

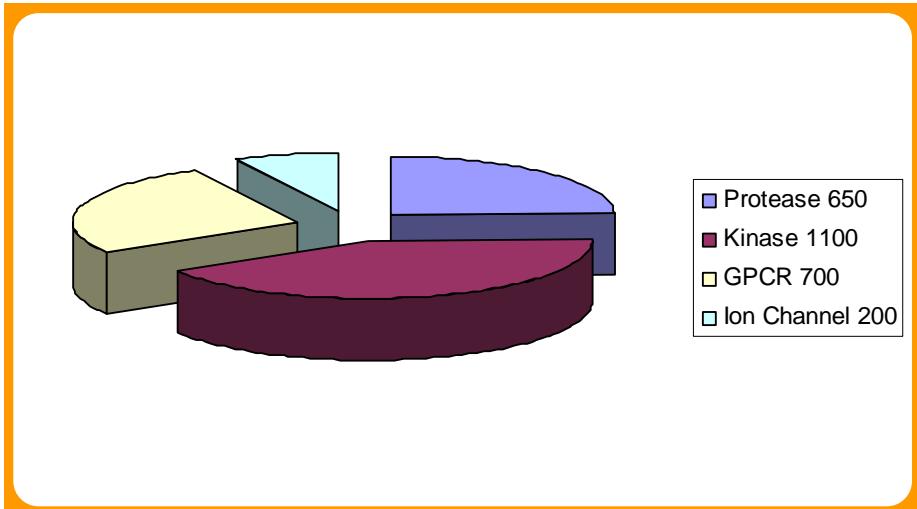
Cathepsin K Inhibitor Development



IASOC
September 16-21, 2006

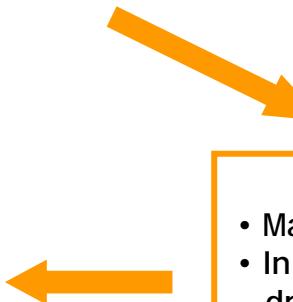
Bertil Samuelsson
Medivir AB, Stockholm

The Drugable Human Genome



- All marketed drugs addresses ~120 different targets
- The top 100 best-selling drugs are directed against only ~45 host proteins
- Very few new innovator targets are successfully developed each year

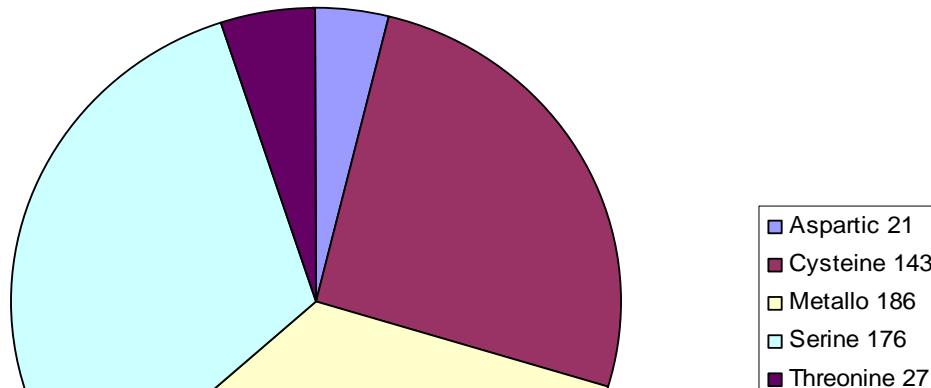
- The new “hot” target class



Proteases:

- Marketed drugs: ~40 from 7 classes
- In Clinical Trials: ~100 investigational drugs from 23 classes

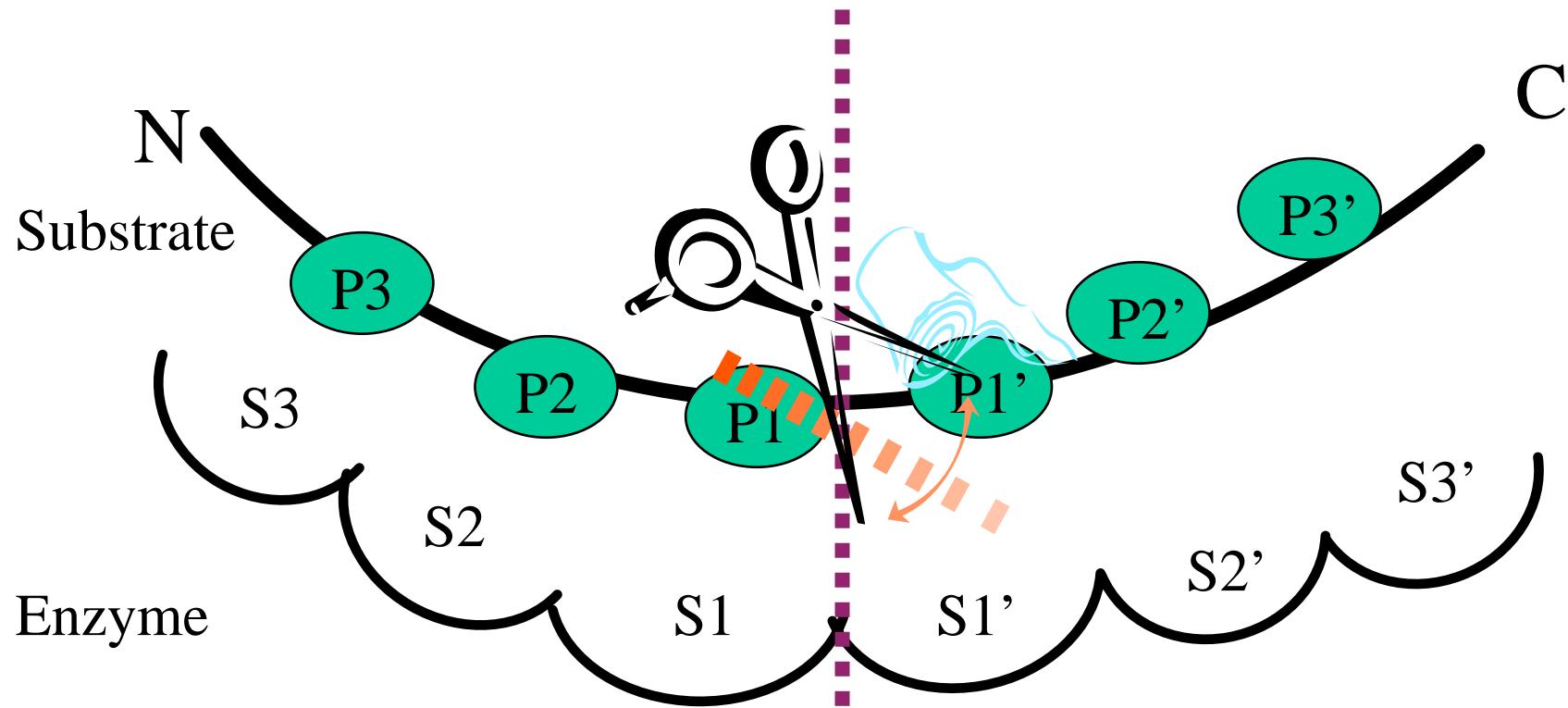
The Distribution of the Human Proteases



The serine, metallo and cysteine
Classes are the largest

"Proteases at work"

Cleaves Bonds Between two Amino Acids - Activates or Inactivates Proteins



Schecter, I. and Berger, A., Biochem. Biophys. Res. Com., **1967**, 27, 157-62.

Cathepsin K Inhibitors - Potential for Changing Treatment Paradigms



Many patients:

- Approx 100 million patients in major pharmaceutical markets have osteoporosis
- Incidence increasing
- Global market \$11 billion by 2008

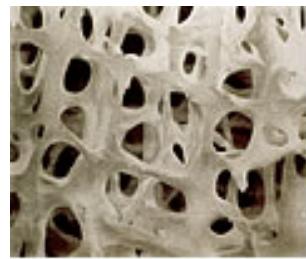
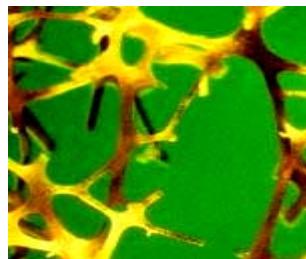
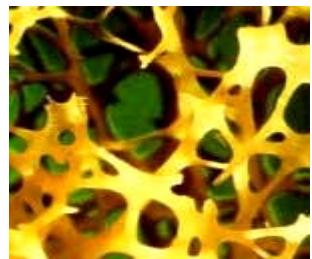
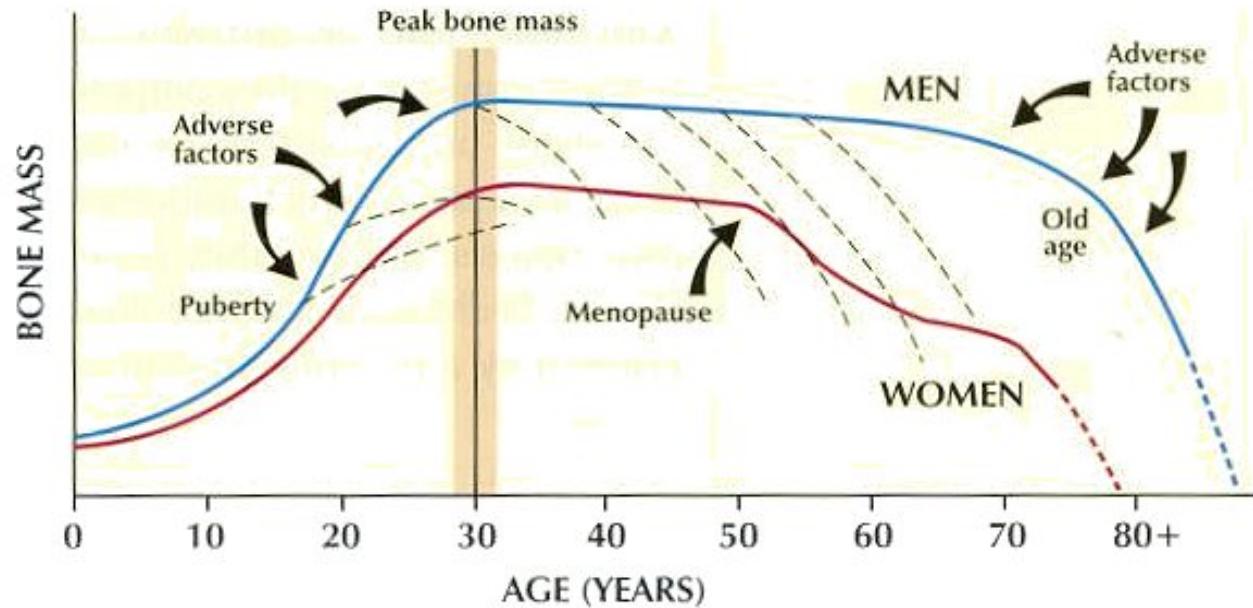
Bisphosphonate issues:

