

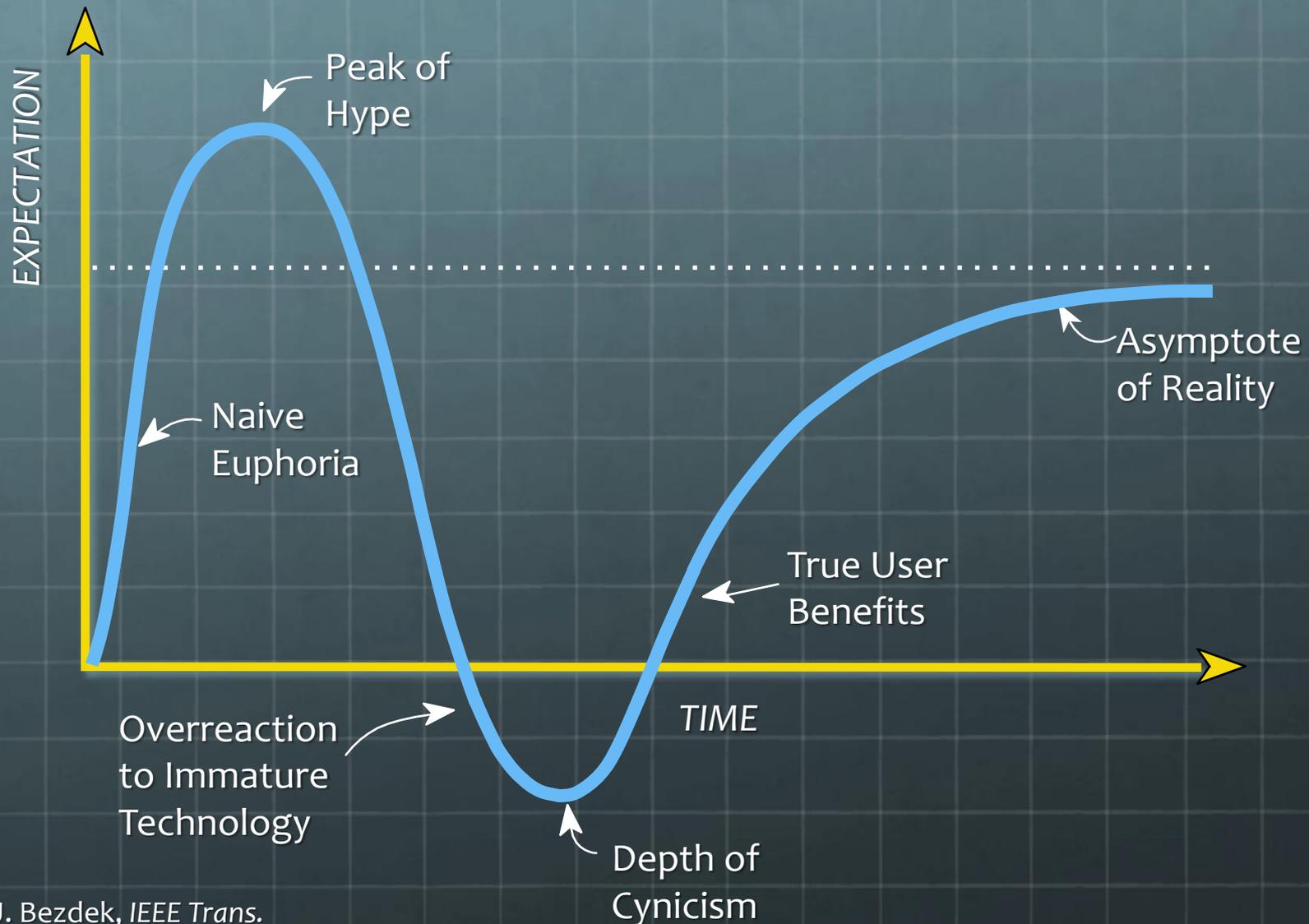
*We don't know a millionth of
a percent about anything.*

-- Thomas Alva Edison



Daumier, "Monsieur Babinet"

Technology Cycle



\$2.50

October 5, 1981

FORTUNE



THE NEXT INDUSTRIAL REVOLUTION

Designing drugs by computer at Merck

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Part 1: SBDD Primer

Targets Which Have Yielded Clinical Candidates With the Help of Structure-Based Drug Design

Therapeutic Area

Targets

Cardiovascular

ACE, Renin, Thrombin, Factor VII, Factor Xa

Glaucoma

Carbonic anhydrase

Inflam / immun

Human neutrophil elastase, P38, IMPDH, ICE, COX2, MMP-X, JAK3

Cancer

Purine nucleoside phosphorylase, Thymidylate synthase, VEGF kinase (KDR), Aurora-2, CDK2, EGF kinase (erbB), Glycinamide ribonucleotide formyl-transferase, HSP90, BTK,

Antivirals

HIV protease, Influenza sialidase (neuraminidase), HCV protease, HCV polymerase, rhinovirus 3C protease, rhinovirus coat proteins

Sepsis

Caspases (broad), secretory PLA2

Diabetes

PPAR-gamma, DPP-IV, Aldose reductase

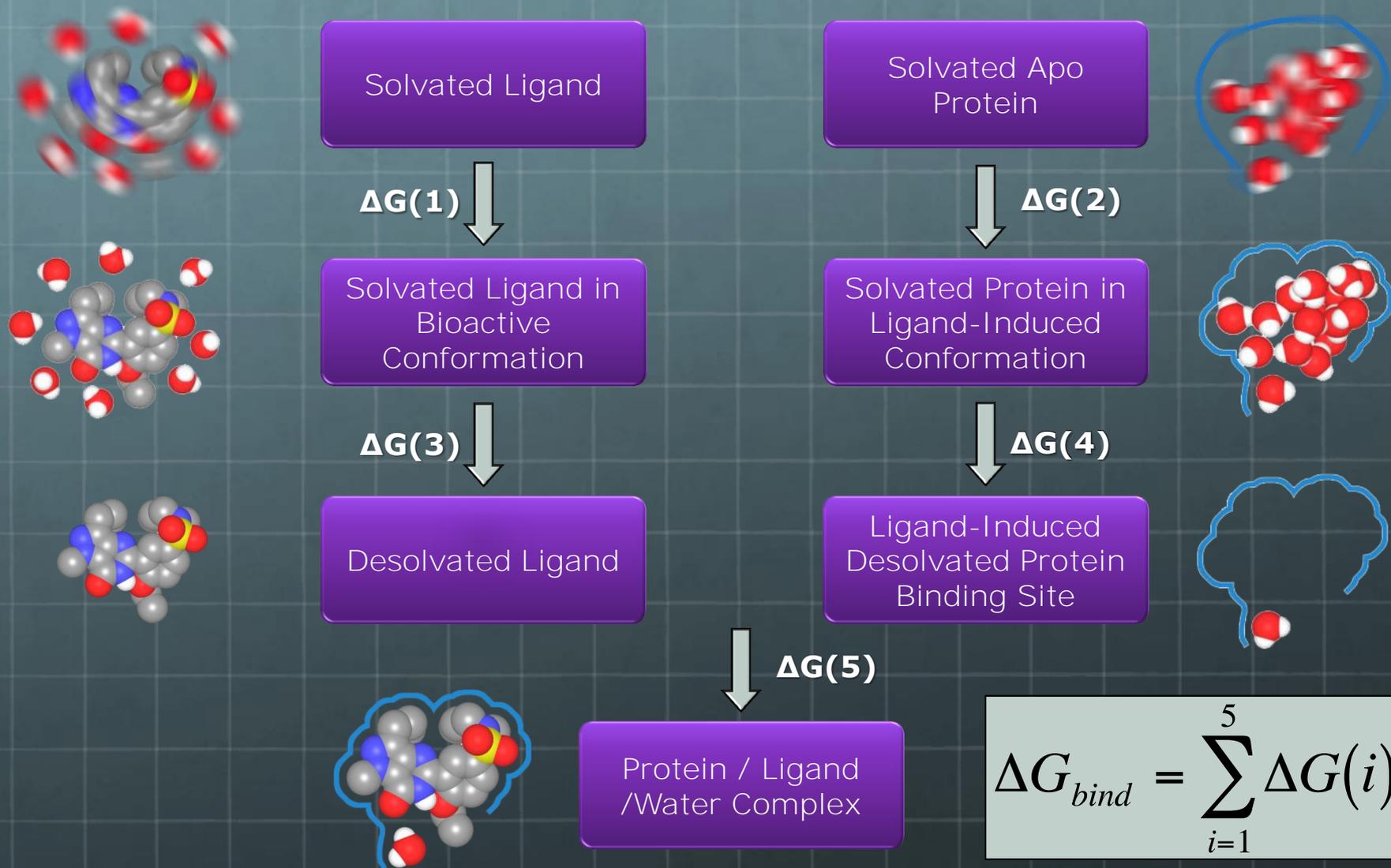
Osteoporosis

Cathepsin K

Various CNS

GSK3 kinase, Acetylcholinesterase, BACE

Thermodynamic Decomposition of Ligand/Protein Binding



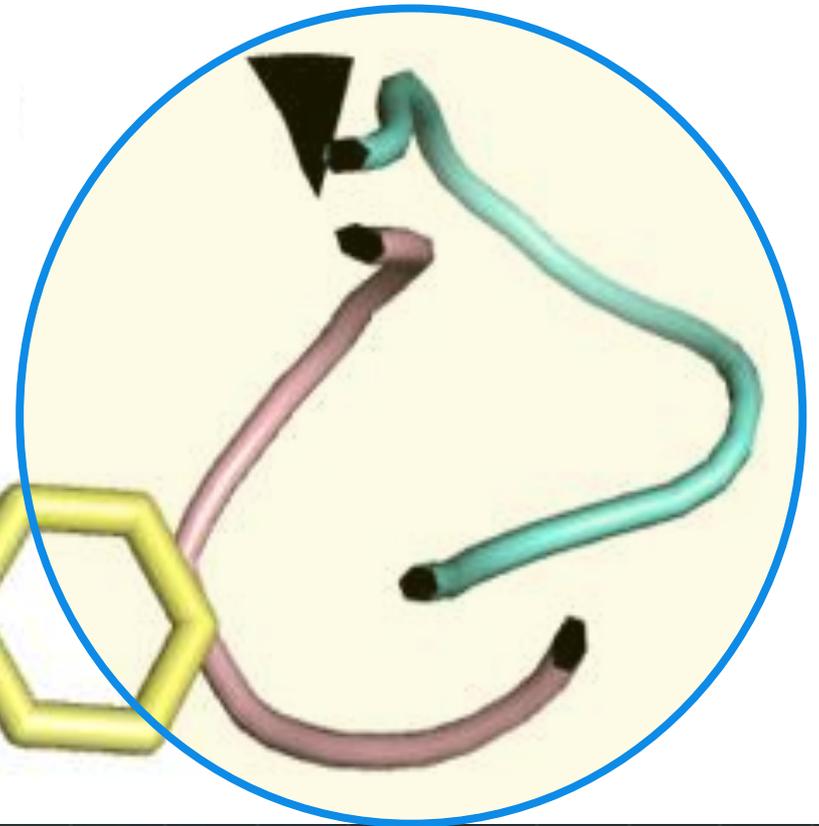
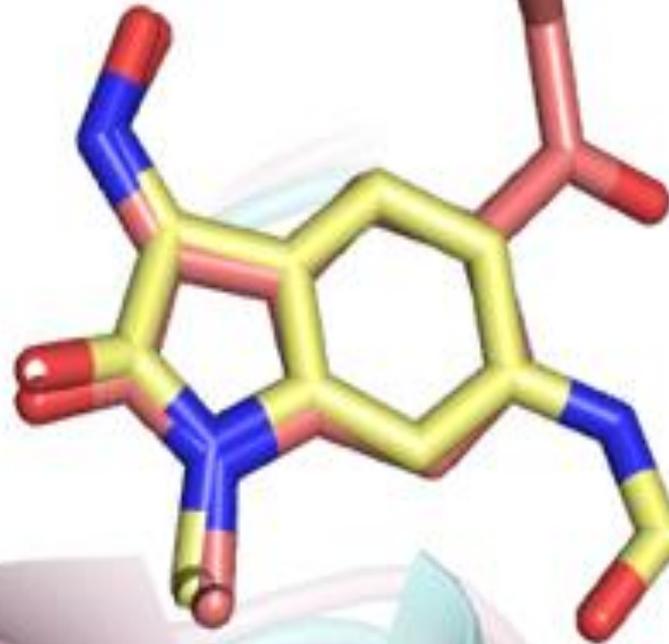
$$\Delta G_{bind} = \sum_{i=1}^5 \Delta G(i)$$

