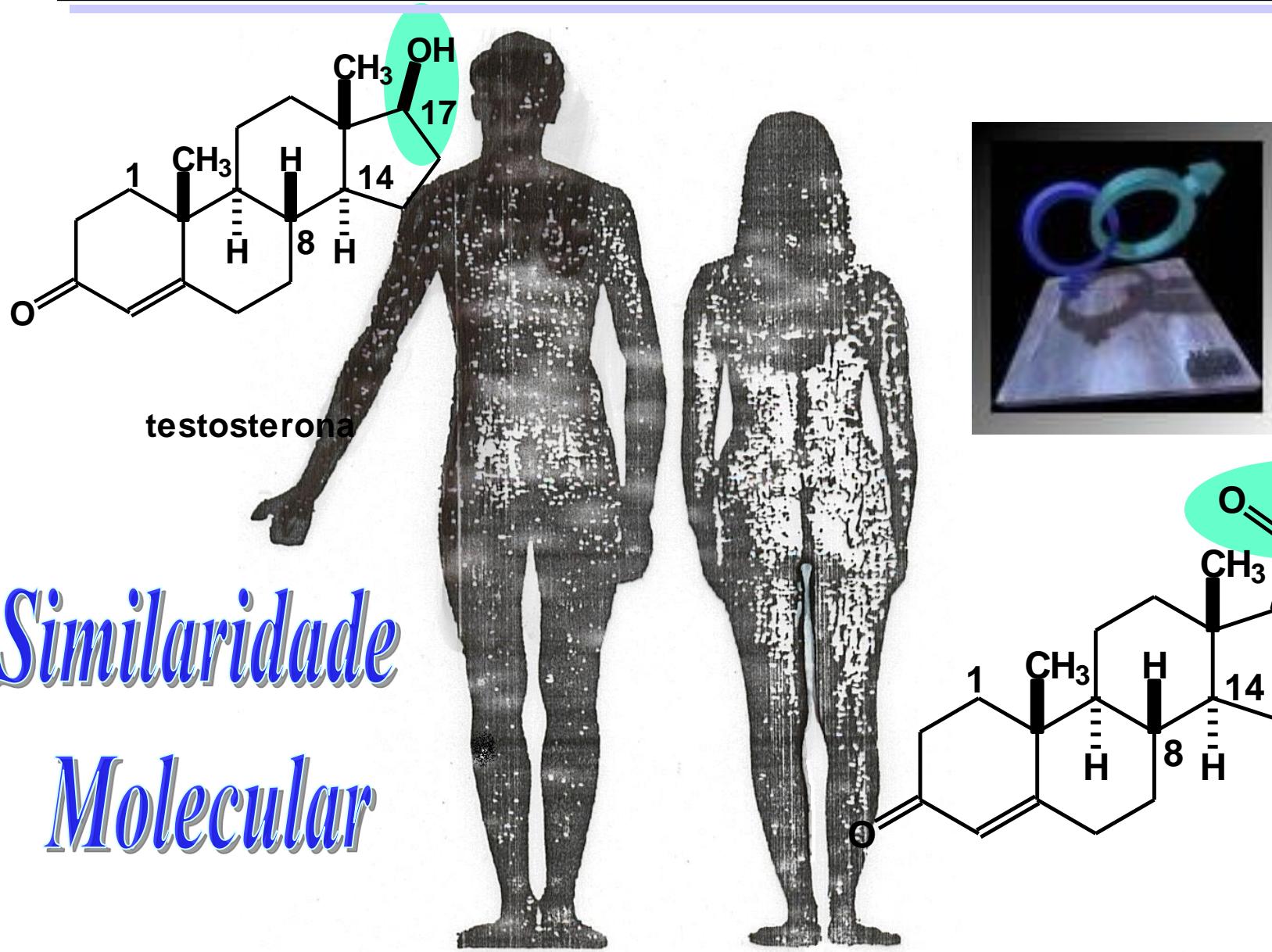




Estrutura do imatinibe complexado no domínio da *h*-ABL-K (código no PDB: 2HYY; Resolução= 2,4 Å) obtida por ancoramento molecular com o programa FlexX (licença # 7512)..
Valor de energia livre de ligação teórica (ΔG_{lig}) de -35,751 kJ/mol.