- Irreversible inhibition of bone build-up
- Raising concern: Necrosis of the jaw

Cathepsin K:

- Selectively expressed in the osteoclasts
- Pivotal role in the degradation of bone matrix
- Attractive new mechanism of action
- Potential for bone forming activity and reduced fracture incidence
- Suitable for broad patient populations and both genders

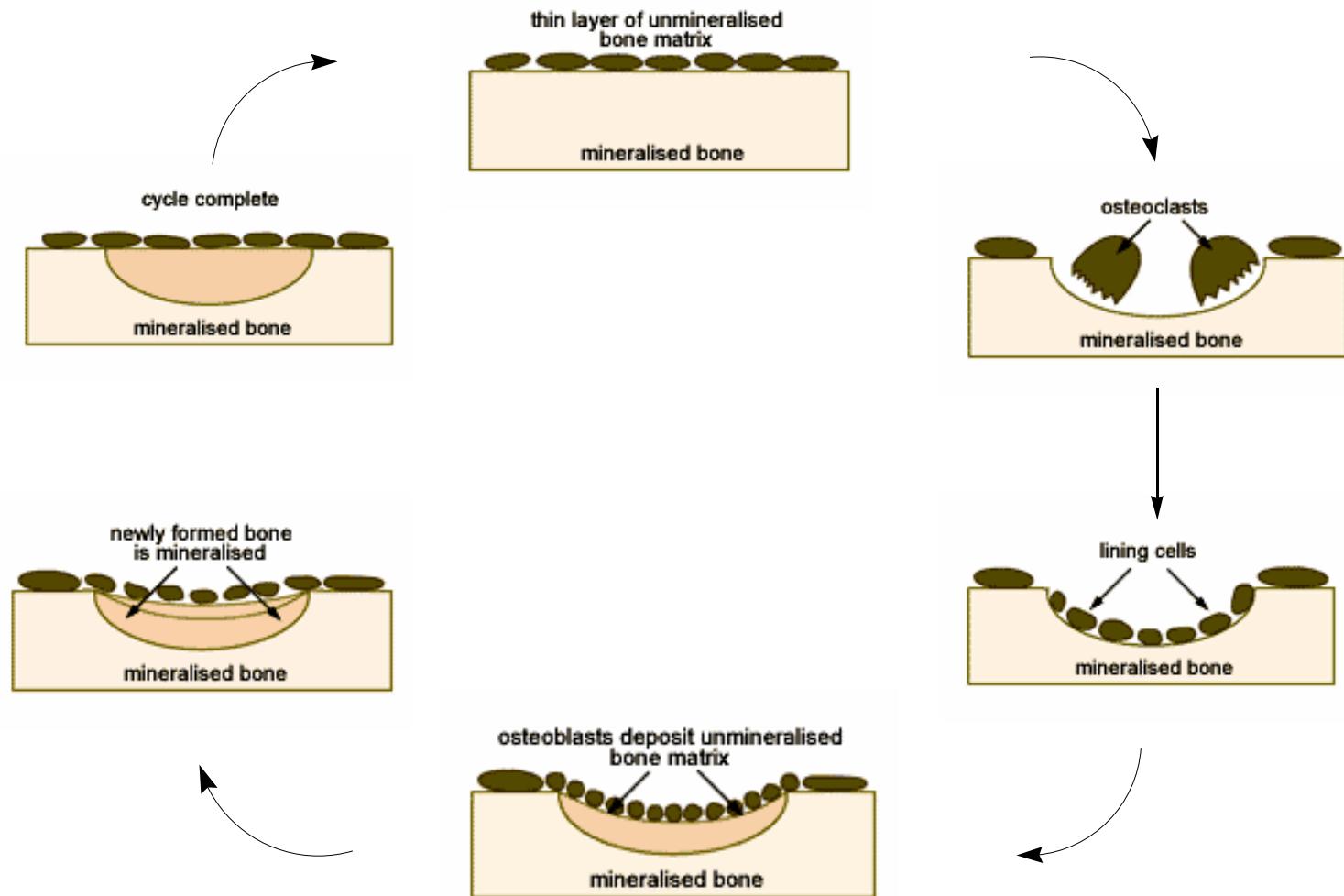
Osteoporosis



Bone Metabolism

- Bone is a form of connective tissue
 - type I collagen fibrils
 - hydroxyapatite crystals
- Unceasing process of remodelling
 - osteoblasts
 - osteoclasts
 - large multinucleated cells
 - originate from hemopoietic stem cells
 - precursor cells released into the bloodstream
 - collect at sites of bone resorption

Bone Remodelling Process



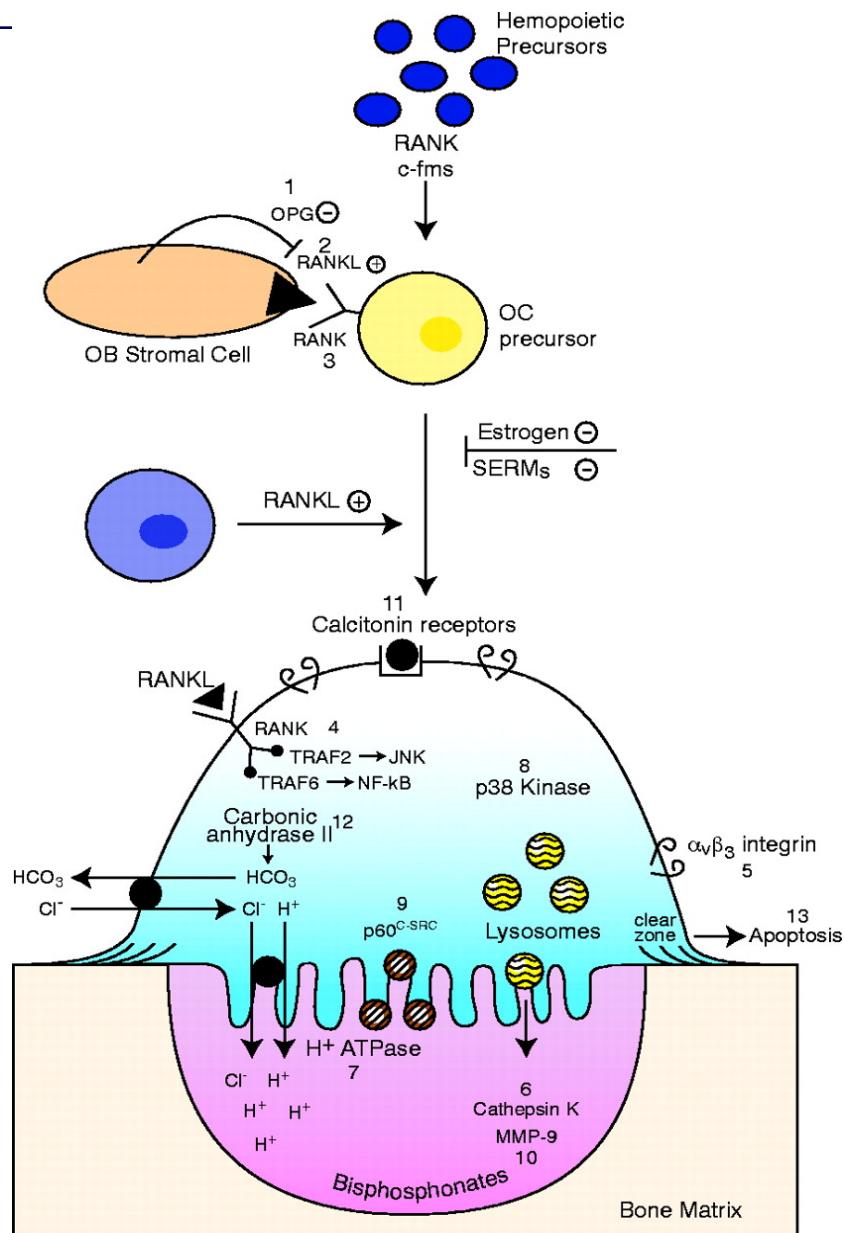
The Molecular Target: Cathepsin K

The osteoclast



Reproduced with Tim Chambers permission

B. Samuelsson
2006-09-11



Cathepsin K

- Cysteinyl proteinase
- Papain superfamily
- 73 % Homology to cathepsin S
- 76 % Homology to cathepsin L
- Highly expressed during bone resorption
- Only known mammalian protease to solubilise collagen by cleavage of type I & II collagen
 - Within triple helix
 - In telopeptide region