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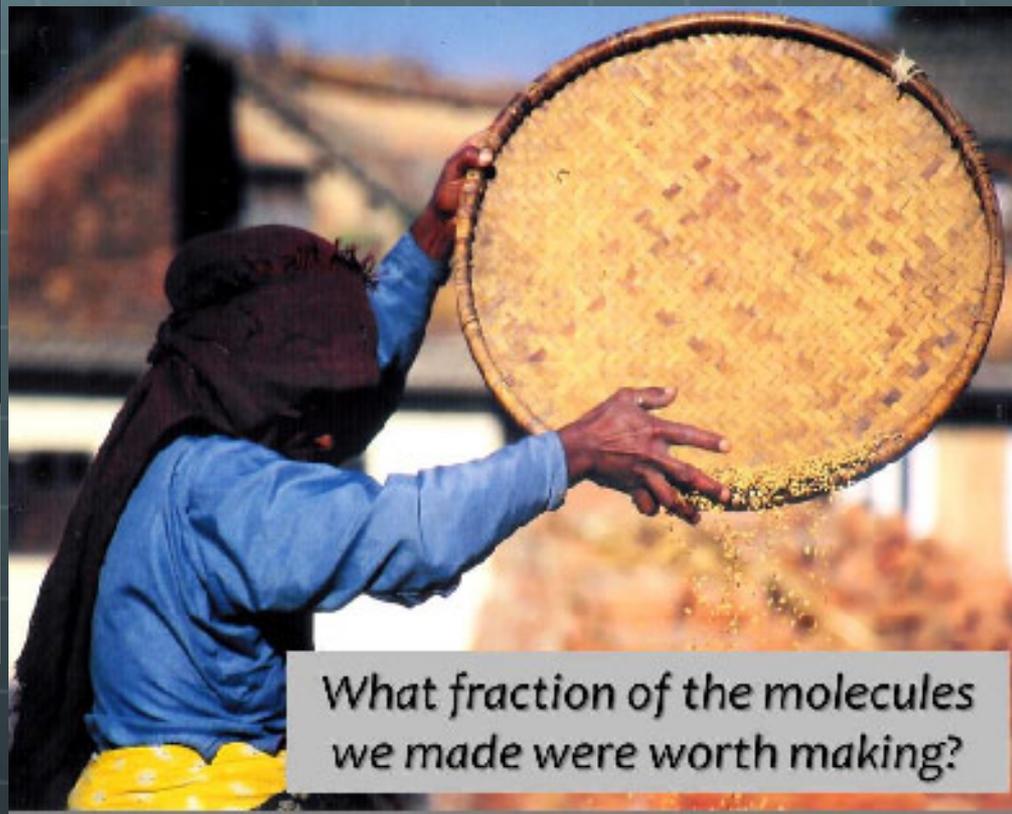
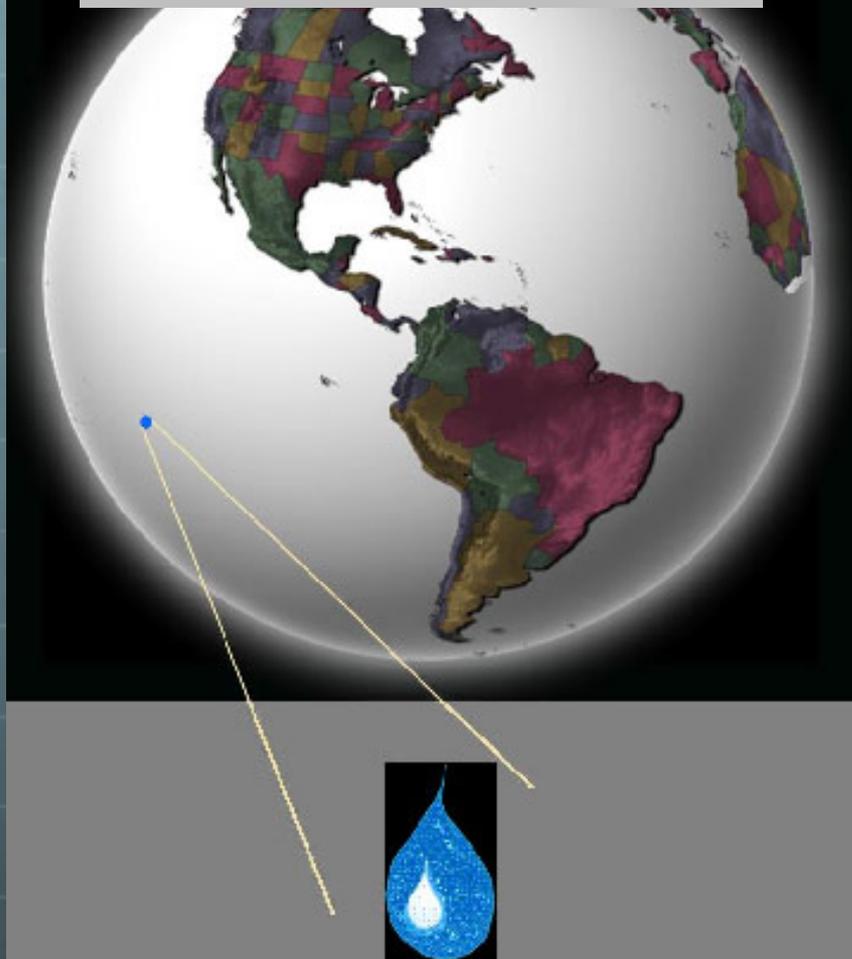
Proteins are Dynamic

Lots of kinase examples - proteins suddenly adopt different conformations and the SAR goes right out the window.

No simple model will ever get this correct.



What fraction of the possible molecules have we made?



What fraction of the molecules we made were worth making?

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SBDD Toolkit

Simple

Hard

Simple filters Empirical data / rules Classification; QSAR Structural mapping Pharmacophores; 3D-QSAR Structure; Dynamics First principles (QM; Stat mech)

Binding ... Selectivity ... Oral Absorption ... BBB ... Protein binding ... Solubility ...
Vss ... Metabolism ... hERG ... CYP inhibition ... CYP reactivity ... Dose ... CYP
induction ... Toxicity ... Production of reactive metabolites ... Tissue
Distribution ... Cell permeability ... Transporter inhibition ... Active
transport ... Gut stability ... Oral half-life ... IV half-life ...

Part 2:
IL-1 β Converting Enzyme
(ICE; Caspase-1)

Observations from Vertex: What Factors Lead to Successful Structure-Based Drug Design?

- Structures available early in each project
 - Willingness and ability to produce protein
- Real-time structures (rapid feedback)
- Experts at interpreting / applying structure
 - Diverse backgrounds, savvy, practical
- Strong links between chem, modeling, x-ray
 - Broad exploration of chemotypes
- Realism about value & limitations of SBDD
 - Don't oversell the technology - use appropriately
- Focus on *drug* design goals
 - Willing to trade *good binding* for *good properties*

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- Focus on *drug* design goals
 - Willing to trade *good binding* for *good properties*

Required heroic biochemical efforts but saved a year+

X-ray structures of >10 of distinct scaffolds

4 modelers with diverse backgrounds, plus savvy crystallographers

Broad exploration of chemotypes; aggressive use of structural info.

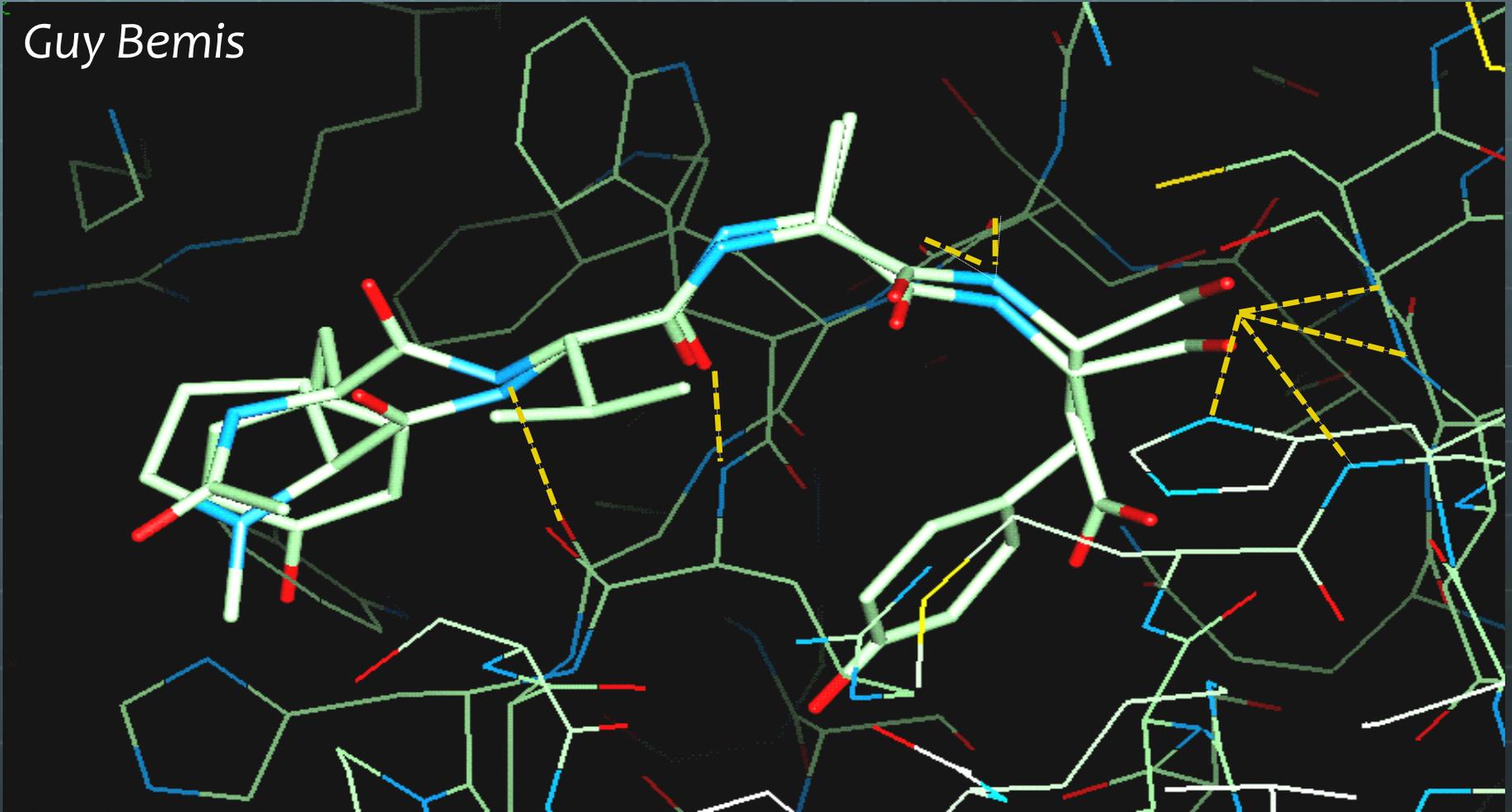
Testing multiple compounds for each scaffold; "bracketing"

Chemistry centered on drug-like cmpds; early focus on PK, whole-blood cell efficacy

