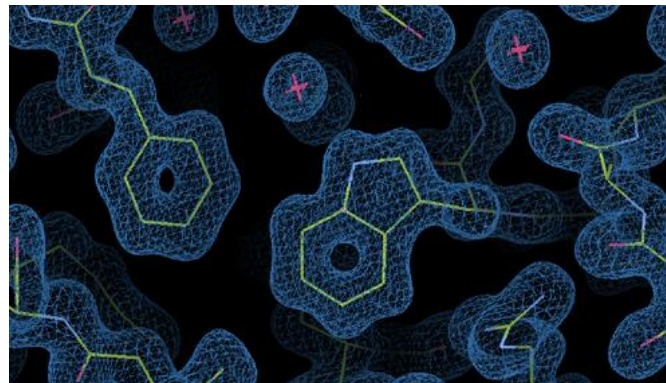


Biologia Estrutural e aplicações

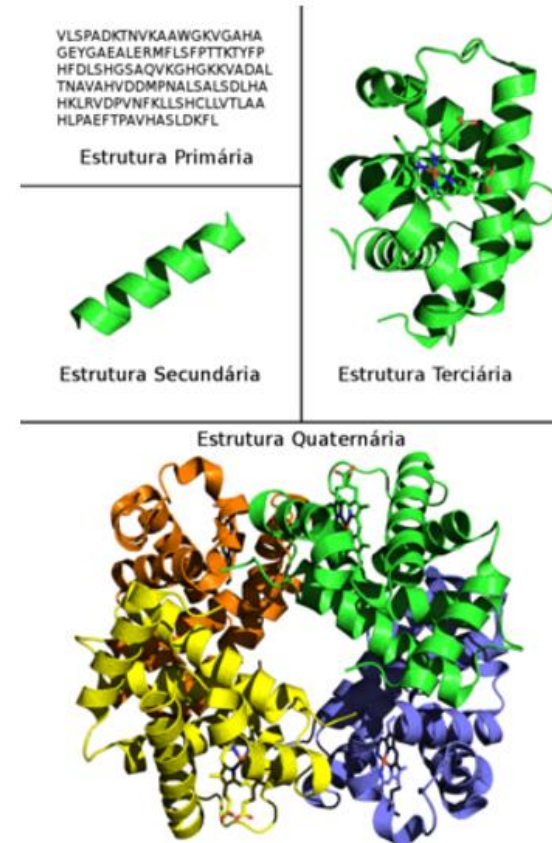
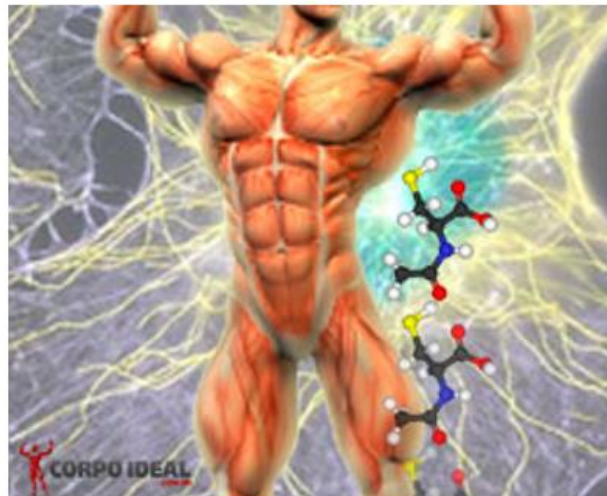
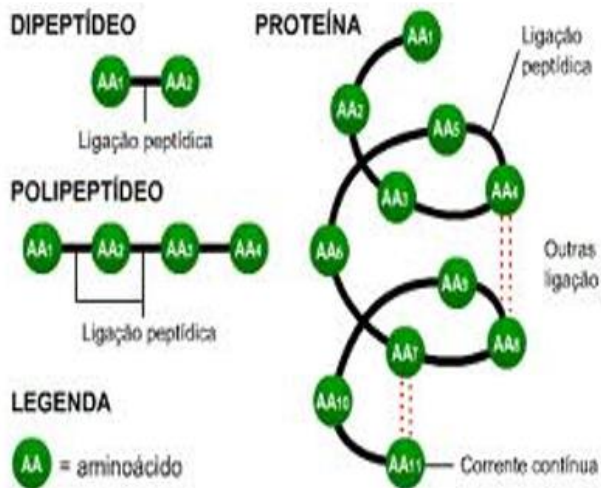
Prof Marcio V B Dias

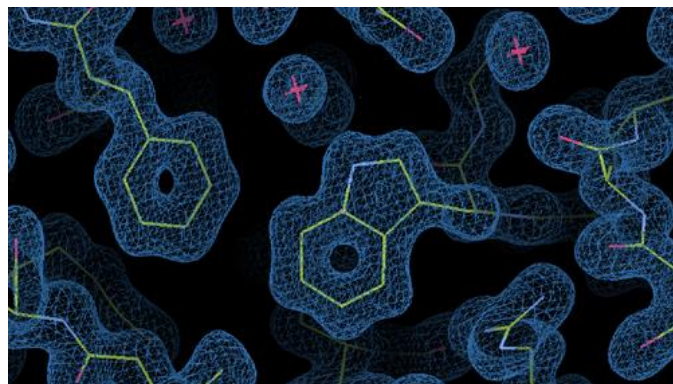
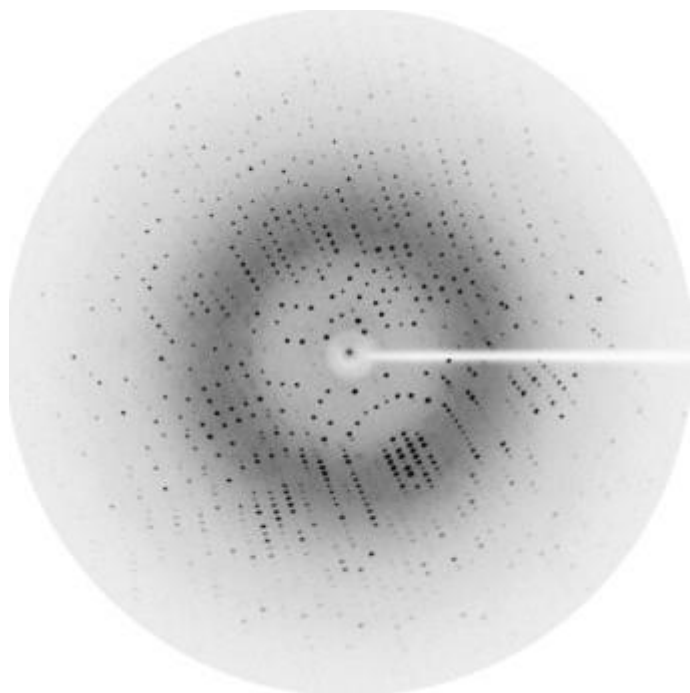
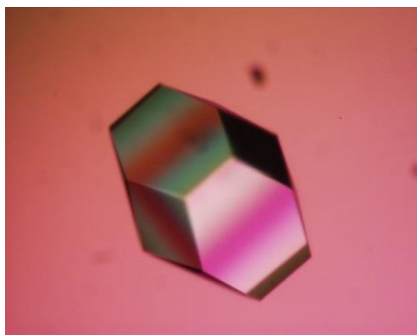
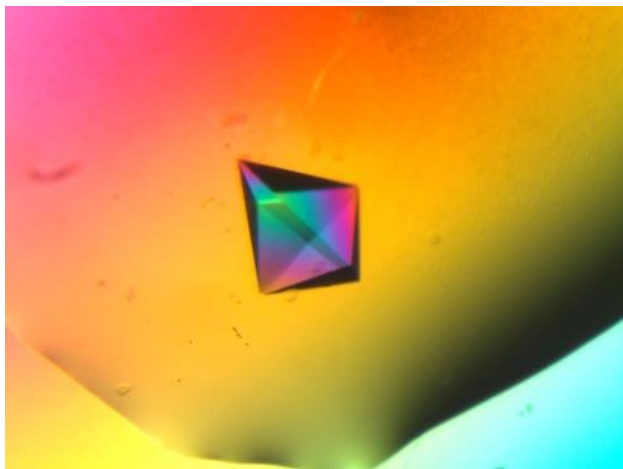
E-mail: mvbdias@usp.br



Por que estudar proteínas???

- ✓ essenciais à estrutura, à função, e ao regulamento celular
- ✓ responsáveis pela homeostasia do organismo
- ✓ inúmeras finalidades: diagnóstica, terapêutica e de prevenção (vacinas)



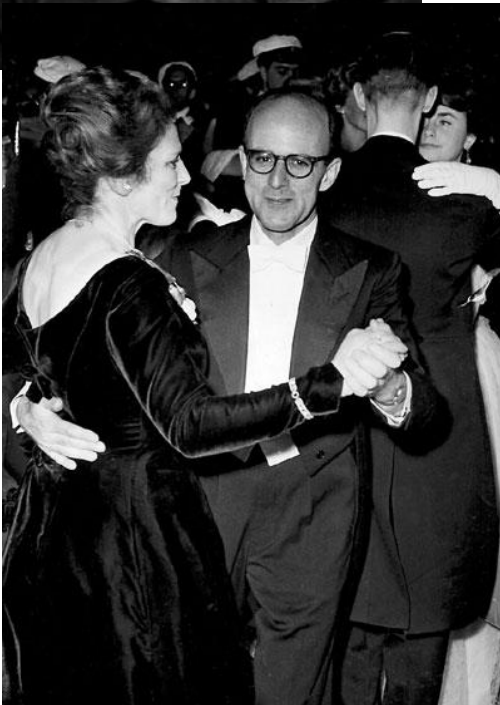




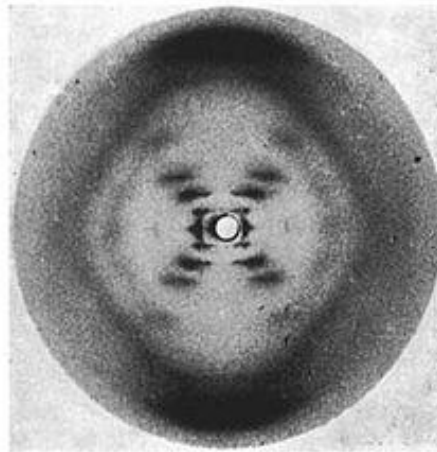
**Dorothy C.
Hodgkin**



Venkatraman Ramakrishnan Thomas A. Steitz Ada E. Yonath



Max Perutz



Rosalind Franklin

Para que cristalizar uma proteína?

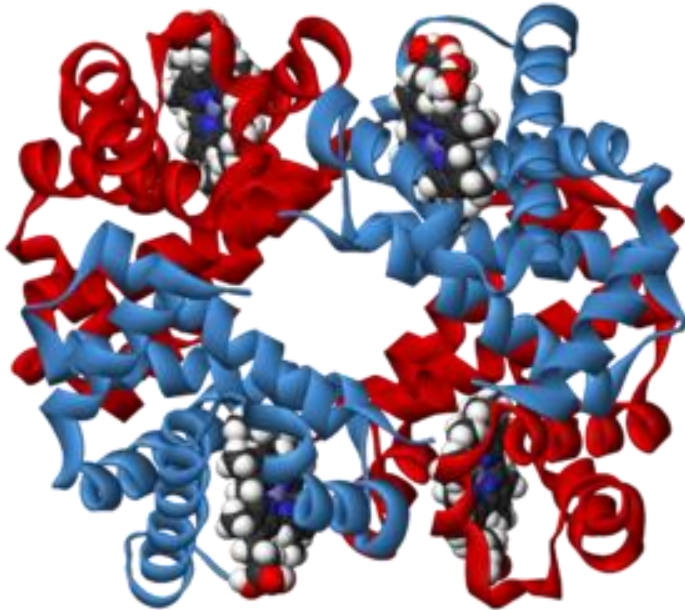
Primeira resposta:

-entender sua estrutura



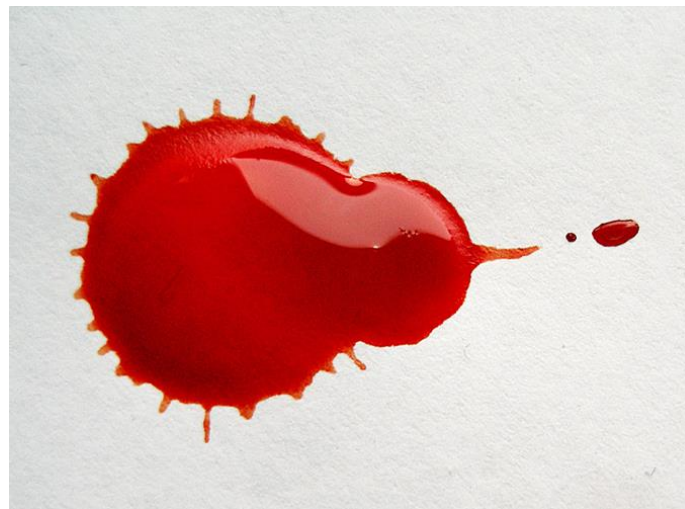
Mas para que entender a estrutura?

- entender o processos biológicos**
 - como transporte de oxigênio**
 - anticorpos**
- interação de proteínas com ligantes**
- mecanismo de catálise**
- desenho de medicamentos**
- regulação alosterica**
- mecanismo de doenças**

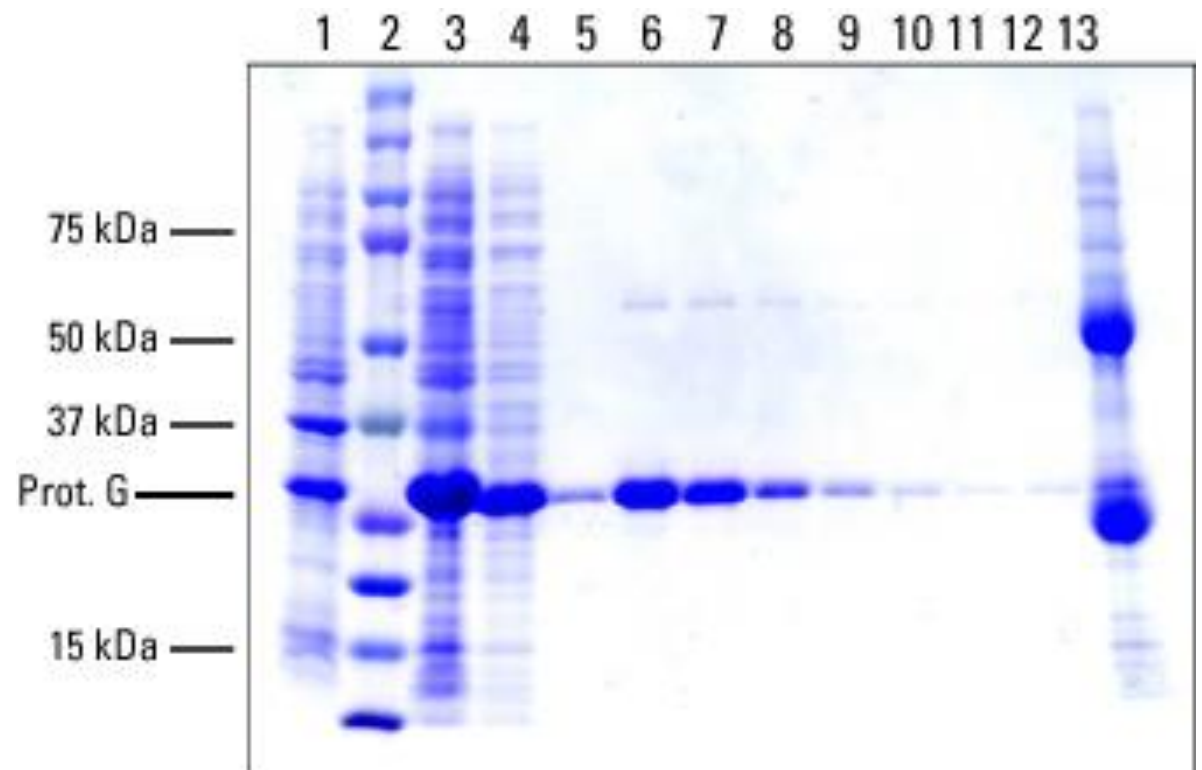


Primeiro Passo

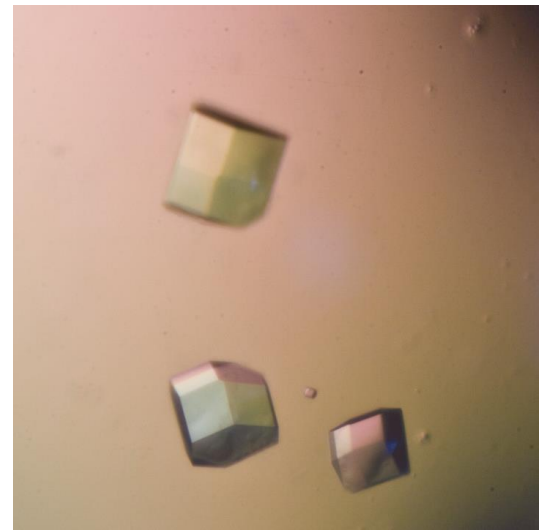
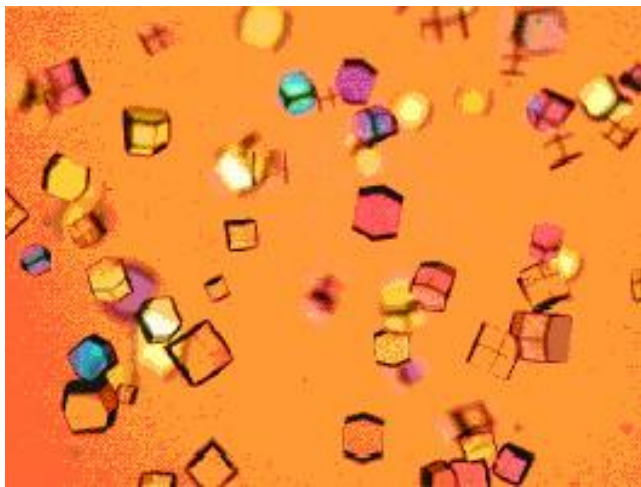
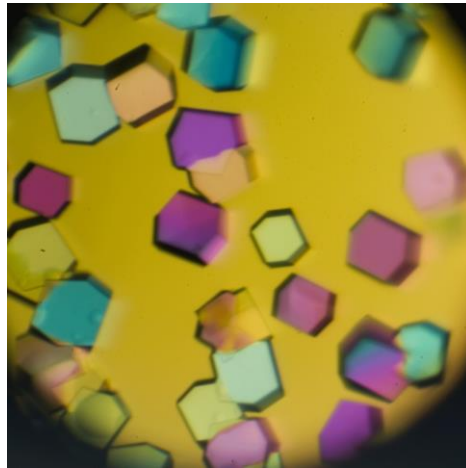
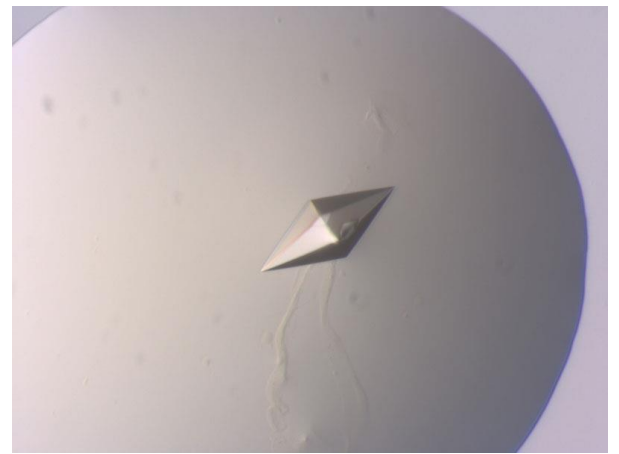
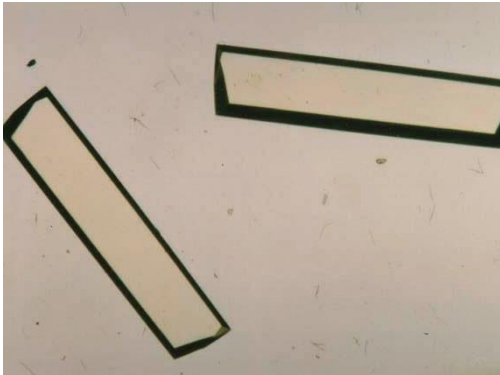
Purificação de proteínas



