

A Química Medicinal

(Planejamento Racional de Novos Fármacos)

XXXVI Semana da Química “Ciência, tecnologia e sociedade: em busca do conhecimento”

unesp Instituto de Química, UNESP – Araraquara, 25-29 de setembro de 2006

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UFRJ

Laboratório de Avaliação e Síntese de Substâncias Bioativas



Universidade Federal do Rio de Janeiro

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6.1 Bioisosterismo: LASSBio-346, LASSBio-501

6.3 Hibridação molecular: LASSBio-756

6.4 Simplificação molecular: LASSBio-294

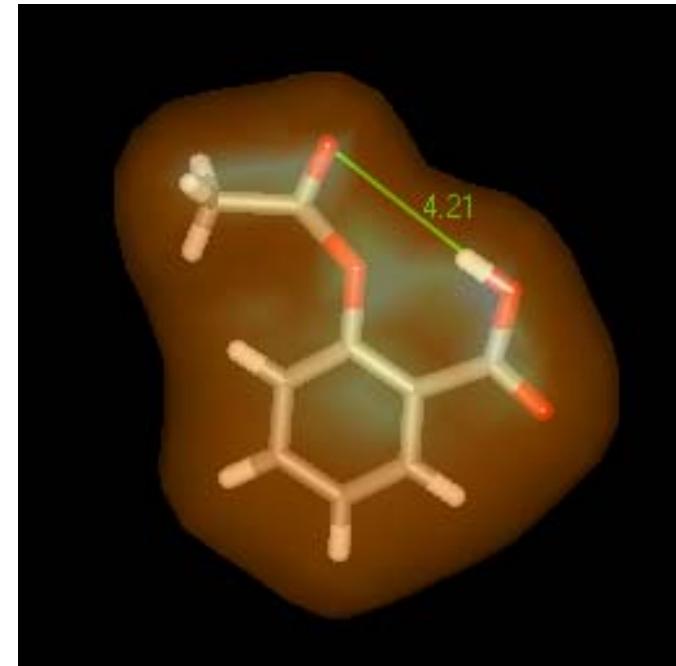
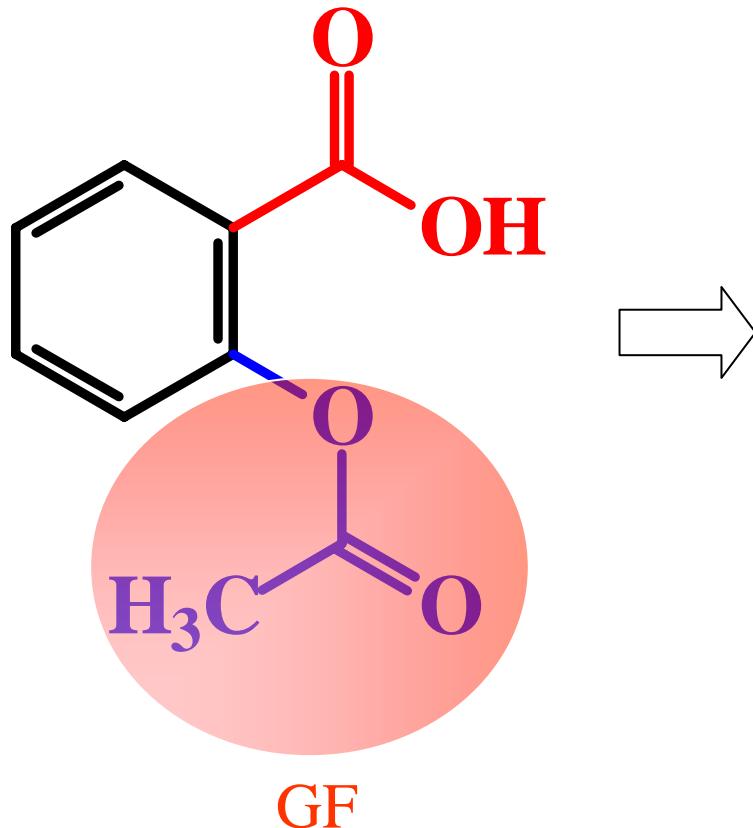
6.5 Desenho de protótipos simbióticos: LASSBio-468

7. Conclusões



Grupamento Farmacofórico do AAS

Grupamento auxofórico



Grupamento farmacofórico

arrangement of atoms (or groups of atoms) responsible for the pharmacological activity of a drug

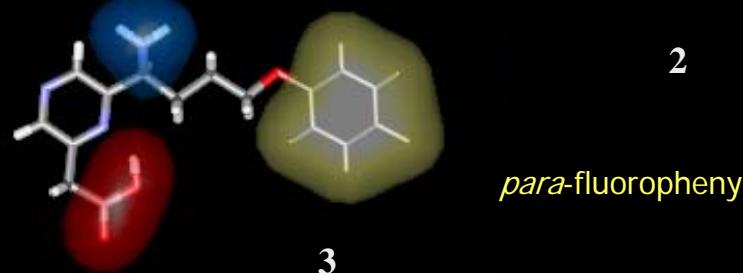
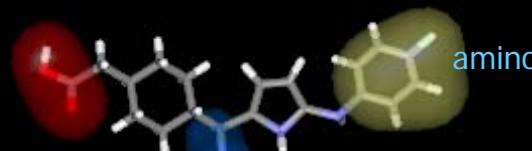
&

is a three-dimensional substructure of a molecule that carries (*phoros*) the essential features responsible for a drug's (*pharmacon*) biological activity

pharmacophore

fluoro-phenyl
amino group
carboxyl

➔ pharmacophore points

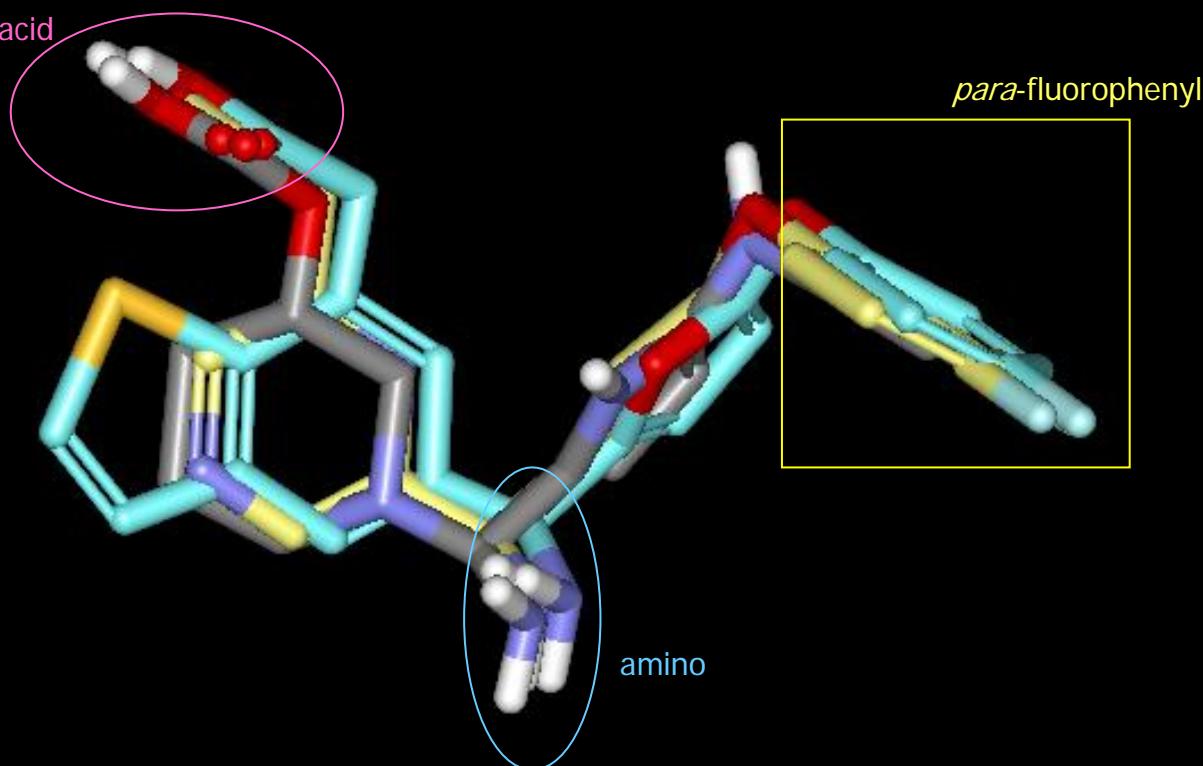


pharmacophore

fluoro-phenyl
amino group
carboxyl

pharmacophore points

carboxylic acid



overlap of the pharmacophoric groups → molecular similarity

Grupamento farmacofórico



230

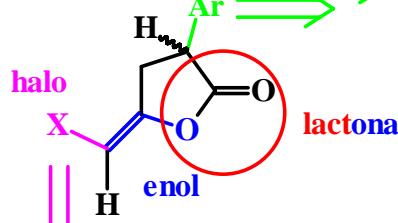
J. Med. Chem. 1986, 29, 230–238

Enol Lactone Inhibitors of Serine Proteases. The Effect of Regiochemistry on the Inactivation Behavior of Phenyl-Substituted (Halomethylene)tetra- and -dihydrofuranones and (Halomethylene)tetrahydropyranones toward α -Chymotrypsin: Stable Acyl Enzyme Intermediate

Michael J. Sofia and John A. Katzenellenbogen*



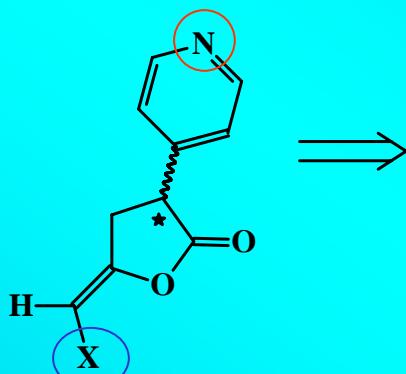
halo-enol-lactona



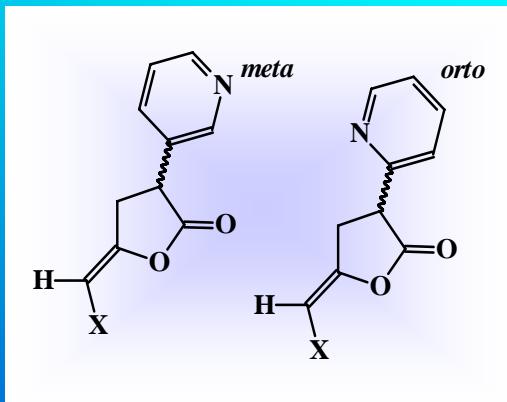
molecular diversity

Inibidor de Protease Suícida

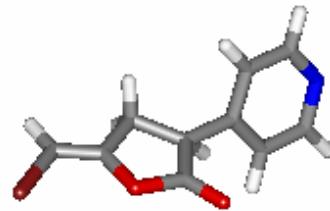
Halo-enol-lactona



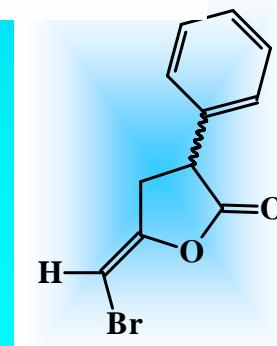
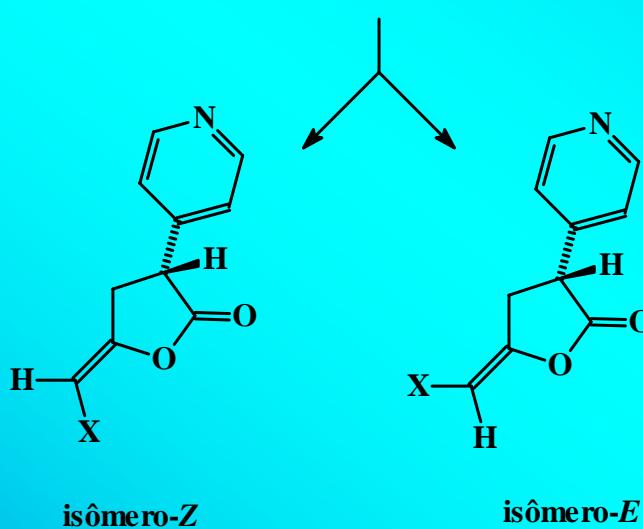
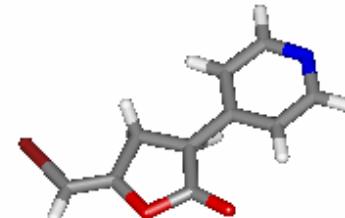
$X = Cl, Br$