The ICE - Chymotrypsin Connection

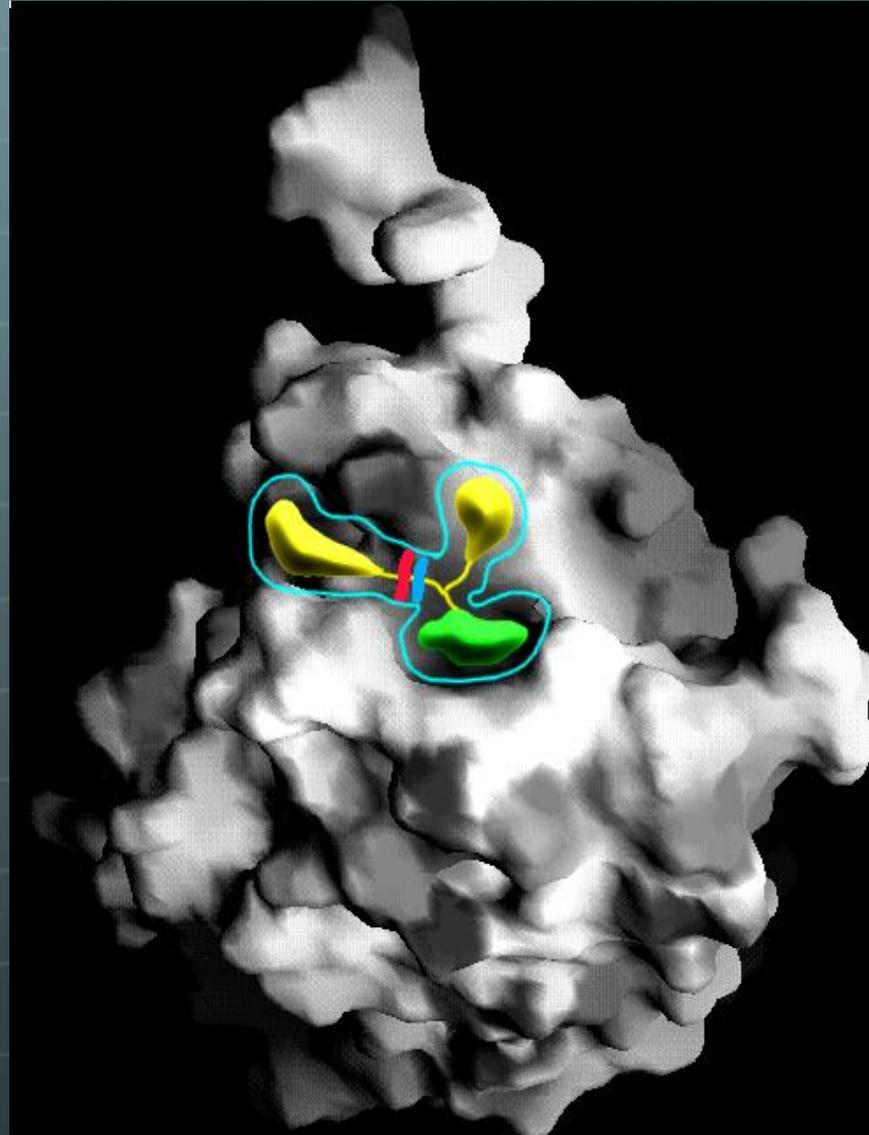
Different global folds - similar ligand recognition motifs

Guy Bemis



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The ICE Active Site Pharmacophore



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ICE: Good or Bad Outcome?

- 🌐 Dev. candidate series designed within 5 wks of xray
- 🌐 First compound synthesized was 20 nM
- 🌐 Sixth compound: decent oral rat clearance and $t_{1/2}$
- 🌐 Development candidate 2 years after that
- 🌐 Efficacious in 280 patient Phase 2A RA study

ICE: Good or Bad Outcome?

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- Development candidate 2 years after that
- Efficacious in 280 patient Phase 2A RA study
- **But:**
 - **Molecules were high-MW acids with poor permeability, poor WB cell activity, low human half-life, and high dose**
 - **Pralnacasan showed fibrosis in dogs after 9 months at very high dose**
 - **Aventis dropped program during Phase 2B RA trial**

ICE: Lessons

- 🌐 Fantastic SBDD effort → blistering speed
- 🌐 Creative insight re: chymotrypsin fold led to breakthrough
- 🌐 Deep understanding of relevant history led to dev candidate scaffold
- 🌐 Deep understanding of protein-ligand recognition motifs led to broad patent claims
- 🌐 But in the end, the molecule was sub-standard and our understanding of the disease biology was inadequate.
- 🌐 So: a SBDD **failure**.

Part 3: Covalency

Covalent Drugs: More Common Than You'd Think

Table 2. Targets, Indications, and Mechanism of Action of Covalently Interacting Small Molecules

mechanism	target	indication	name of drug or representative drug ^a	reacting functionality	reversibility	dose (mg) ^b	
acylation	serine-type D-Ala-D-Ala carboxypeptidase	bacterial infection	amoxicillin ^c	β -lactam	irreversible	100–500	
		obesity	orlistat	lactone	reversible	360	
		Alzheimer's	rivastigmine	carbamate	reversible	6–12	
	β -lactamase	bacterial infection	clavulanate ^c	β -lactam	irreversible	500	
		prostaglandin endoperoxidase synthase	pain	aspirin	ester	reversible	1000
	vitamin K epoxide reductase (warfarin-sensitive)	anticoagulant	warfarin	coumarin			2–10
		enol-acyl carrier protein reductase	bacterial infection (tuberculosis)	isoniazid	hydrazide ^d	irreversible	300
	aldehyde dehydrogenase		alcoholism	disulfiram	disulfide	irreversible	500 ^e
	alkylation	UDP-N-acetylglucosamine-1-carboxyvinyltransferase	bacterial infection	fosfomycin	epoxide		3000
			bacterial infection (tuberculosis)	D-cycloserine	amine ^d		> 250
metal/metalloid binding	GABA-AT ^f aromatase	epilepsy	vigabatrin	amine ^d	irreversible	3000 ^e	
		breast cancer	exemstane ^e	methyl boronic acid	irreversible	25	
disulfide bond formation	H ⁺ /K ⁺ ATPase	multiple myeloma	bortezomib	boronic acid	reversible	3	
		gastroesophageal reflux disease	omeprazole ^e	sulfenamide	irreversible	20	
(seleno-enzyme)	P2Y12 purinoceptor antagonist	platelet aggregation inhibitor	clopidogrel	thiol	irreversible	75	
		thyroxine 5'-deiodinase (type 1)	hyperthyroidism	propylthiouracil	thiourea		450
hemiketal formation	serine protease	viral infection	VX-950 (1q)	ketoamide	reversible	n/a	
Michael addition	ribonucleoside diphosphate reductase	cancer	gemcitabine ^e	vinyl ketone		≥ 150 –000 ^h	
		thymidylate synthase	cancer	floxuridine ^e	unsaturated amide	reversible	0.1–0.6 (mg/kg)/d
	ErbB1/2 ^g	cancer (NSCLC)	HKI-272 (1t)	unsaturated amide	irreversible	n/a	
	5- α -reductase	benign prostatic hyperplasia	finasteride ^e	unsaturated amide ^d	reversible	5	
Pinner reaction	MAO-B	Parkinson's disease ⁱ	selegiline ^e	acylenic imine ^d	irreversible	1	
		diabetes	osteoporosis	vildagliptin	nitrile	reversible	100
cathepsin K ^g	odanacatib			nitrile	reversible	10–50 ^j	

^a Prodrugs are indicated in italics. ^b As determined from the FDA label or other medical references. ^c Because of the large number of drugs developed for these targets, one representative drug is indicated in the table. ^d Indicates functionality covalently modified by the cofactor. ^e Estimated dose. ^f Approved in Canada, U.K., and Mexico. ^g Under clinical investigation. ^h Dose = 1000 mg/m² weekly. The average body surface area of a person is approximately 1.5–2 mm². ⁱ Several irreversible MAO inhibitors are on the market for the treatment of depression. ^j Weekly dose used in the clinical trial "MK0822 (Odanacatib) Late Phase II Dose-Finding Study" described at www.clinicaltrials.gov.

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Source: Potashman, Michele H., and Mark E. Duggan. "Covalent Modifiers: An Orthogonal Approach to Drug Design." *Journal of Medicinal Chemistry* 52, no. 5 (2009): 1231-46.

Aspirin MOA Finally Revealed

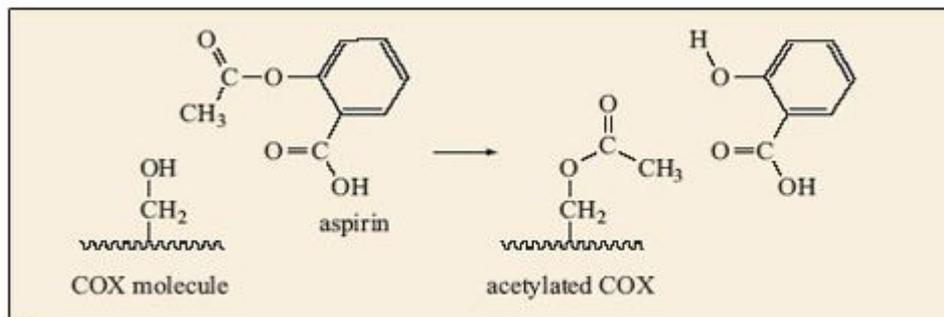
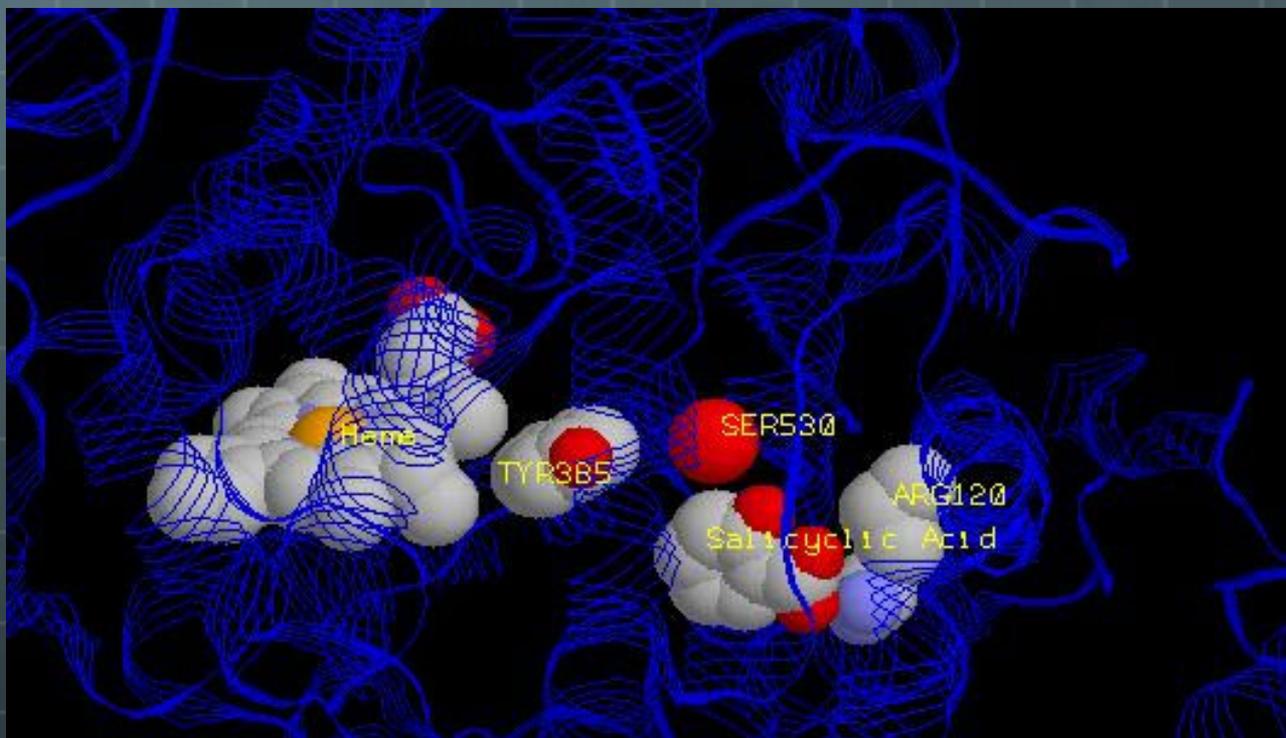