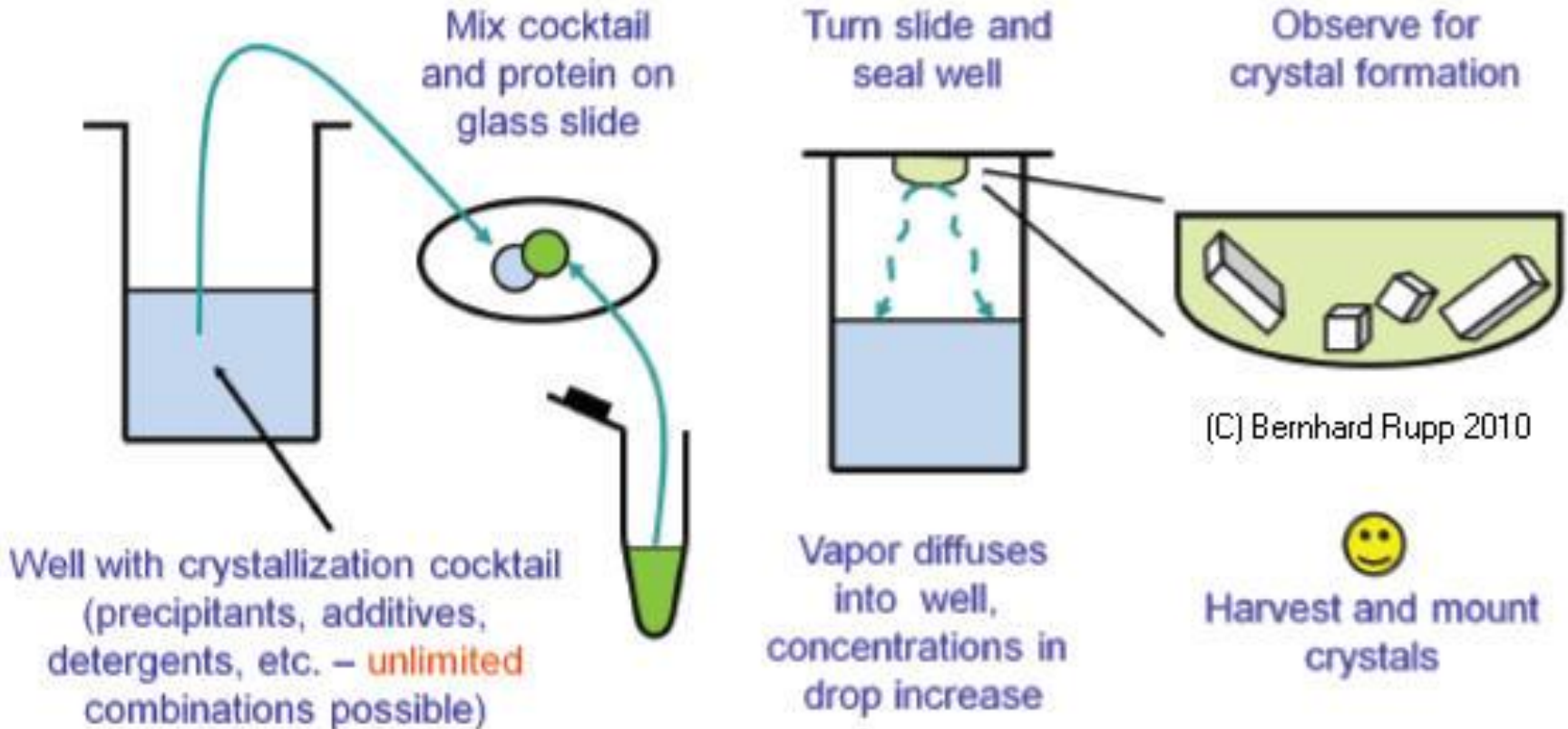
Purificação de proteínas baseado em afinidade



Cristais de proteína

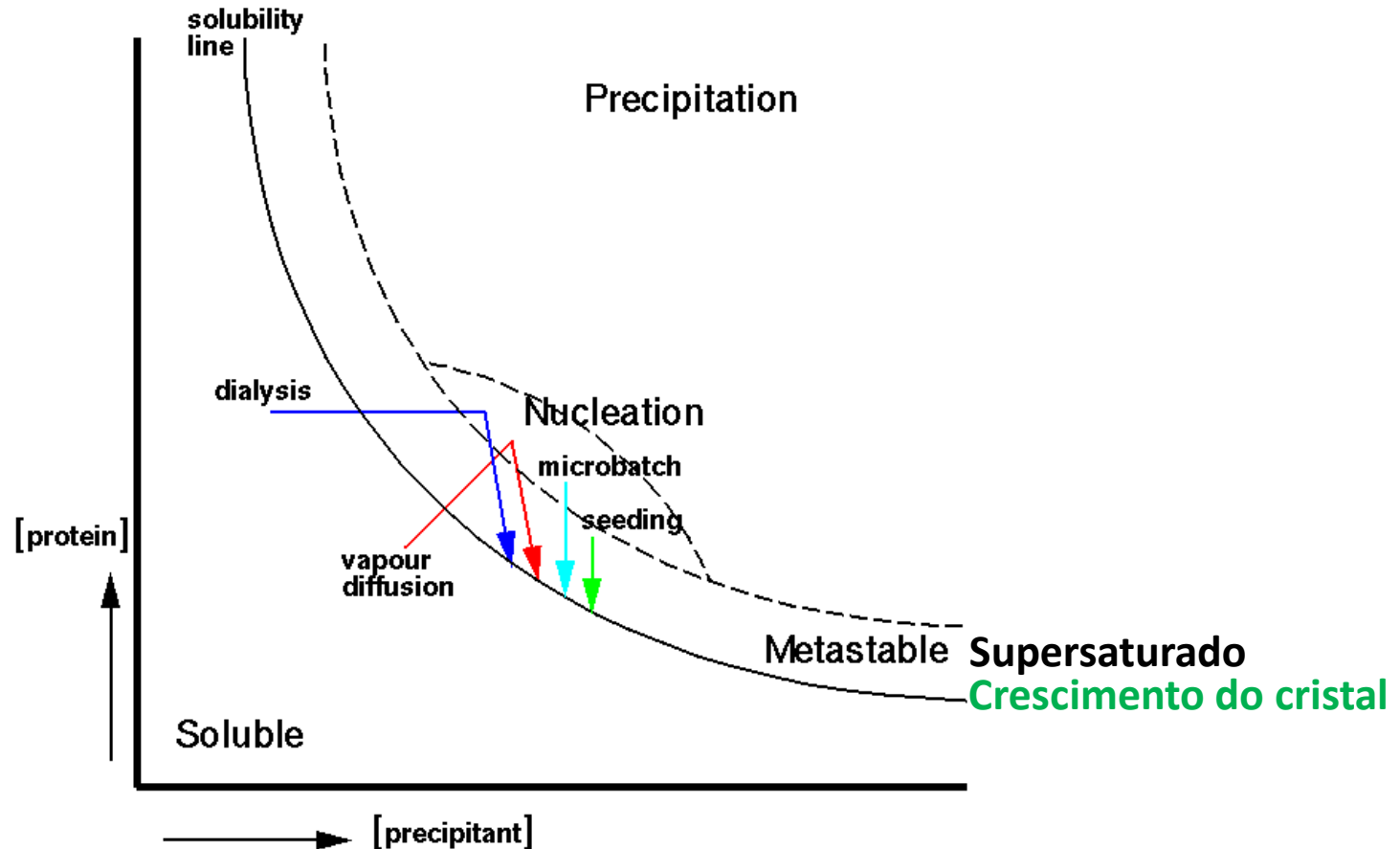


O Processo de Cristalização



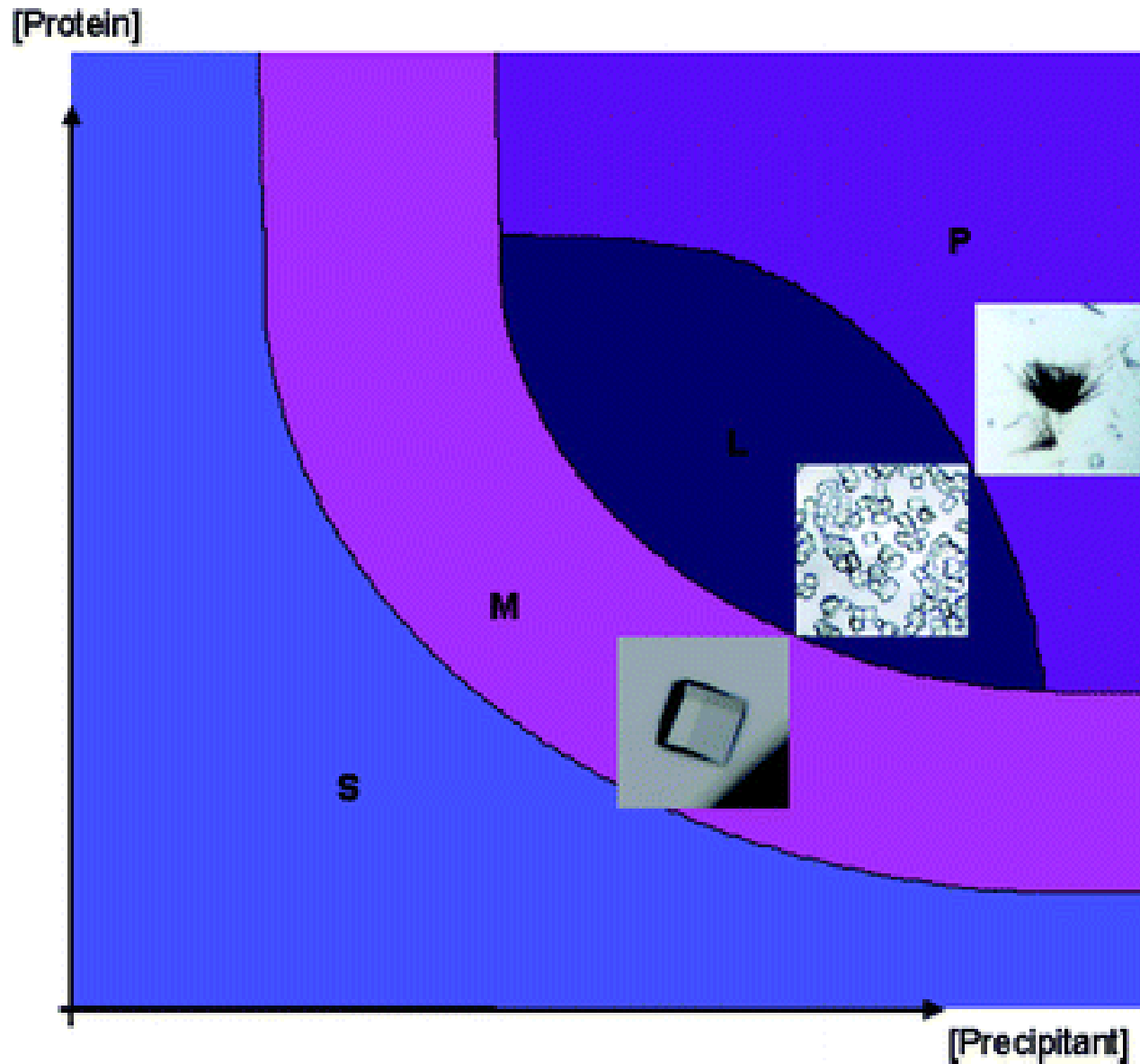
Como obter um cristal de Proteína?

Diagrama de fase

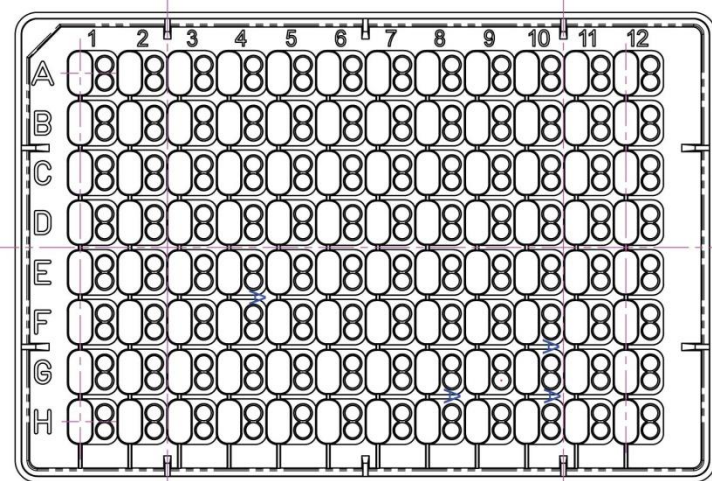
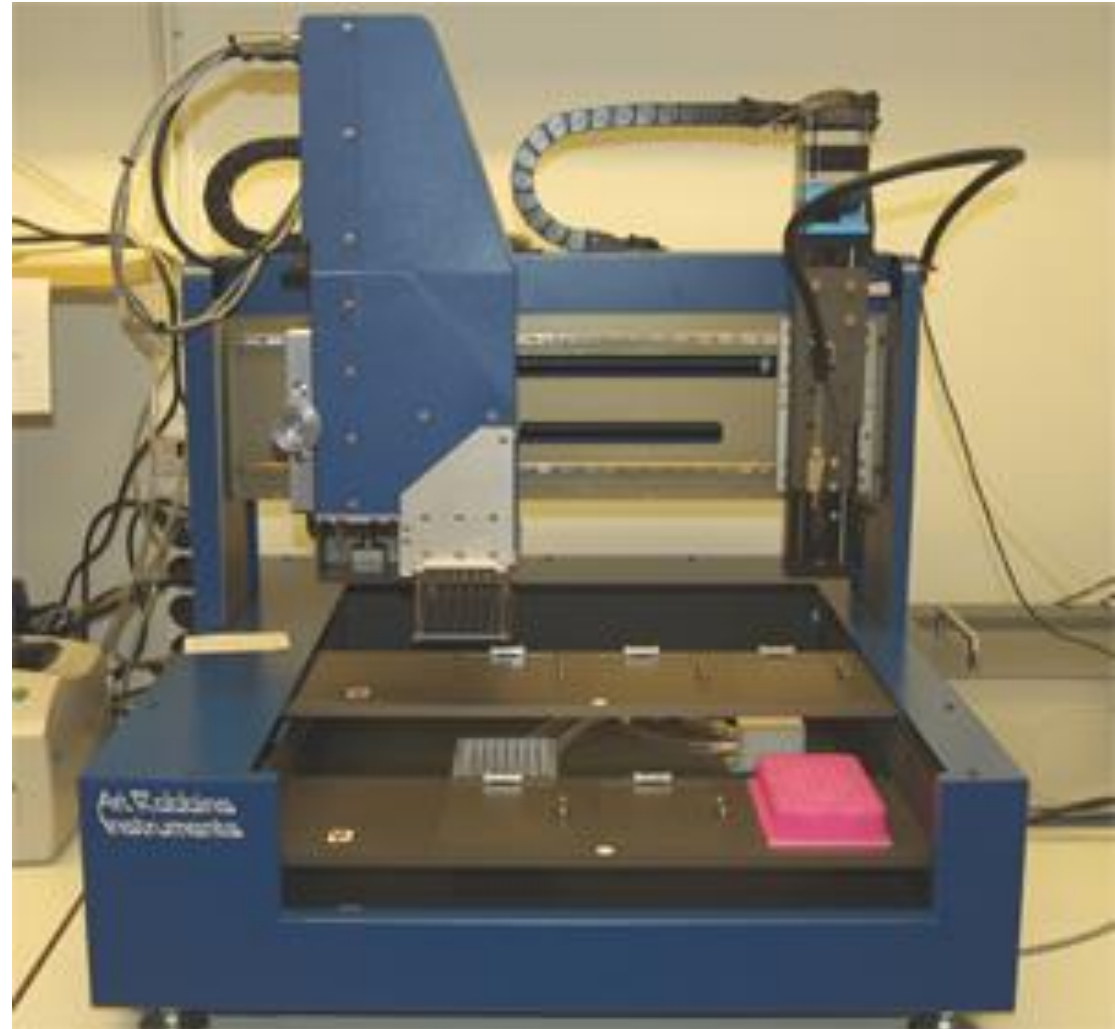
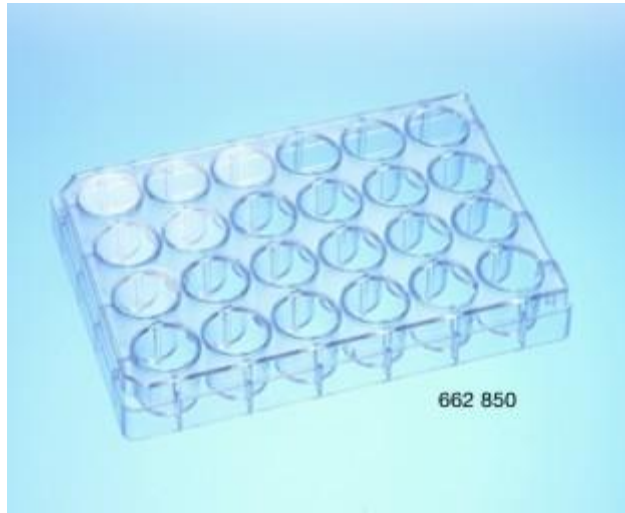


Protein Solubility Phase Diagram and Crystallisation

Diagrama de fase



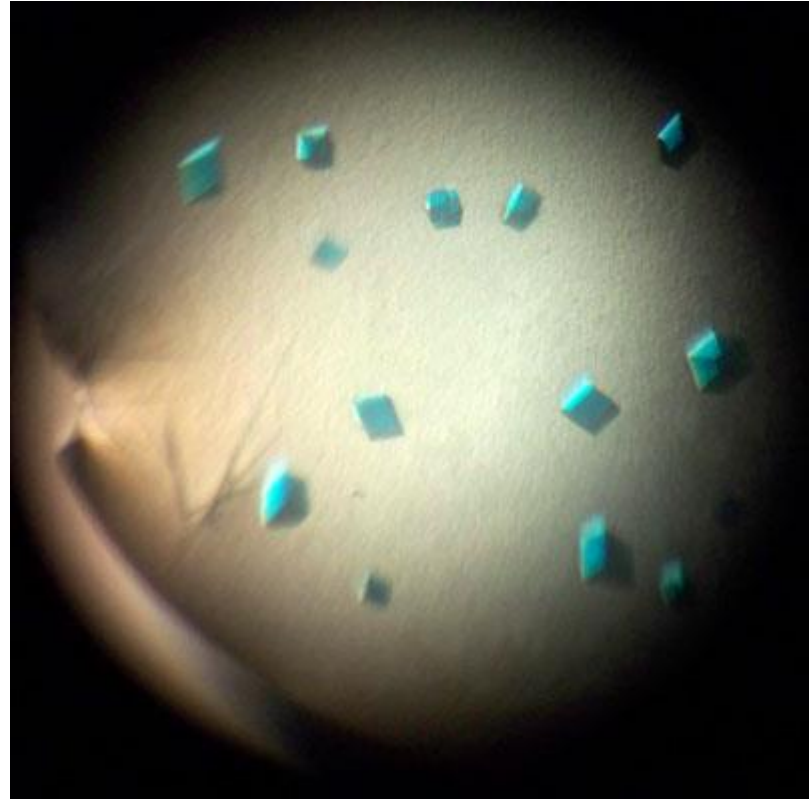
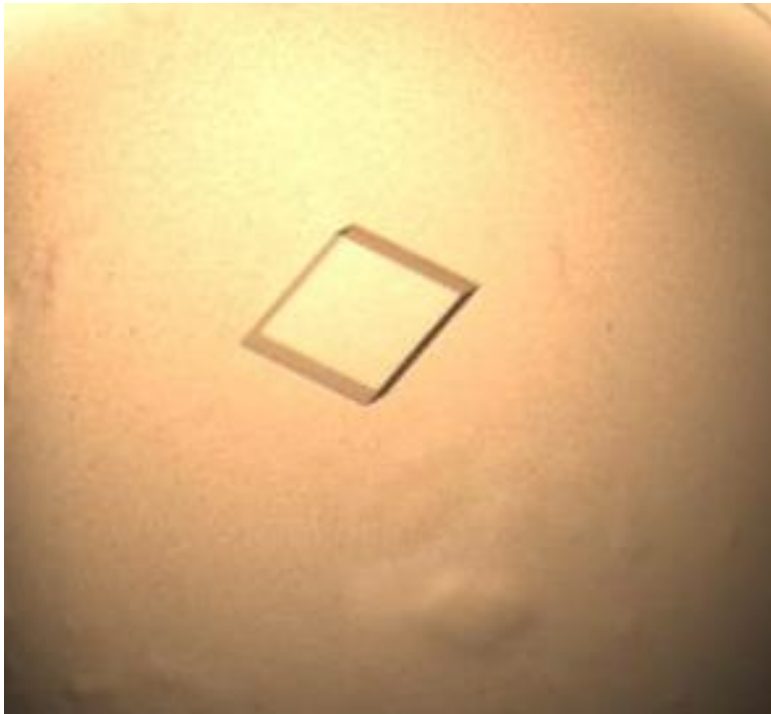
Cristalização rototizada