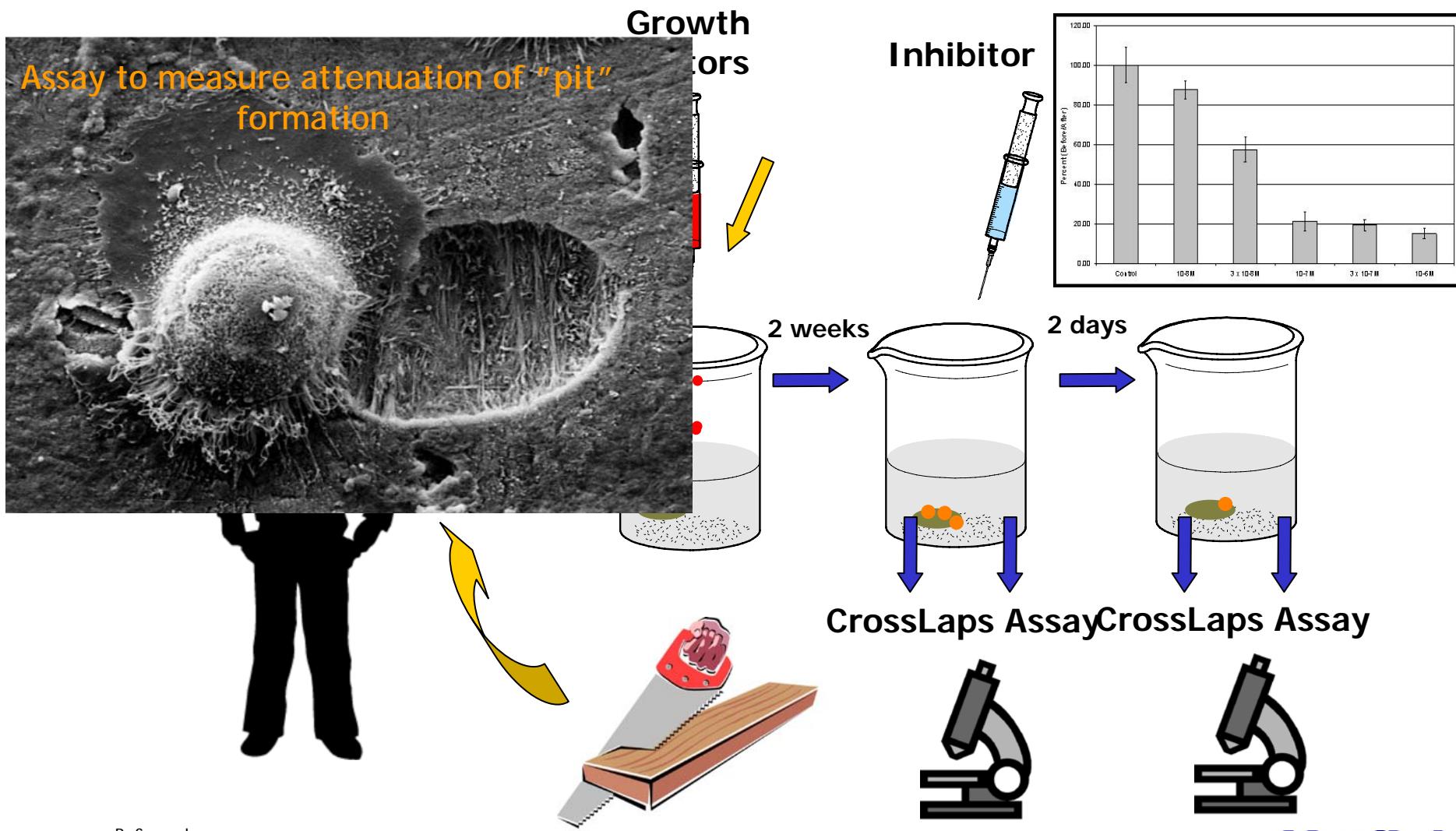
Role of Cathepsin K in Bone Remodelling

- Cathepsin K genetic deficiency leads to pycnodysostosis
 - Antisense inhibition of cathepsin K mRNA
 - Blocks protein translation
 - Leads to dose dependent inhibition of osteoclast pit formation
 - Cathepsin K knockout mice have been shown to develop osteopetrosis
 - Animal studies have shown that inhibition of cathepsin K leads to a reduction in bone resorption in cynomolgus monkeys in a dose dependent manner
-
- Cathepsin K is a critical enzyme in bone resorption
 - Cathepsin K inhibitors are developed to provide therapy for bone disorders such as osteoporosis

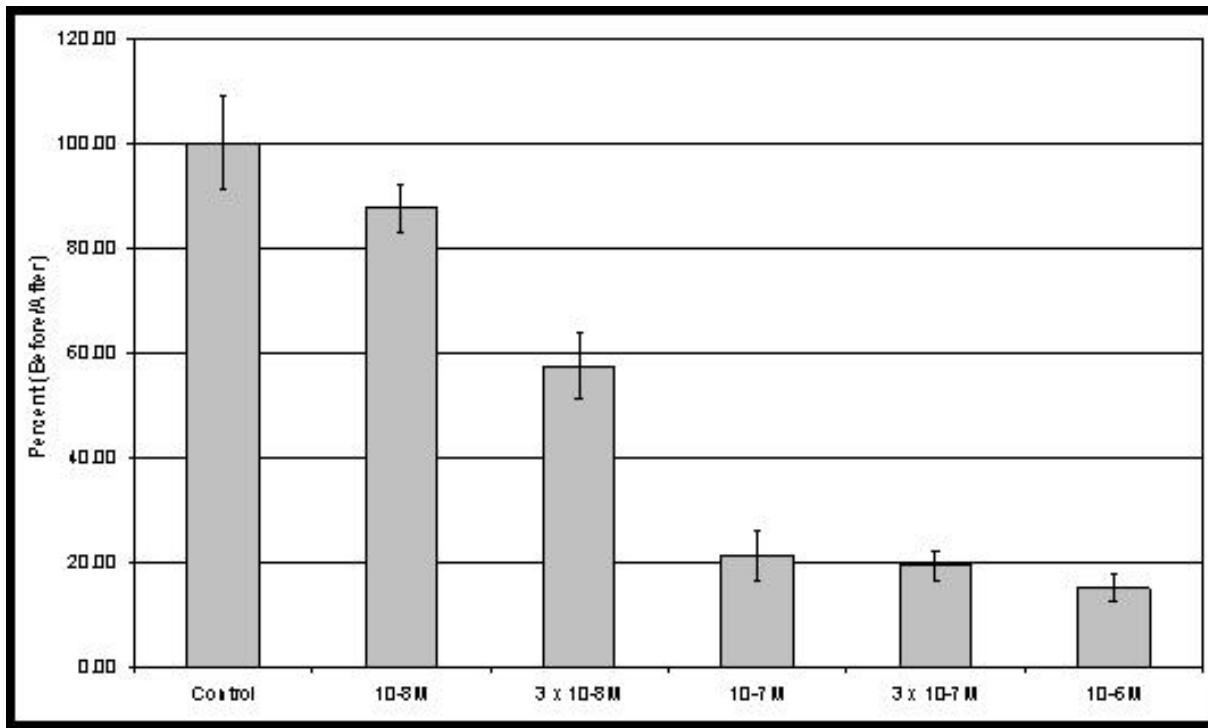
C1 Family – Selectivity is Important

Cathepsin	Tissue Expression	mouse phenotype
B	Ubiquitous	Reduced apoptosis
L	Ubiquitous	Defective CD4 selection, hair loss, increased mortality
V	Thymic	-
K	Osteoclasts	Pycnodynostosis
S	Antigen-presenting cells	Defective antigen presentation
H	Ubiquitous	-
W	CD8+ T cells	-
F	Macrophages, Ubiquitous	-
C	Myeloid cells, Ubiquitous	Hyperkeratosis, periodontitis
O	Ubiquitous	-
Z	Ubiquitous	-