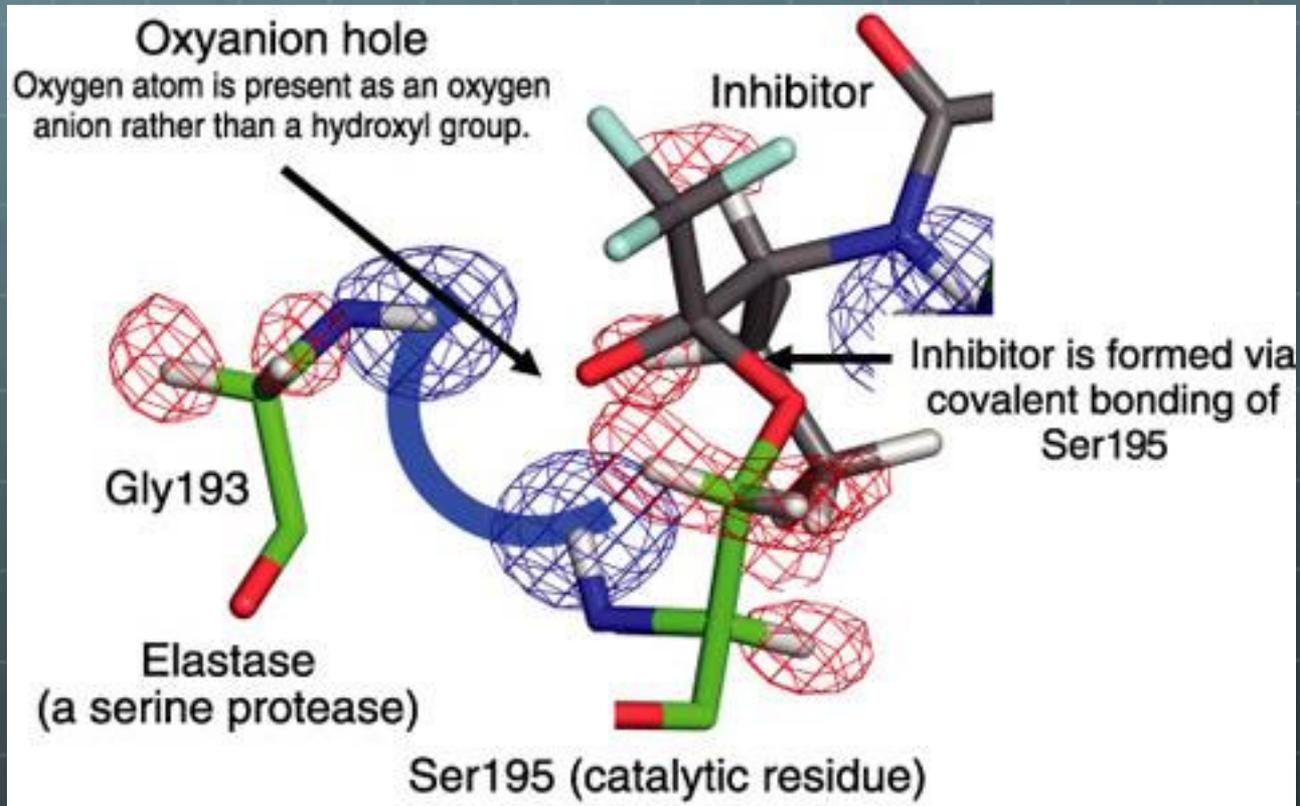
Figure 16: The acetyl group in aspirin reacts with an alcohol group inside the COX cavity.

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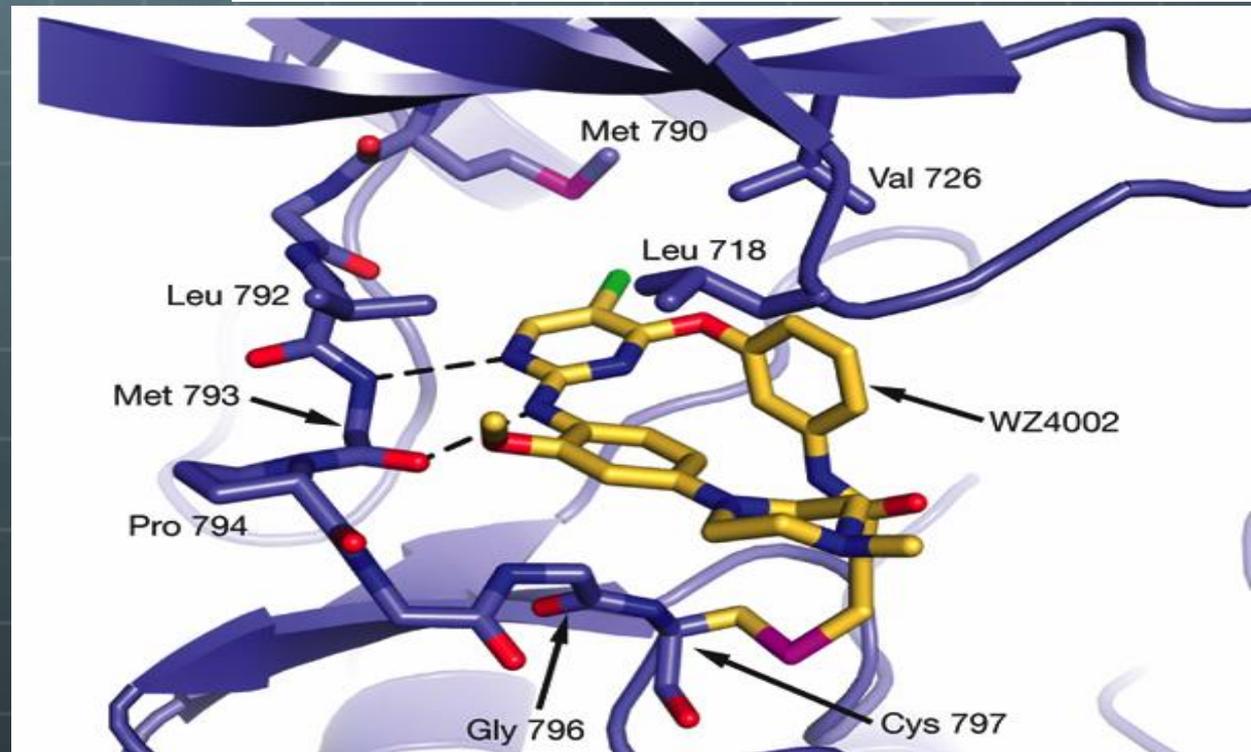
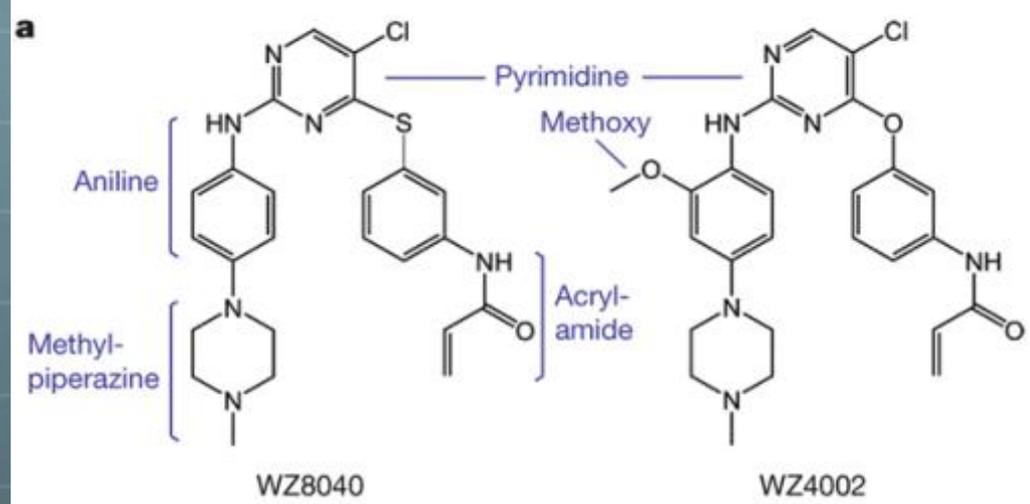
Covalent Serine Protease Inhibitors



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Irreversibles Don't Have to Use Catalytic Residues

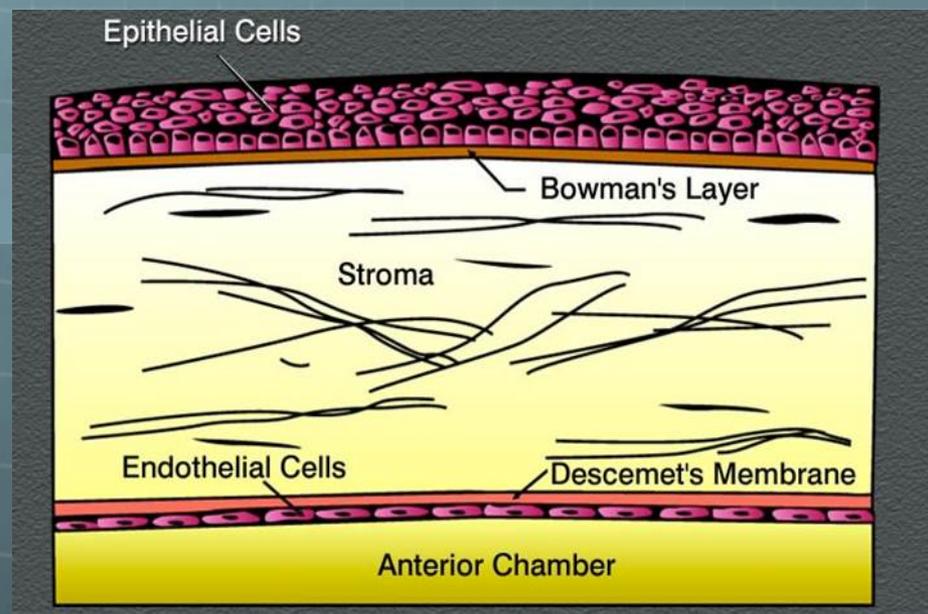
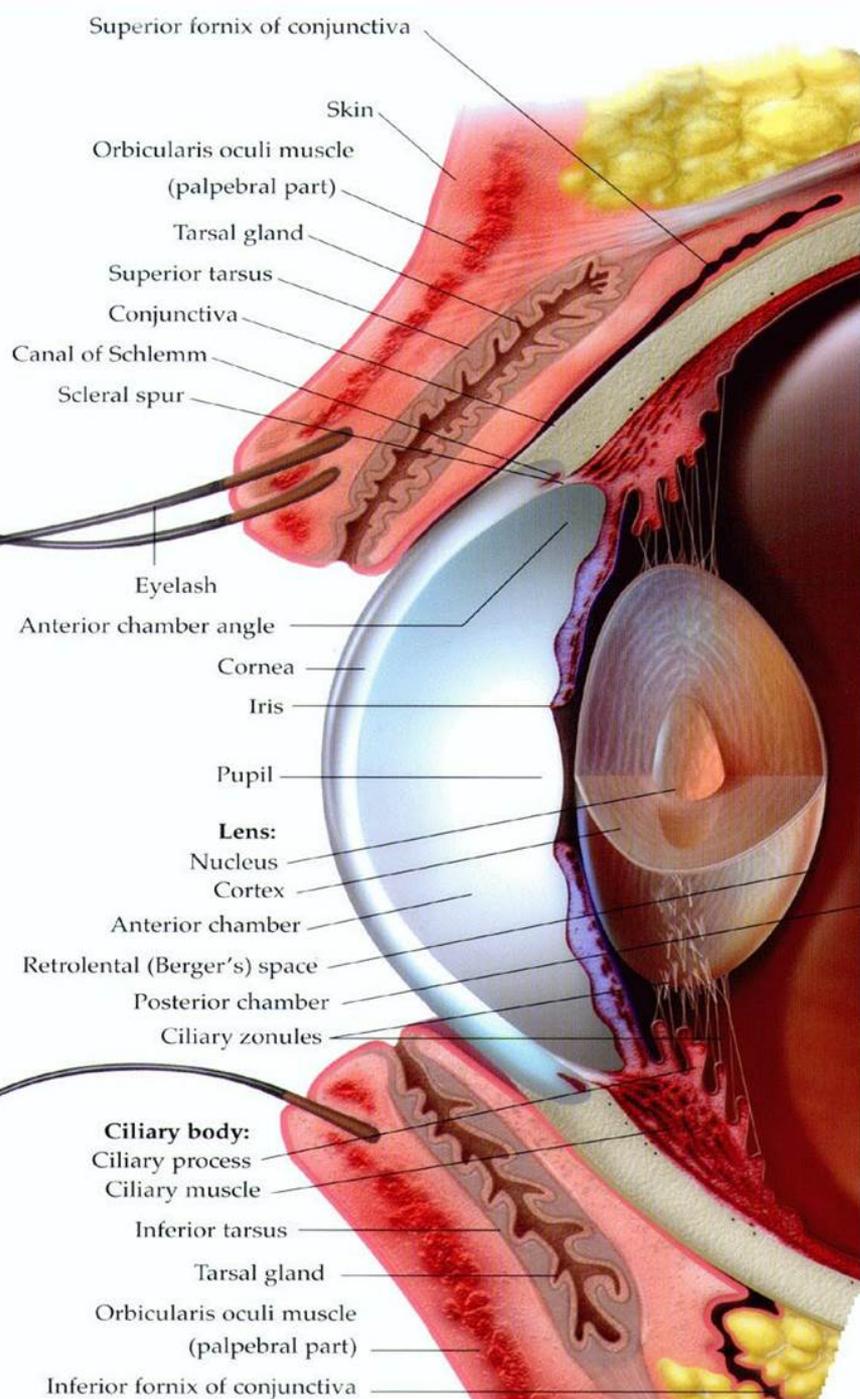
- 🌐 Epidermal growth factor receptor (EGFR) kinase inhibitors
- 🌐 Acrylamide moiety reacts with conserved cysteine
- 🌐 Discovered by screening against mutants resistant to other EGFR inhibitors



Courtesy of Macmillan Publishers Limited. Used with permission.
Source: Zhou, Wenjun, Dalla Ercan, et al. "Novel Mutant-Selective EGFR Kinase Inhibitors Against EGFR T790M." *Nature* 462, no. 7276 (2009): 1070-4.