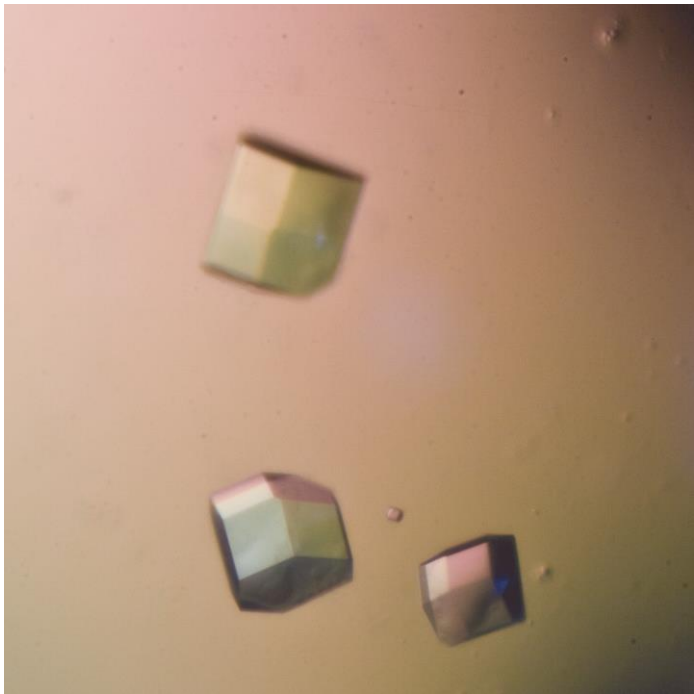
Diferentes Kits de Cristalização

- Tampão
- Sal
- Agente precipitante

Diferenciar Cristal de Proteína com de Sal



Montagem dos cristais



Sírius

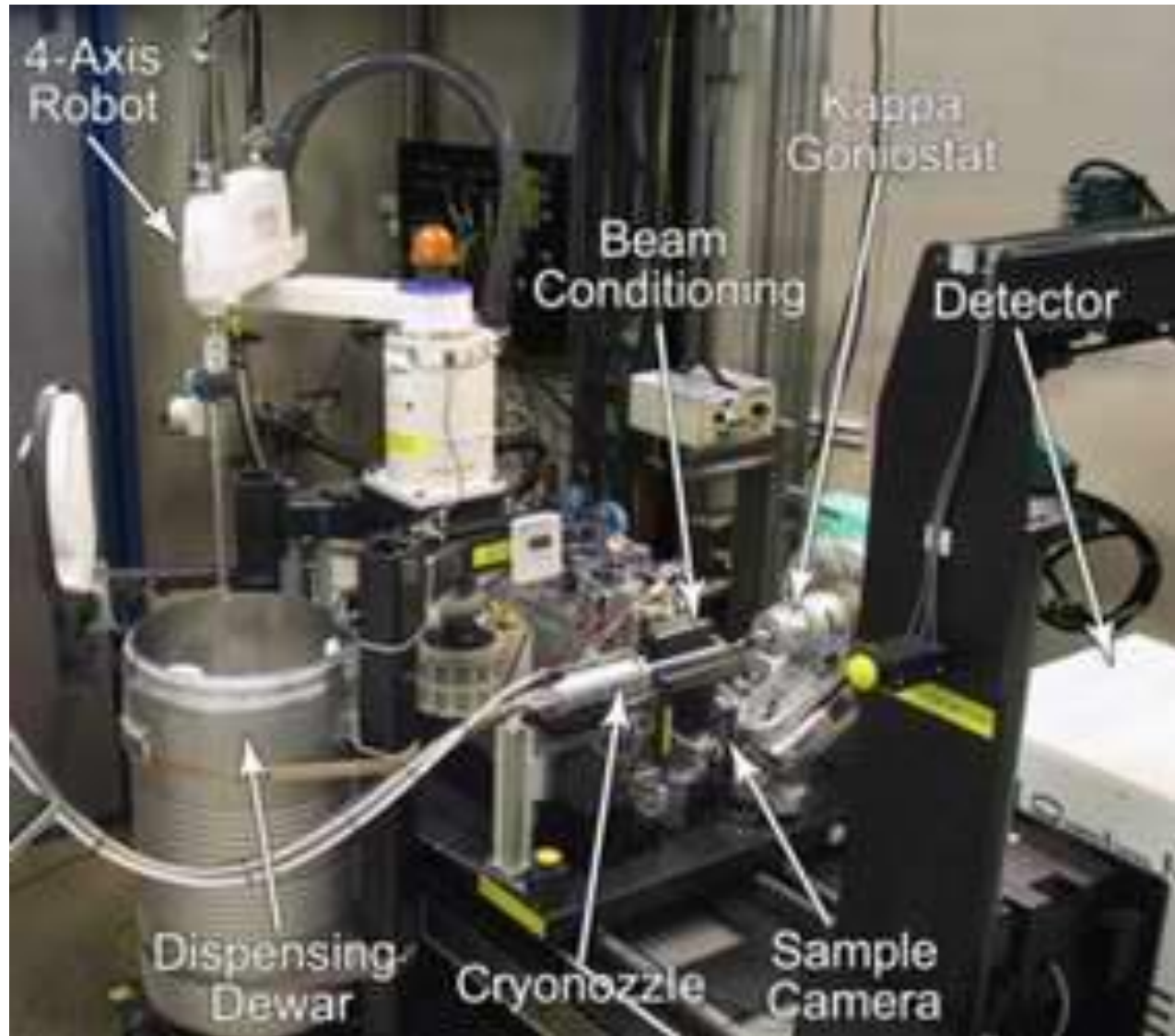


https://www.youtube.com/watch?v=lbxOSSUkgv0&list=RDCMUcYhTgGdeaBbbZ-h_LVJ_6bw&start_radio=1&rv=lbxOSSUkgv0&t=549

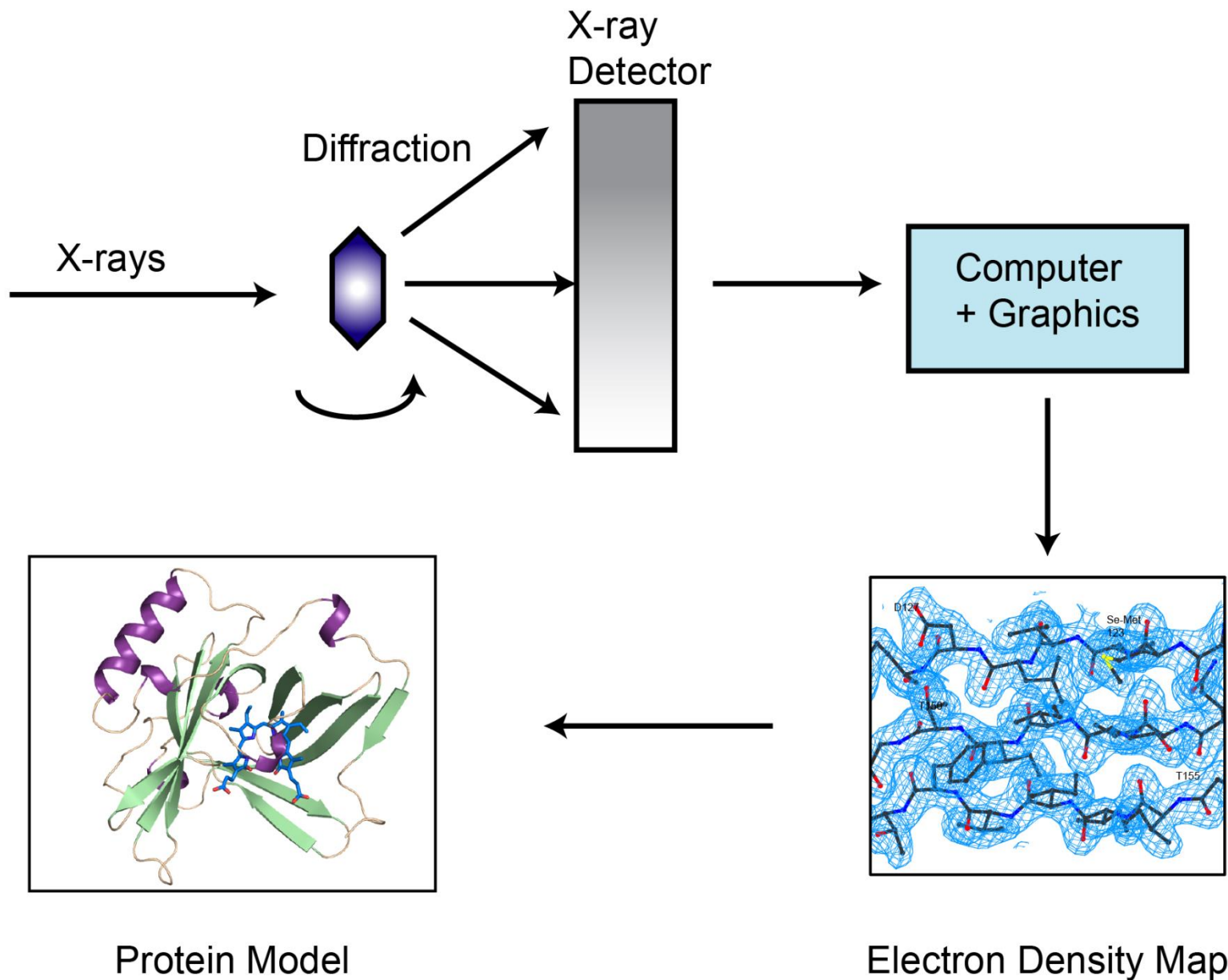
MX2

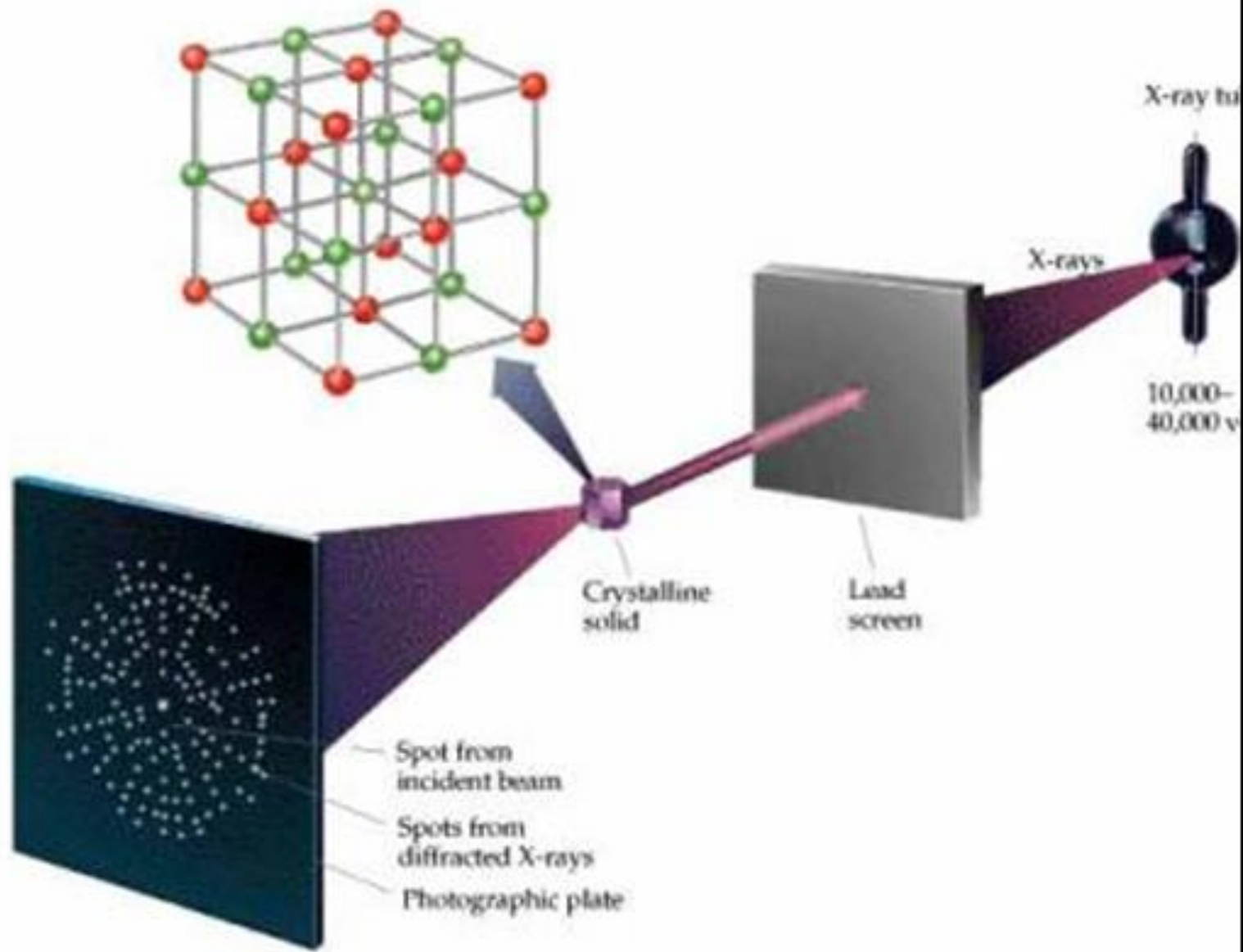


Instrumentação de uma estação de coleta

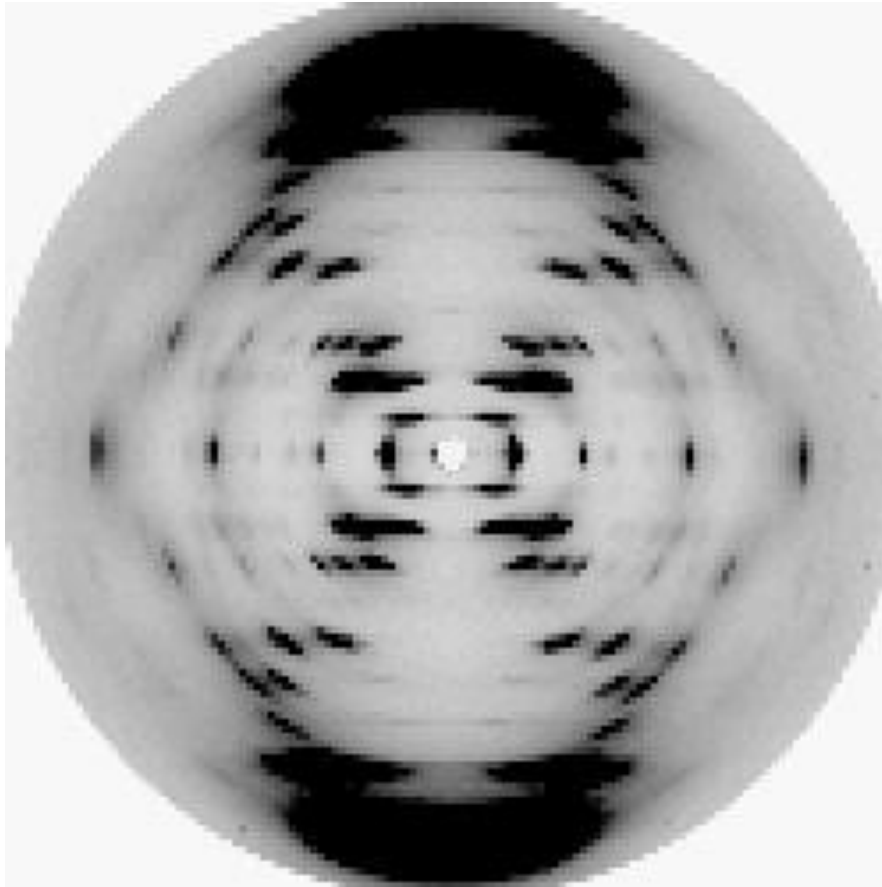


Overview of the X-ray Crystallographic Method





Padrão de difração para o DNA



Padrão de difração de proteína

automar (V1.0.0) - /e/data/lyso/c/lyso_001.mar1200 (1200 pixels & 437 > 65535)

File Options Image Options Edit Page Show Page Log files Set defaults Help

Show Page ...