Reconhecimento Molecular-II

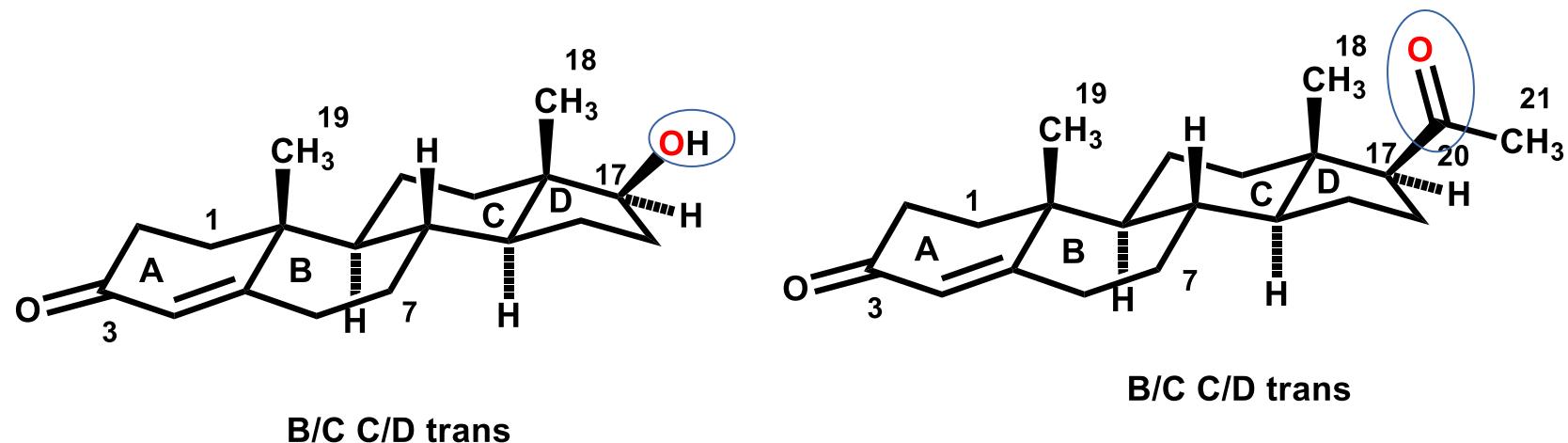
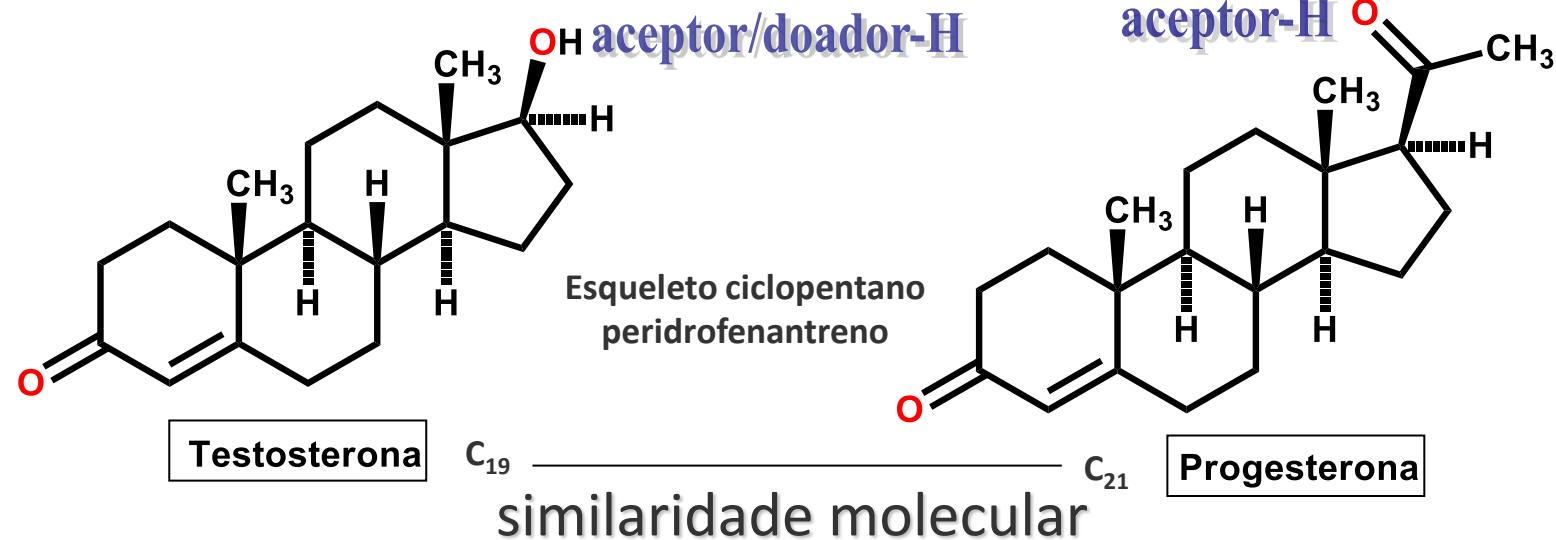