Part 4: Four SBDD Drugs

Glaucoma



The epithelium - Covers the surface of the cornea, is about 5-6 cell layers thick.

Bowman's membrane - Very difficult to penetrate.

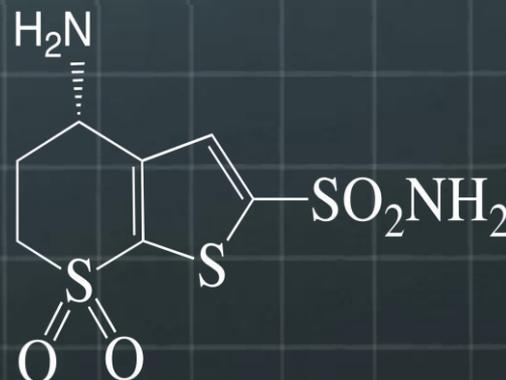
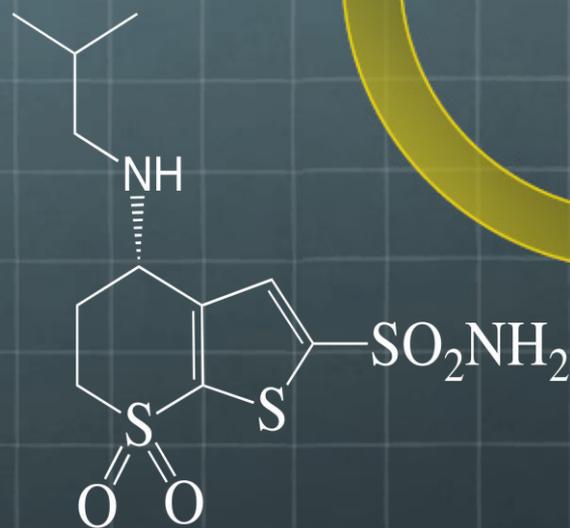
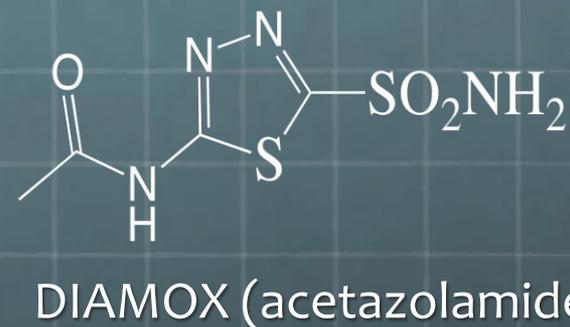
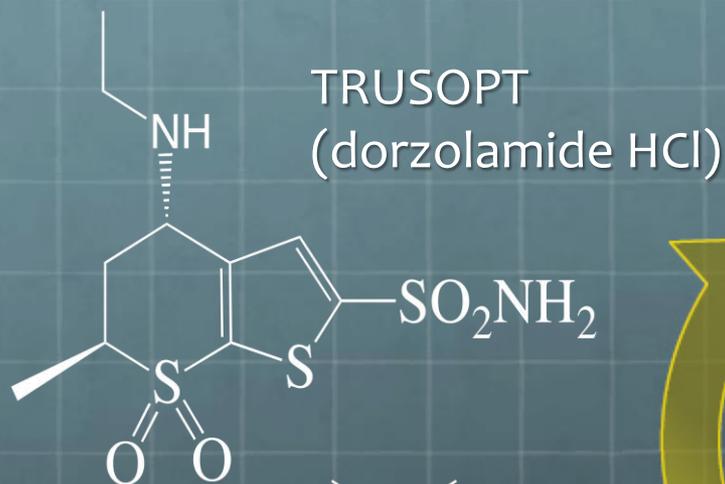
The stroma - The thickest layer, composed of tiny collagen fibrils that run parallel to each other, this precision formation gives the cornea its clarity, strength, elasticity, and form.

Descemet's membrane - A thin but strong sheet of tissue that acts as protection against infection and injuries. It is composed of collagen fibers (different from those of the stroma).

The endothelium - Essential in keeping the cornea clear. It pumps this excess fluid out of the stroma, which has the danger of swelling with water.

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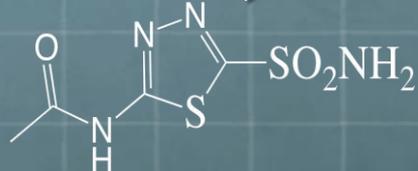
First Crystallography-Based Drug Design Example (Merck)



J. Med. Chem. **30**, 591-599 (1987)
J. Med. Chem. **32**, 2510-2522 (1989)
J. Med. Chem. **37**, 1035-1054 (1994)

A Struggle of Biblical Proportions?

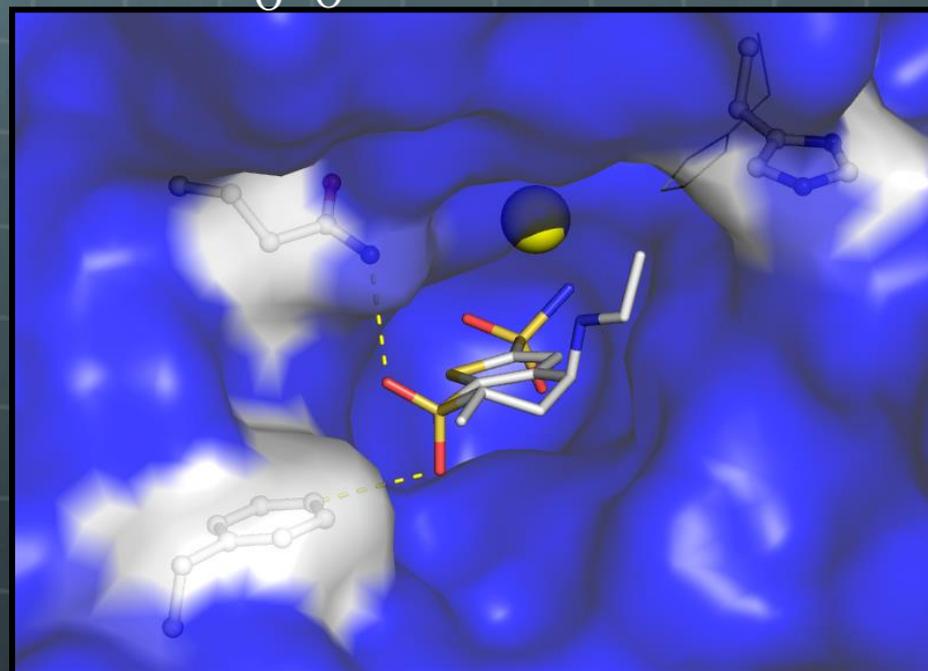
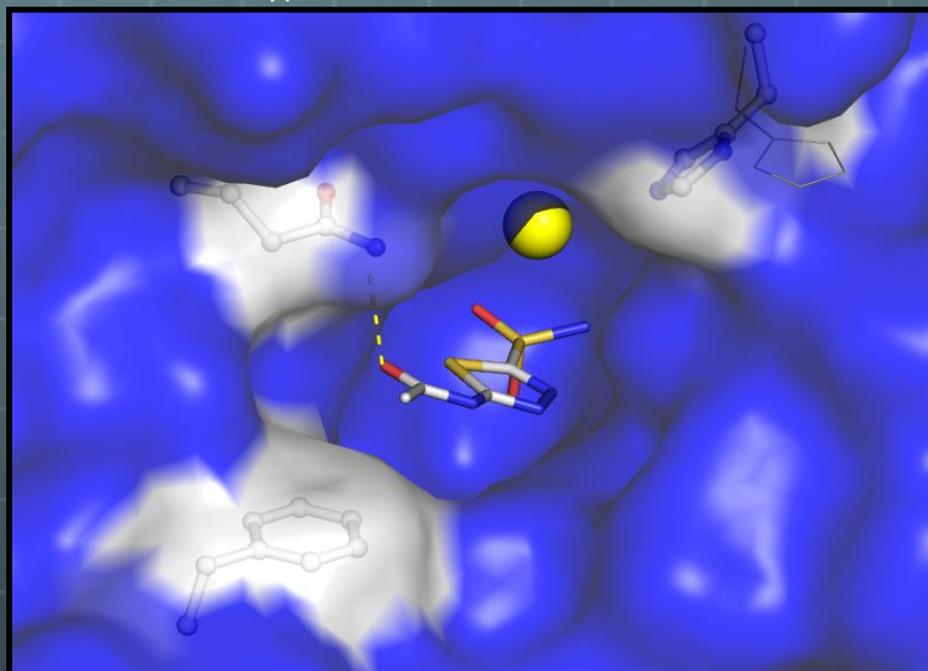
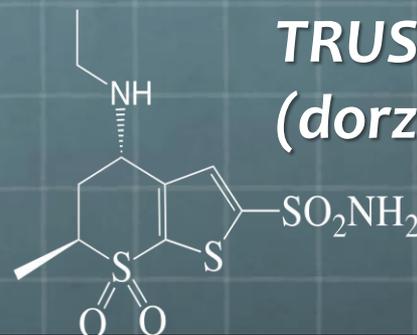
DIAMOX
(acetazolamide)



40 YEARS



TRUSOPT
(dorzolamide HCl)



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1YDA, Nair et al
Biochemistry 34, 3981-3989 (1995)

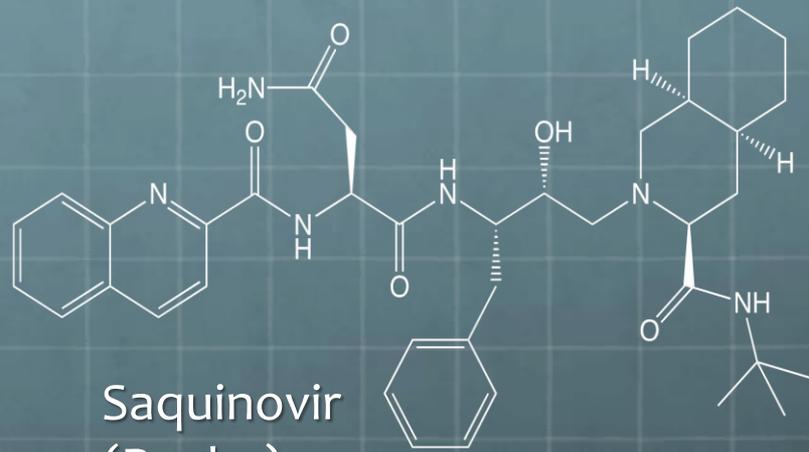
1CIL, Smith et al
Protein Sci. 3, 118-125 (1994)