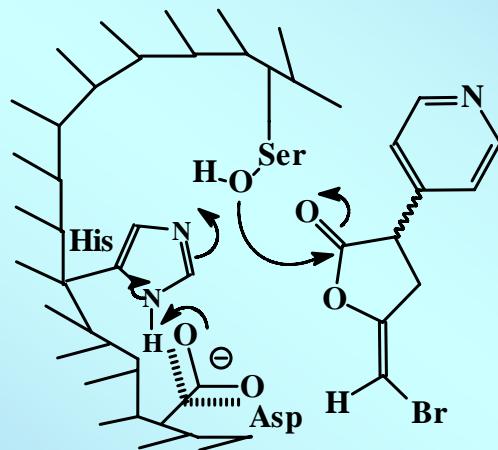
Isômero *cis*



Isômero *trans*



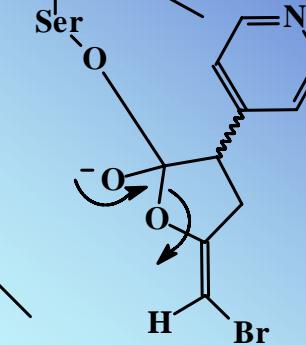
Nu = SerOH



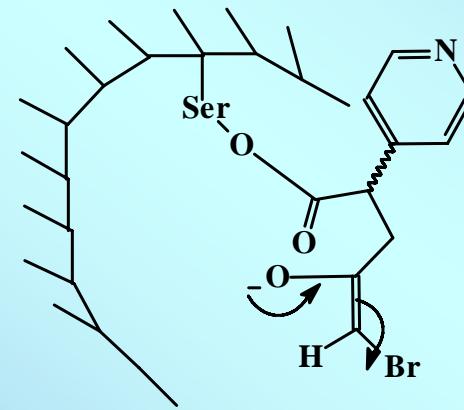
Enz-Nu = ataque
nucleofílico

"Pull-push effect"