Display Setup Text

Plot: Index

First image: 1

Last image: 6

Min. resol.: 100

Max. resol.: 2.402

Edit input file

Autoupdate parameters

Run Step

Peak Search Edit

Index Edit

Predict Edit

Strategy Edit

automar

Integrate Edit

Join Edit

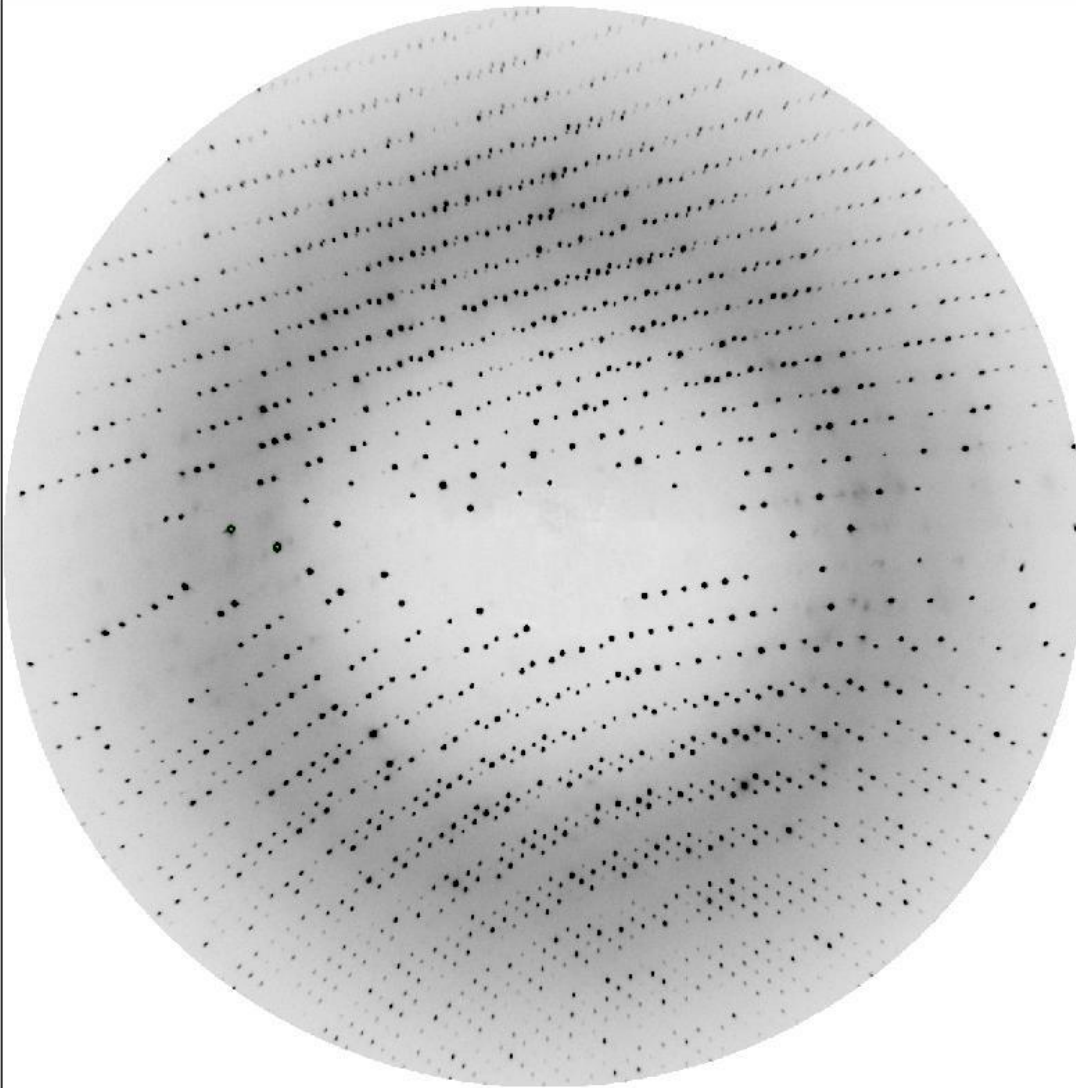
Scale Edit

Truncate Edit

Convert Edit

Run Selection

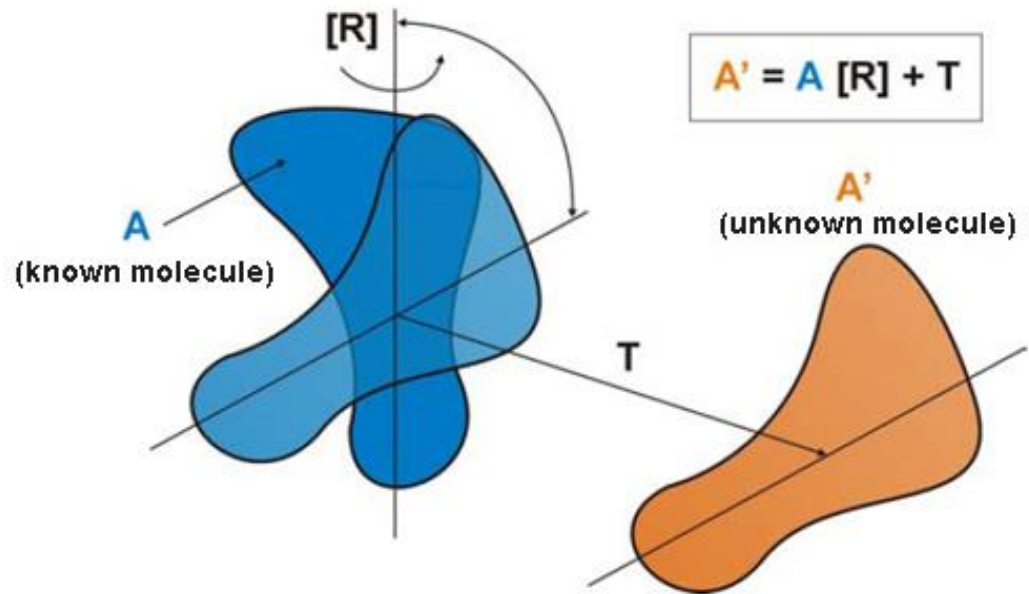
Run on localhost



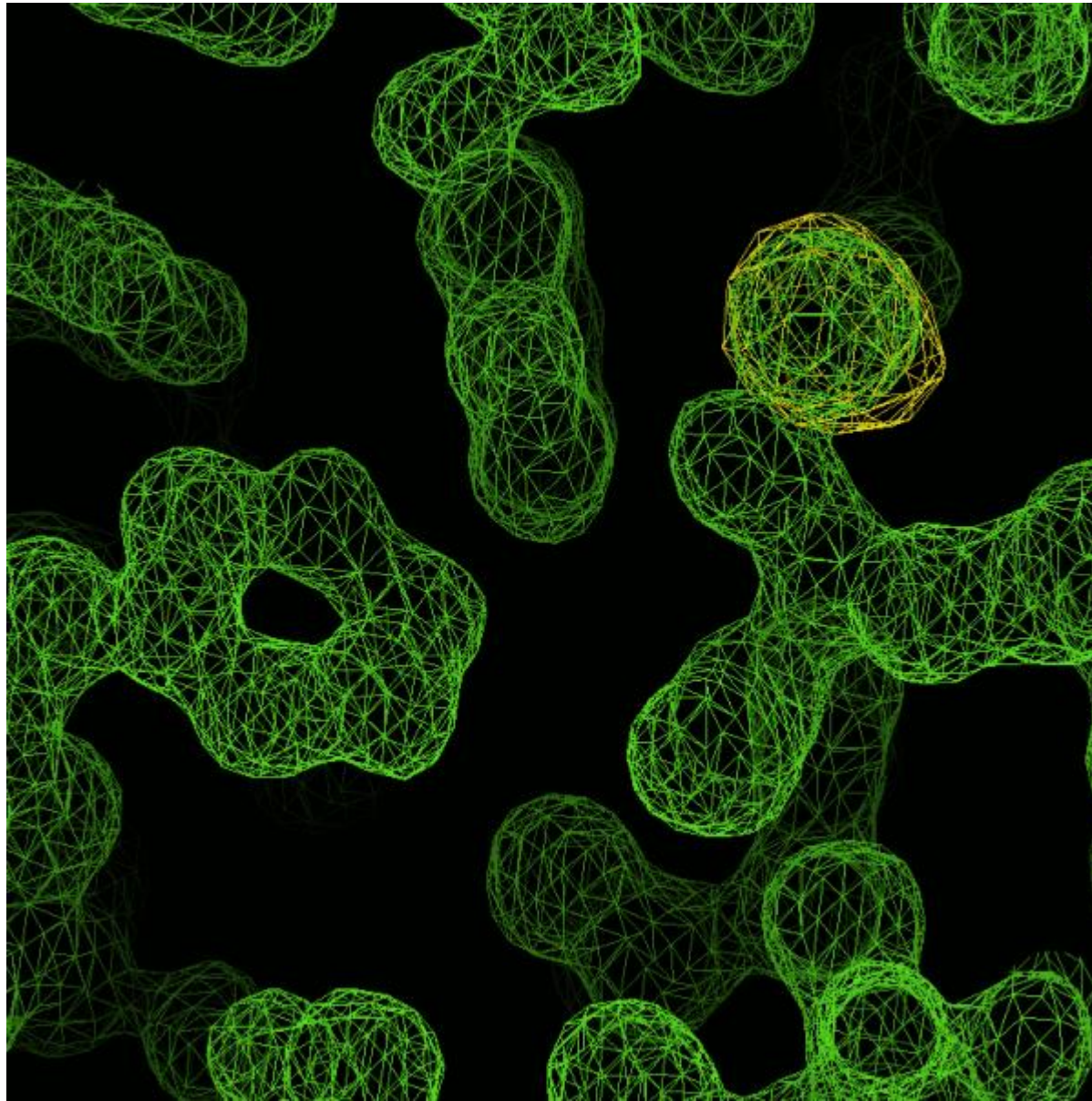
Métodos de determinação da fase

-Substituição molecular

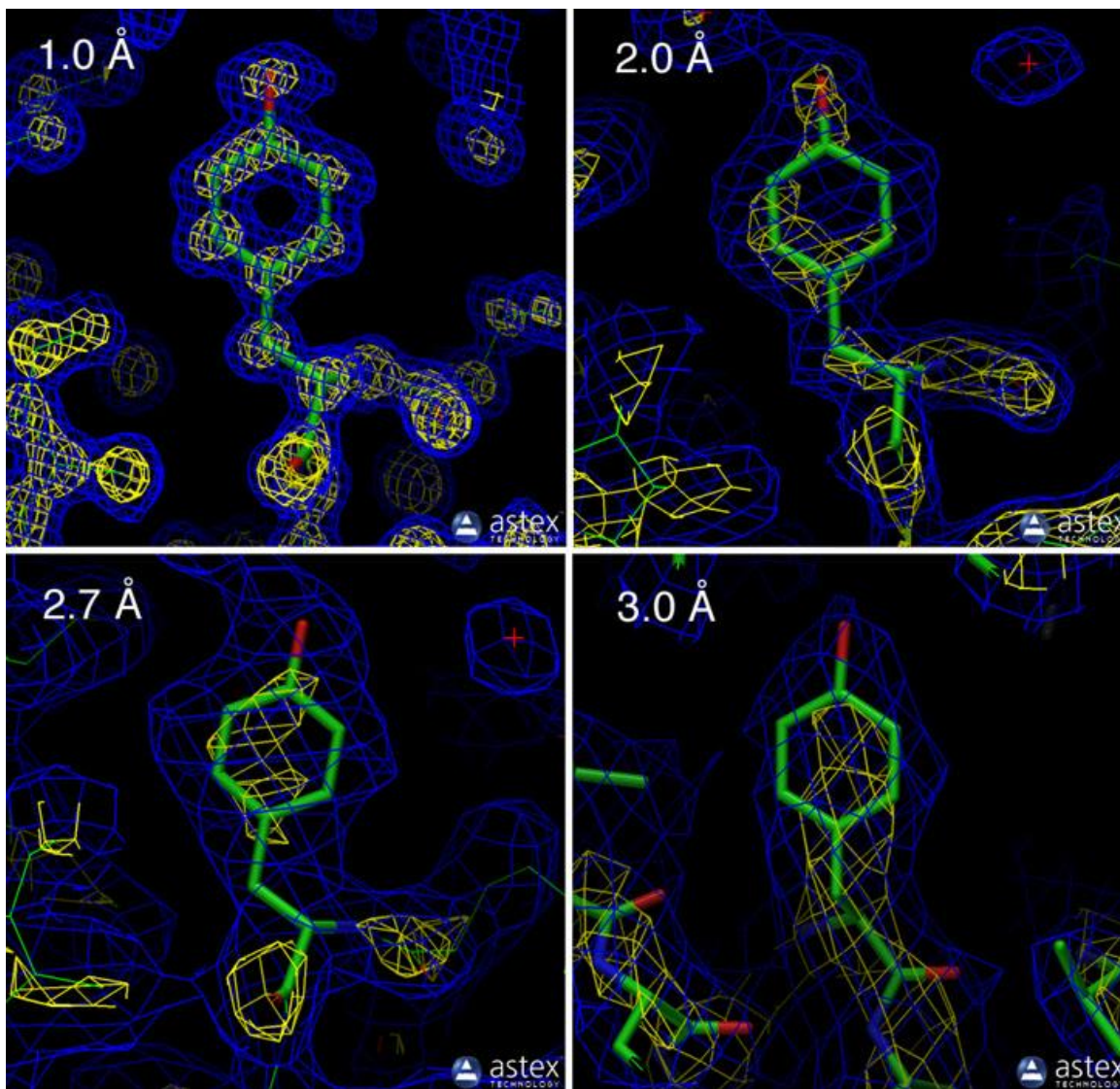
-Métodos experimentais



Mapa de densidade eletrônica

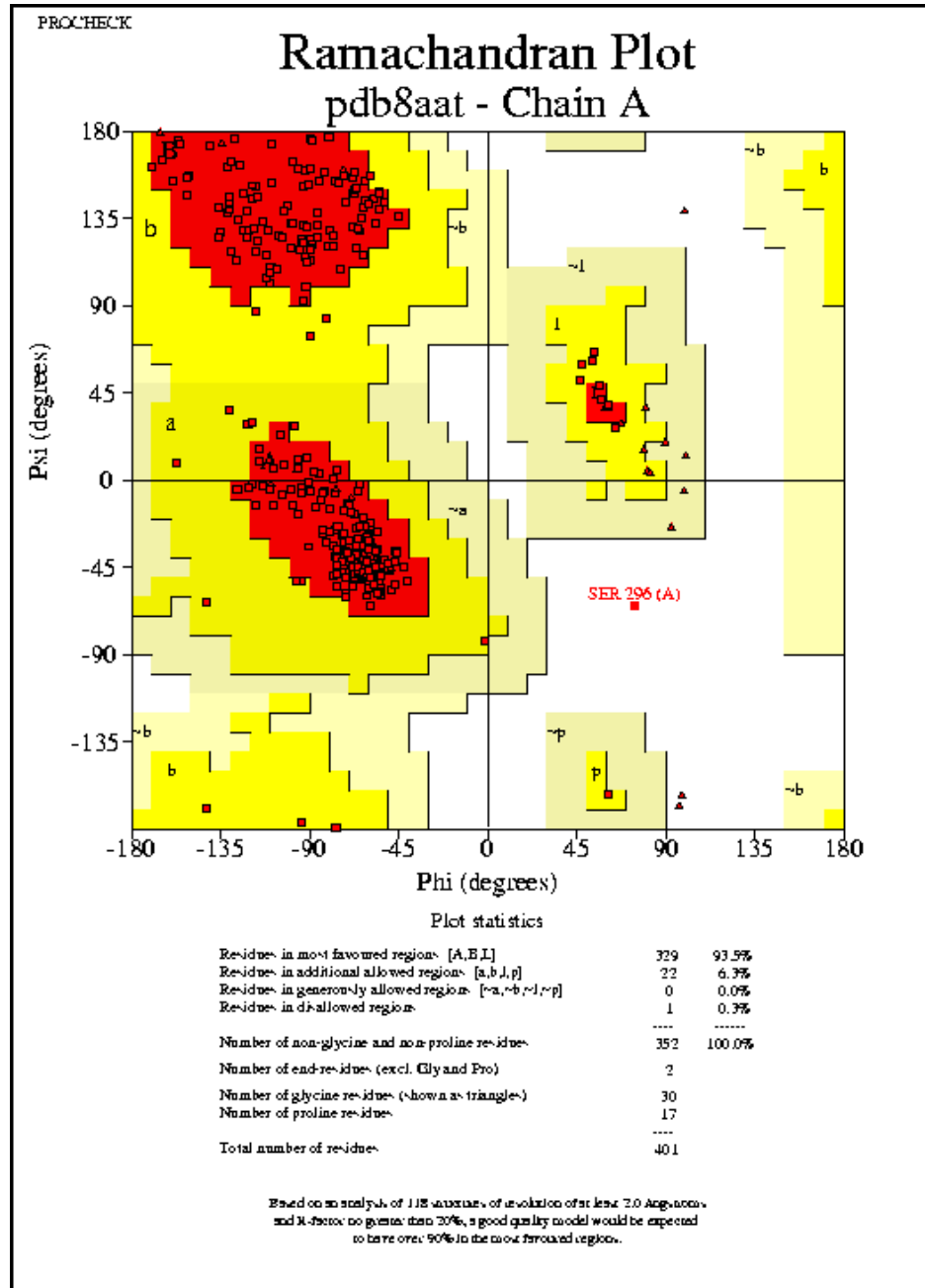


Resolução estrutural



Análise da qualidade da estrutura

Baseado em restrições conformacionais

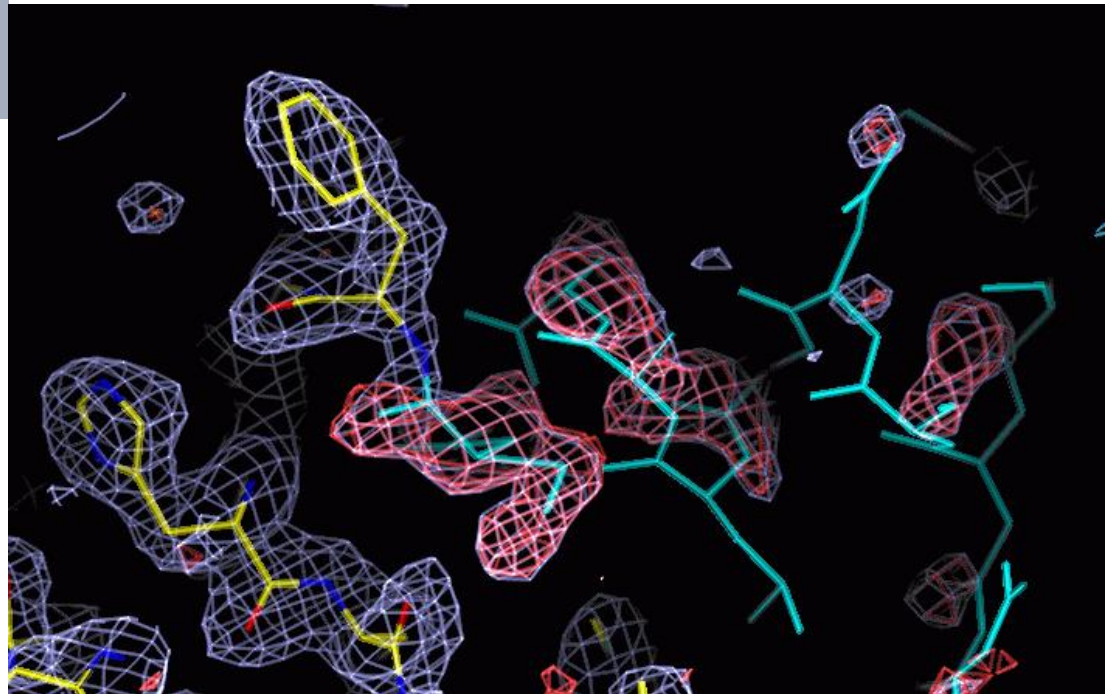


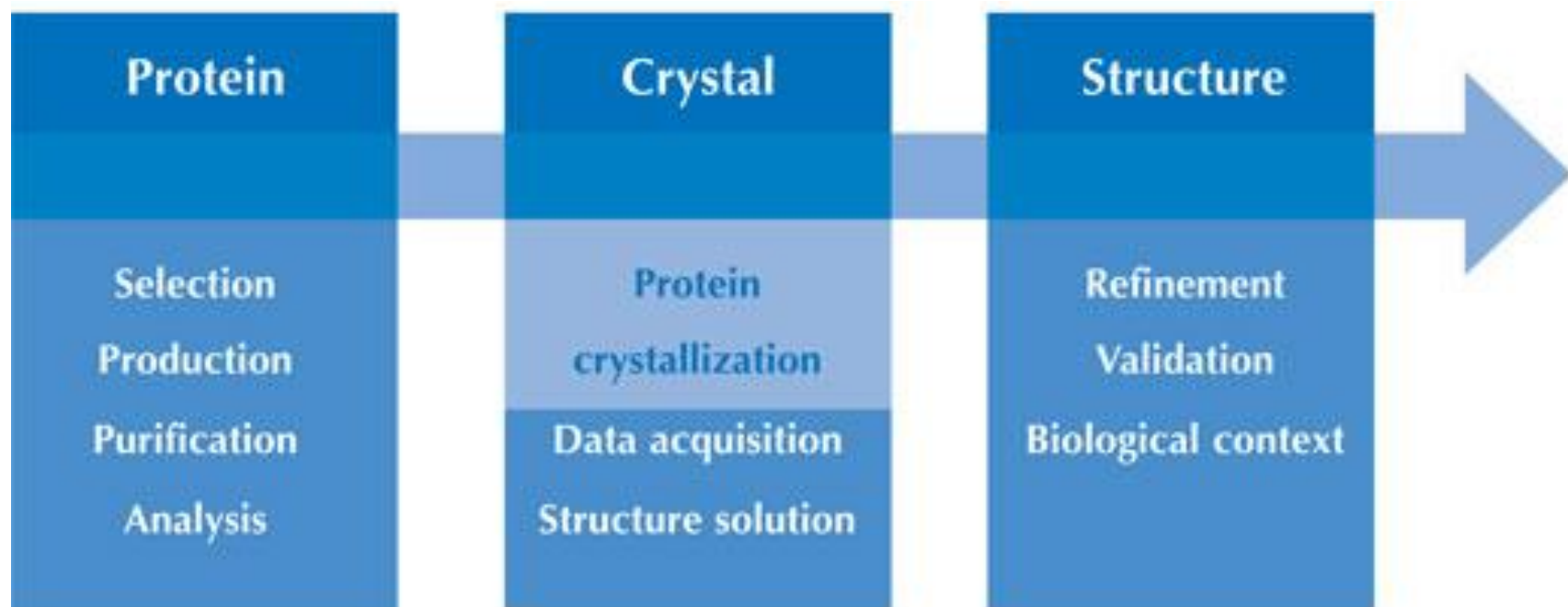
Fatores importantes para averiguar a qualidade de uma estrutura cristalográfica:

- Resolução
- Fator-R e R-free
- Qualidade estereoquímica

Refinamento cristalográfico

Coot





Exemplos de aplicação da biologia estrutural:

- Planejamento de fármacos baseado em estrutura**
- Melhoramento em biocatálise**

Planejamento de fármacos e Desenvolvimento



Como os fármacos são descobertos e desenvolvidas?