Development of a Human Osteoclast Resorption Assay - Key Cell-based test

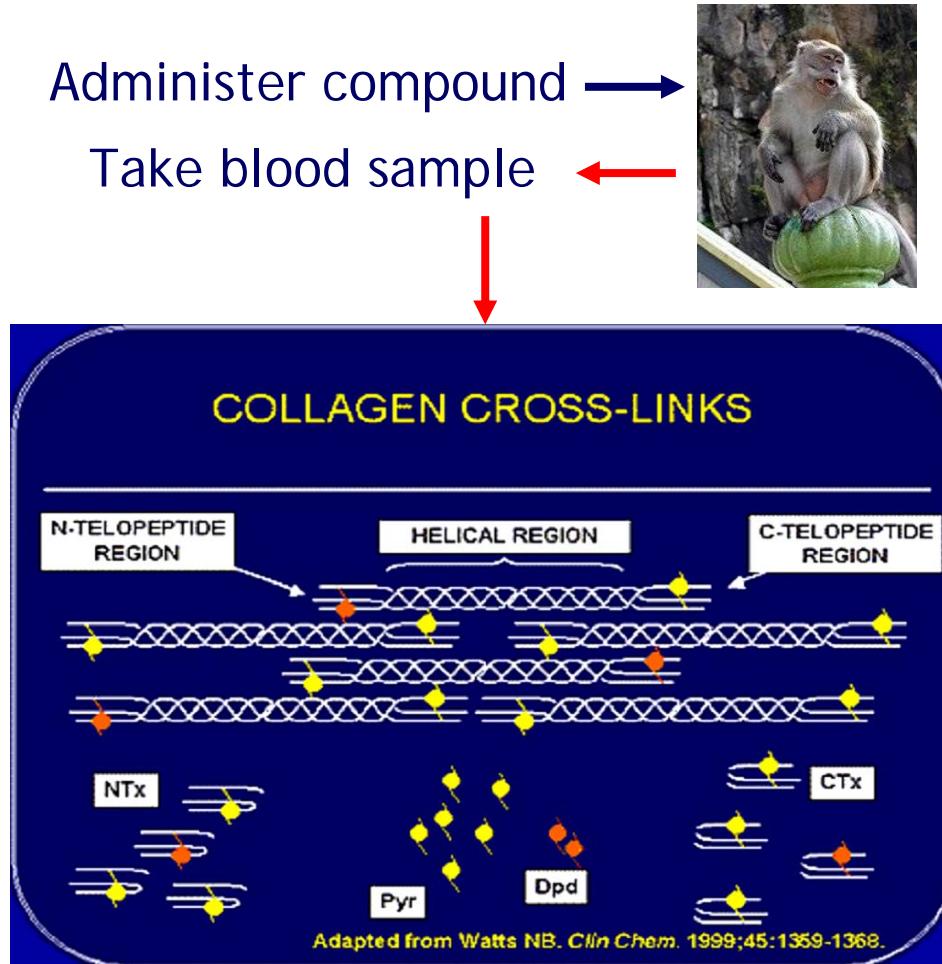


Human Osteoclast Assay



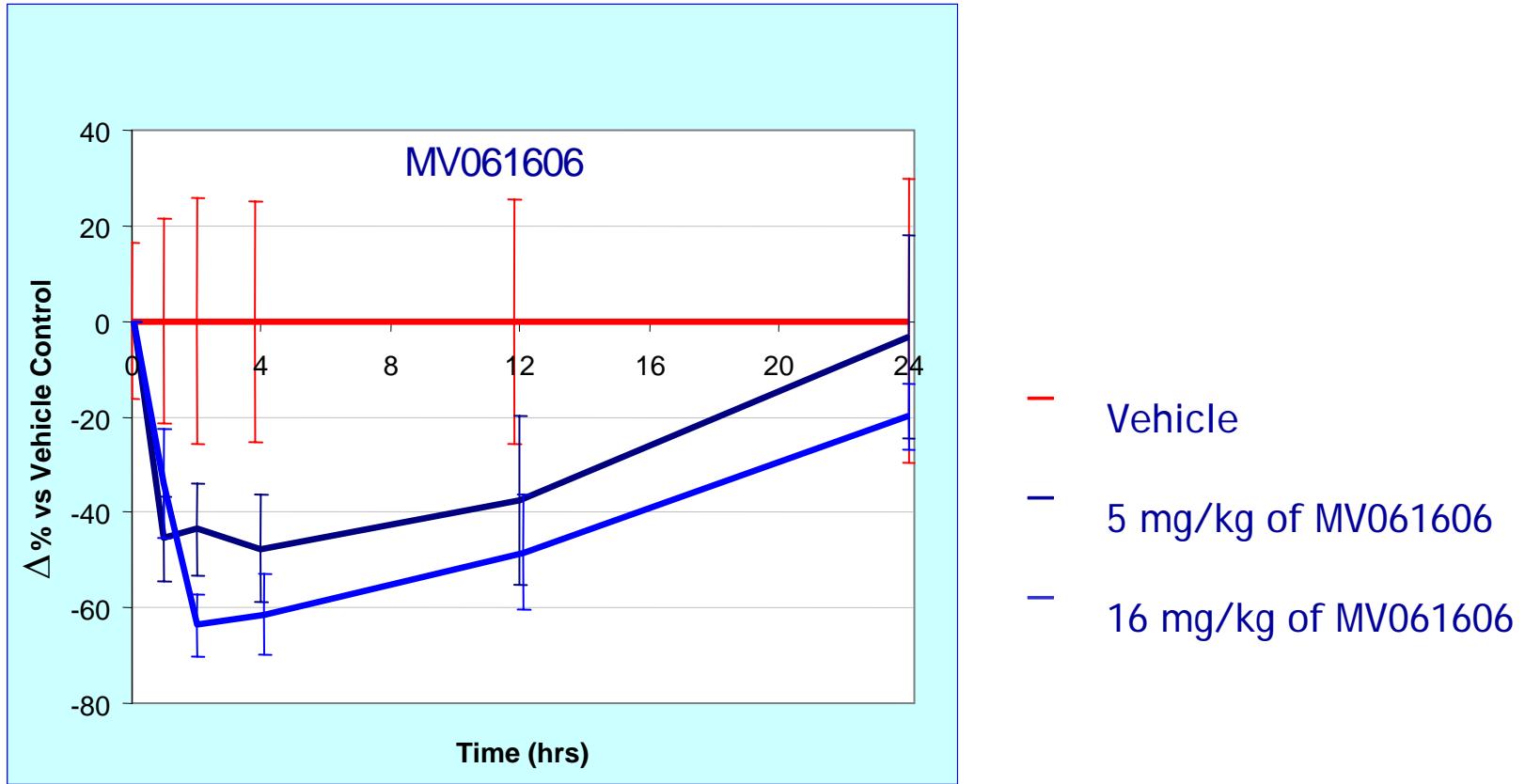
- In-house inhibitors exhibit excellent potency in the human osteoclast resorption assay, coupled with high potency in the isolated enzyme assay and selectivity against related enzymes

In vivo efficacy can be Demonstrated in Cynomolgus: CTx is a bone resorption biomarker



CTX Measurement in intact Monkeys

- Oral dosing in cynomolgus (5 mg/kg and 16 mg/kg)

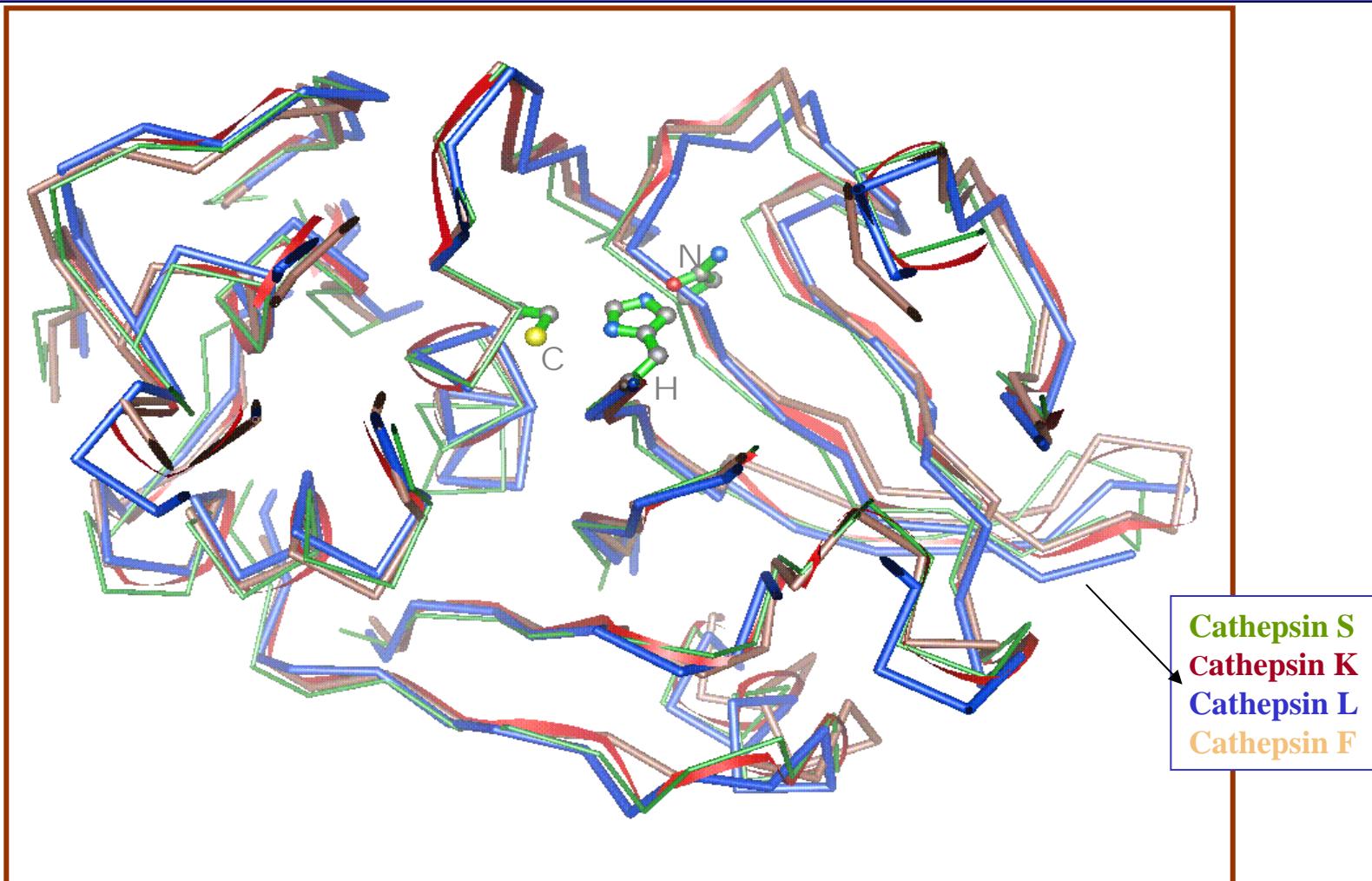


In vivo efficacy demonstrated in monkeys based on biomarkers

Evolution of the Cathepsin K Inhibitor Program at Medivir

- Novel leads provided by screening Medivir's compound libraries and RAPID™
- RAPiD & FOCUS
 - RAPiD generic library
 - >120,000 FRET compounds
 - Abz-B-C-D-E-Tyr(NO₂)-Asp-NH₂
 - Natural/non-natural amino acids, dipeptide mimetics
 - Substrate and inhibitor recognition sequence information obtained
 - Selectivity against other related enzymes
 - Initial SAR
- To date > 2500 inhibitors synthesised in-house providing advanced SAR
 - Fully reversible binding
- X-ray crystal structure information and SBDD have driven synthesis
- Early focus of *in vitro* and *in vivo* DMPK has ensured drug-like properties of the inhibitors developed