Protease

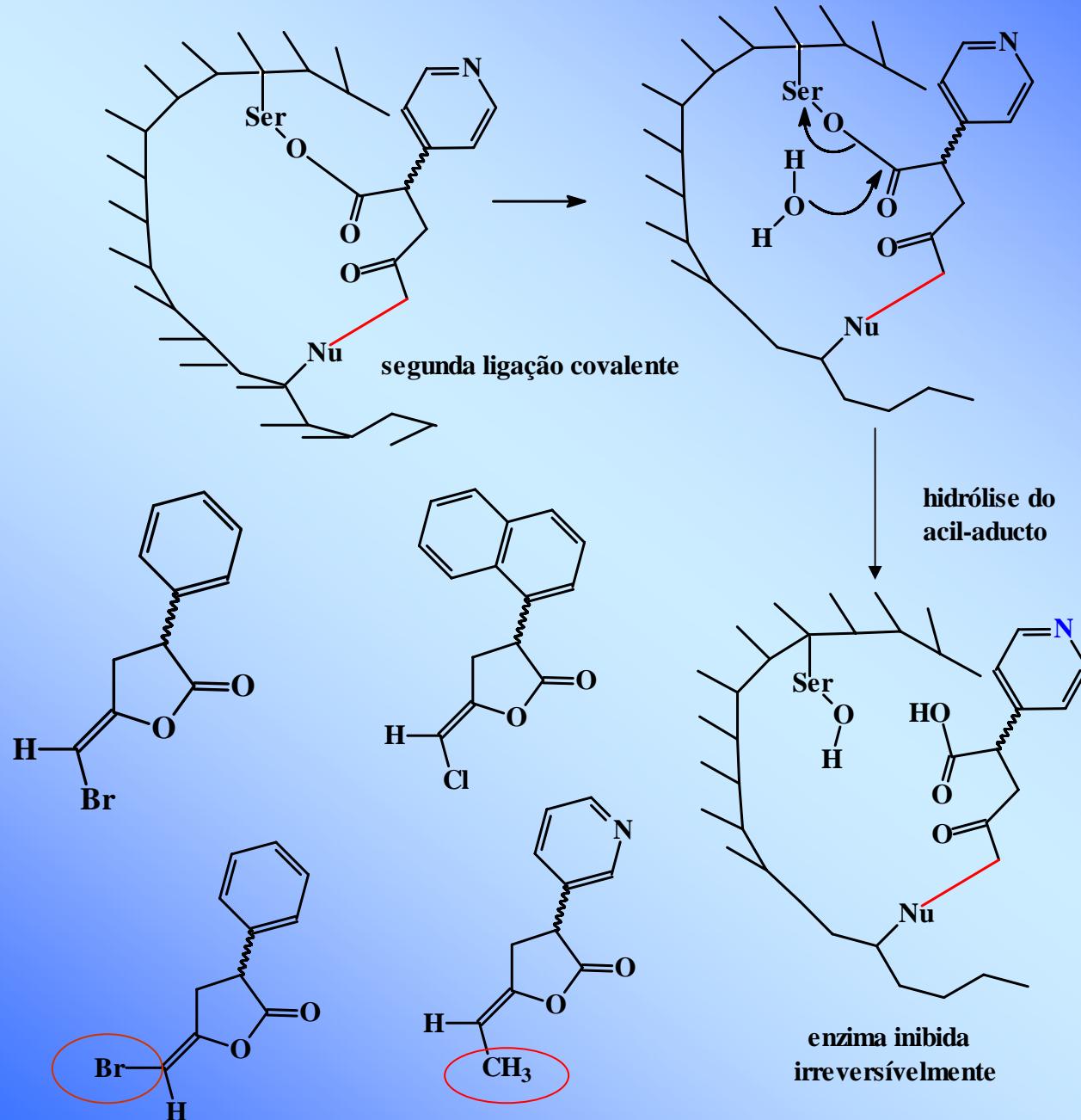


Estado de Transição

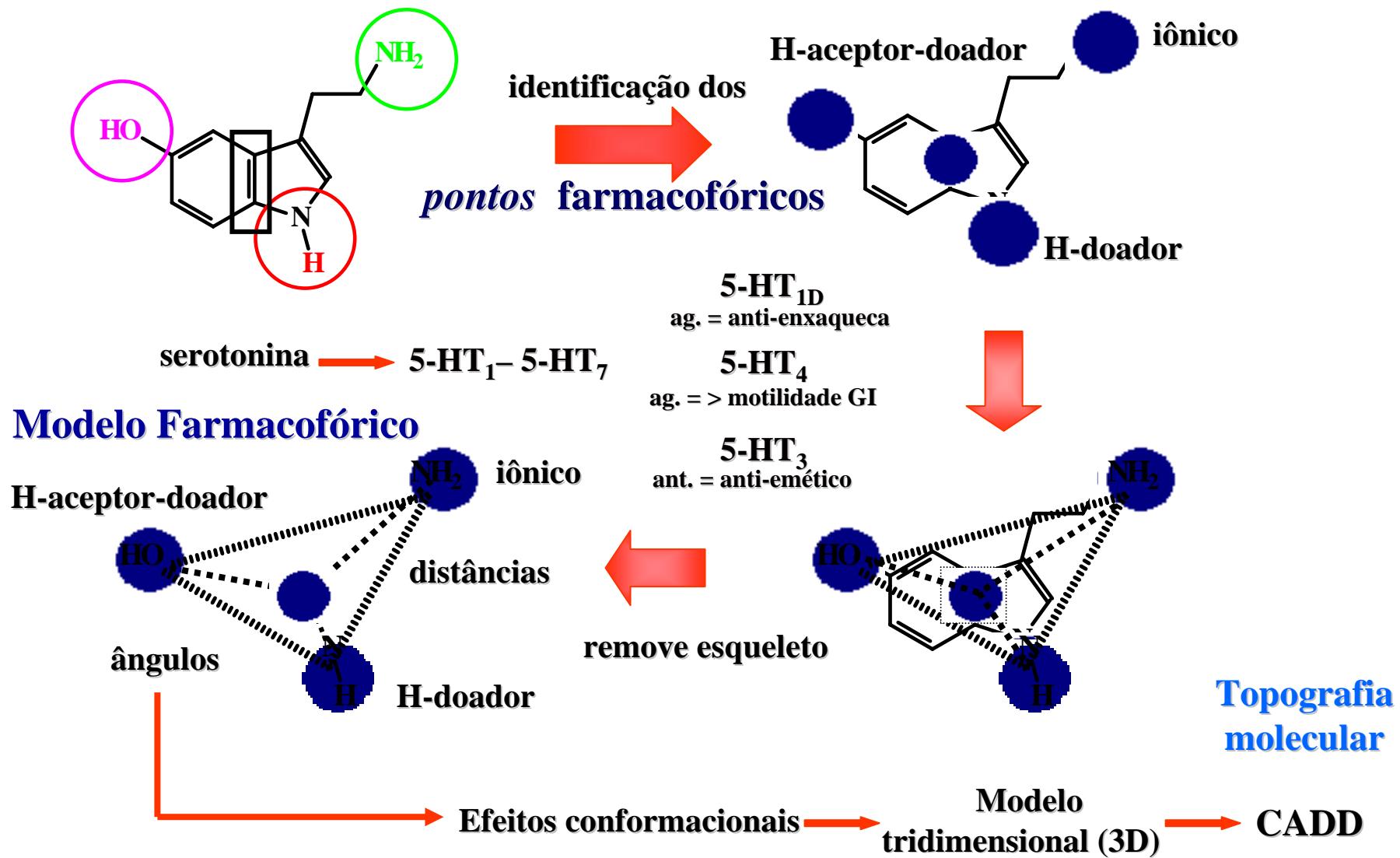


ADUTO
ACIL-ENZIMA

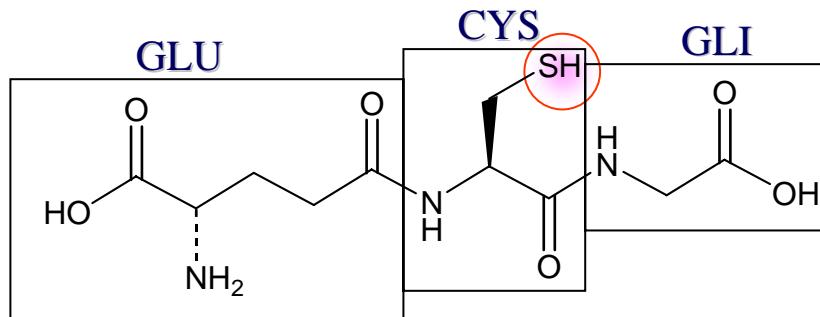
α -halocetona



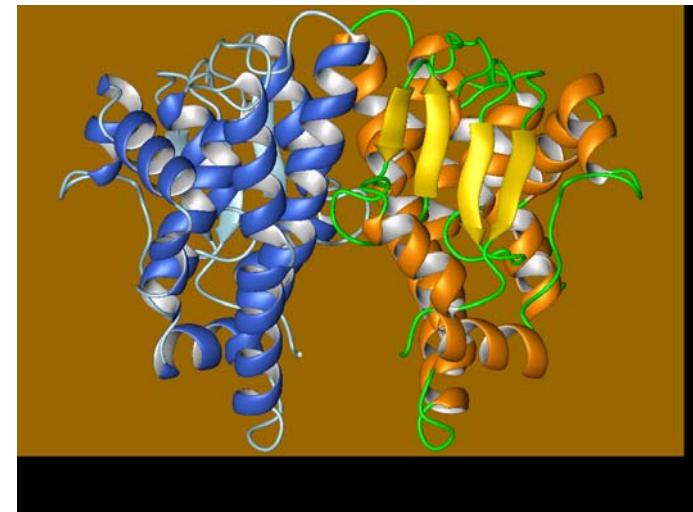
Construção de Modelo Farmacofórico 2D/3D



Grupamento toxicofórico



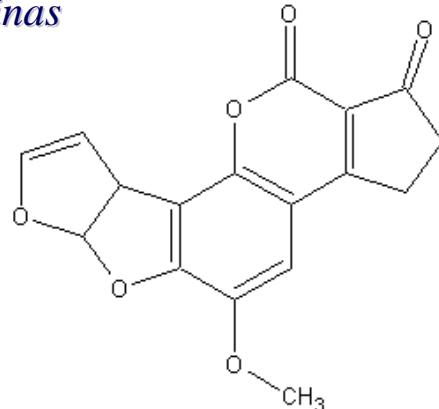
Glutatião = (Nu⁻) bionucleófilo



Toxicófilo/toxicofórico:

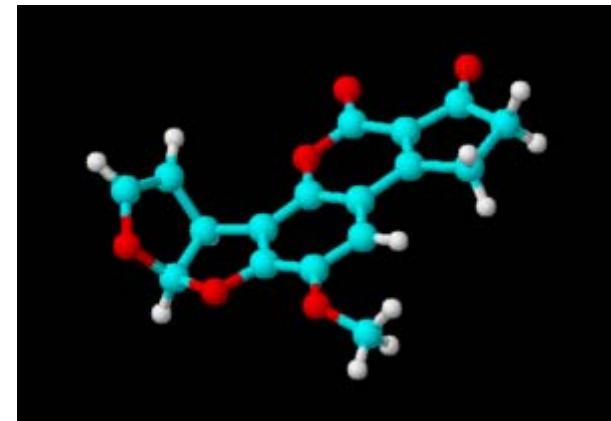
é o grupamento ou a sub-unidade estrutural de uma substância responsável pelas propriedades tóxicas.

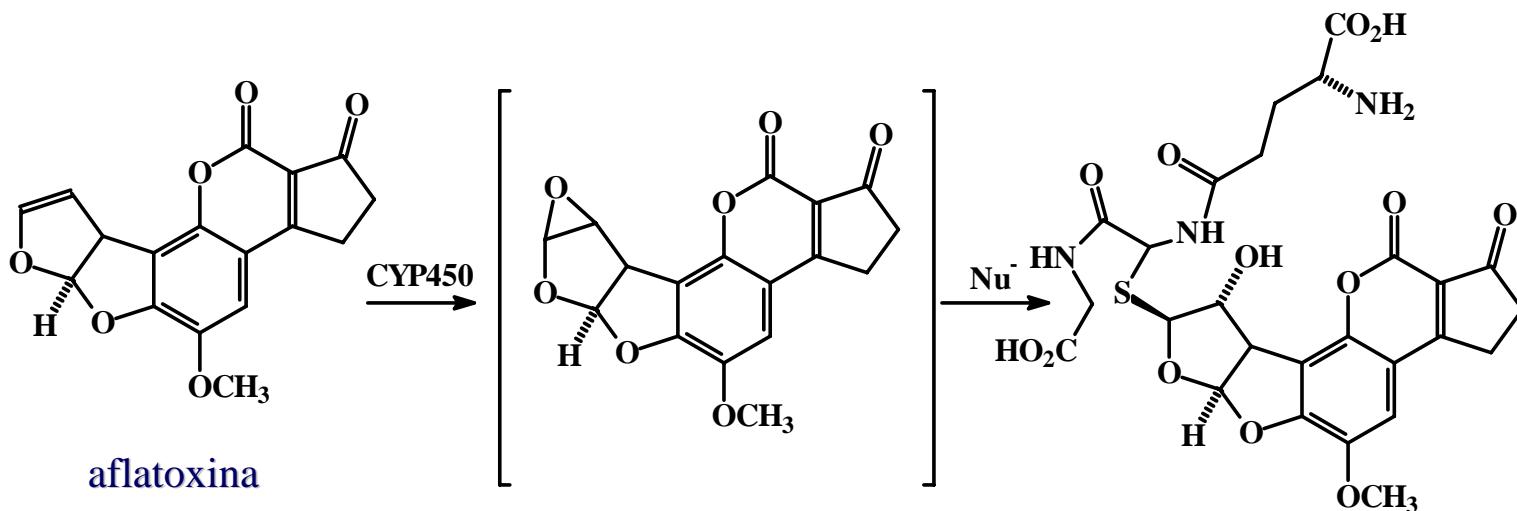
Micotoxinas



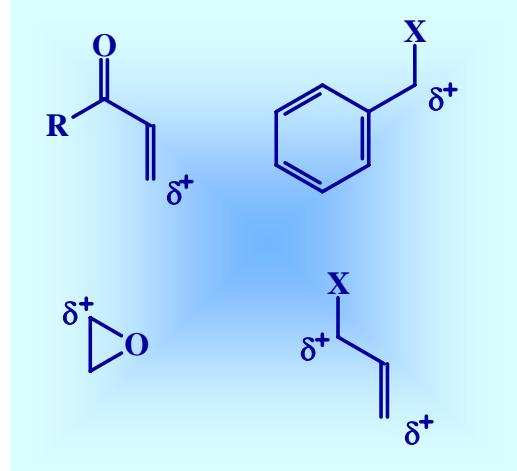
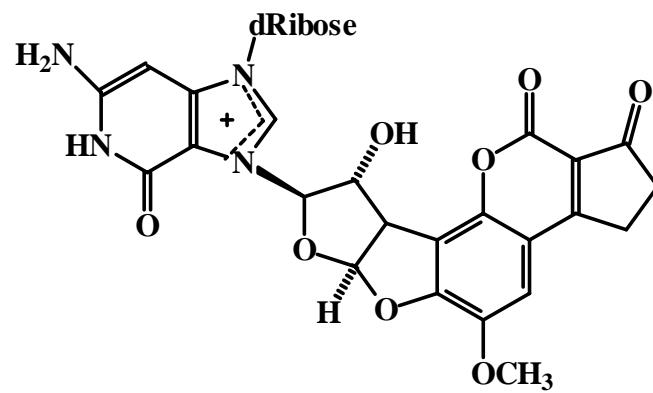
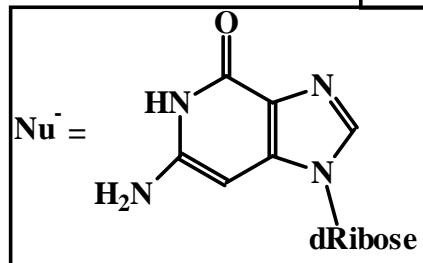
Aspergillus sp

Aflatoxina (B1/B2/G1/G2)





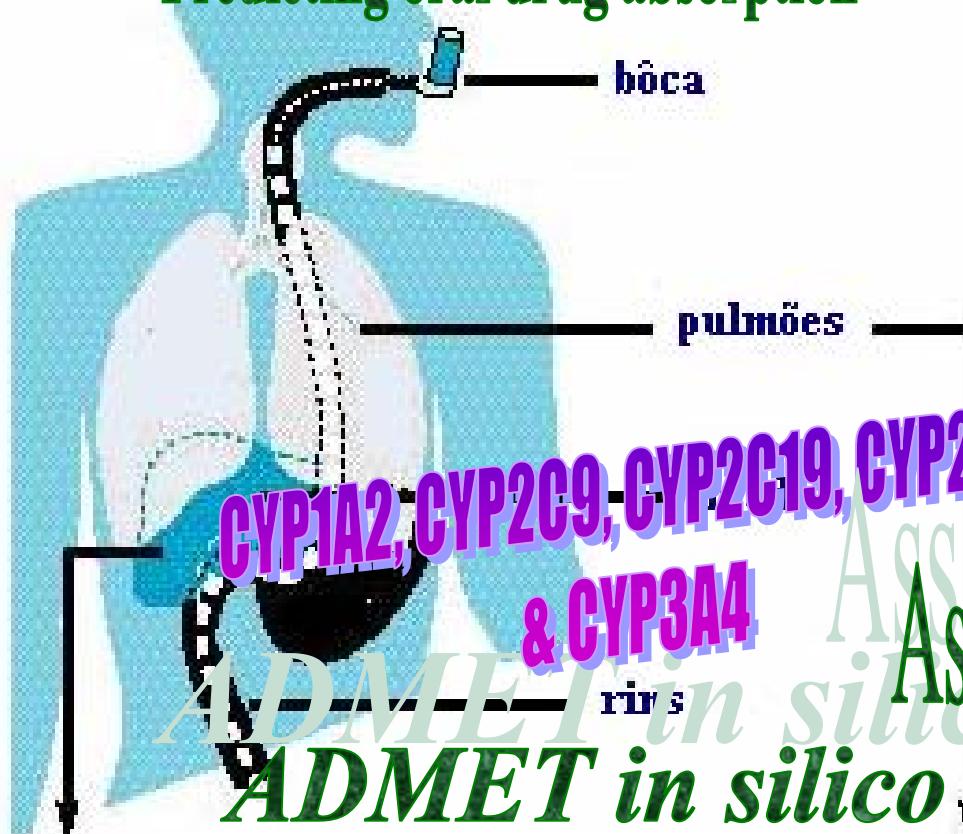
Nu^- = glutatião-transferase



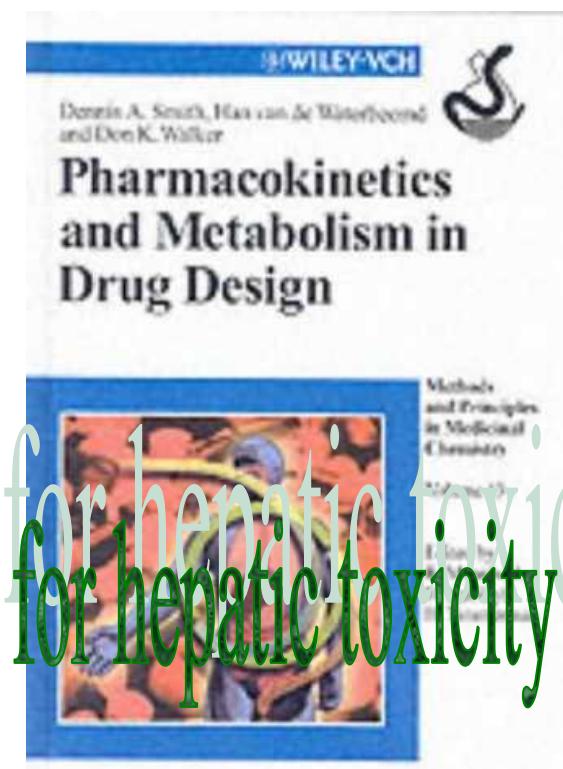
Fase Farmacocinética

Predicting oral drug absorption

Predicting oral drug absorption



ADME*



* absorção, distribuição, metabolismo & eliminação

Parte 2

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6.3 Hibridação molecular: LASSBio-756