Trabalhando em conjunto

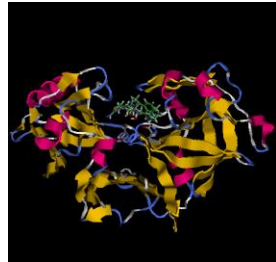
- **Biologia estrutural**
- **Bioquímica**
- **Química medicinal**
- **Toxicologia**
- **Farmacologia**
- **Biofísica química**
- **Tecnologia de informação**

Processo de descoberta e planejamento de fármacos

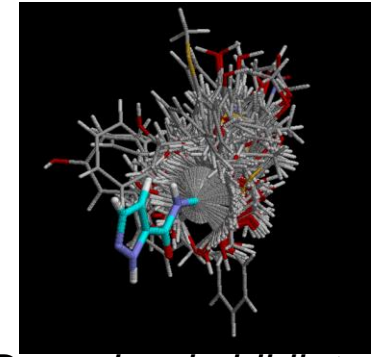
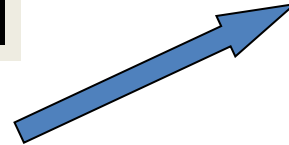


Banco de dados de sequências gênicas

Raio-X ou Homologia



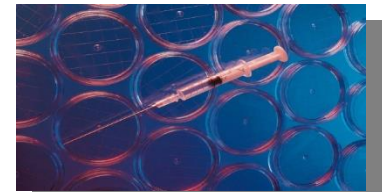
Med Chem/Combichem



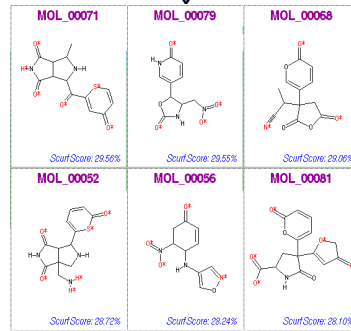
Desenho de bibliotecas



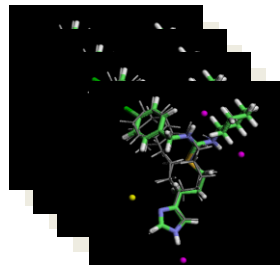
Bibliotecas de compostos



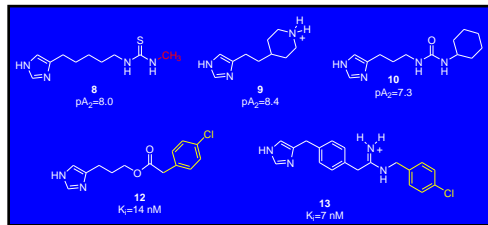
Screening



Desenho de modelos



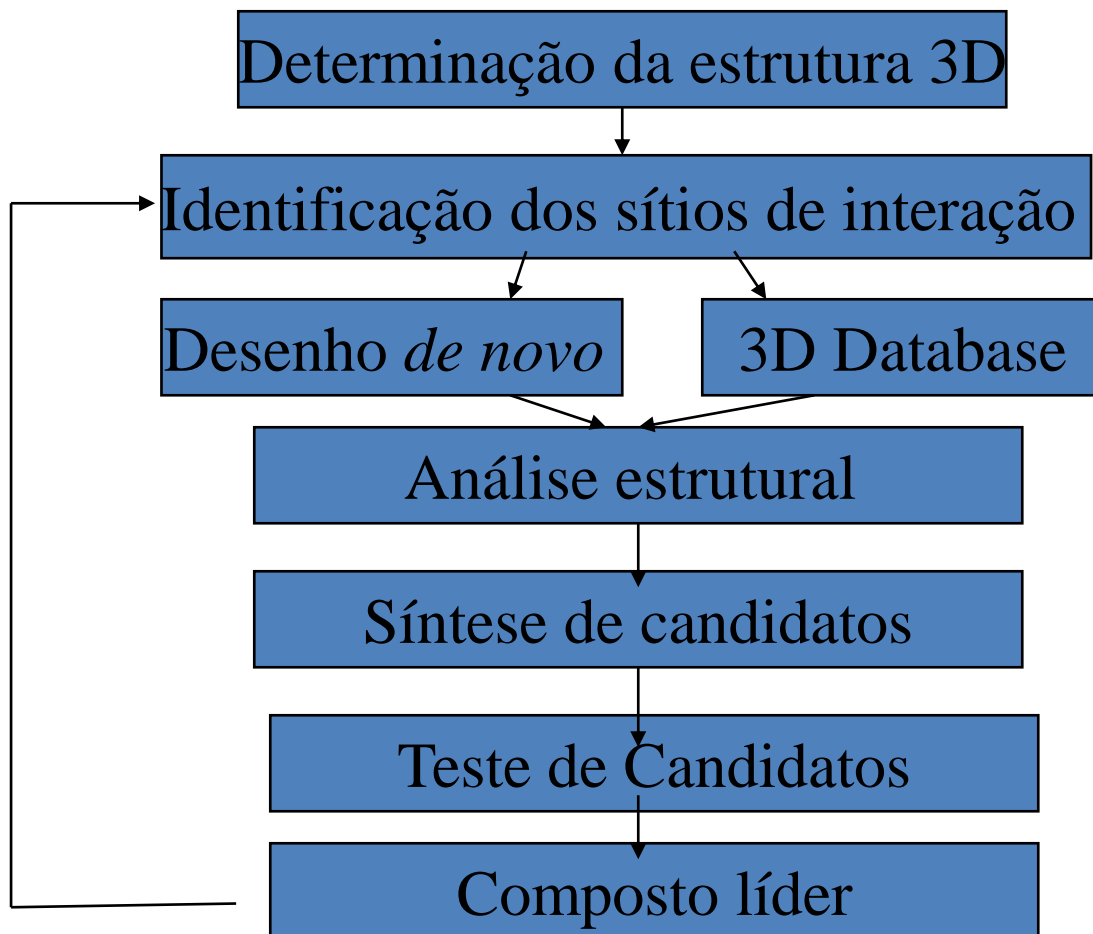
Modelo farmacofórico



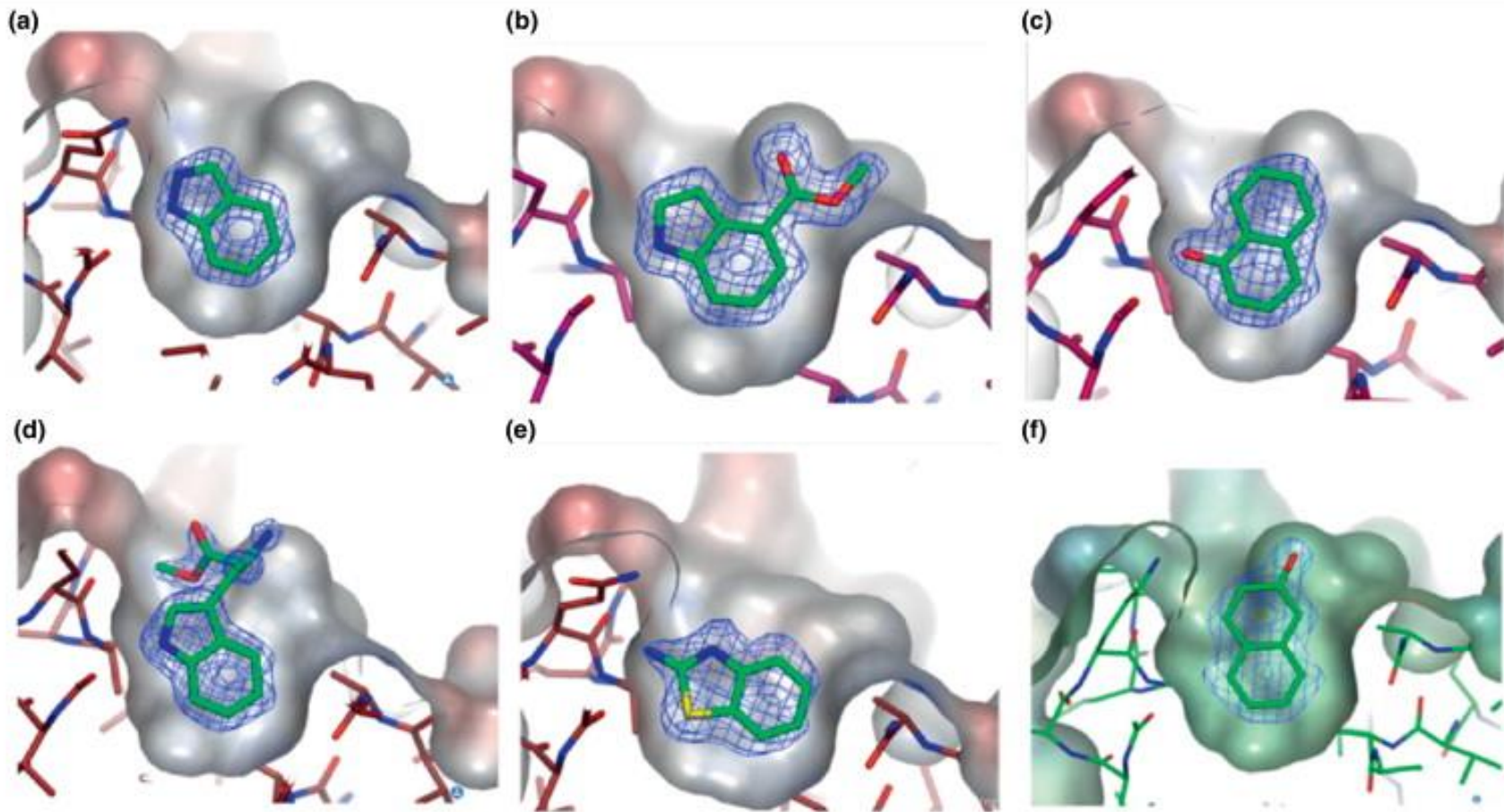
Dados de modo de ligação



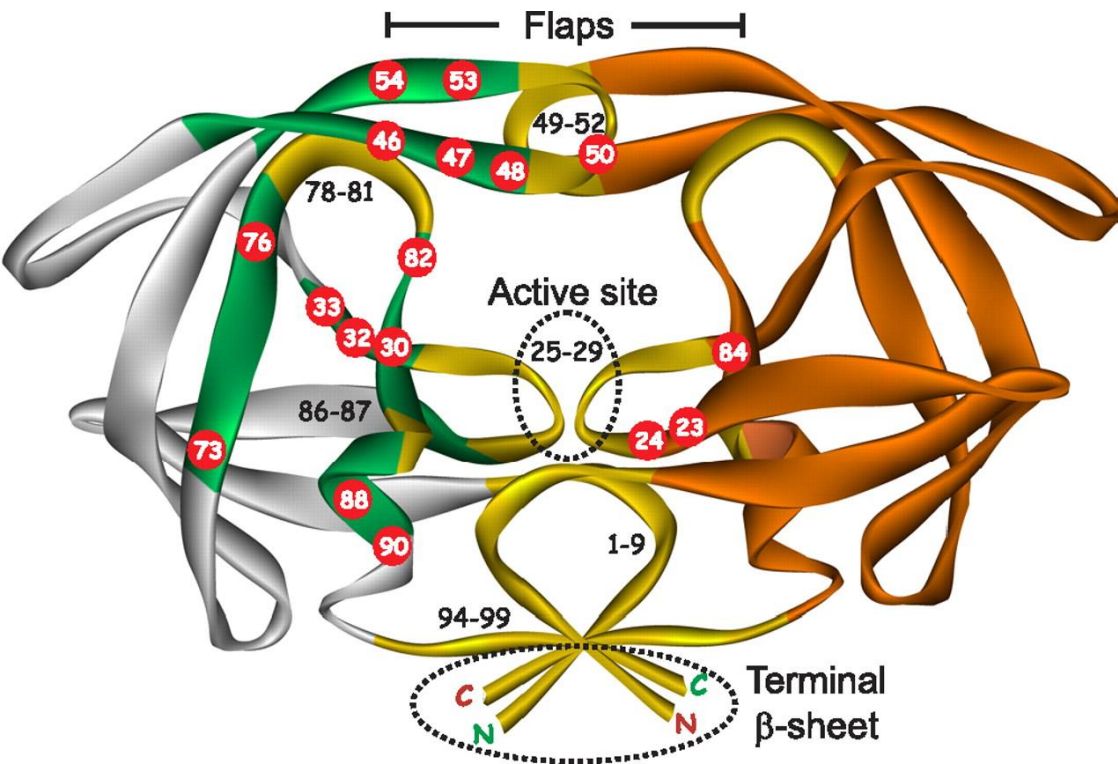
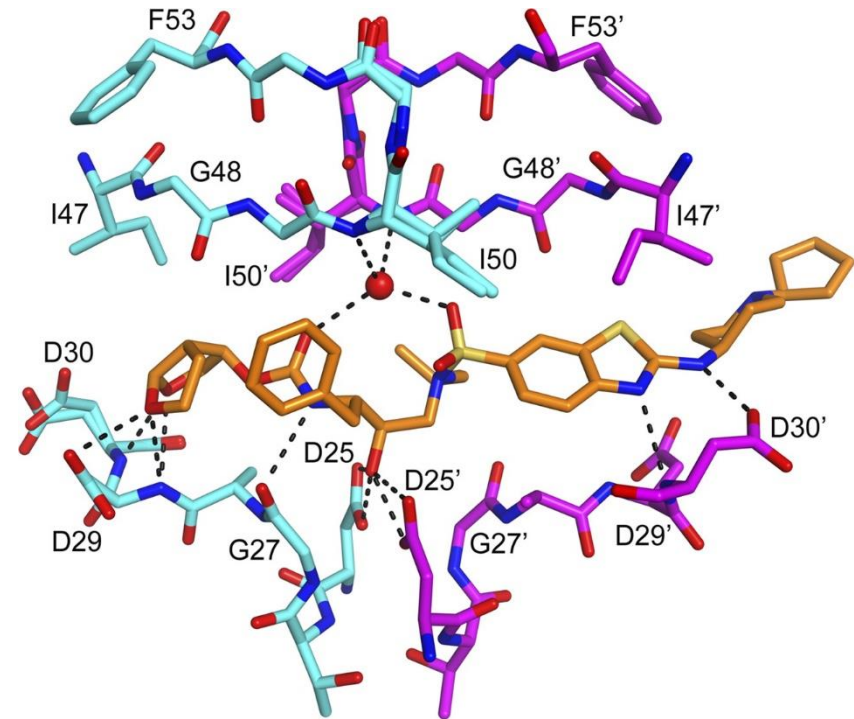
Planejamento de fármacos baseado em estrutura



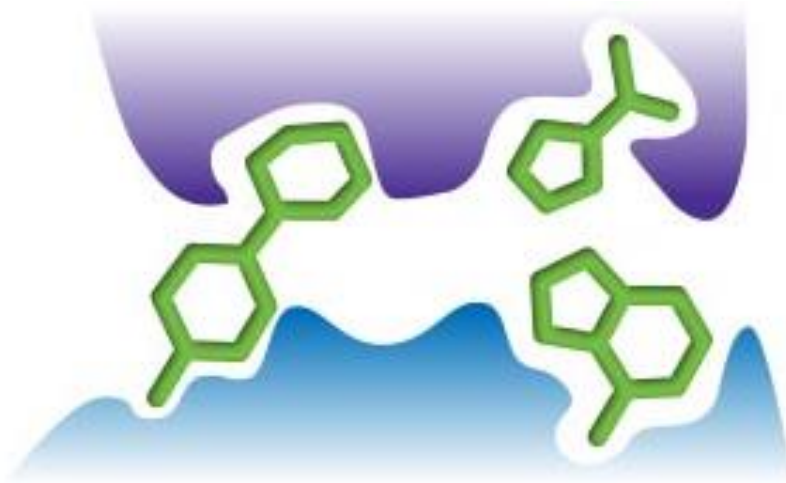
Descoberta ou desenho de moléculas que interagem com alvos bioquímicos de estrutura tridimensional conhecidas