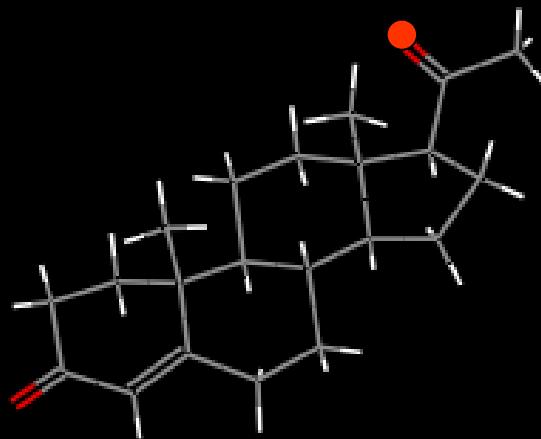
*Similaridade
Molecular*

progesterona

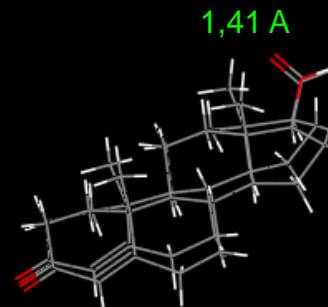


Reconhecimento Molecular

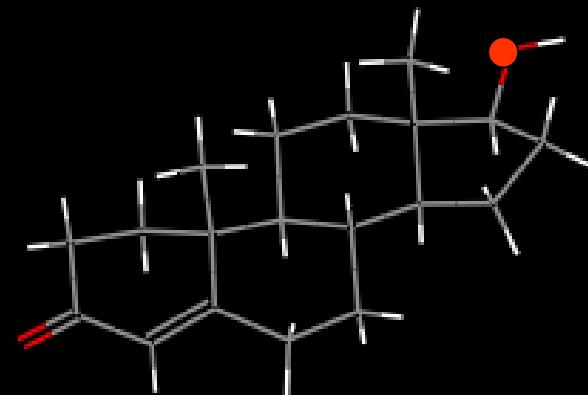




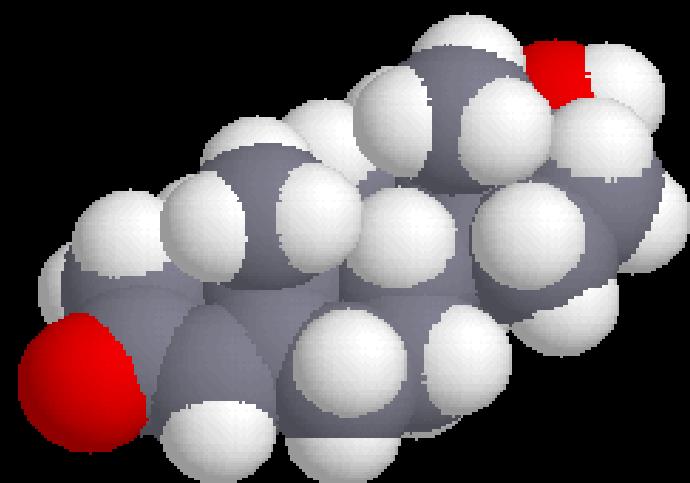
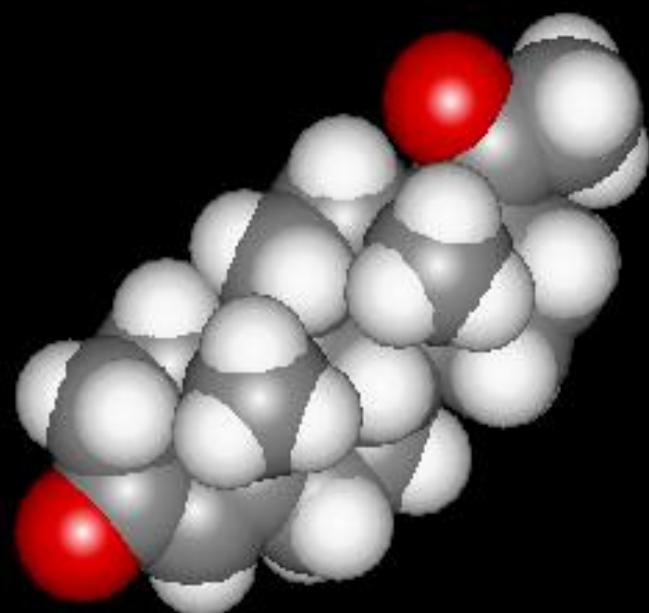
progesterona