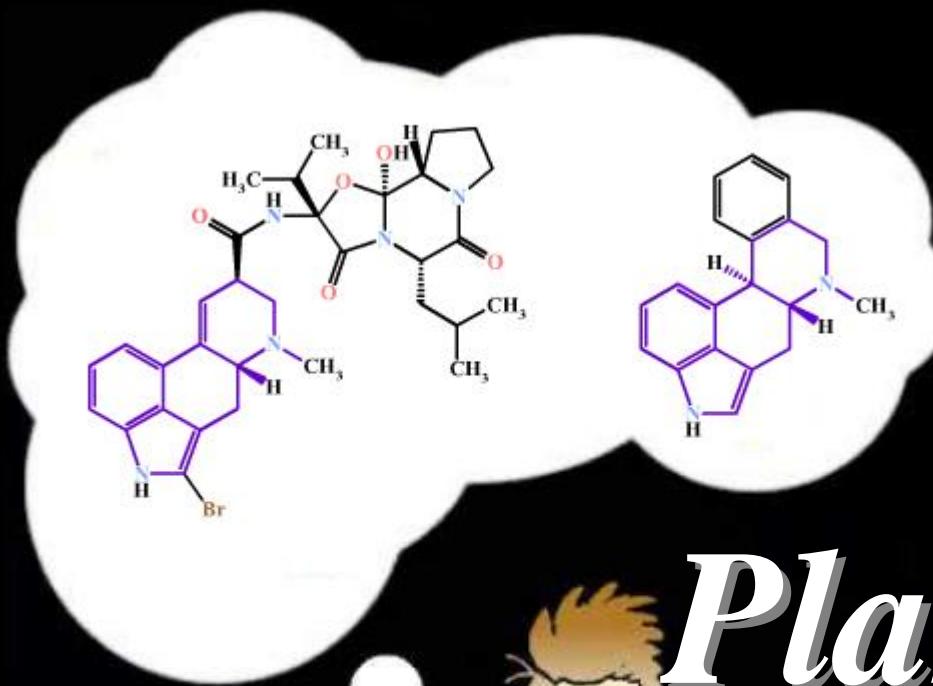
6.4 Simplificação molecular: LASSBio-294

6.5 Desenho de protótipos simbióticos: LASSBio-468

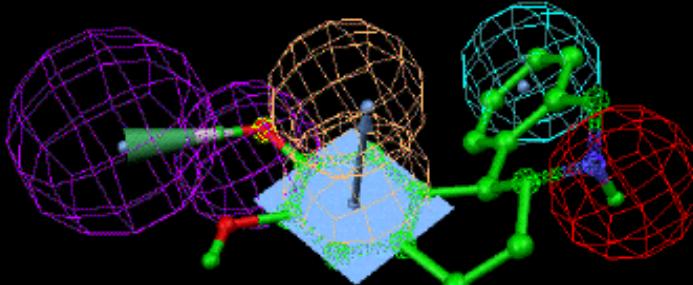
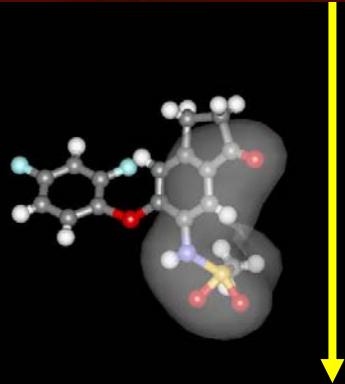
7. Conclusões



Fármacos Inteligentes

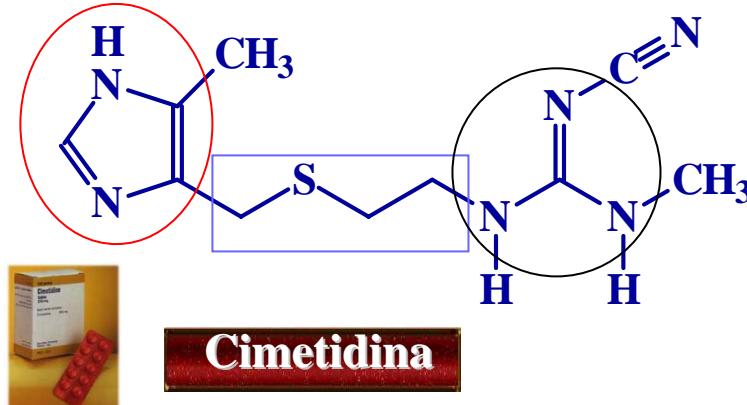


*Planejamento
racional*

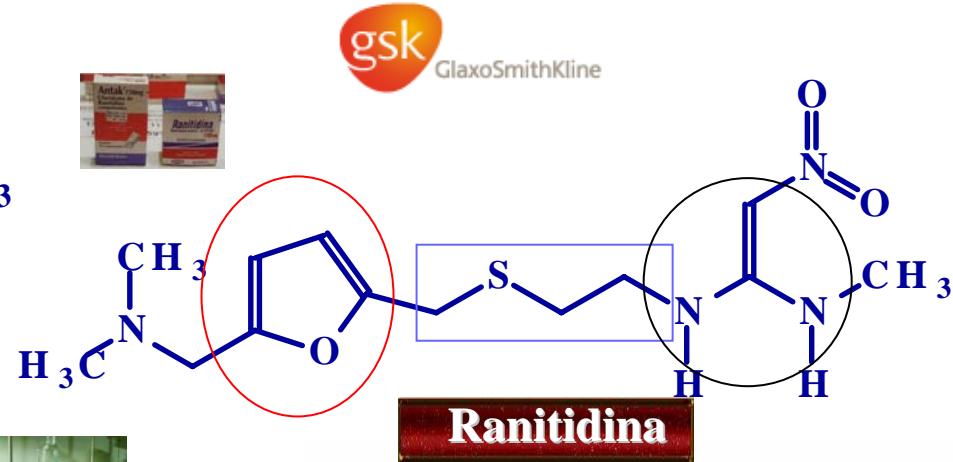


m e d s h e m Química Medicinal

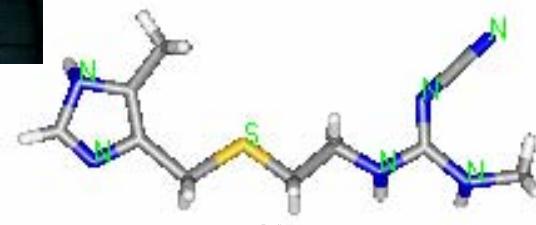
Atualmente, os fármacos, capazes de atuarem em **qualquer alvo-terapêutico**, são *descobertos* por planejamento racional.



Robin Ganellin *et al.*, 1974
 US 3950333 1974, 1976 - SK&F
Brit. J. Pharmacol. **53**, 435 (1975).

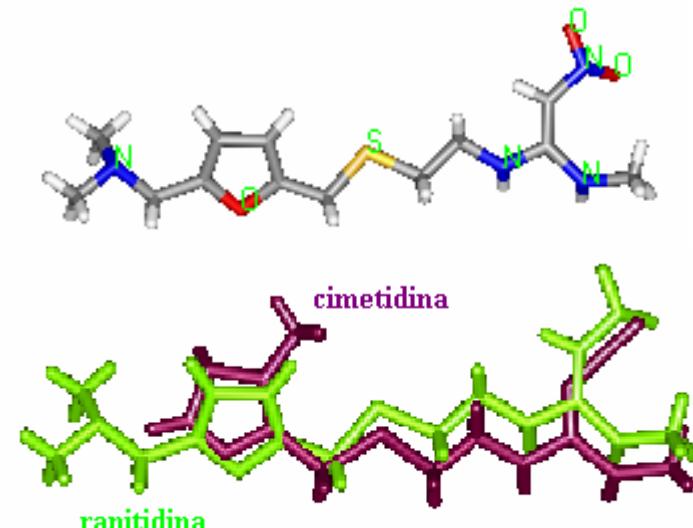


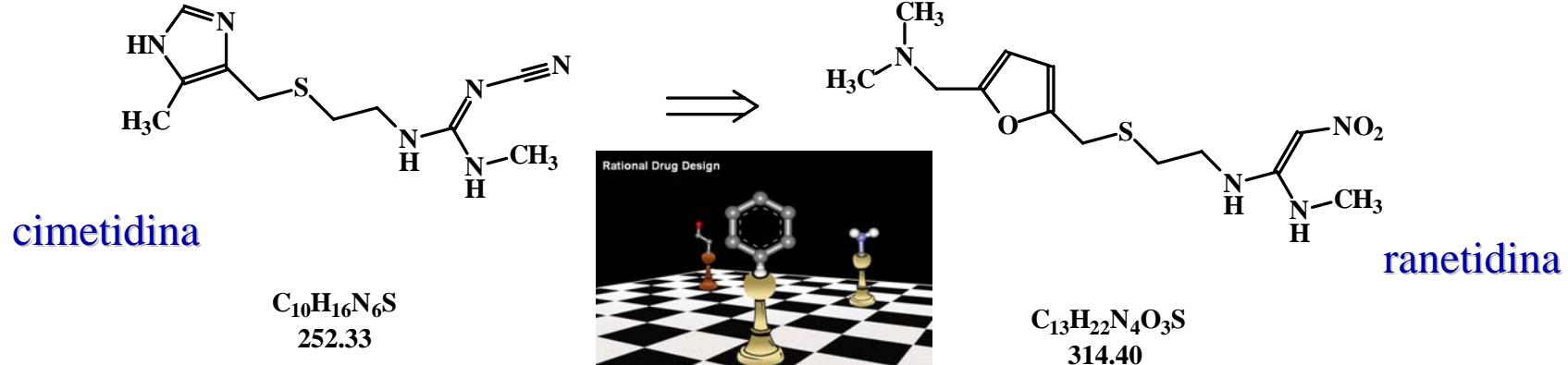
Barry J. Price *et al.*, 1978
 US 4128658 1978 - Allen & Hanburys
Brit. J. Pharmacol. **66**, 464 (1979)