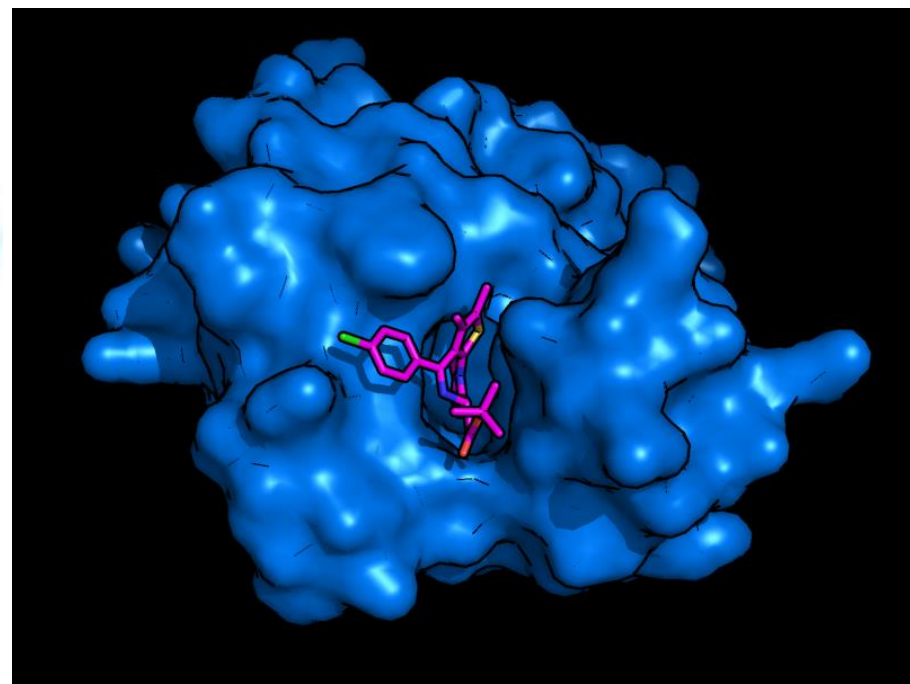
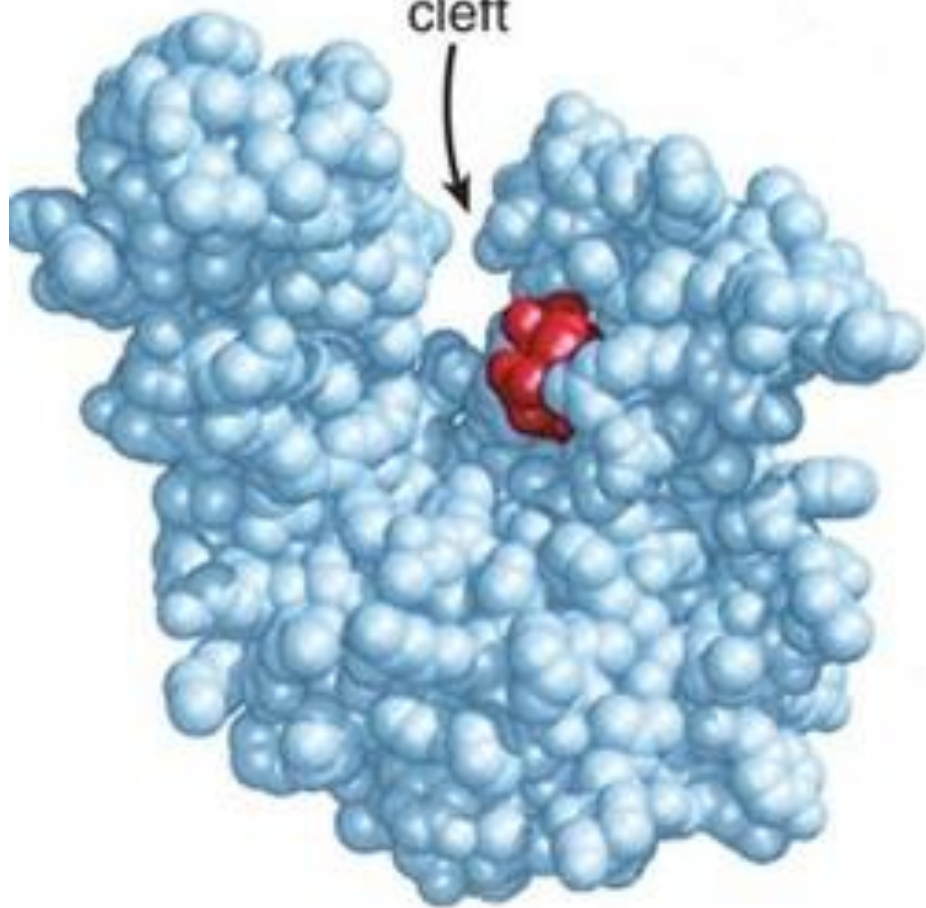
HIV protease – um exemplo clássico do processo de desenho de fármacos baseado em estrutura

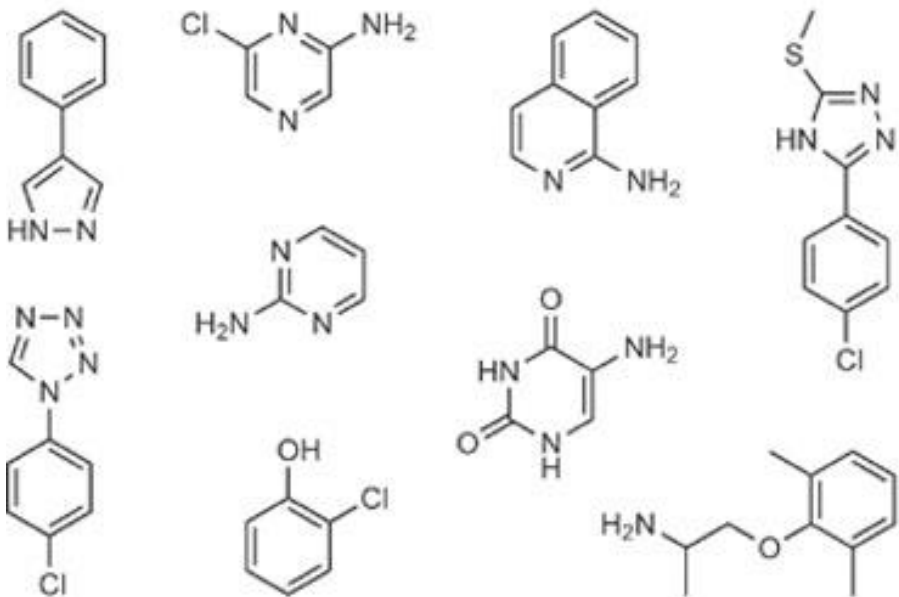


Um exemplo moderno da aplicação da estrutura de proteínas – planejamento de fármacos baseado em fragmentos

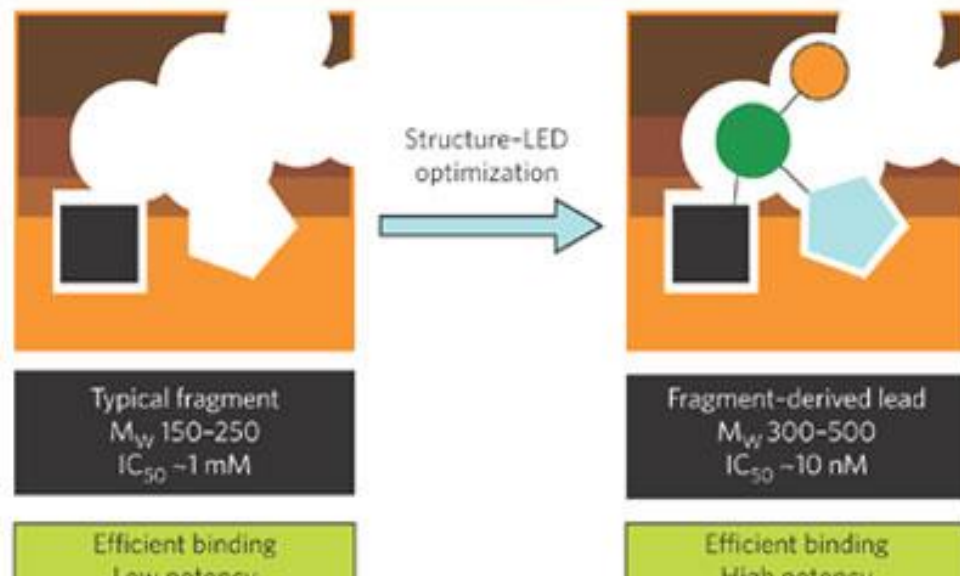


Active-site
cleft



a

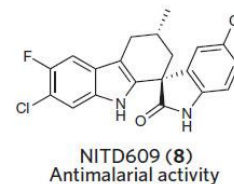
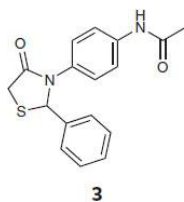
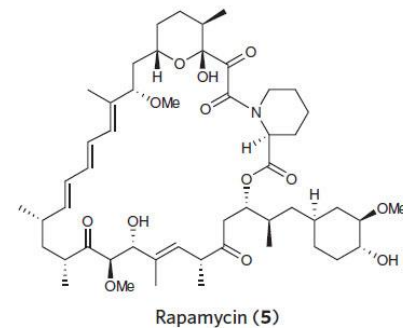
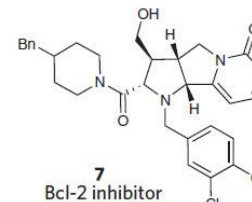
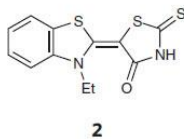
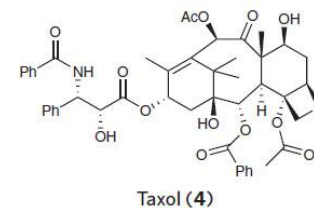
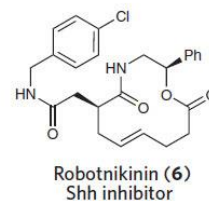
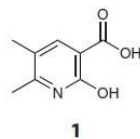
Fragment-based drug discovery



Typical compounds in CVLs:
low structural complexity

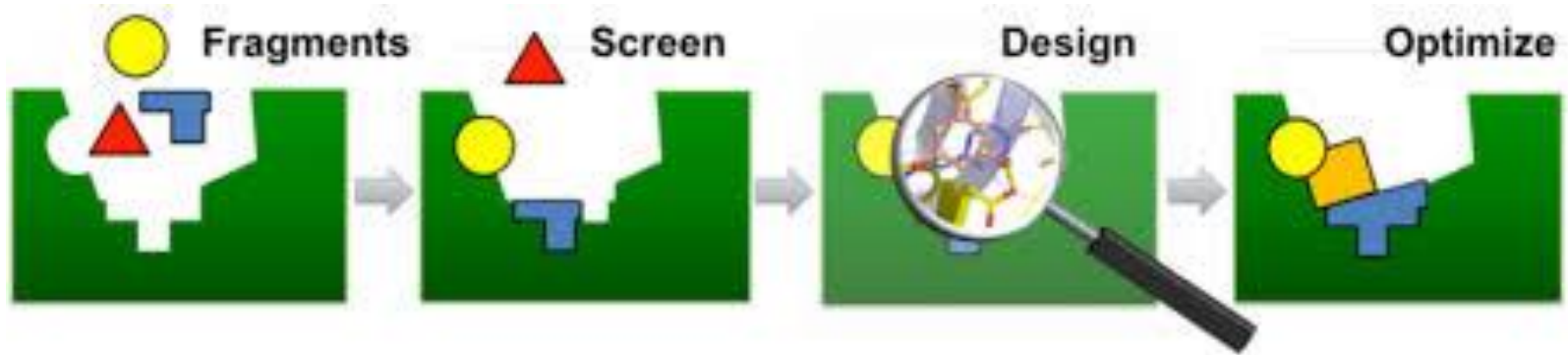
Examples of bioactive compounds:
intermediate structural complexity

Examples of NP-derived drugs:
high structural complexity



Increasing structural complexity

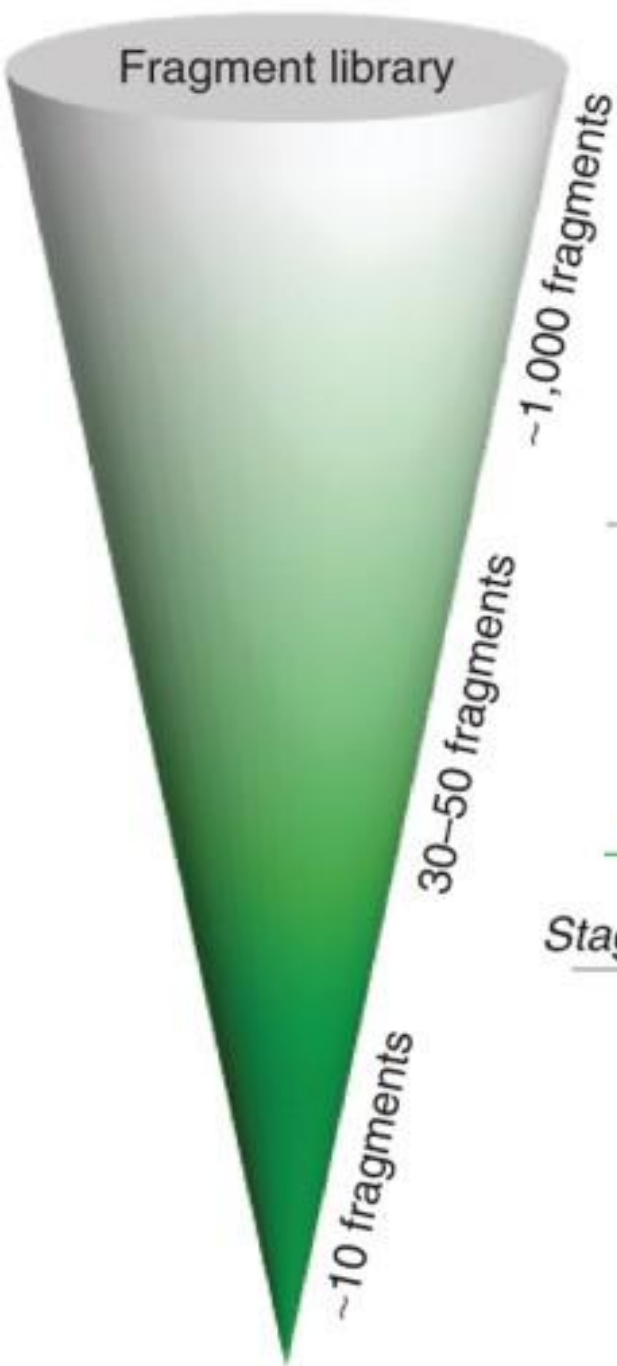
How we know which fragments are the best?



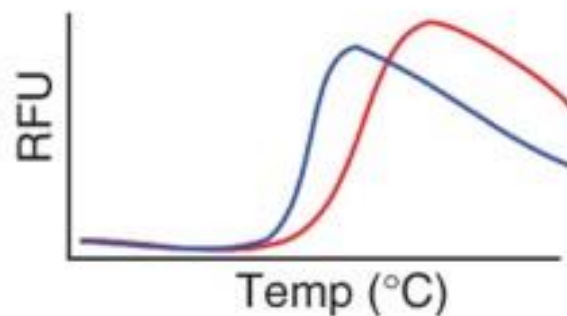
-1000 compounds (about 300 da)

How to screen?