Comparison of catalytic domains

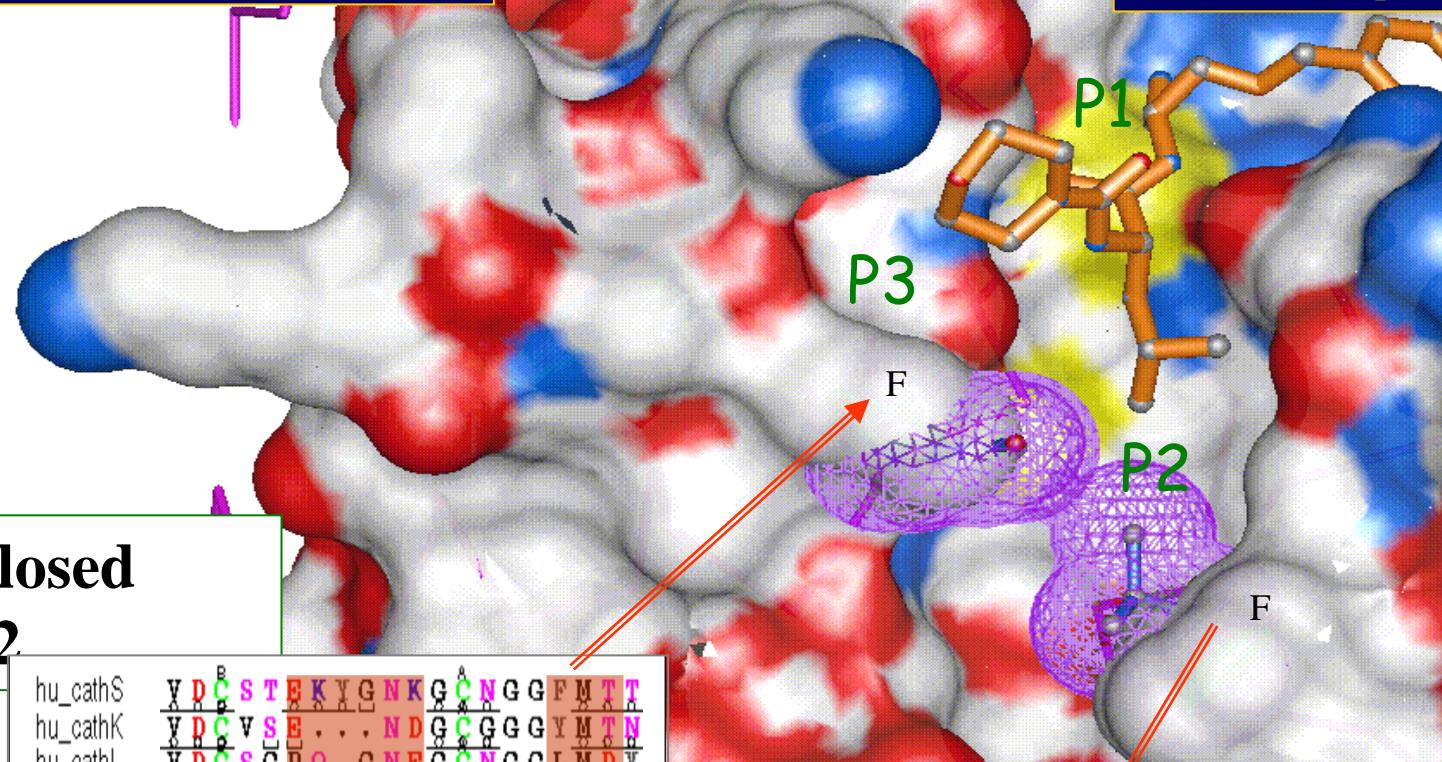


Comparison of the primary sequences Cathepsins S,K,L and F

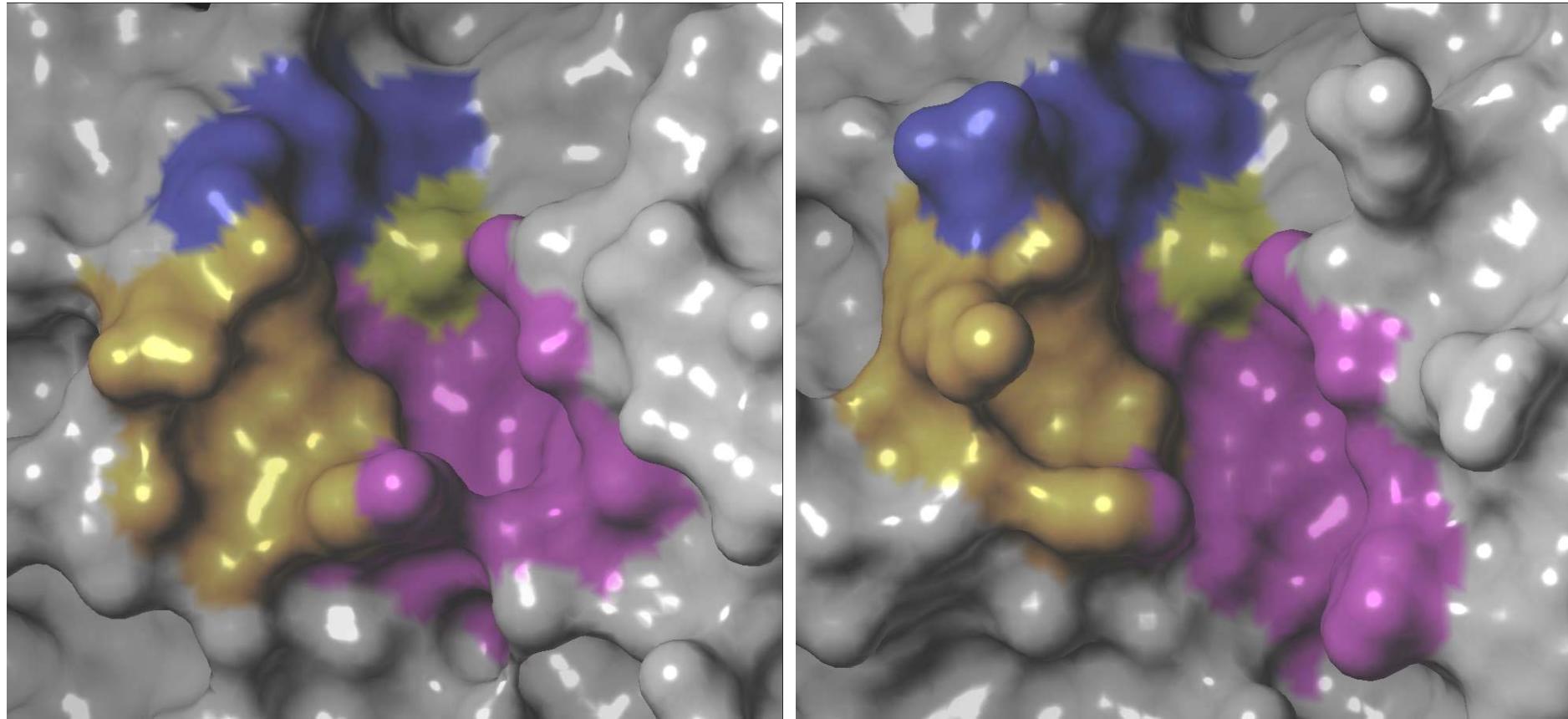
Cathepsins : basis for selectivity

Cathepsin K

Surface Cathepsin S



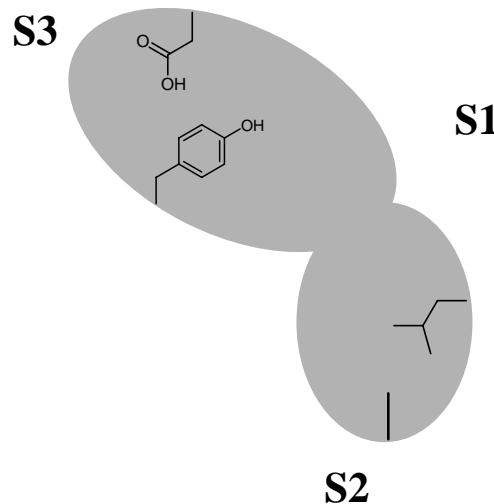
Visualization of Cathepsin K and S Active Sites



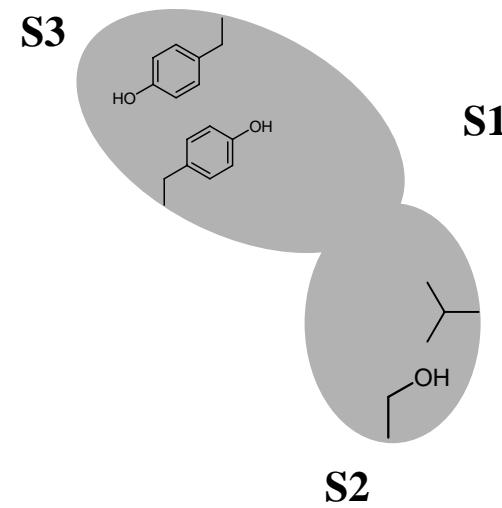
Cathepsin K Enzyme

Amino Acid Residue	S3		S2	
	61	67	134	160
Human	Asp	Tyr		
Rat	Tyr	Tyr	Ala	Leu
			Ser	Val

Human



Rat



- Species differences need to factored in

Cathepsin K selectivity was factored in from the start

RAPiD combinatorial library and
FOCUS software provide SAR and selectivity data from protease pre-screens



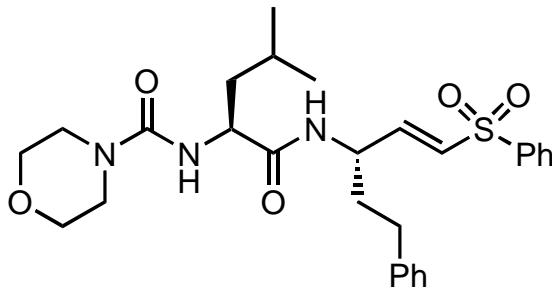
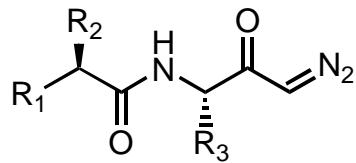
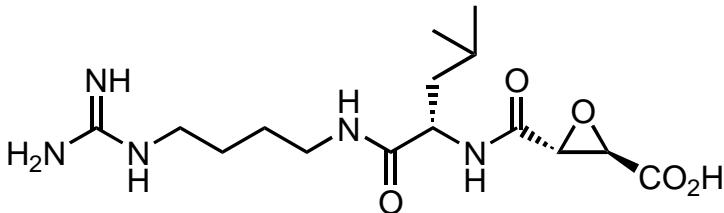
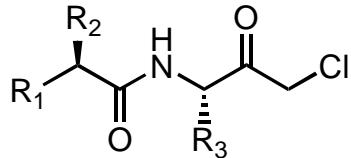
Cathepsin S



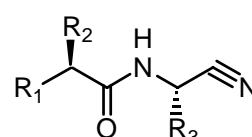
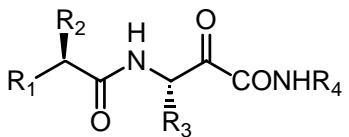
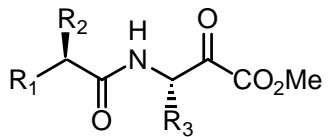
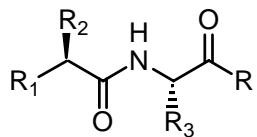
Cathepsin K