**Inovação
terapêutica**

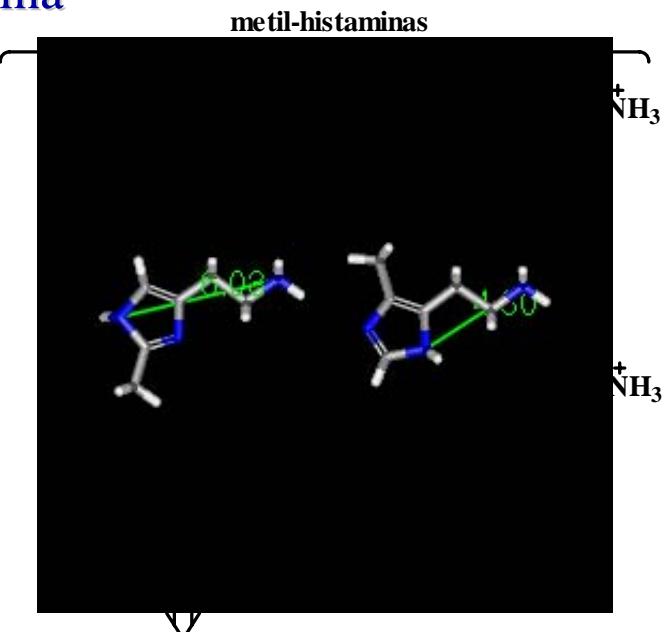
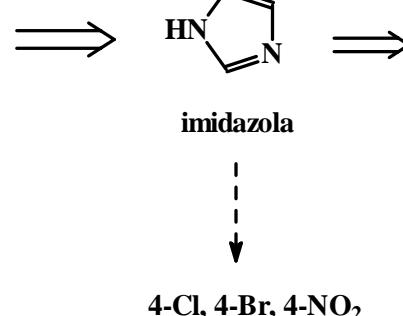
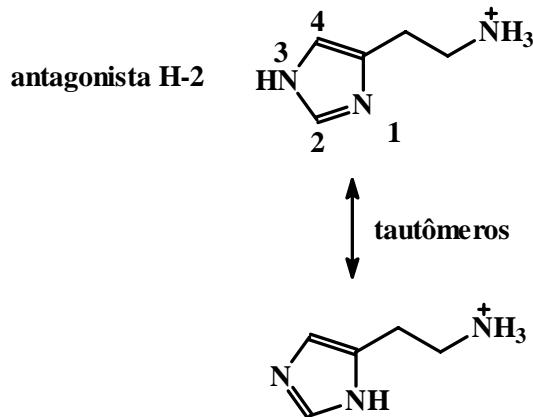
*similaridade
molecular
me-too*



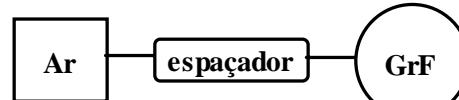


A gênese da cimetidina

A procura do protótipo:

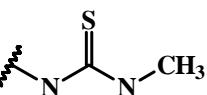
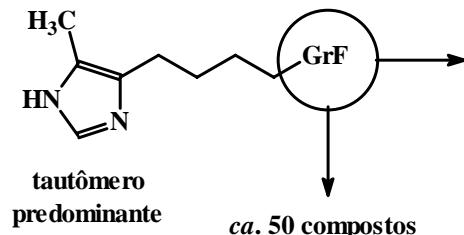


metil-alquil-imidazola



Ar = arila, heteroarila
GrF = grupamento funcional

N-metil-tiouréia



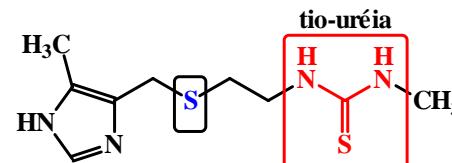
bioisosterismo

ant/agonista
& CYP450

clássico

$\text{CH}_2 \longrightarrow \text{S}$

Ensaiada em 700 pacientes com
úlcera duodenal



metiamida

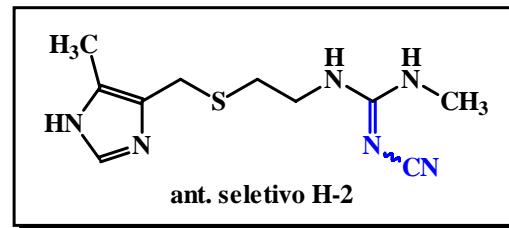
ant. seletividade H-2
& <<CYP450, p.o.

ensaios clínicos
granulocitopenia

bioisosterismo
clássico

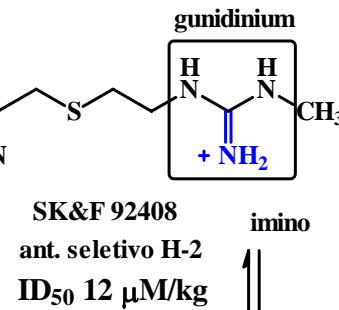


Cimetidina
 $ID_{50} 1,4 \mu\text{M}/\text{kg}$

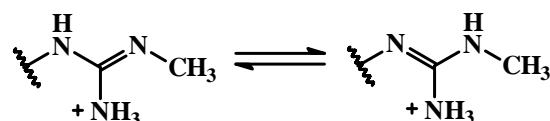


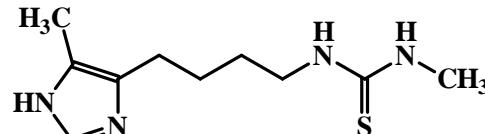
A gênese da cimetidina

bioisosterismo



amino

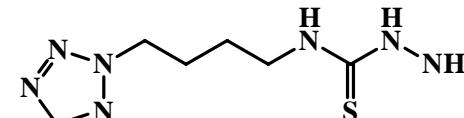




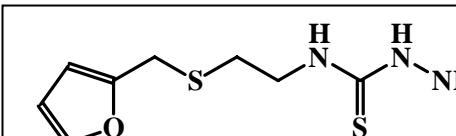
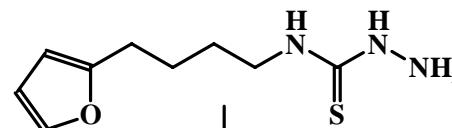
burinamida

Allen & Hanburys Ltd.

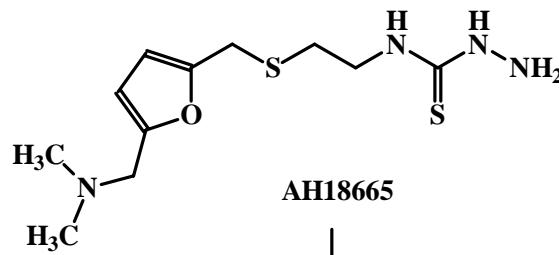
análogos da burimamida



AH15475

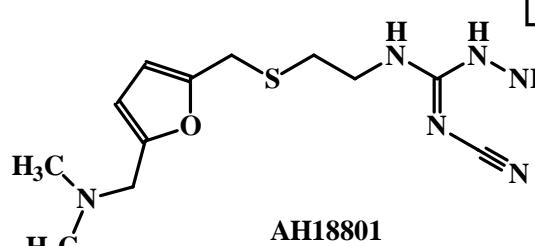


AH18166



AH18665

ciano-guanidila



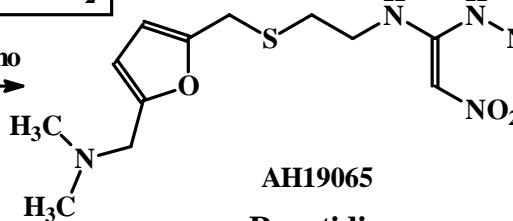
AH18801

não cristalino

mesma potência da cimetidina

=NCN → =CHNO₂

bioisosterismo



AH19065

Ranetidina

nitro-vinila



cristalino



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A new era of logical drug design

The research program leading to cimetidine also represented a revolution in the way pharmaceuticals are developed. Traditionally, the development of a new drug would often depend on the fortuitous discovery of a plant or microbial extract that showed some of the required biological activity. Using that first extract as a lead, many similar compounds would be made and tested for pharmacological effectiveness. In many cases, the researchers did not know how the drug worked, so finding an optimal compound was difficult.

The development of cimetidine was radically different: it was one of the first drugs to be designed logically from first principles. SK&F's multidisciplinary research team first looked at the physiological cause of acid secretion. They confirmed that a molecule found in the body called histamine triggers the release of acid when it binds to a specific receptor (now called the H₂-receptor) in the stomach lining. Their aim was to find a molecule that successfully competed with histamine in combining with the receptor, but then blocked, rather than stimulated, acid release. Such a molecule was called a histamine H₂-receptor antagonist and represented a new class of drugs.

Using a step by step analysis of structural and physical properties, the team made a series of histamine-based molecules, which were then tested for antagonist activity using carefully designed pharmacological assays. Today, this approach of rational drug design underpins the discovery programs of many major pharmaceutical companies.





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& Nominate

Landmark designation

The American Chemical Society and The Royal Society of Chemistry designated the discovery of histamine H₂-receptor antagonists, which led to the development of the anti-ulcer drug Tagamet®, as an International Historic Chemical Landmark in ceremonies at SmithKline Beecham's research facilities (now GlaxoSmithKline) at Harlow, United Kingdom, on November 24, 1997, and King of Prussia, Pennsylvania, on February 27, 1998. The text of the plaques at the two sites reads:

Pioneering work by scientists in the laboratories of this company led to the first clinically effective inhibitor of gastric acid secretion. The worldwide introduction of cimetidine (Tagamet) revolutionized the treatment of peptic ulcers by dramatically reducing the need for surgical intervention. The work is recognized as the classic example of the systematic modification of a natural messenger substance (histamine) to create a therapeutically useful blocking agent. Effective commercialization of this discovery was greatly facilitated by the subsequent investigation and design of novel synthetic routes, which led to the development of an efficient chemical manufacturing process.

The Royal Society of Chemistry's Historic Chemical Landmarks can be found at
<http://www.rsc.org/lap/publicat/landmarks.htm>.

Designed by [MSK Partners](#), Hunt Valley, Maryland.