-biophysical techniques, as NMR, MS, ITC, crystallography, TS, BLI, SPR

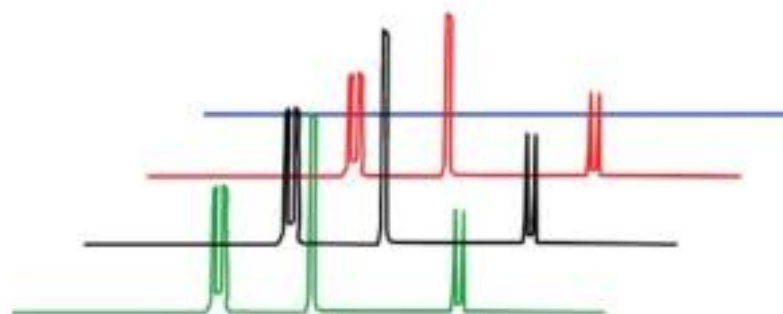


Stage 1: preliminary screening



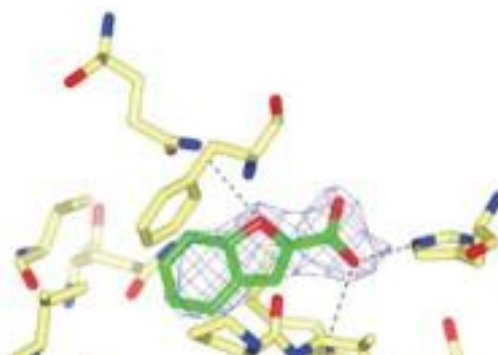
Differential scanning fluorimetry

Stage 2: validation



Ligand-observed NMR spectroscopy

Stage 3: characterization

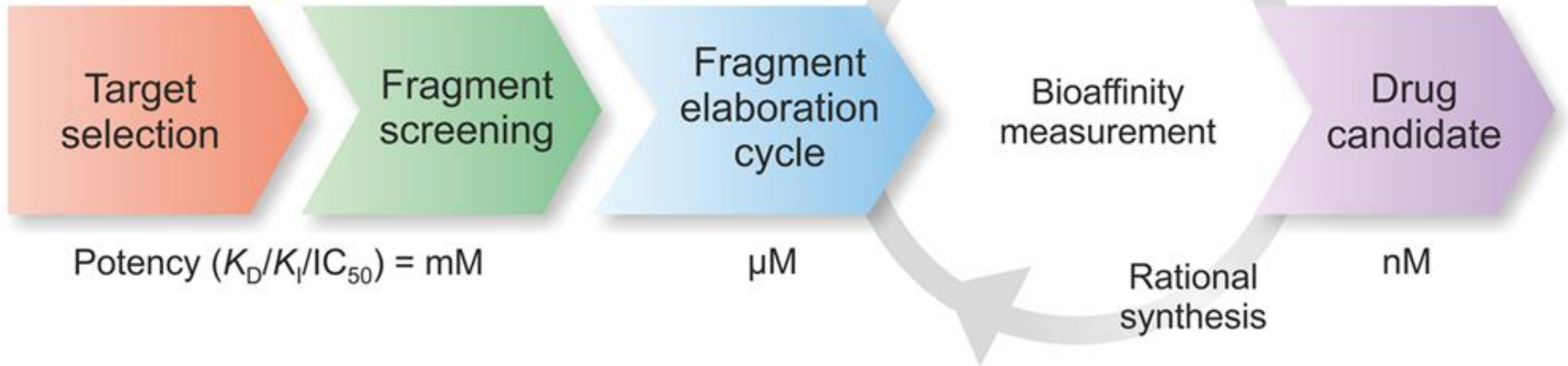


Isothermal titration calorimetry

X-ray crystallography

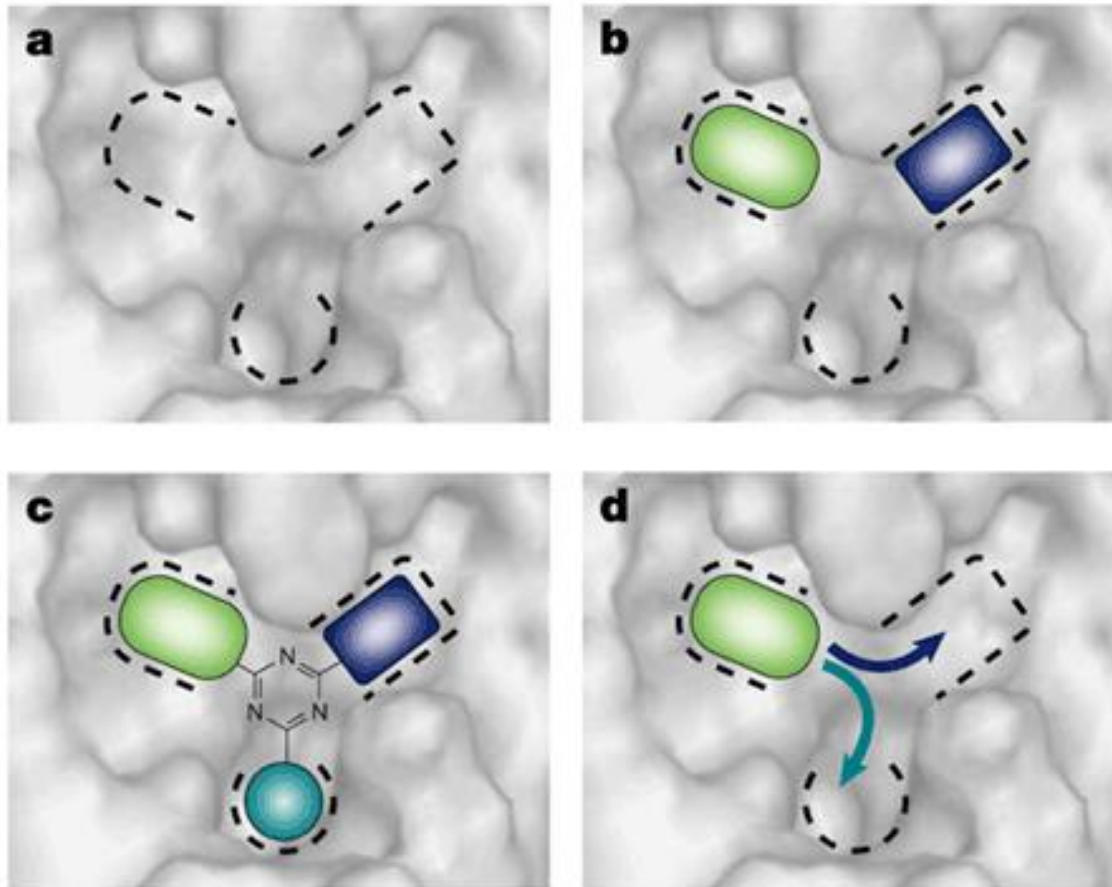
FBDD pipeline

Library design

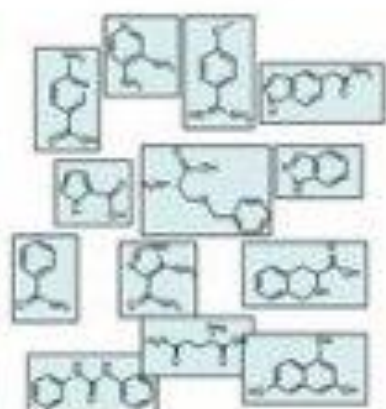


FBDD biophysical toolkit

Strategies of FBDD



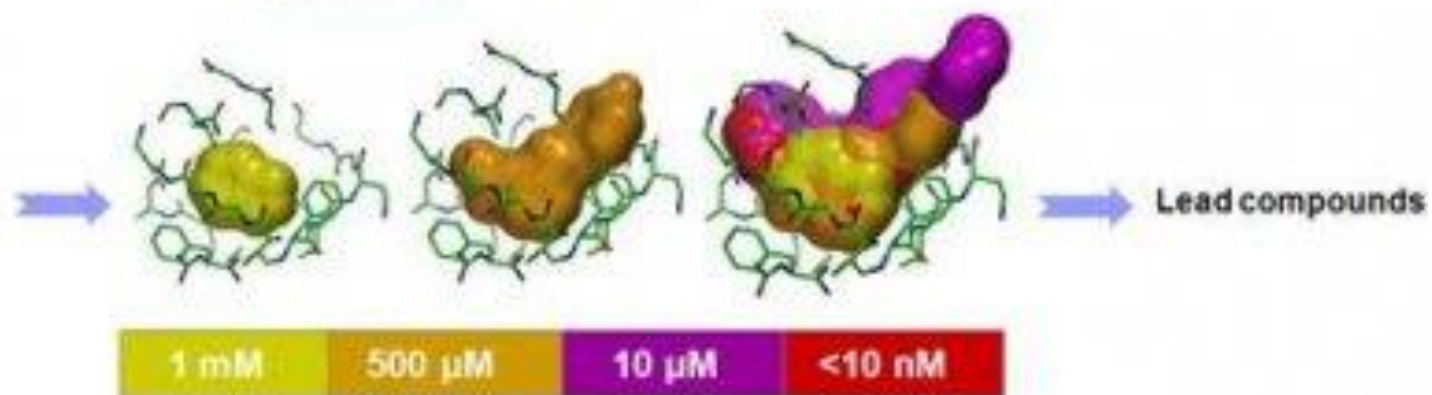
Fragment Hits

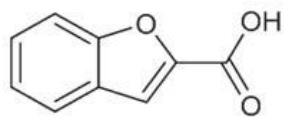


Parallel synthesis of analogues – SAR

Molecular modelling - docking

Biochemical and cellular validation

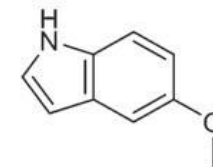
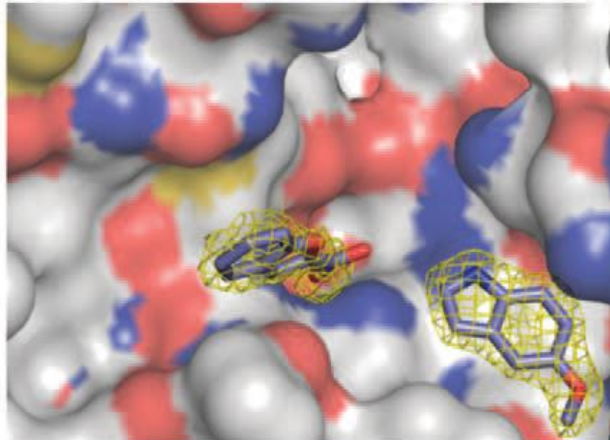




5

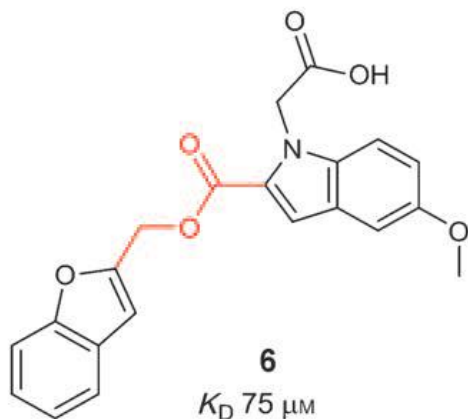
K_D 1000 μM
LE 0.34

a)



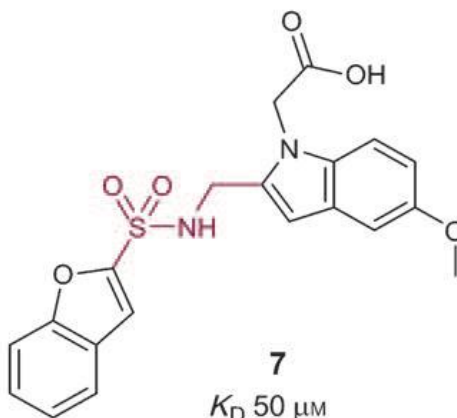
1

K_D 1100 μM
LE 0.36



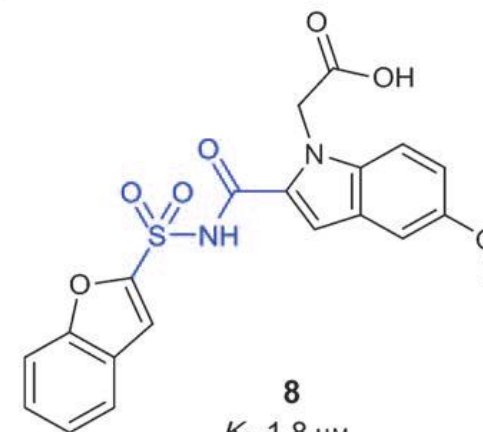
6

K_D 75 μM
LE 0.20



7

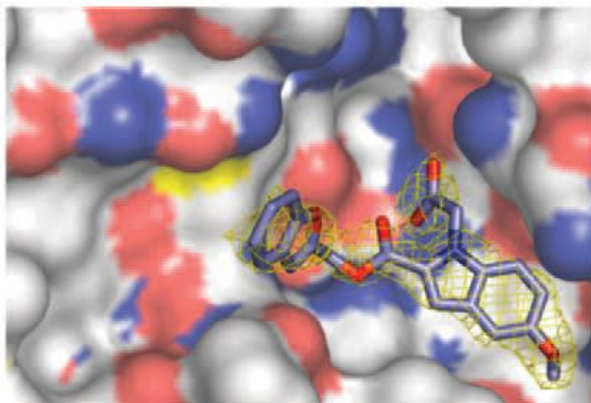
K_D 50 μM
LE 0.20



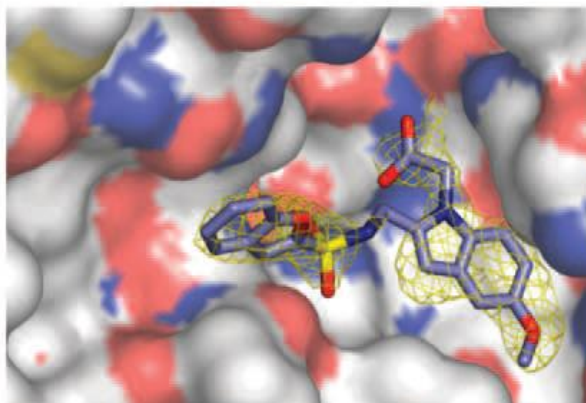
8

K_D 1.8 μM
LE 0.26

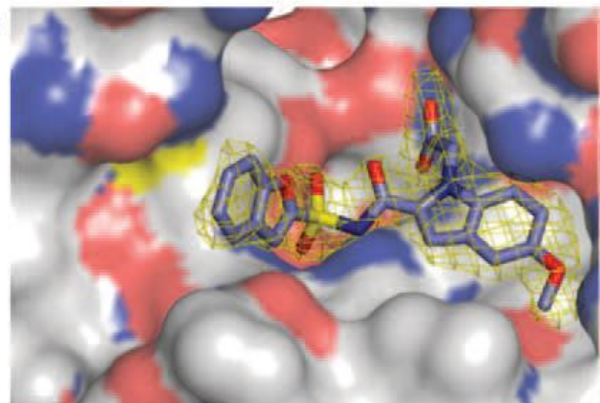
b)

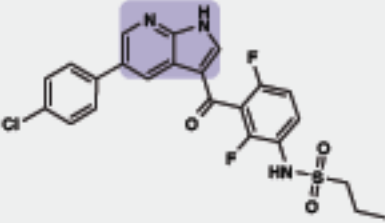
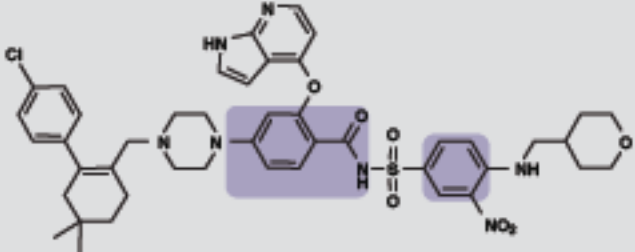
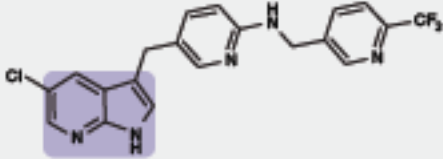
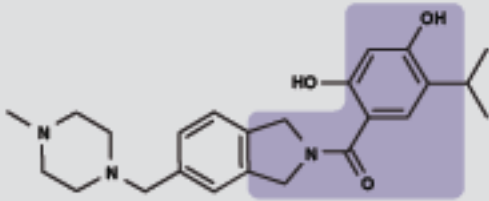


c)



d)




Structure*	Drug	Drug Target	Status
	Vemurafenib	BRAF-V600E	Approved
	Venetoclax	BCL-2	Approved
	Pexidartinib	FMS, KIT and FLT3-ITD	Phase III
	Onalespib	HSP90	Phase II

*shading indicates initial fragment

<https://www.technologynetworks.com/drug-discovery/articles/advances-in-fragment-based-drug-discovery-303059>

Review Article | Published: 15 July 2016


Twenty years on: the impact of fragments on drug discovery

Daniel A. Erlanson , Stephen W. Fesik, Roderick E. Hubbard, Wolfgang Jahnke & Harren Jhoti

Nature Reviews Drug Discovery **15**, 605–619(2016) | [Cite this article](#)

Perspective | Published: June 2009

The rise of fragment-based drug discovery

Christopher W. Murray & David C. Rees 

Nature Chemistry **1**, 187–192(2009) | [Cite this article](#)

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Biologia estrutural e biocatálise

Biocatálise é a aplicação de enzimas e microrganismos em química sintética, e usar esses biocatalisadores naturais para novas propostas: aplicação em que enzimas NÃO foram evoluídas para tal.



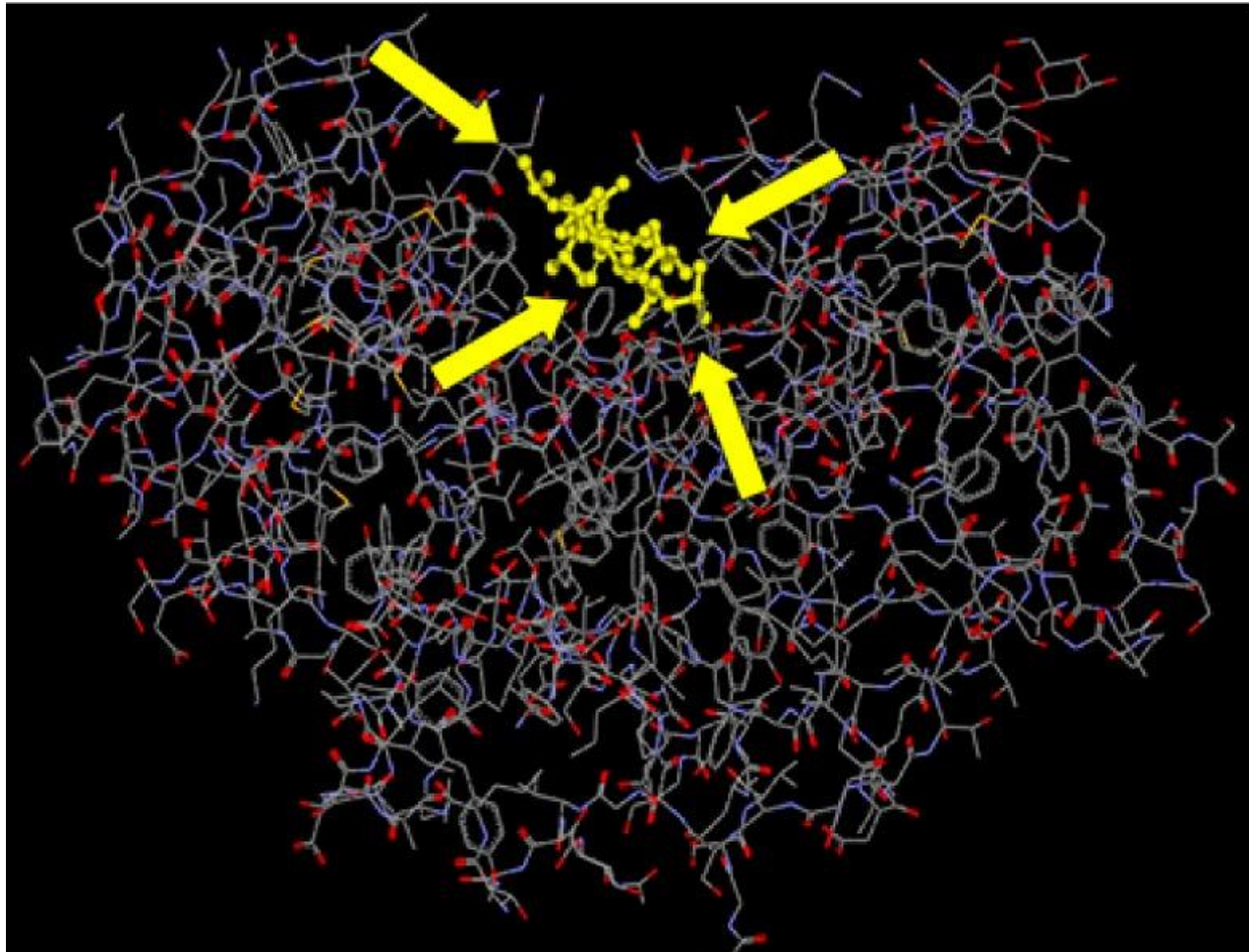
Francis Arnould – Pioneira em biocatálise
Prêmio Nobel em 2018

A biocatálise tem 2 objetivos principais:

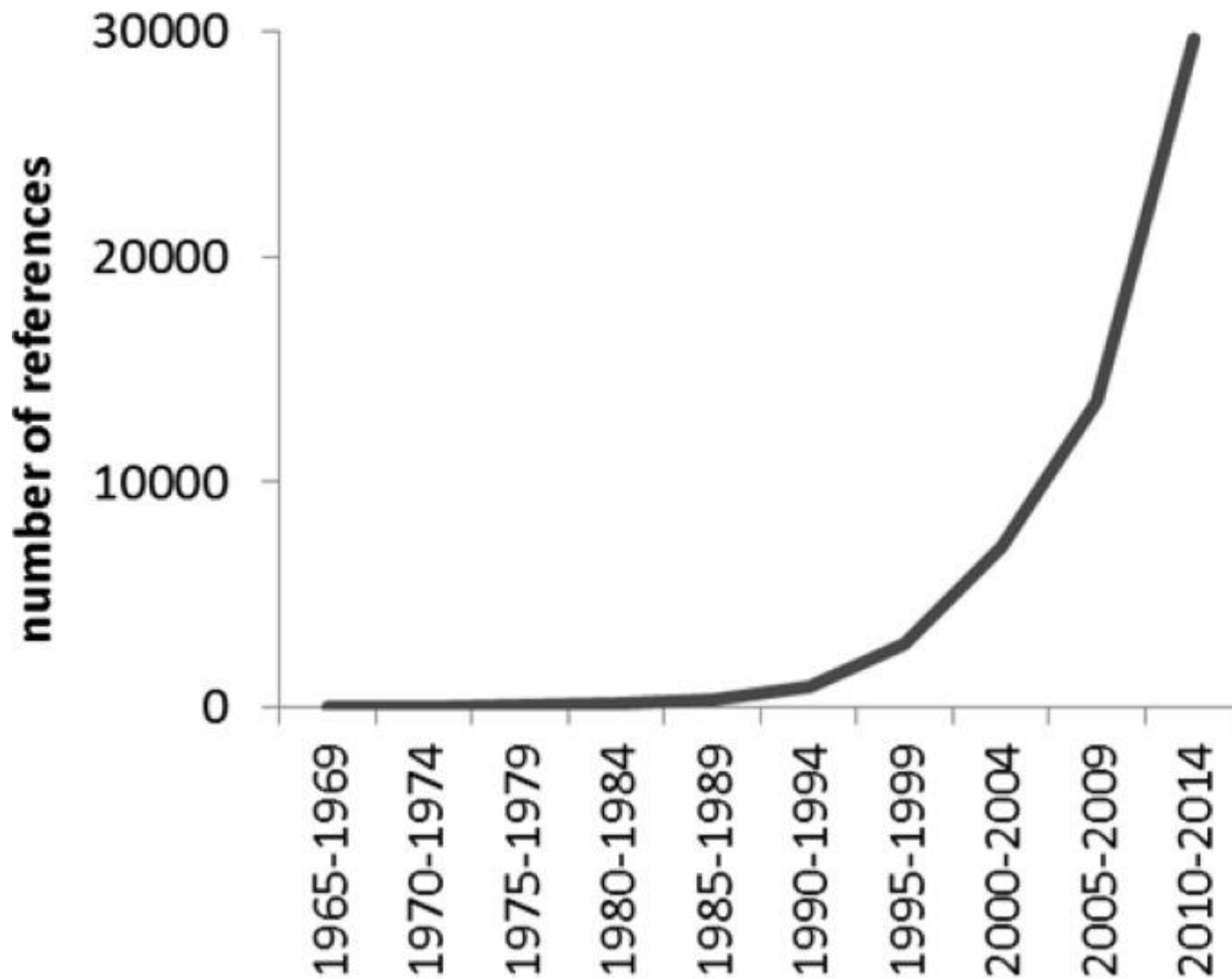
- **Acelerar reações químicas ou transformações;**
- **Utilizar da “química verde” diminuindo a quantidade de compostos tóxicos despejados no ambiente**