Exemplar Classical Cysteine Protease Inhibitors

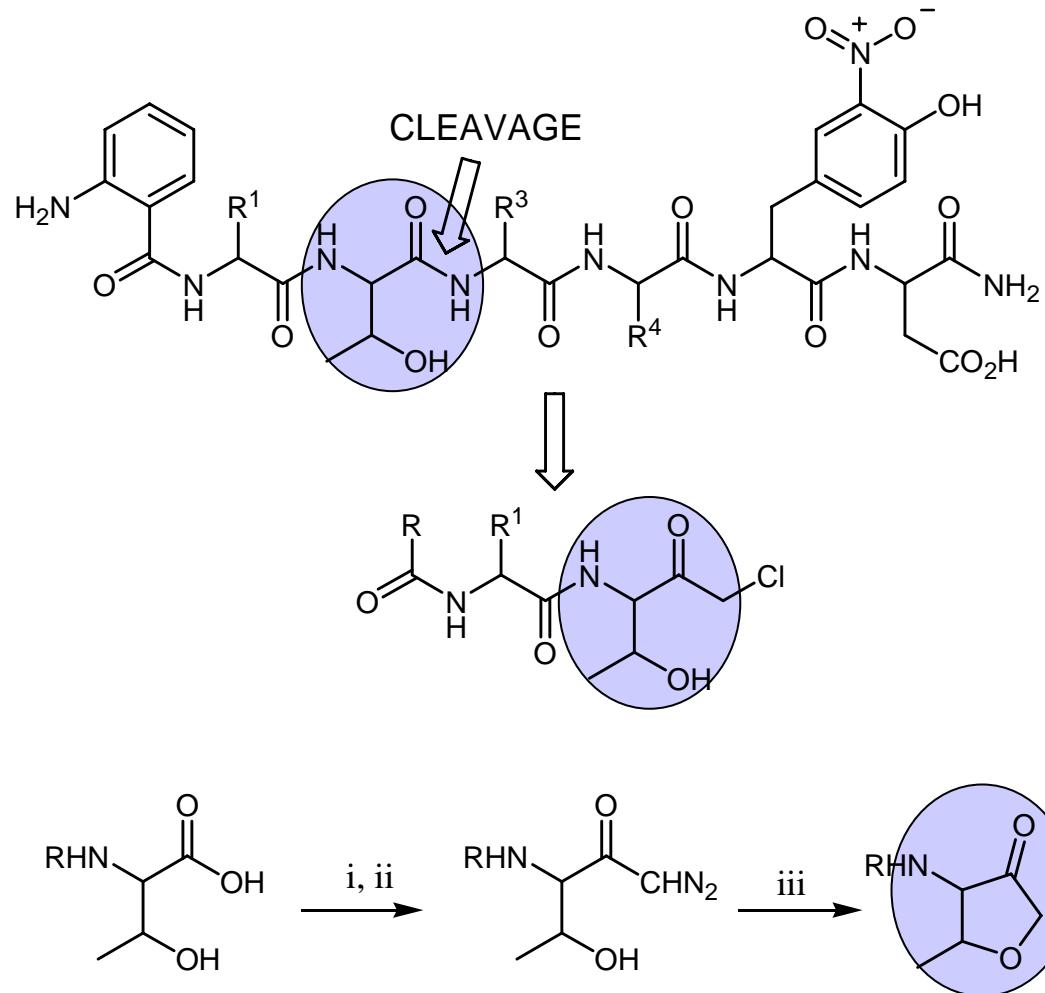
Irreversible



Reversible



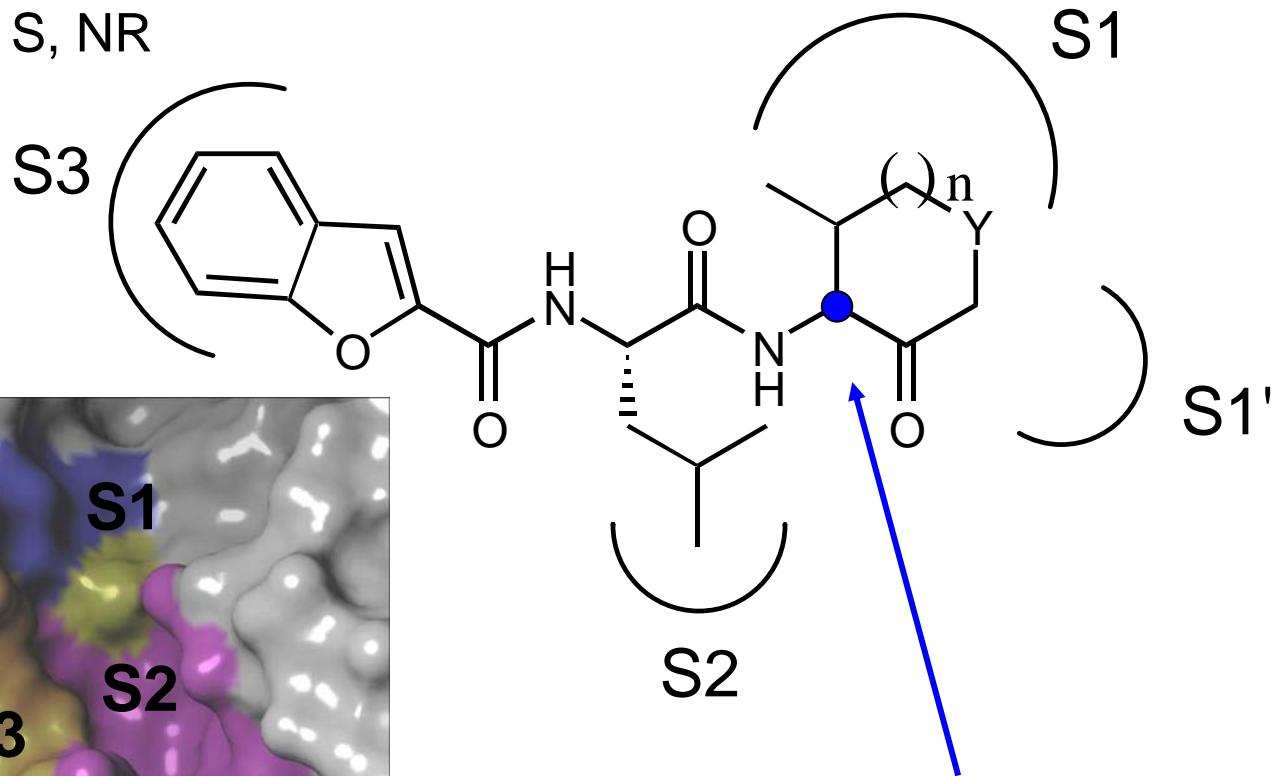
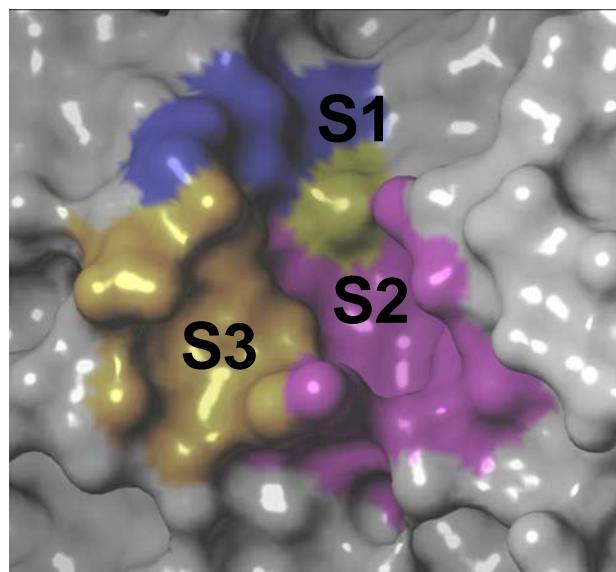
Cathepsin K Inhibitor Design



SAR Modifications

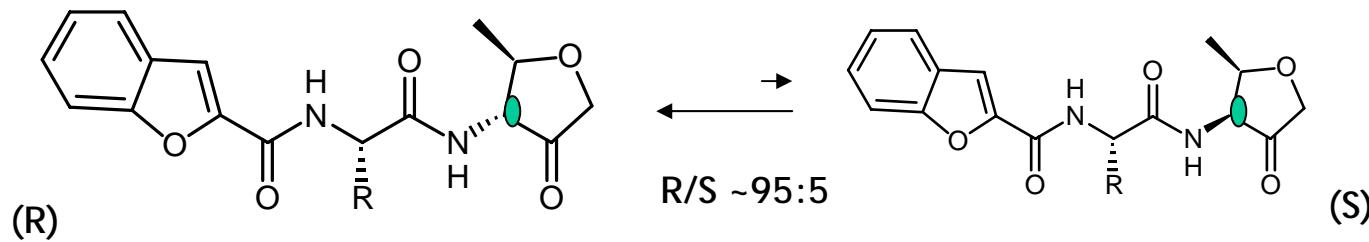
$n = 0, 1, 2 \text{ or } 3$

$Y = O, S, NR$



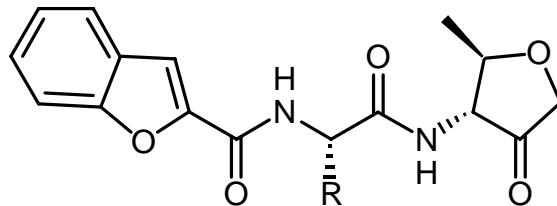
Potential development issue: chirally labile position

The (S)-Isomer is the More Active



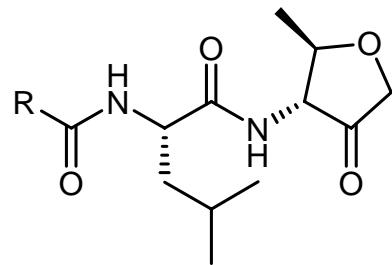
	R =	Cath K (K_i μM)	Cath L (K_i μM)
(S)		0.004	0.52
(R)		0.020	0.75