1,41 Å



testosterona

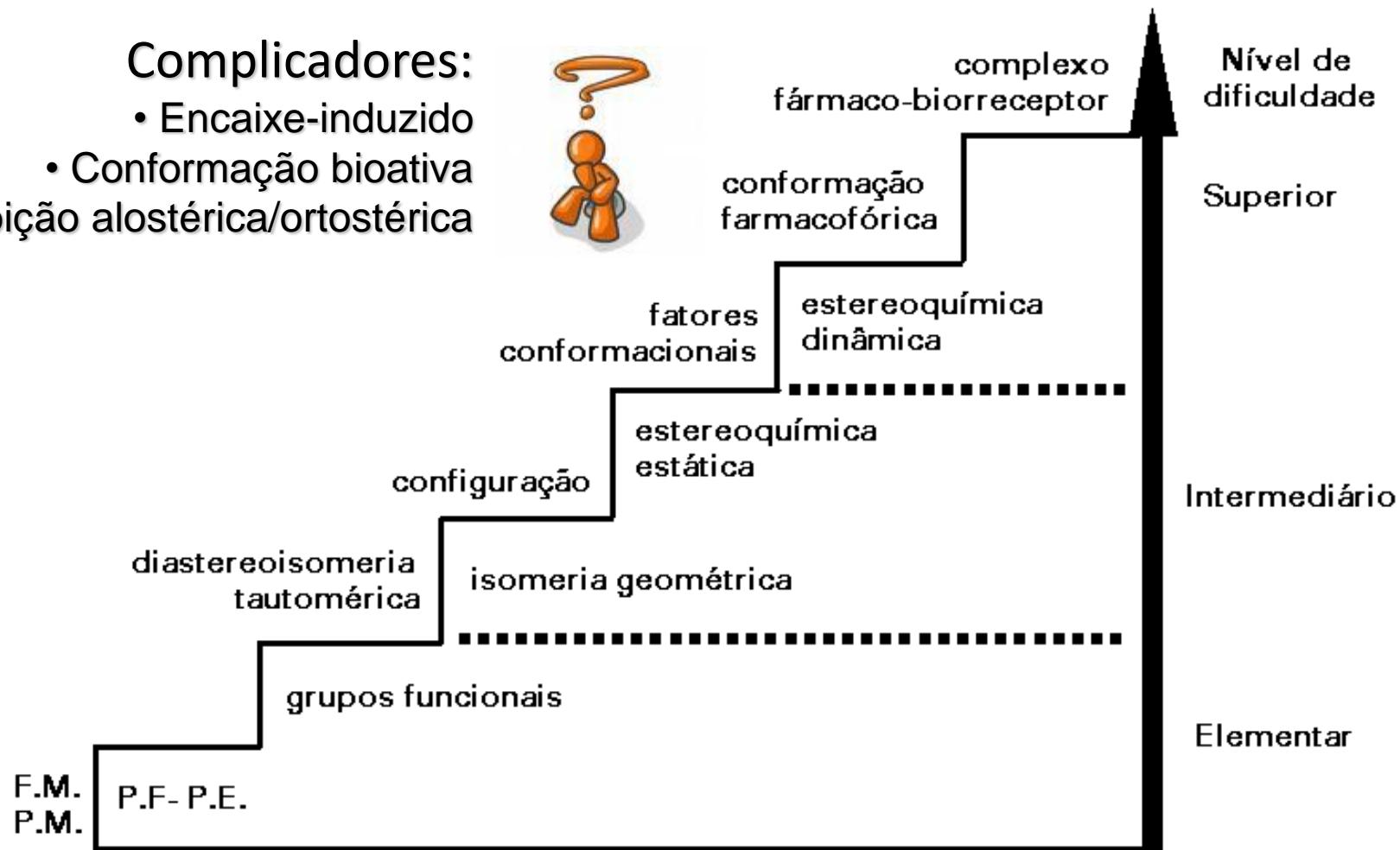




Nível hierárquico da descrição da complementaridade F-R

Complicadores:

- Encaixe-induzido
- Conformação bioativa
- Inibição alostérica/ortostérica

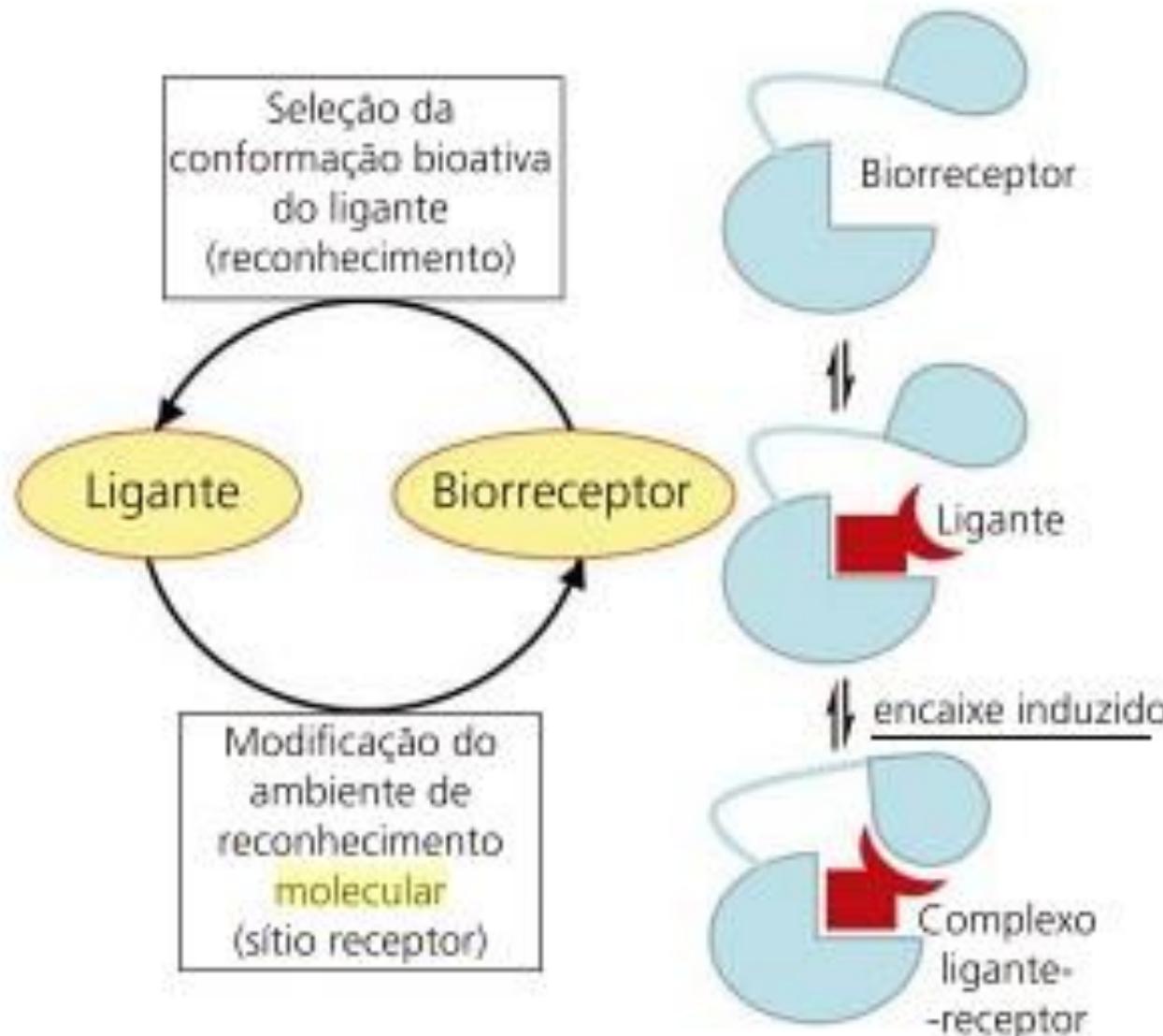


Propriedades Estruturais

bioligante- ligante
composto protótipo



O encaixe induzido F-R



Altera a energia livre do sistema