Carbonic Anhydrase: Lessons

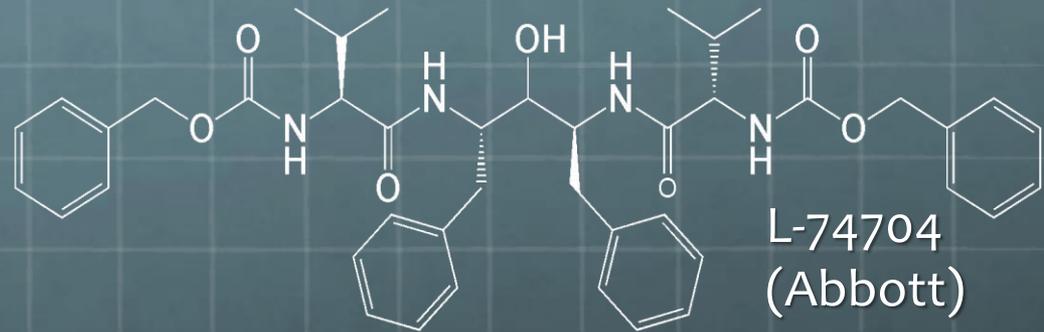
-  When working on validated targets, “stay the course”
-  SBDD can be used to optimize physical / biological properties
-  Conformational analysis is critical

HIV

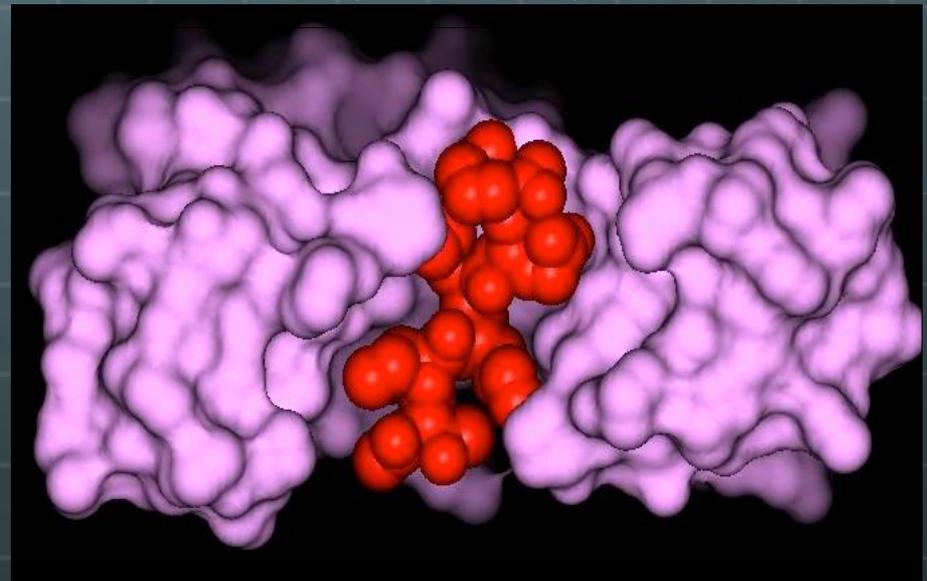
HIV Protease: Prototypes, Circa 1992



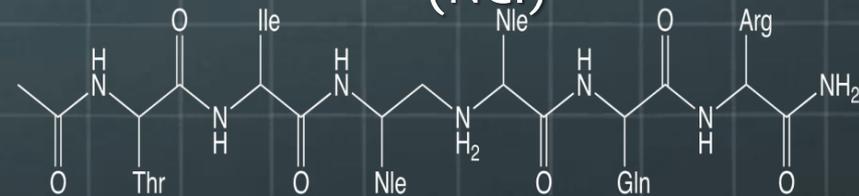
Saquinovir
(Roche)



L-74704
(Abbott)

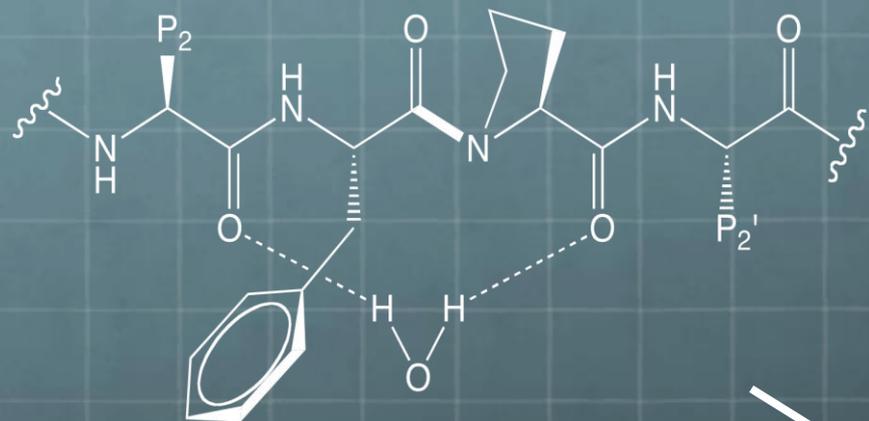


MVT-101
(NCI)

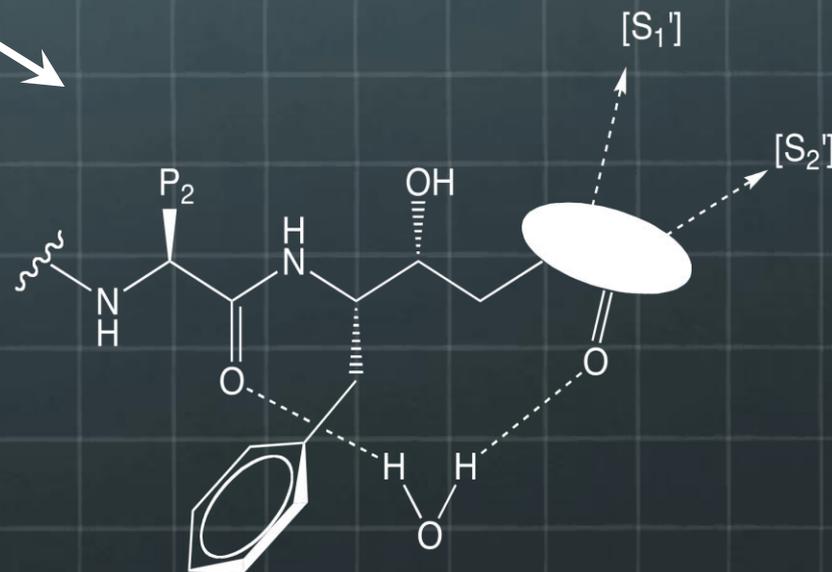


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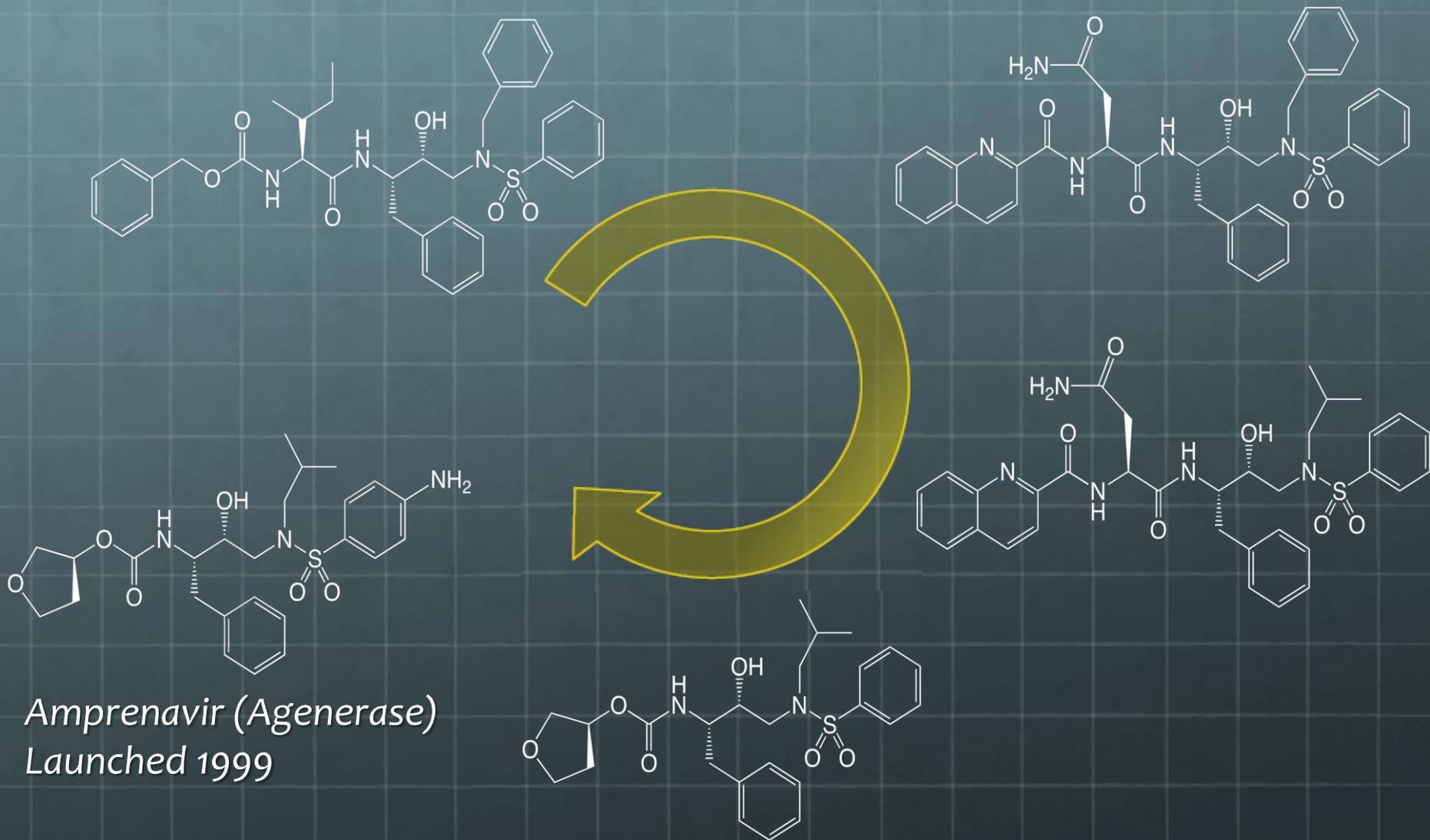
Novel Scaffolding → Simpler Molecules?