P2 Modifications



R =	Cath K (K_i μM)	Cath L (K_i μM)
	0.3	> 100
	0.02	0.75
	2.12	9.41
	0.82	4.24

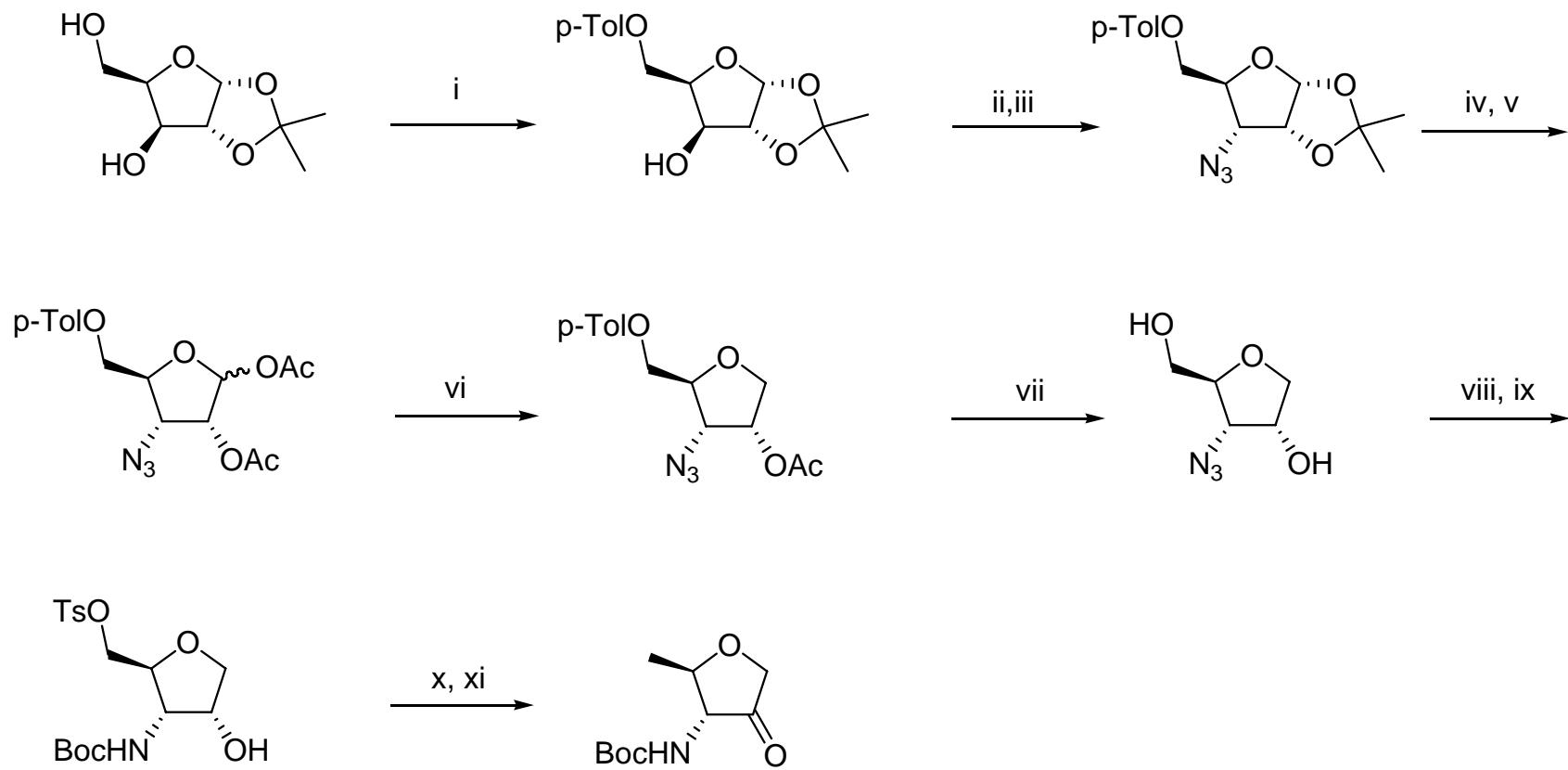
P3 Modifications



R =	Cath K (K_i μM)	Cath L (K_i μM)
	9.5	0.35
	0.13	0.67
	7.6	0.76
	0.02	0.75

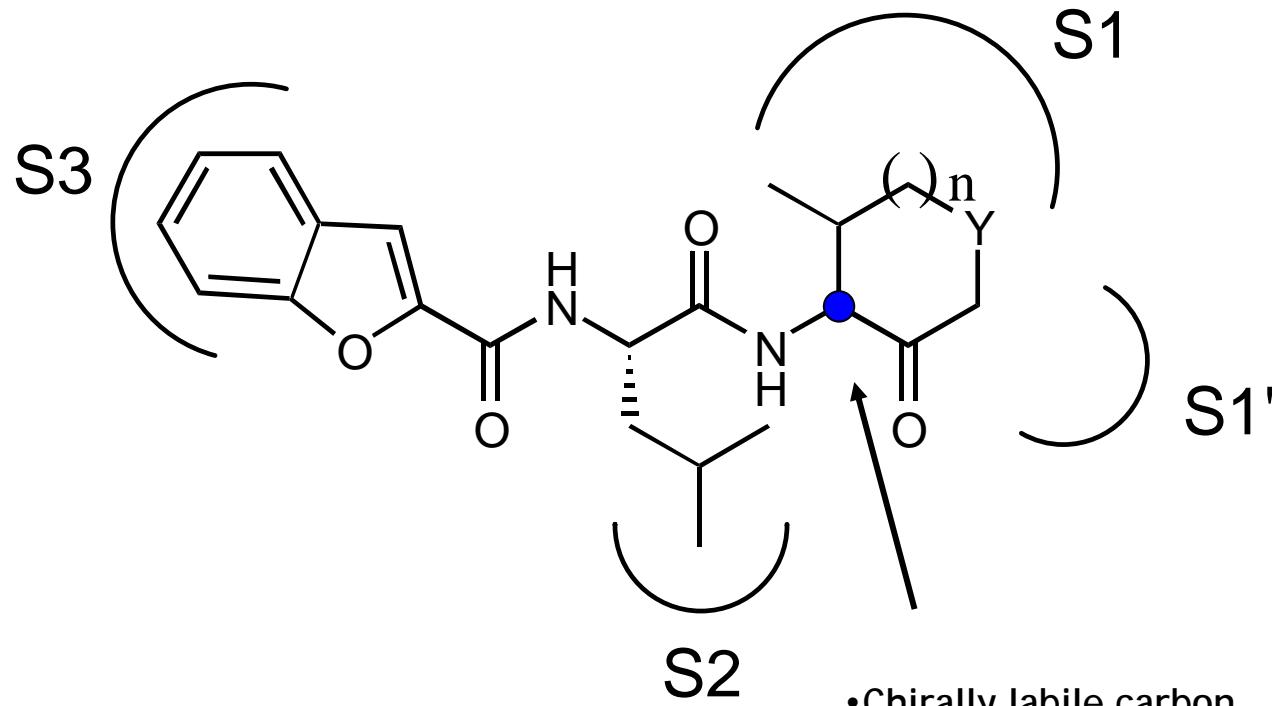
The benzofuran P2 capping group provides good potency and selectivity

Synthesis of the P1 Furanone Template



i. *p*-TolCl, pyridine, 72 % yield; ii. Tf₂O, pyridine, ; iii. NaN₃, 30 % yield over two steps; iv. 75% HCO₂H;
v. Ac₂O, pyridine, 74 % yield over two steps; vi. TMSOTf, Et₃SiH, 50 % yield; vii. K₂CO₃, MeOH, 80 %
yield; viii. H₂/Pd/C, Boc₂O, ix. TsCl, pyridine, 95 % over two steps; x. LiAlH₄; xi. Dess Martin
Periodinane, 70 % yield over two steps

Design of New P1 groups

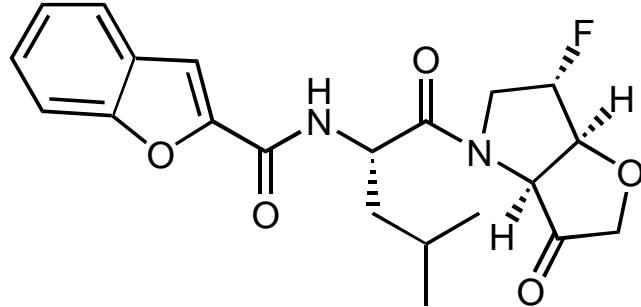


$n = 0, 1, 2 \text{ or } 3$

$Y = O, S, NR$

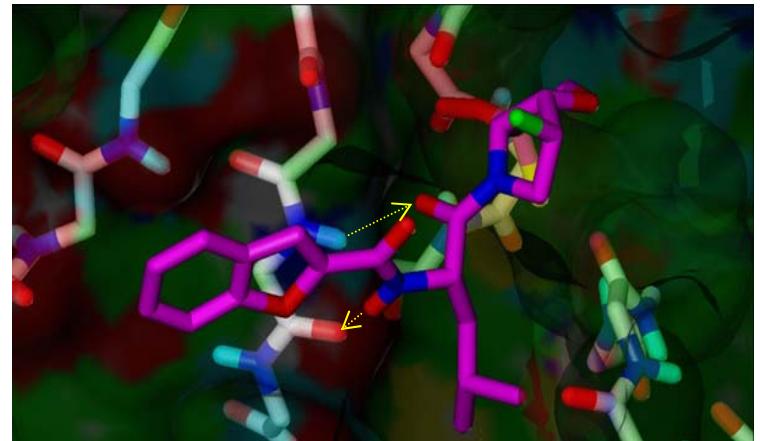
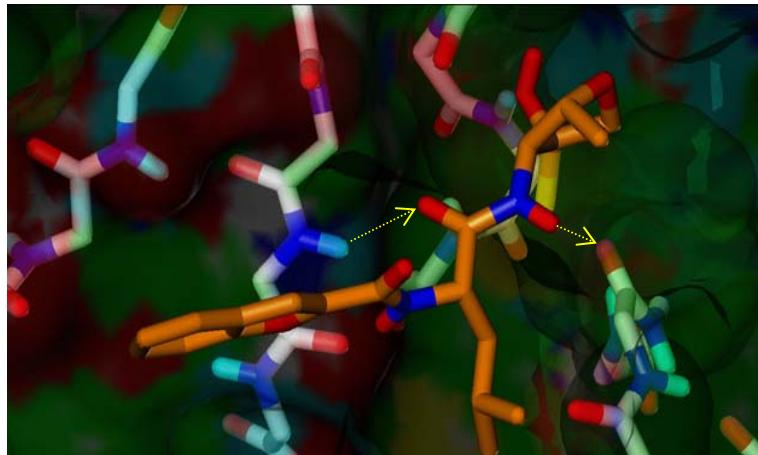
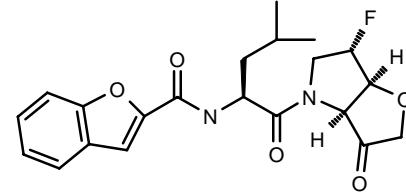
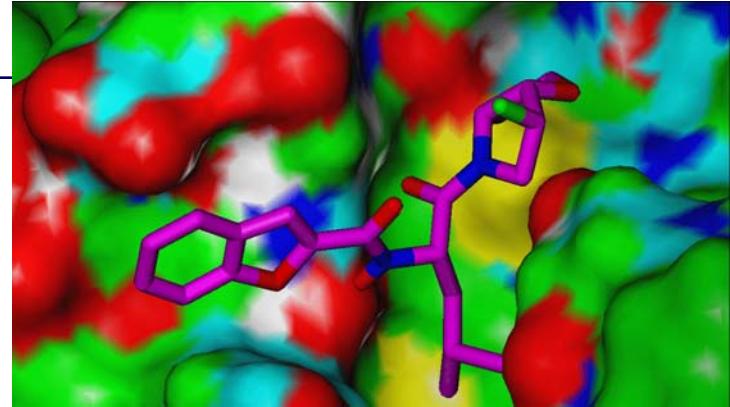
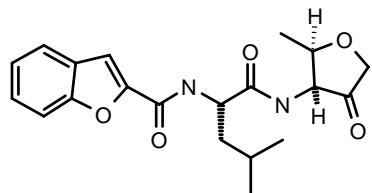
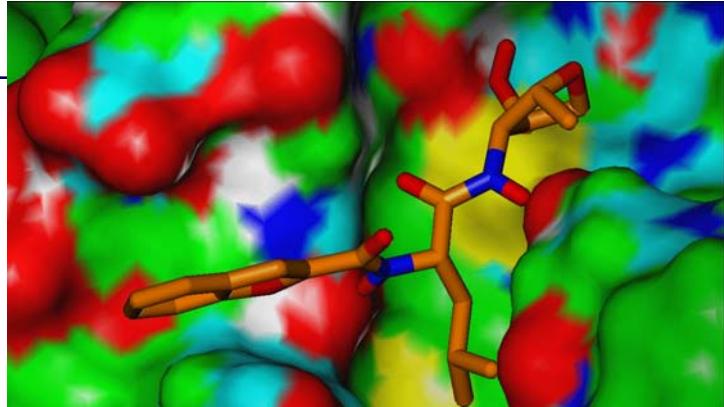
- Chirally labile carbon
- Chirally stable P1 templates desirable
- Should retain potency/DMPK properties