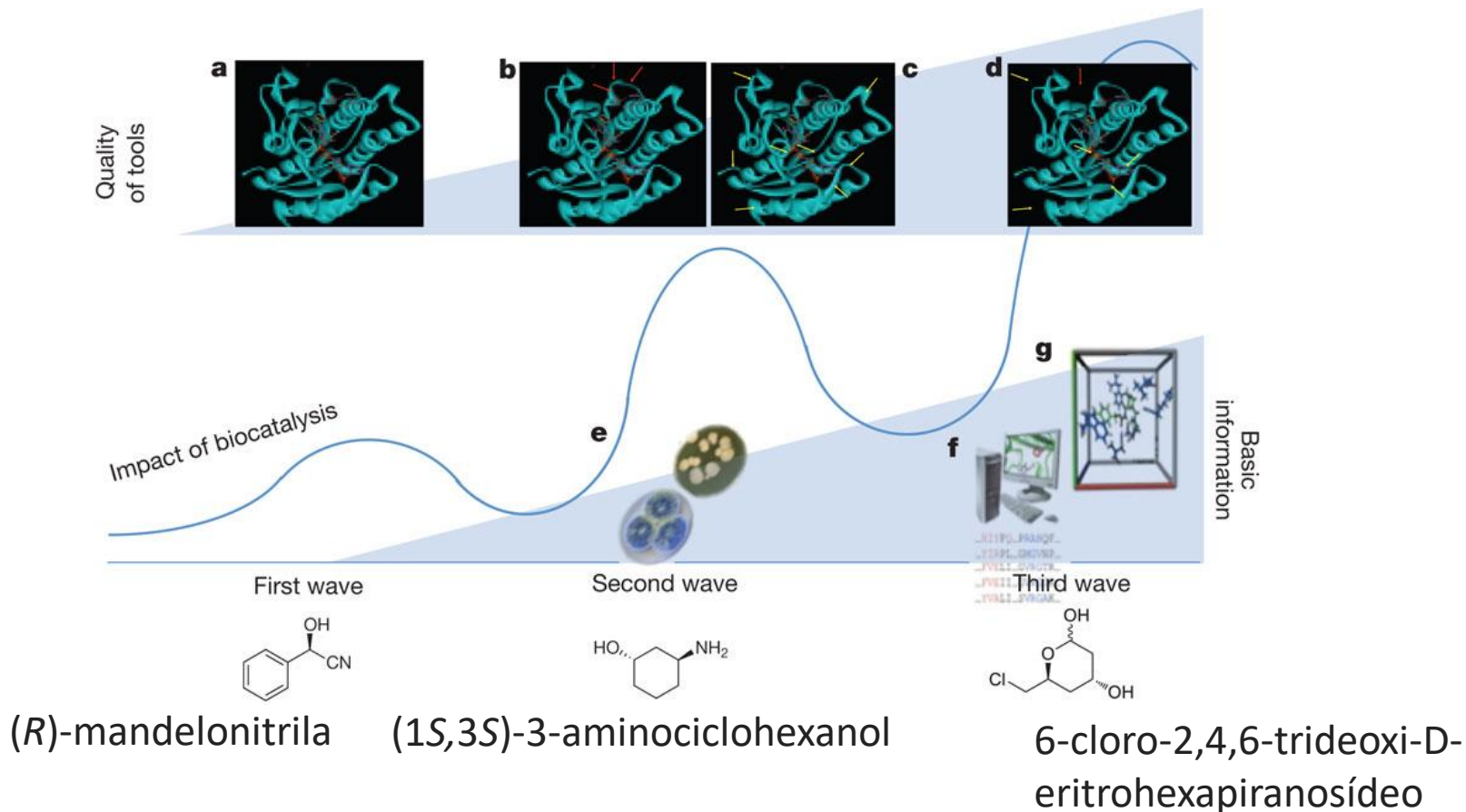
A biologia estrutural traz informação sobre os pontos de contato de substratos com a proteína



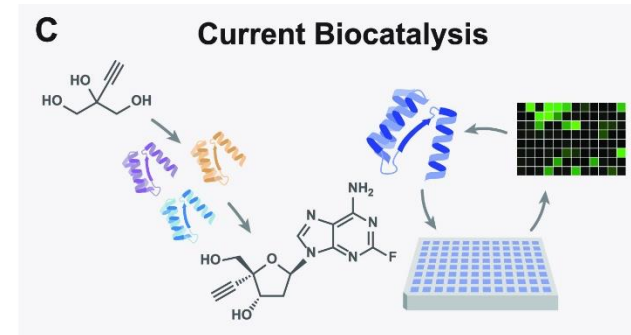
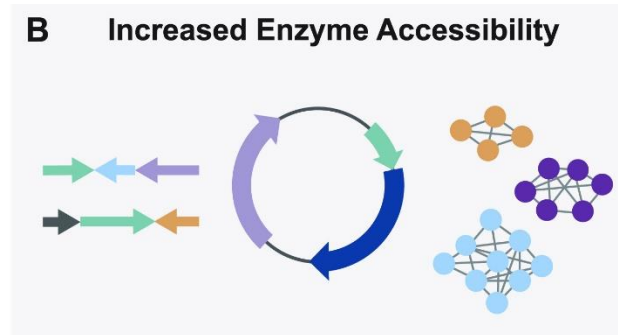
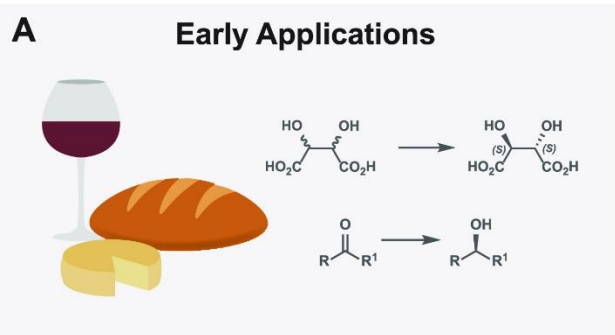
A biocatálise é um tópico de amplo interesse nos últimos anos pela indústria farmacêutica

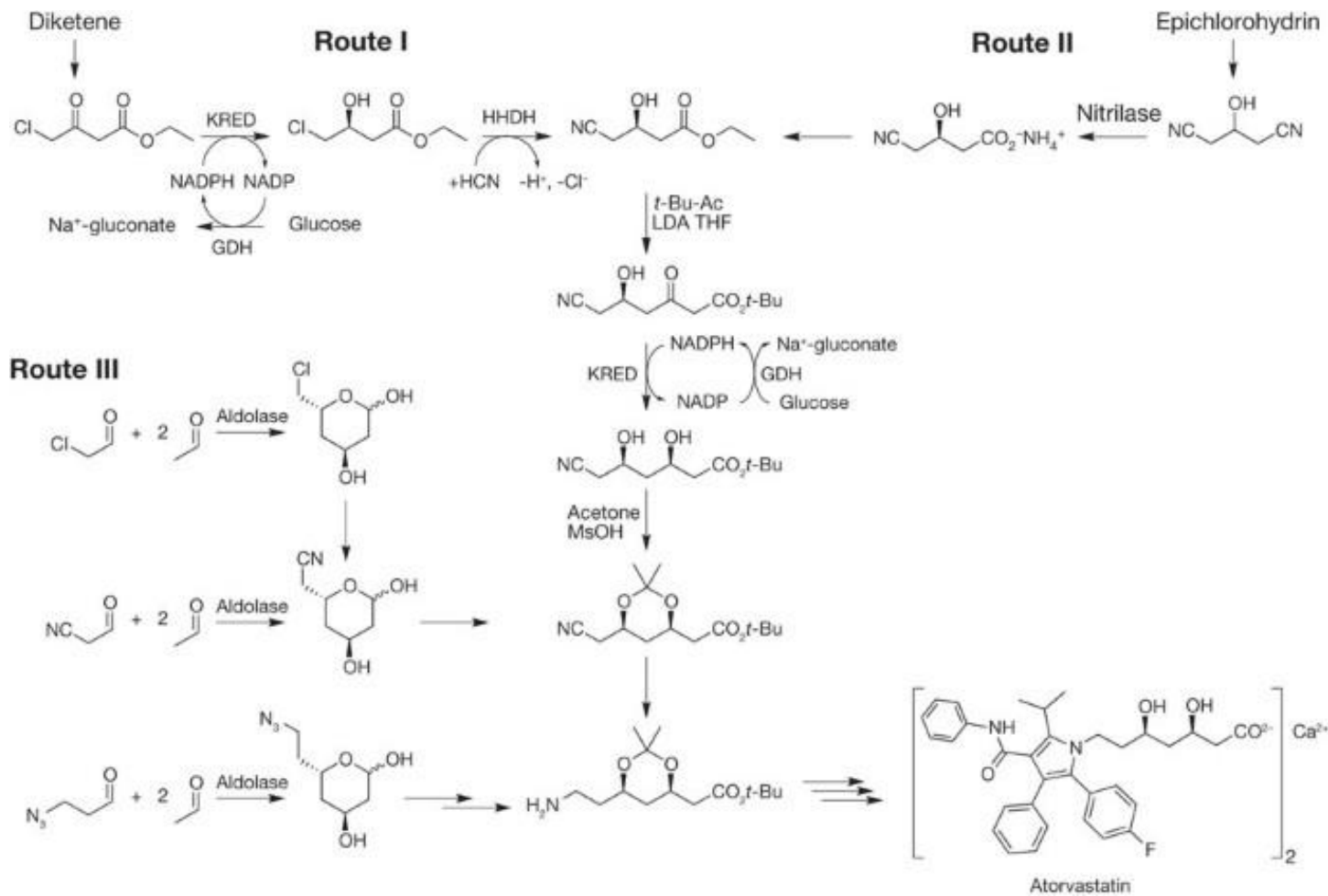


The evolution of enzyme discovery and protein engineering strategies used to identify desired catalysts.



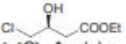
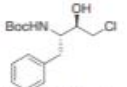
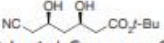
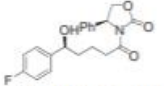


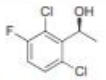
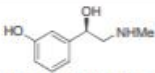
Evolução da aplicação da biocatálise





Alguns exemplos da aplicação de ceto-oxidoredutases na indústria

Table 1 | Recently developed biocatalytic processes in the pharmaceutical industry

Product	Technology (company, reference)	Product	Technology (company, reference)
KREDs			
 Ethyl (S)-4-chloro-3-hydroxybutanoate (1)	Various protein engineering approaches, including directed evolution, to increase activity, stability and coenzyme specificity. Initial intermediate for atorvastatin (Lipitor) manufacture on the industrial scale. (Kaneka ^{B1} , Codexis ^{B2})	 Tert-butyl (2S,3R)-4-chloro-3-hydroxy-1-phenylbutan-2-yl-carbamate (10)	Recombinant expression and also protein engineering for activity. Intermediate for atazanavir (Reyataz). (Bristol Myers Squibb ^{B2} ; Codexis, WO/2011/005527)
 Tert-butyl 6-cyano-3,5-dihydroxyhexanoate (2)	Protein engineering for activity and stability to 30% organic substrate (the hydroxyketone substrate and a tert-butyl acetoacetate impurity, both liquids). Intermediate for atorvastatin. (Codexis, US/2011/7879585; Pfizer, US/2008/0118962)	 (4S)-3-[(5S)-5-(4-fluorophenyl)-5-hydroxy-pentanoyl]-4-phenyl-1,3-oxazolidin-2-one (11)	Protein engineering for activity and solvent stability. Intermediate for ezetimibe (Zetia, Vytorin). (Codexis, WO/2010/025085)
 (R)-3-hydroxythiolane (R3HT) (3)	Directed evolution primarily for enantioselectivity. Intermediate for sulopenem-type antibiotics (since discontinued). (Codexis ^{B3} , WO/2009/029554)	 D-pseudoephedrine (12)	Protein engineering to overcome product inhibition. The ketone substrate is racemic at C ₂ and racemizes during the reaction to give high-e.e. product. The product of the reaction is a generic API. (Daiichi ^{B4})
 (S)-1-(2,6-dichloro-3-fluorophenyl)-ethanol (DCFPE) (4)	Directed evolution of a variant identified from the R3HT (3) project for increased activity. Note that the ketone is hindered owing to the ortho chloro-substituents. Raw material for crizotinib (Xalkori). (Codexis, WO/2009/036404)	 (R)-phenylephrine (13)	Protein engineering for activity and tolerance to aminoketone substrate. The product of the reaction is a generic API. (Codexis, WO/2011/022548)

Engineering the third wave of biocatalysis

U. T. Bornscheuer¹, G. W. Huisman², R. J. Kazlauskas^{3,4}, S. Lutz⁵, J. C. Moore⁶ & K. Robins⁷

Over the past ten years, scientific and technological advances have established biocatalysis as a practical and environmentally friendly alternative to traditional metallo- and organocatalysis in chemical synthesis, both in the laboratory and on an industrial scale. Key advances in DNA sequencing and gene synthesis are at the base of tremendous progress in tailoring biocatalysts by protein engineering and design, and the ability to reorganize enzymes into new biosynthetic pathways. To highlight these achievements, here we discuss applications of protein-engineered biocatalysts ranging from commodity chemicals to advanced pharmaceutical intermediates that use enzyme catalysis as a key step.

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Outlook

State-of-the-Art Biocatalysis

Joshua B. Pyser, Suman Chakrabarty, Evan O. Romero, and Alison R. H. Narayan*



Cite This: *ACS Cent. Sci.* 2021, 7, 1105–1116



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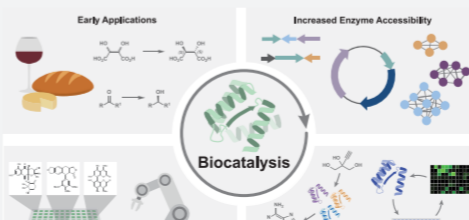


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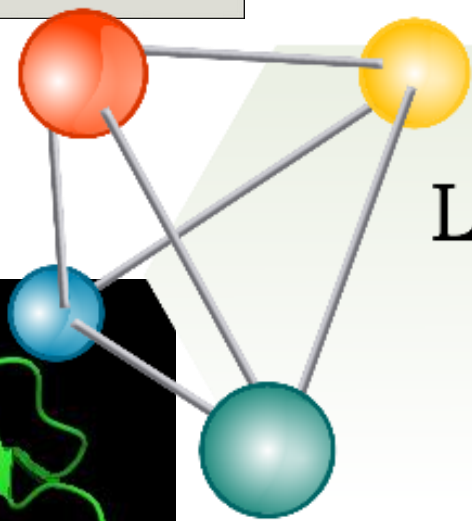
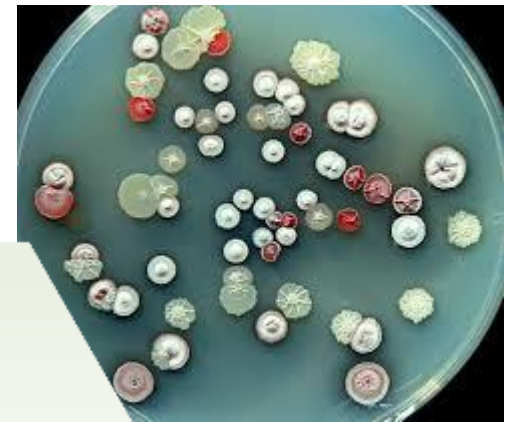
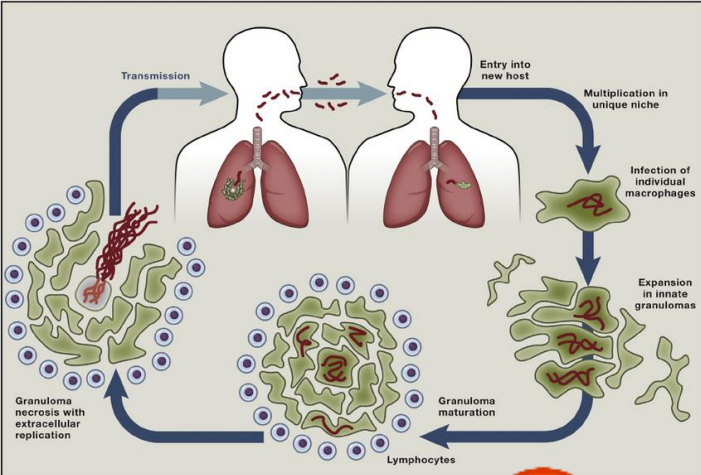
ABSTRACT: The use of enzyme-mediated reactions has transcended ancient food production to the laboratory synthesis of complex molecules. This evolution has been accelerated by developments in sequencing and DNA synthesis technology, bioinformatic and protein engineering tools, and the increasingly interdisciplinary nature of scientific research. Biocatalysis has become an indispensable tool applied in academic and industrial spheres, enabling synthetic strategies that leverage the exquisite selectivity of enzymes to access target molecules. In this Outlook, we outline the technological advances that have led to the field's



1980s and 1990s, initial
are based, extended the
is of unusual synthetic
to the manufacture of
Examples include the
synthesis of diltiazem

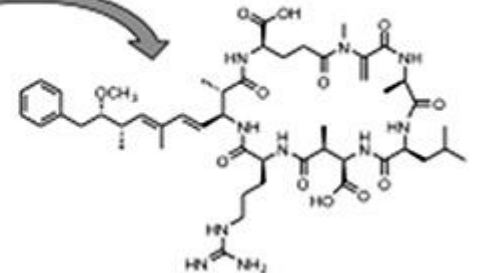
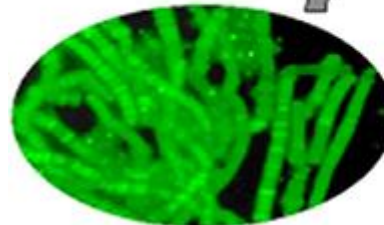
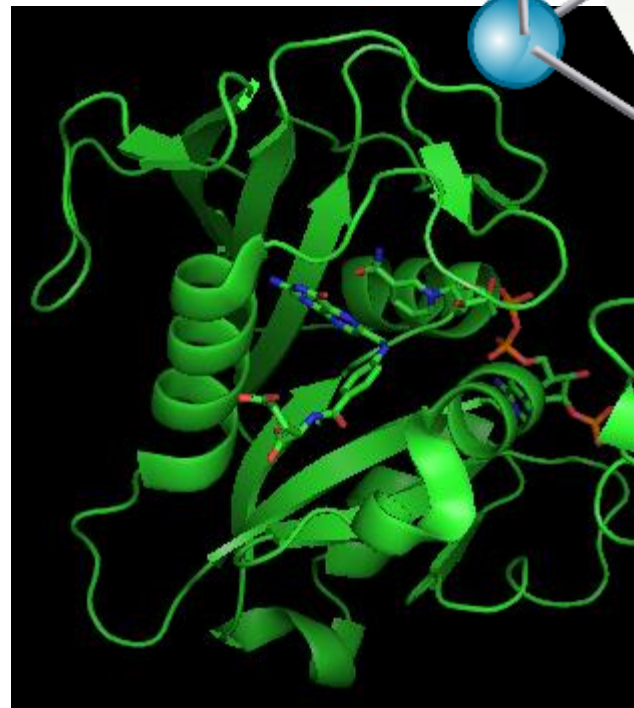
Consideração final sobre o uso da biologia estrutural:

- **Desenho de fármacos – aumentar pontos de interação para ganho de afinidade e inibição**
- **Biocatálise – aumentar pontos de interação ou promiscuidade para aumentar eficiência de catálise ou atividade contra substratos alternativos.**



Laboratório de Biologia Estrutural Aplicada

Divisão Fármacos



Exercício:

1) COVID-19 mudou drasticamente nosso modo de vida nos últimos 18 meses, levando à morte de um grande número de pessoas. Embora hoje tenhamos vacina, ainda há poucas moléculas que estão em estudos clínicos para o tratamento desta doença. Assim, a biologia estrutural tem sido extremamente importante neste momento e em um ano e meio de pandemia, o número de estruturas para proteínas depositados no “Protein Data Bank” é bastante expressivo, incluindo na presença de potenciais inibidores. Vocês, como futuros biomédicos, podem trabalhar na interpretação de resultados estruturais. Assim, entrem no PDB, código de acesso **7D1M** e analise as interações do composto CG376 com a *main protease de SARS-CoV-2*, indicando quais são essas interações (hidrofóbicas e interações de hidrogênio) que garantem a afinidade desse composto. Tente encontrar pelo menos uma modificação no composto que poderia aumentar sua afinidade pela proteína.

Para facilitar: Entre em “<https://www.rcsb.org/>” digite **7D1M** (em procura)/abaixo da estrutura vai em **3D View/ligand interactions/components/polimer/add representation/line|label/** Observe os amino ácidos que interagem com o ligante.

1) Biocatálise tem se tornado uma importante abordagem para a síntese de novos compostos utilizando técnicas menos agressivas ao meio ambiente (química verde) e mais baratas. Portanto para que uma enzima possa ser utilizada nessas técnicas, modificações são necessárias. Baseado no que se sabe sobre as proteínas de organismos termofílicos, que modificações poderiam ser propostas para aumentar a estabilidade de proteínas a altas temperaturas? Como a biologia estrutural poderia auxiliar nesse processo para uma enzima que não apresenta similaridade com outras já resolvidas?