Fármacos Inteligentes

Super-Super Drug

otimização de protótipo natural



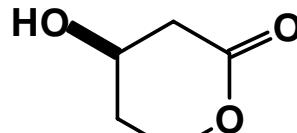
Jan., 1987: Lipitor®
US\$ 1 billion

2002: US\$ > 7.0 billions

* A. M. Thayer, CE&N, Nov. 12, 2002

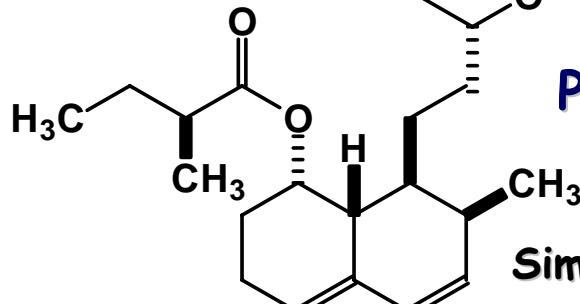
atorvastatina





Super-fármaco

* A. M. Thayer, CE&N, Nov. 12, 2002

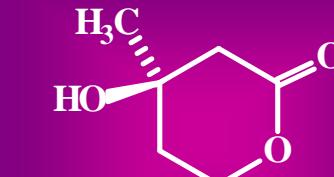


Protótipo natural



1975 - Compactina

A. Endo, J. Antibiot. 1979, 32, 652
Monascus ruber



Mevalo-lactone

HMG-CoA reductase

ácido pirrol-heptanóico

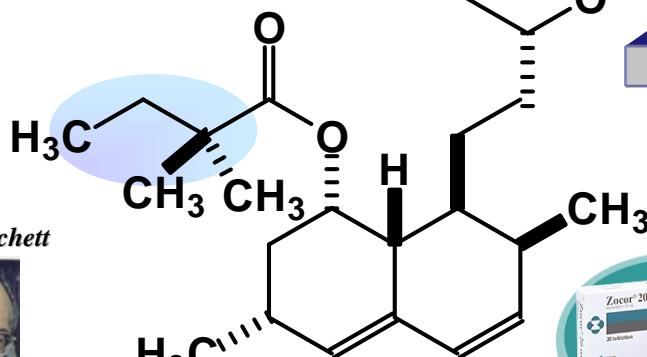


Pfizer

Bruce Roth



γ -lactona



Arthur Patchett



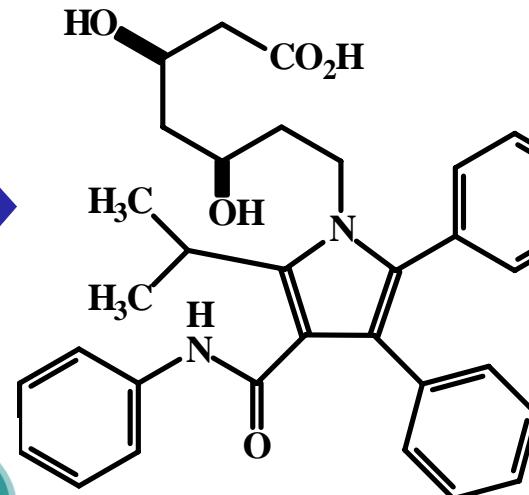
J. Med. Chem. 1986, 29, 849

J. Med. Chem. 2002, 45, 5609

Simvastatin
(Zocor)
MK-733
1981

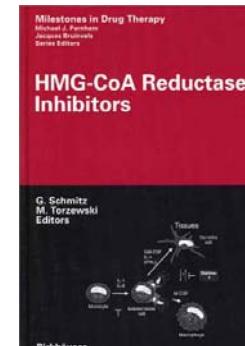


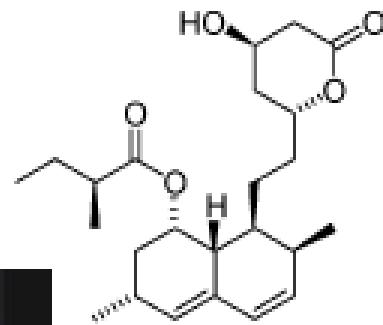
MERCK



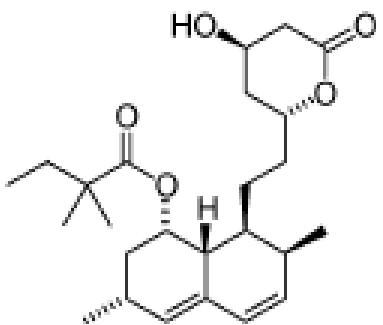
atorvastatina

$C_{33}H_{35}FN_2O_5$

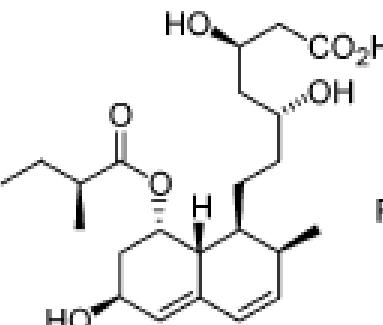


**Me-too**

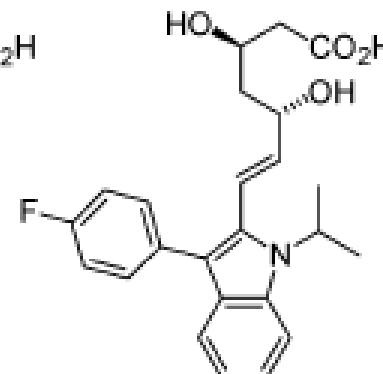
Lovastatin (15)
(Mevacor®)



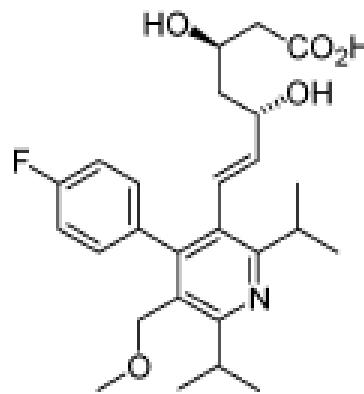
Simvastatin
(Zocor®)



Pravastatin
(Pravachol®)

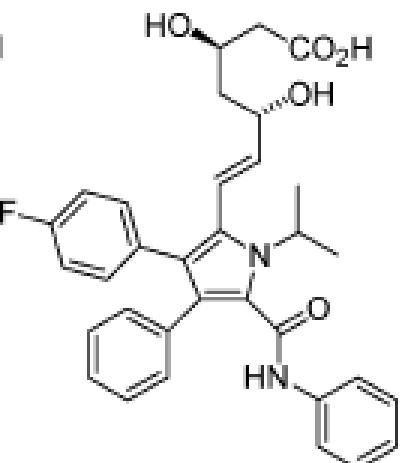


Fluvastatin
(Lescol®)

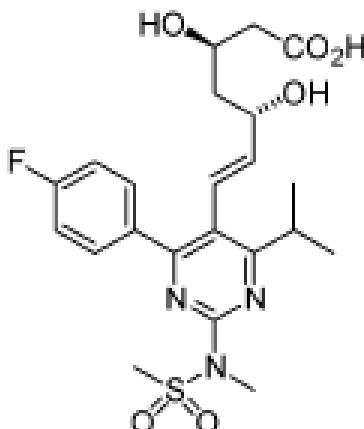


Cerivastatin
(Baycol®)

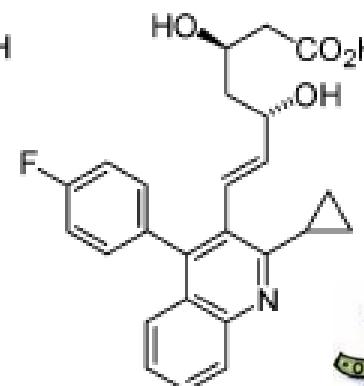
Note: removed from market



Atorvastatin
(Lipitor®)



Rosuvastatin (36)
(Crestor®)



Pitavastatin (37)
(Livalo®)



Características estruturais comuns aos cinco fármacos mais vendidos no mundo em 2004:

- Possuem apenas 7 elementos químicos: C,H,O,N,S,F,Cl;
- Todos possuem heteroátomos;
- Todos são multicíclicos (< cinco anéis);
- 80% têm unidades aromáticas e são heterocíclicos;
- Têm apenas 10 centros estereogênicos;

Totalizam 111 C's

1 C = US\$ 280 mi;

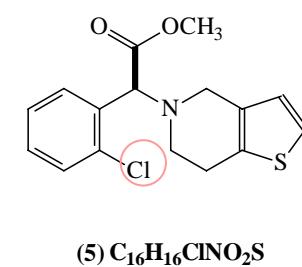
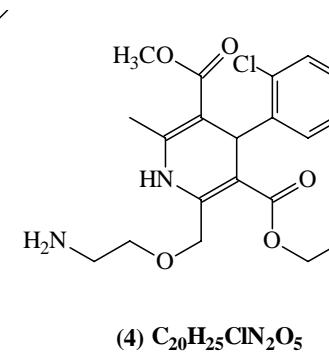
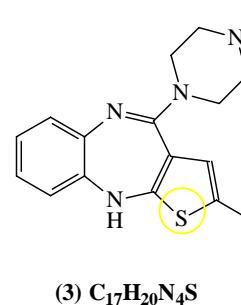
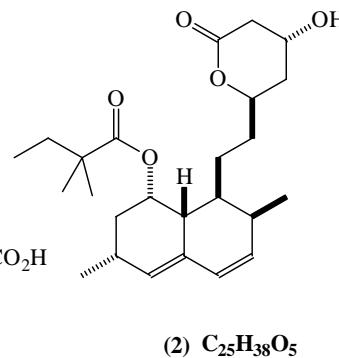
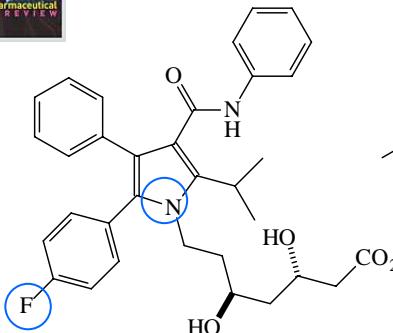
30,9&