- 🌐 Preserve the interactions with catalytic Asps
- 🌐 Maintain the hydrogen bonds to the flap water
- 🌐 Design a scaffold which can reach S₁' and S₂'
- 🌐 Design a scaffold with minimal binding strain

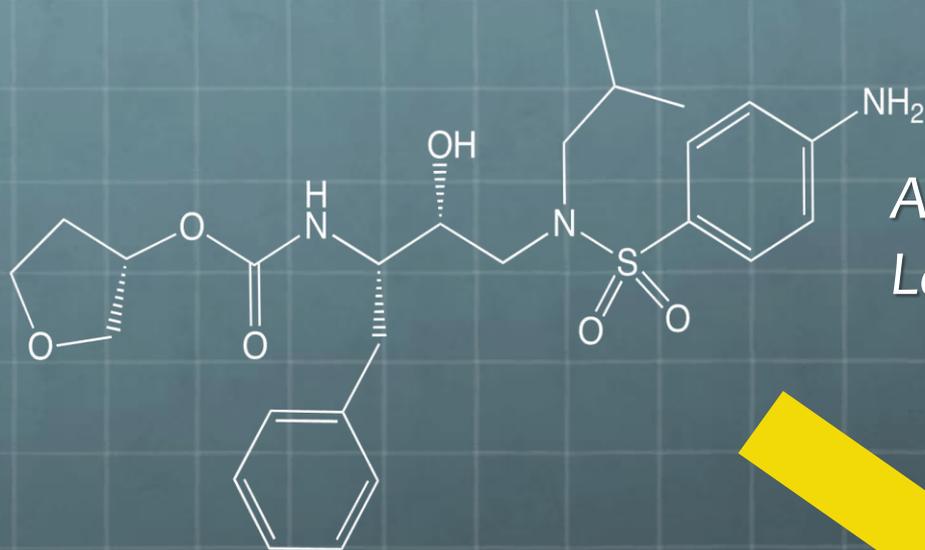


5 Chemists, 12 Months, 204 Compounds

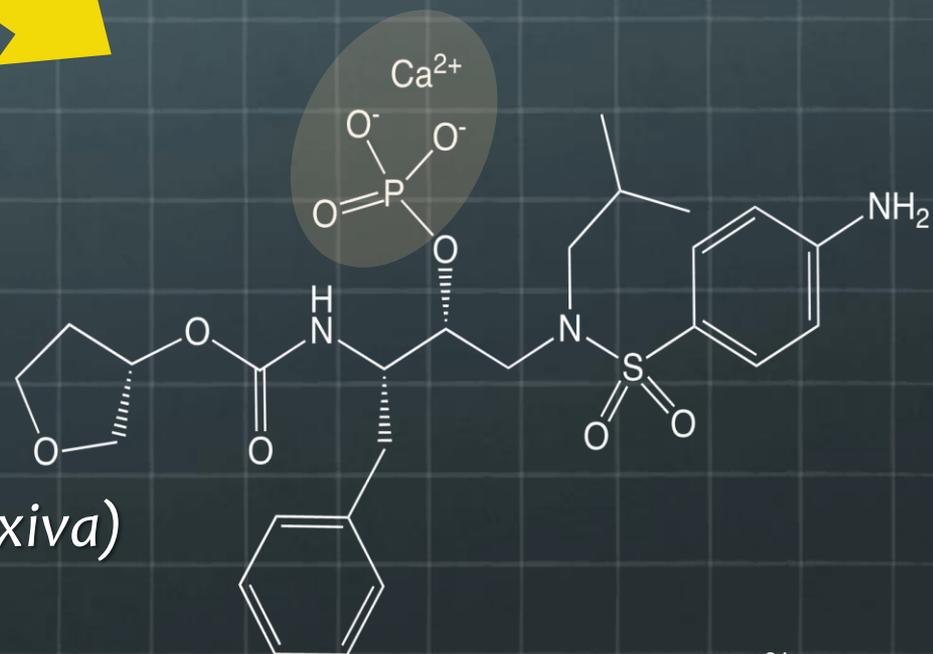


D'Oh!

Roger Tung



*Amprenavir (Agenerase)
Launched 1999*



*Fosamprenavir (Lexiva)
Launched 2003*

HIV-Protease: Lessons

- 🌐 Conformational analysis is incredibly powerful
- 🌐 SBDD can help optimize physical properties
- 🌐 Sometimes the marketing guys are right
- 🌐 Pay attention to formulation early

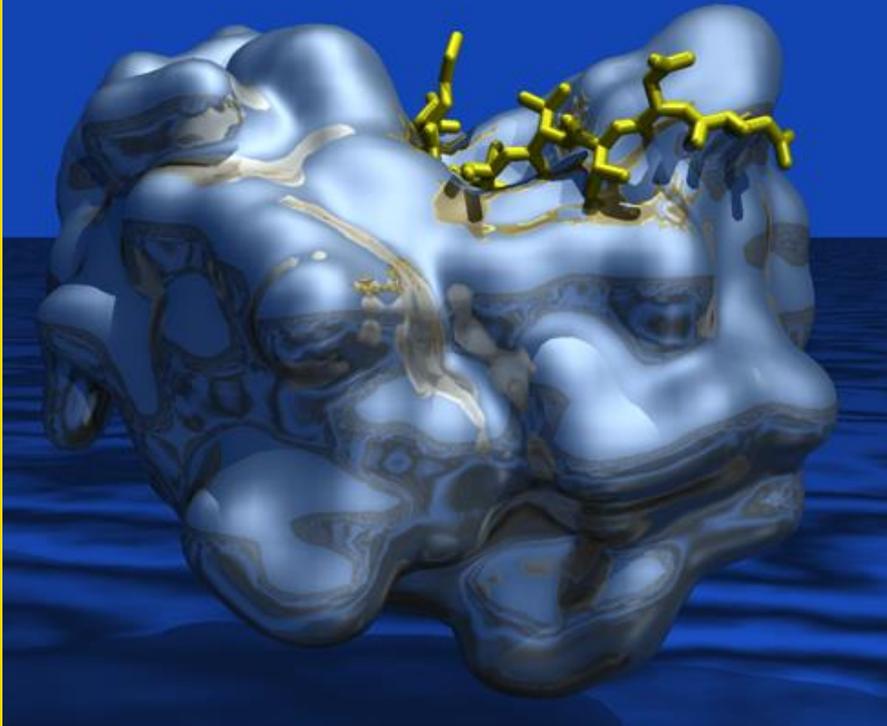
HCV

Hepatitis C Infection

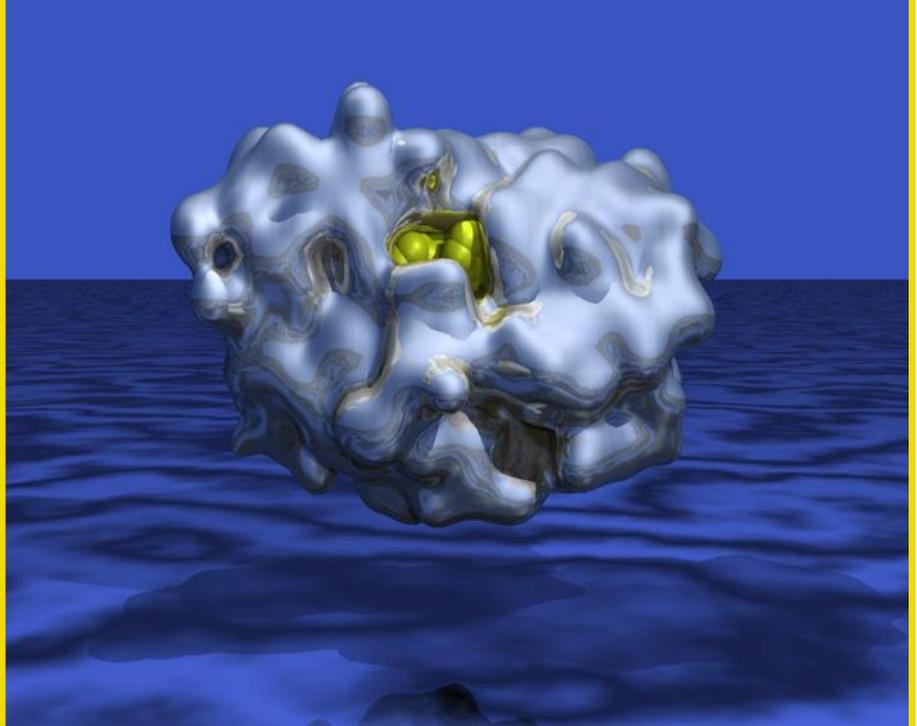
- 🌐 Infects ~200 million people worldwide
- 🌐 Progresses to cirrhosis in 20-30% of cases
- 🌐 Progresses to hepatocellular carcinoma in 1-3% of cases
- 🌐 Responsible for ~10,000 death / yr in US
- 🌐 PEG IFN- α + Ribavirin <50% effective

HCV Protease: A Dinner Plate

HCV NS3 • 4A Protease with NS5A-5B Substrate



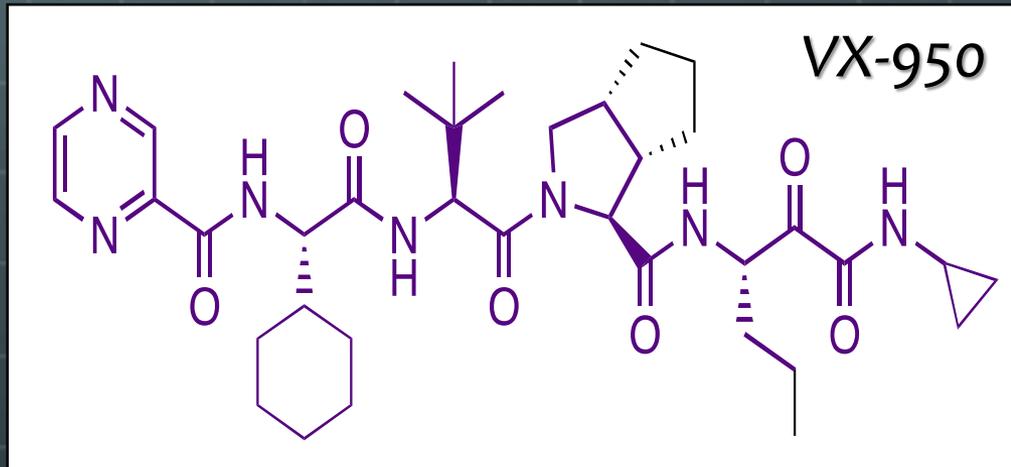
HIV Protease with bound Agenerase®



“Several loops found in other chymotrypsin family proteases are missing from HCV. These loops normally play a critical role in defining the shapes of the non-prime-side substrate-binding pockets. The absence of these loops in HCV-PR renders the binding groove **relatively featureless**, and this constitutes a **challenge for drug design efforts**. It is therefore anticipated that **structural information** for enzyme-inhibitor complexes **may be crucial** for optimization of potent, drug-like inhibitors.”

HCV: Telaprevir

- Efficacy surrogate: high ratio of liver concentration to IC_{50}
 - $[C_{liver}] > 10X IC_{50}$
 - Fa more important than %F
- High $[C_{liver}]$ compared to other organs or tissues
 - Minimize potential for systemic toxicity
 - Some plasma exposure required to cover extra-hepatic sources of virus and for drug load monitoring in patients



HCV Program Strategy

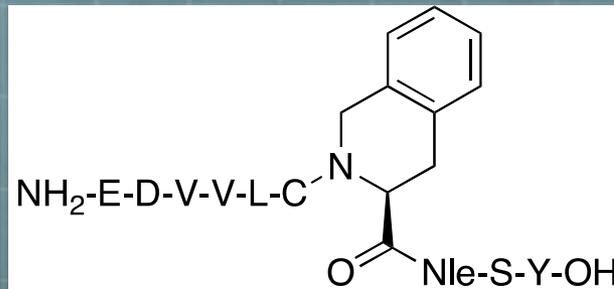
- Efficacy surrogate: high ratio of liver concentration to IC_{50}
 - $[C_{liver}] > 10X IC_{50}$
 - F_a more important than %F
- High liver concentrations are generally desirable compared to other organs or tissues
 - Minimize potential for systemic toxicity
 - Some plasma exposure required to cover extra-hepatic sources of virus and for drug load monitoring in patients

Perni, R.B. et. al. *Bioorg. Med. Chem. Lett.* **2004**, 14, 1939-1942

Y. Yip et. al. *Bioorg. Med. Chem. Lett.* **2004**, 14, 251-256

F. Victor et. al. *Bioorg. Med. Chem. Lett.* **2004**, 14, 257-261

Truncating the Decapeptide Substrate Mimic

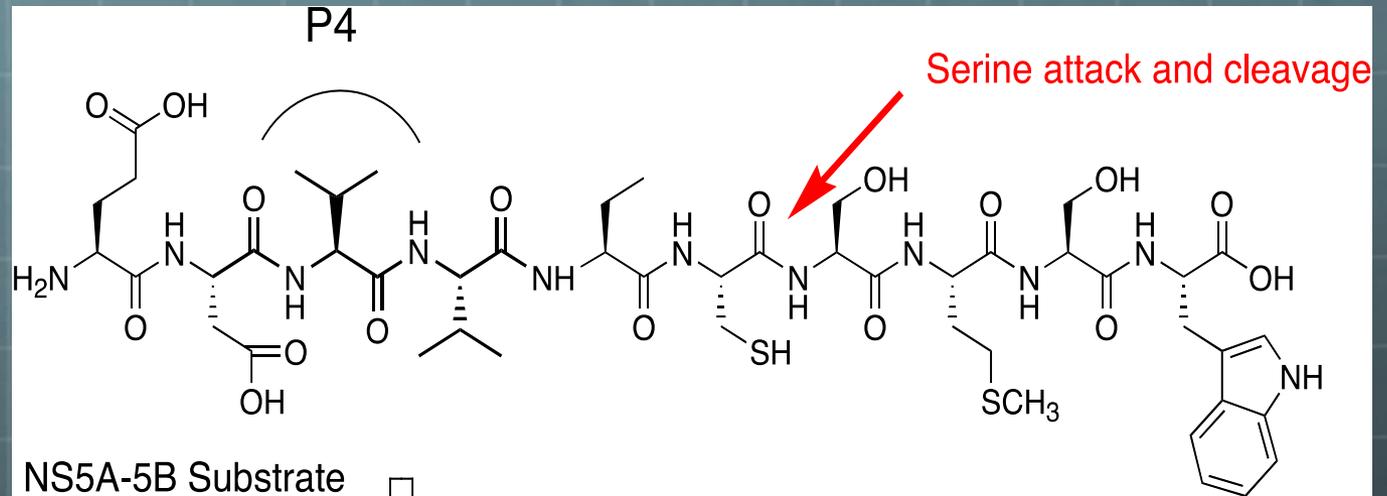


K_i (uM)