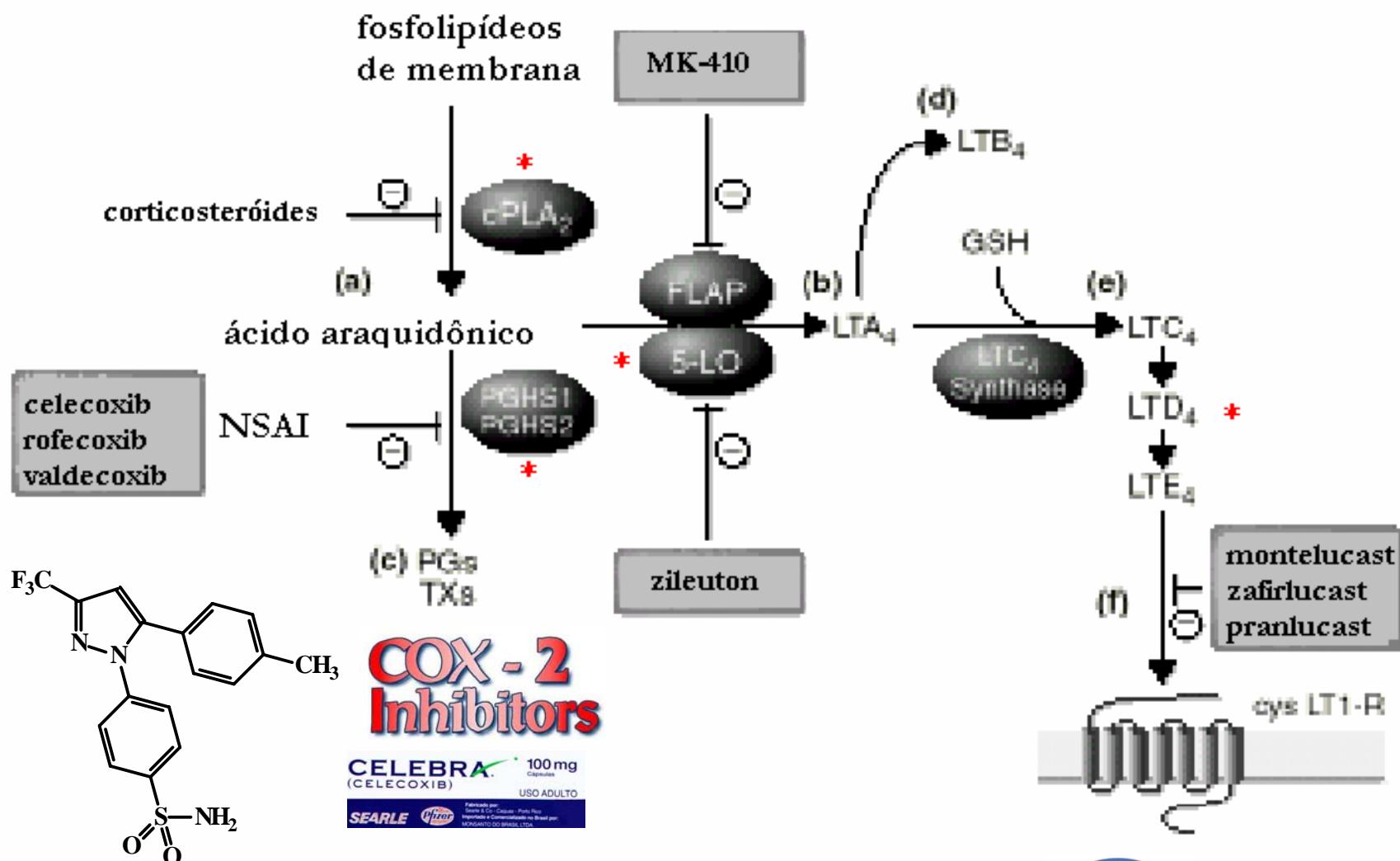
Fonte:



C&EN December 5, 2004
 Volume 82, Number 49
 S. Class pp. 18-29

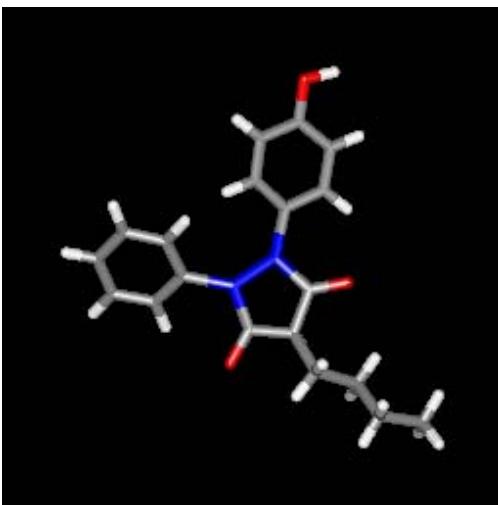
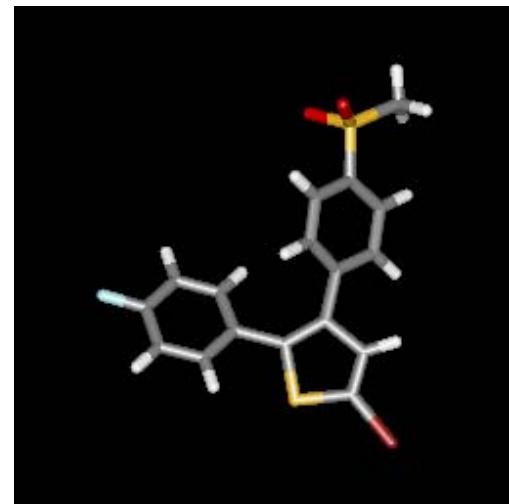
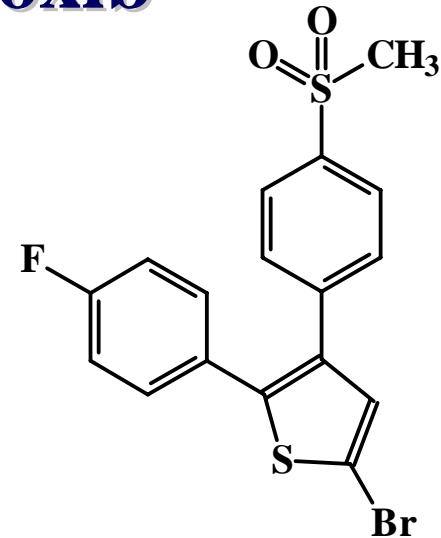
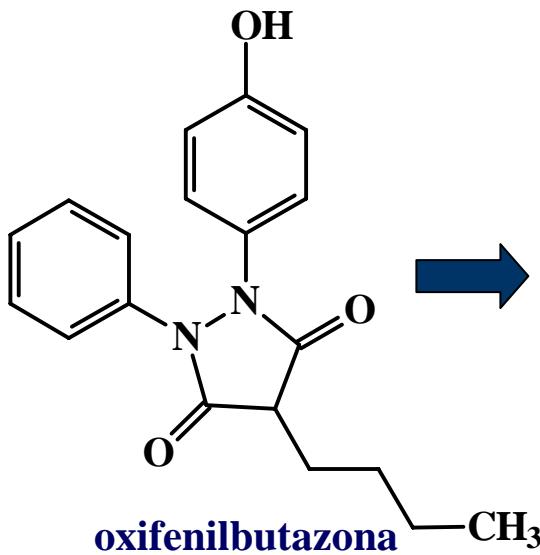
em US\$ bilhões, nos últimos 12 meses;
 & estima-se que o mercado mundial, em 2004, tenha sido ca. US\$ 505 bilhões



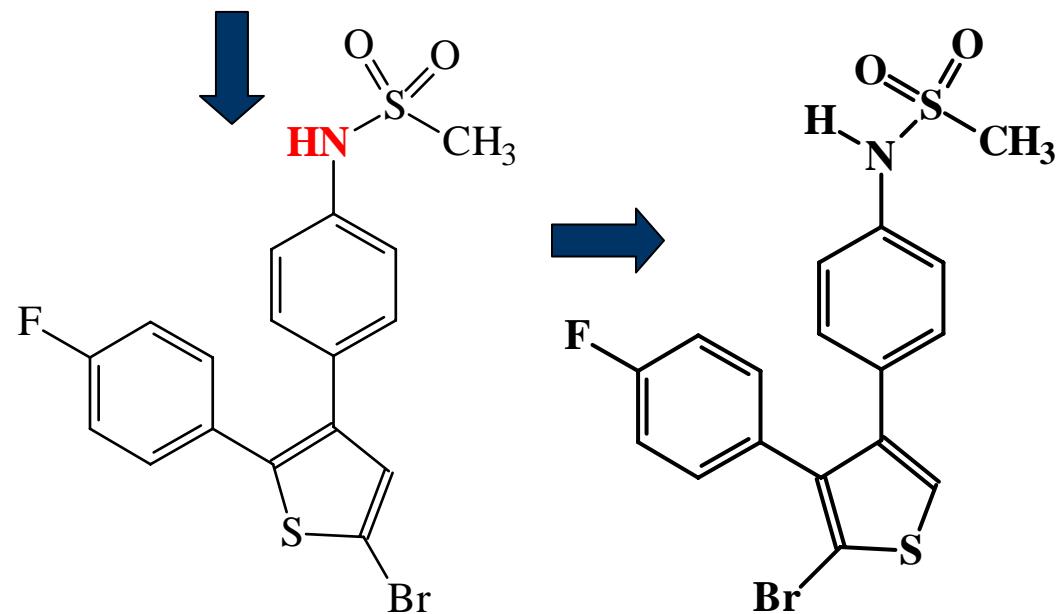


Gênese do Celecoxib

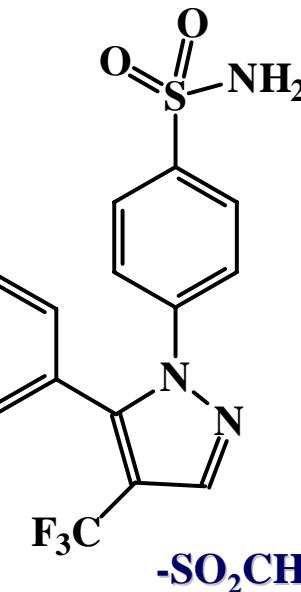
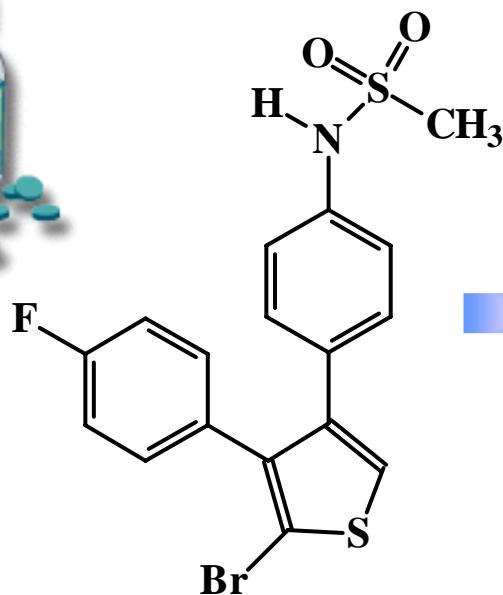
Gênese do Celecoxib



Pfizer



Gênese do Celecoxib



Gans (DuPont) 1990
Vida-média = 12 dias !
(ADME)

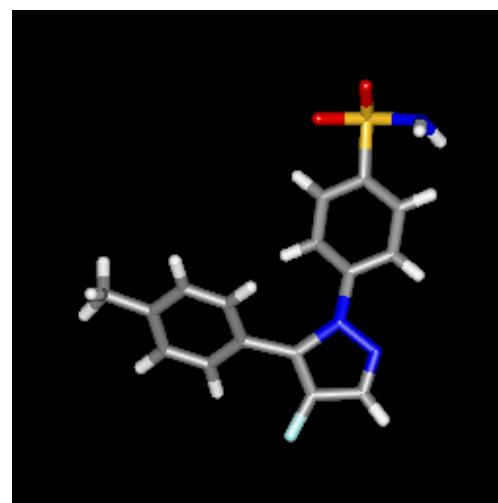


celecoxib (Celebra®)

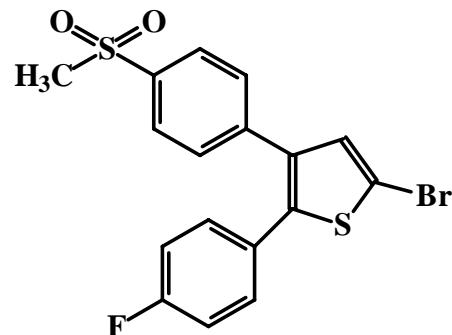
US\$ 797 milhões no
3 trimestre 2004 (>14%)