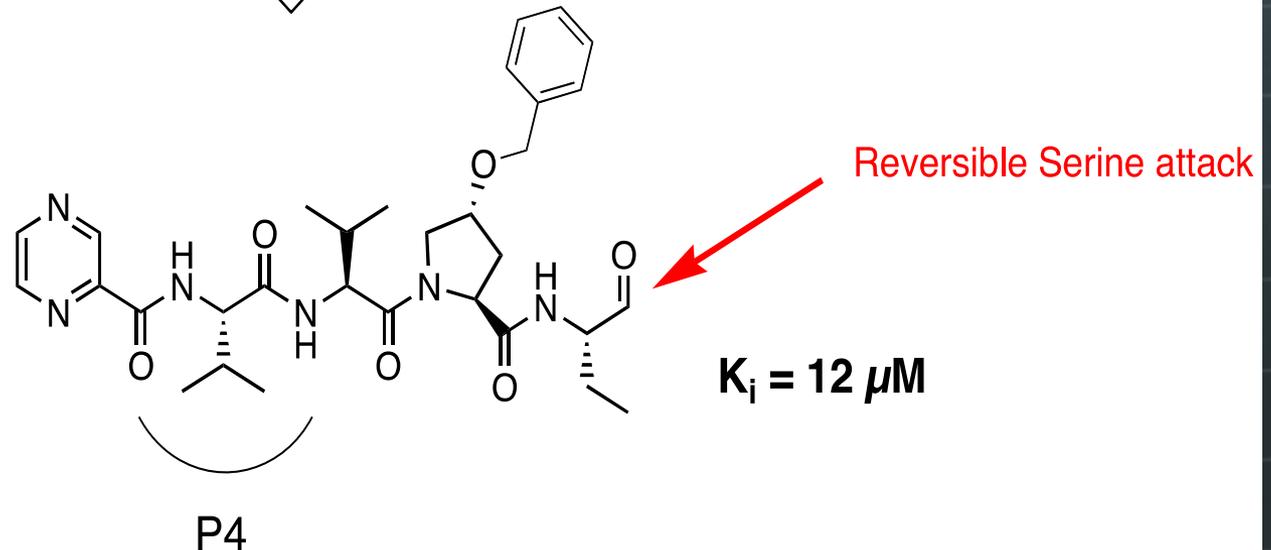
H-Glu-Asp-Val-Val-Leu-Cys-Tic-Nle-Ser-Tyr-OH	0.34
H-Glu-Asp-Val-Val-Leu-Cys-Tic-Nle-Ser-OH	27
H-Glu-Asp-Val-Val-Leu-Cys-Tic-Nle-OH	17
H-Glu-Asp-Val-Val-Leu-Cys-Tic-OH	14
H-Asp-Val-Val-Leu-Cys-Tic-Nle-Ser-Tyr-OH	4.4
H-Val-Val-Leu-Cys-Tic-Nle-Ser-Tyr-OH	79
H-Val-Leu-Cys-Tic-Nle-Ser-Tyr-OH	500
H-Leu-Cys-Tic-Nle-Ser-Tyr-OH	2000

Inhibitor Evolution

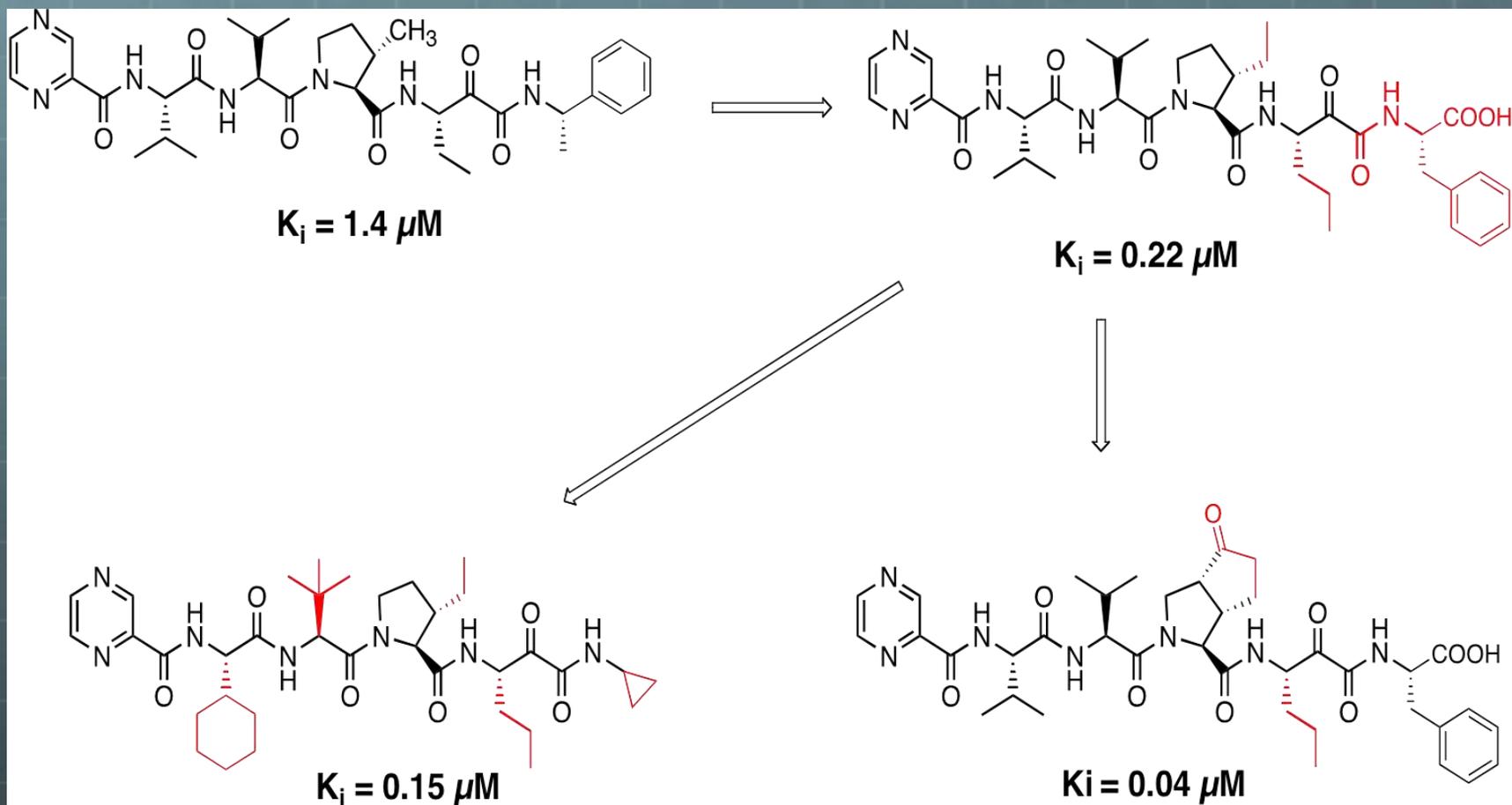
Substrate



Inhibitor



Multi-Subsite Optimization

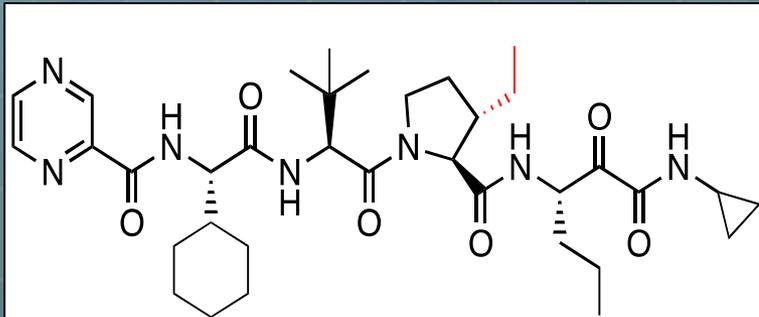


Perni, R.B. et al. *Bioorg. Med. Chem. Lett.* **2004**, 14, 1939-1942

Y. Yip et al. *Bioorg. Med. Chem. Lett.* **2004**, 14, 251-256

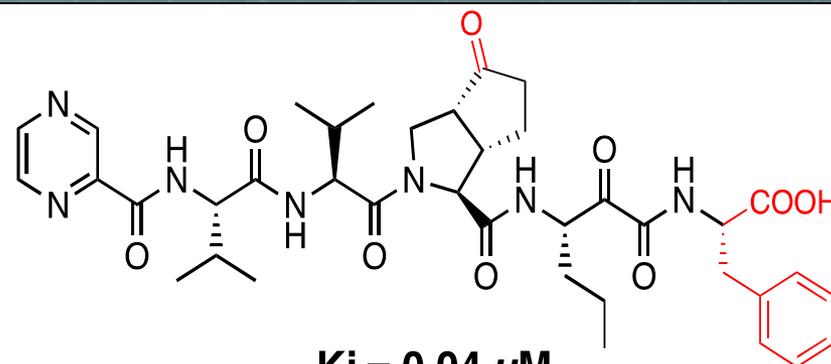
F. Victor et al. *Bioorg. Med. Chem. Lett.* **2004**, 14, 257-261

The Finish Line



$K_i = 0.15 \mu\text{M}$

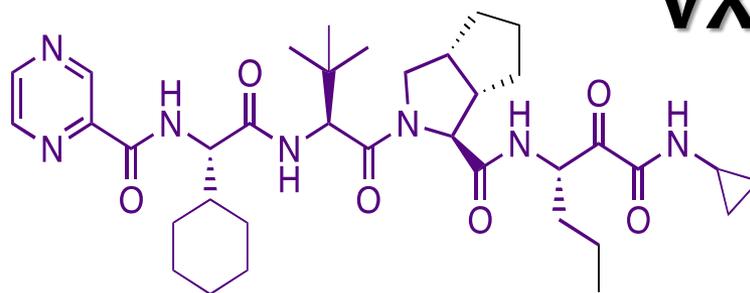
Replicon $\text{IC}_{50} = 0.45 \mu\text{M}$



$K_i = 0.04 \mu\text{M}$



Two-step binding mode:
conformational flip of both
enzyme and ligand at the
catalytic machinery



VX-950

$K_i = 44 \text{ nM}$ (15 min pre-incubation)

$K_i^* = 7 \text{ nM}$

Replicon $\text{IC}_{50} = 0.35 \mu\text{M}$

How Do You Know You're Done?

A “good drug” --

- Serves an important need
- Has enough potency, bioavailability, and safety
- Is novel
- Can be made (formulation, synthesis, stability, ...)

“Every design balances--connects--dozens of values, like a conceptual mobile, and the weights of those values, their relative utility or attractiveness, are **changing constantly**.”

“At some point you have to **shoot the engineers** and ship.”

“A great design attracts applications, and in doing so necessarily makes its creators look **short-sighted and slightly dumb**.”

HCV-Protease: Lessons

- 🌐 Stick with validated targets even if hard – but be realistic about timelines
- 🌐 Consider how drugs of various mechanisms can be combined
- 🌐 Consider the target organ in your design
- 🌐 If you're first, chances are that a better drug will come along quickly. That's OK – don't worry about looking dumb later!
- 🌐 Have a vigorous 2nd generation plan

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20.201 Mechanisms of Drug Actions
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