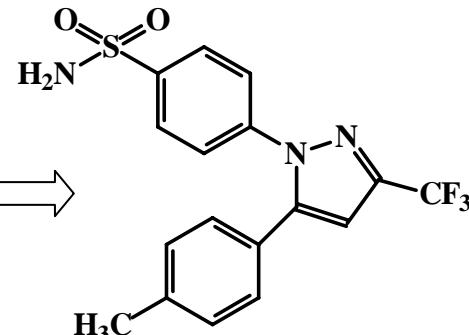
CELEBRA® 100 mg
(CELECOXIB)
Fabricado por:
União Química - Porto Alegre
Importado e Comercializado no Brasil por:
SEARLE Pfizer MONSANTO DO BRASIL LTDA



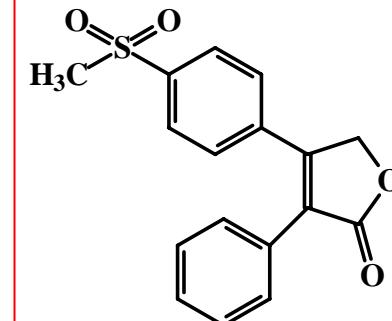
Similaridade Molecular: “me-too”



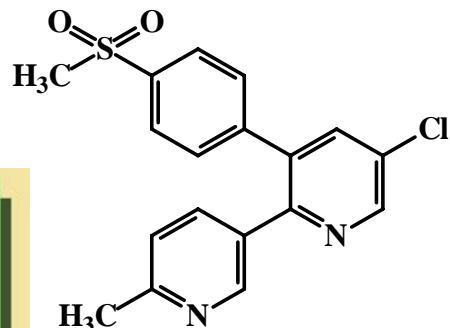
$C_{17}H_{12}BrFO_2S_2$
411.30



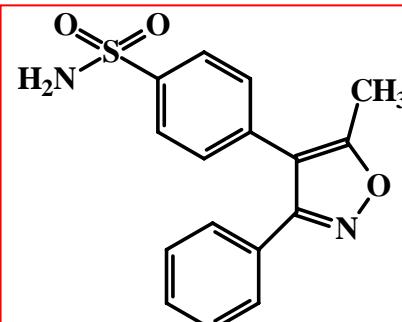
$C_{17}H_{14}F_3N_3O_2S$
381.37



$C_{17}H_{14}O_4S$
314.35



2002
 $C_{18}H_{15}ClN_2O_2S$
358.84

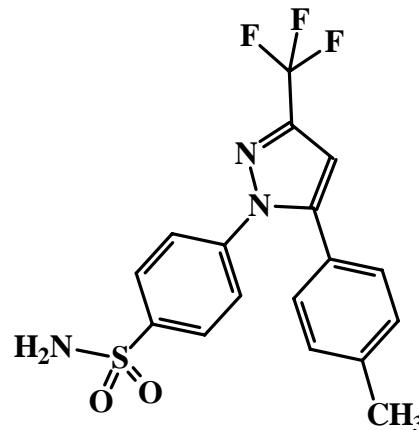


$C_{16}H_{14}N_2O_3S$
314.35

PM 314- 411 uma
anéis: 3
doador-H: 0-1
aceptor-H: 2-4

vendas US\$ 12,8 bilhões no
3 trimestre 2004
(agosto-outubro)

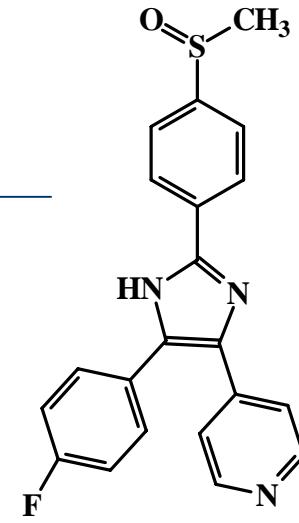
Molecular similarity recognition



$\text{C}_{17}\text{H}_{14}\text{F}_3\text{N}_3\text{O}_2\text{S}$
381

COX-2 inhibitors

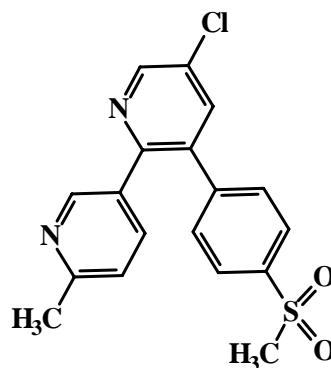
terphenyl system



$\text{C}_{21}\text{H}_{16}\text{FN}_3\text{OS}$
377

SB203580

P38 MAP kinase inhibitor



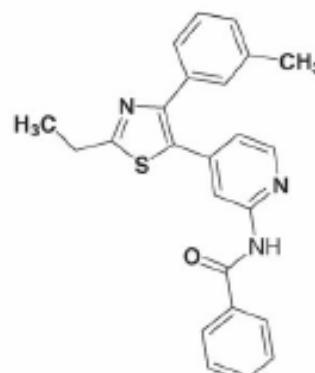
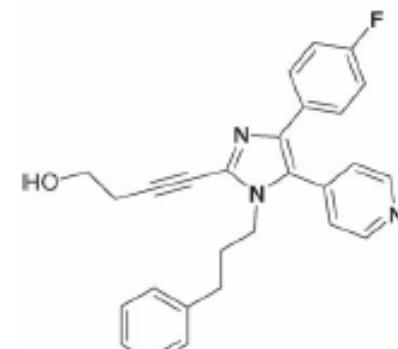
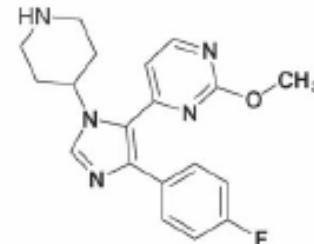
Coxib's act at p38 MAPK level?



p38 MAP kinase inhibitors: A future therapy for inflammatory diseases

Ruth J. Mayer*, James F. Callahan

GlaxoSmithKline Pharmaceuticals, Respiratory and Inflammation CEDD, P.O. Box 1539, King of Prussia, PA 19406, USA



No Published Structures

VX-702
(Vertex)

SCIO-469
(Scios)

SB-681323
(GlaxoSmithKline)

Available online at www.sciencedirect.com

Bioorganic & Medicinal Chemistry Letters 15 (2005) 3506–3509



The molecular basis for coxib inhibition of p38 α MAP kinase

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Carlos M. R. Sant'Anna^{a,c} and Eliezer J. Barreiro^{a,b,*}

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^bDepartamento de Farmacologia Básica e Clínica, Instituto de Ciências Biomédicas, Universidade Federal do Rio de Janeiro (UFRJ), Rio de Janeiro, RJ, 21941-590, Brazil

^cDepartamento de Química, ICE, Universidade Federal Rural do Rio de Janeiro (UFRJ), Seropédica, RJ 23851-970, Brazil

Received 11 April 2005; revised 20 May 2005; accepted 26 May 2005

Abstract—In this work, we present the results of two combined approaches, molecular docking and comparative molecular field analysis (CoMFA), to propose how the selective cyclooxygenase-2 inhibitor celecoxib could act as a p38 mitogen-activated protein (MAP) kinase inhibitor. The docking analysis revealed why celecoxib has a less favorable binding energy ($\Delta G = -12.4$ kcal/mol) than the selective p38 MAP kinase (p38 MAPK) inhibitor, SB203580 ($\Delta G = -22.2$ kcal/mol). The CoMFA results revealed unfavorable steric effects that can be related to the predicted lower p38 MAP kinase inhibitory activity of celecoxib. Additionally, FlexX and CoMFA results also suggested that etoricoxib, another selective COX-2 inhibitor, could inhibit p38 MAP kinase.

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Binding of celecoxib & SB203580 with p38 MAPK

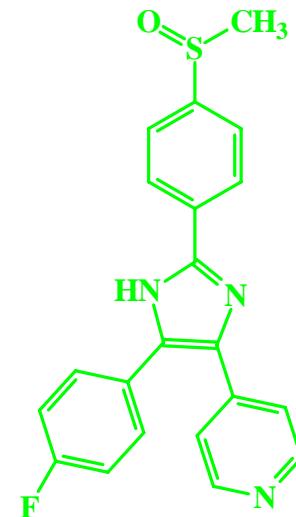
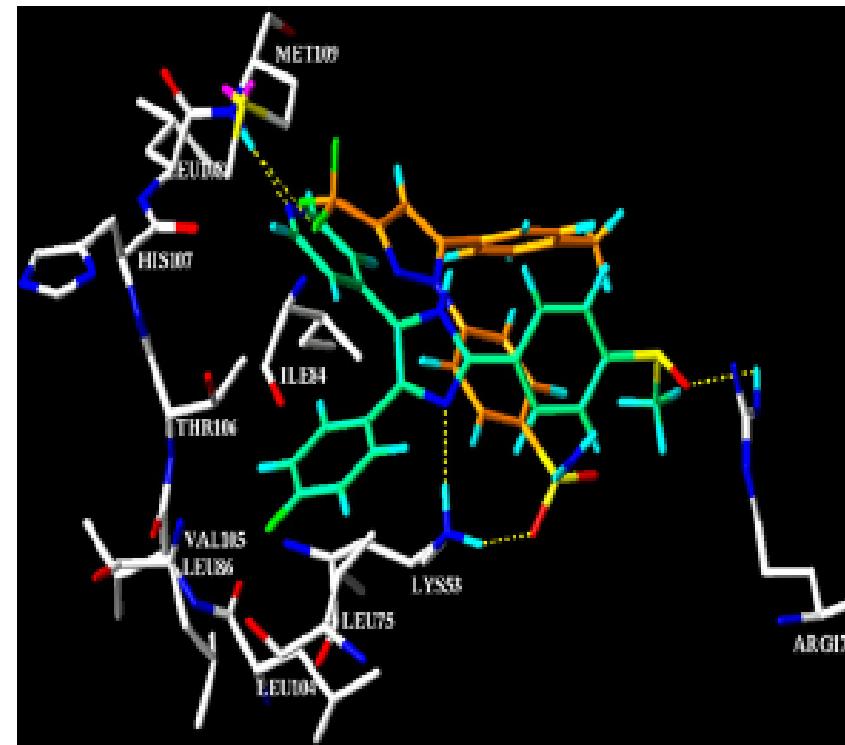
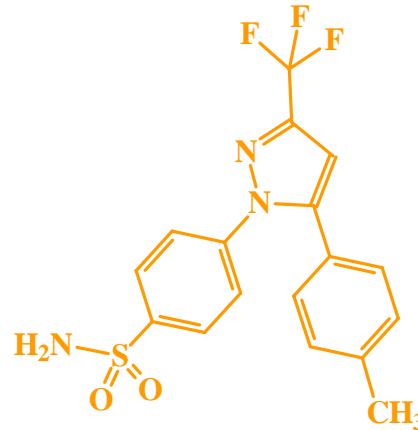


Figure 2. Probable binding conformation of celecoxib and its alignment in the binding site of p38 MAPK. FexX docking shows most important amino acid of p38 MAPK to interact with celecoxib. For celecoxib the carbon atoms are shown in orange (C), blue (N), red (O), and green (halogen). For SB203580, the carbon atoms are shown in green-blue (C), blue (N), red (O), and green (halogen).