Example of Non-epimerizable P1 Groups



- A P1 fused 5:5 ring system designed
- Maintains the preferred (*S*)-chirality
- Chirally stable

Will it Bind? - Docking of Methyl Furanone & Fluorobicycle Ligands to Cathepsin K

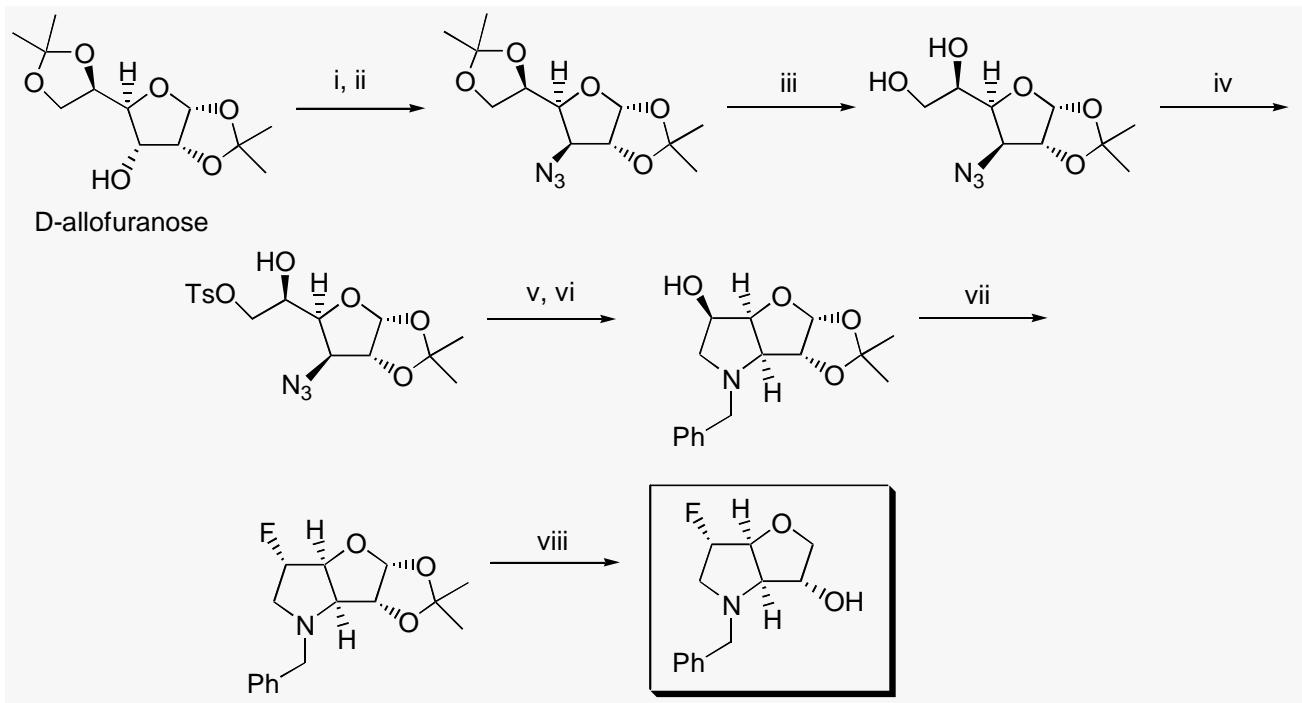


- The P1-P2 amide N-H hydrogen bond to the enzyme is lost while the overall placement and direction of the amide remains similar

Overlay of Bound Conformation of Methyl-Furanone and Fluorobicycle P1s

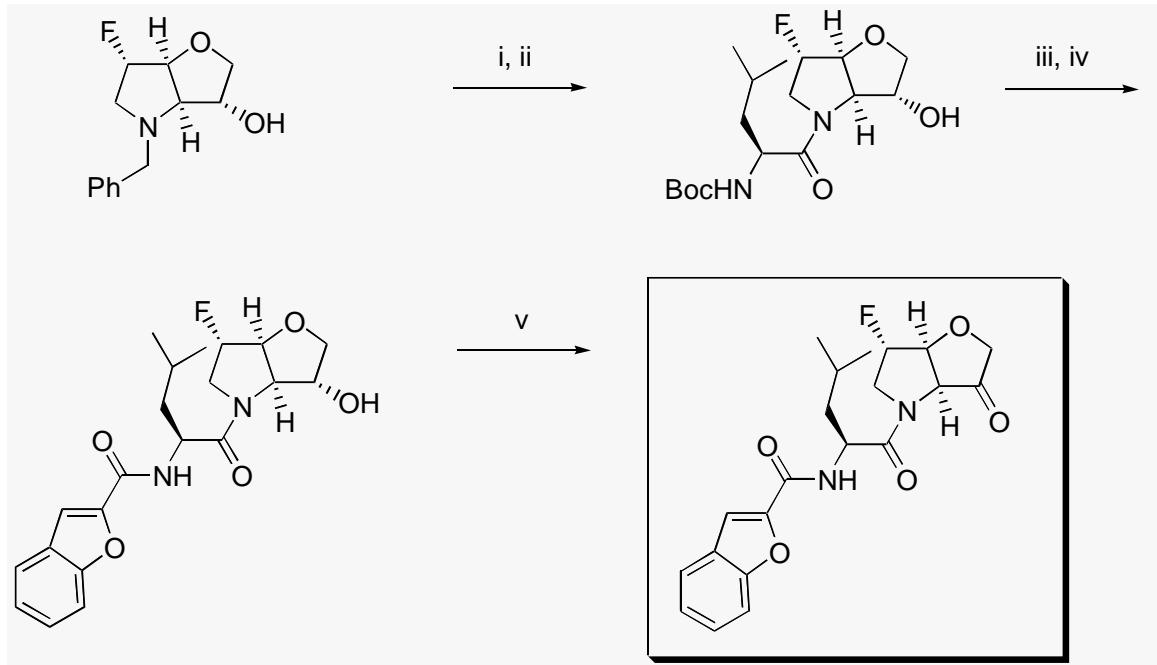


Synthesis of Novel P1 Bicyclic Template



- i. Tf_2O , pyridine; ii. NaN_3 , 60 % yield over two steps; iii. 70 % AcOH , Δ , 80 % yield; iv. $p\text{-TsCl}$, pyridine, 72 % yield; v. 10 % Pd on C, H_2 ; vi. PhCHO , NaCNBH_3 , 74 % yield over two steps; vii. Deoxofluor, pyridine, 65 % yield; viii. a. TFA, H_2O ; b. Ac_2O , pyridine; c. HBr , AcOH ; d. LiAlH_4 , 62 % yield over four steps

Synthesis of a Cathepsin K Inhibitor



i. 10 % Pd on C, H₂; ii. Boc-LeuOH, WSC, HOBr, NMM, 85 % yield over two steps; iii. TFA, H₂O; iv. 2-Benzofurancarboxylic acid, WSC, HOBr, NMM, 80 % yield over two steps; v. Dess-Martin periodinane, 65 % yield

- High potency displayed: K_i value of 2.6 nM
- > 100 fold selectivity over related Cathepsins
- Good overall DMPK properties
- Very promising inhibitor class

Clinical Pipeline of Cathepsin K Inhibitors

	Merck	P&G/Sanofi-Aventis*	GSK	Novartis	Eli Lilly
Launched	Fosamax Bisphoshonate	Actonel Bisphoshonate	Boniva Bisphoshonate	Zometa Bisphoshonate	Evista SERM
Pipeline	L-001037536 Cathepsin K, P I		Relacatib Cathepsin K, P I	Aclasta Bisphosphonate, P III	
	c-3578 α -v / β -3 antagonist, P I		NPS-423562 PTH, P I	AAE-581 Cathepsin K, P II	
			NPS-2143 PTH, P I	AFG -495 Cathepsin K, suspended	

* Sanofi-Aventis: ongoing preclinical cat K program

- Several pharmaceutical companies with a major osteoporosis franchise are now developing cathepsin K inhibitors

Acknowledgements

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Soyfur Miah
Shirley Rahman

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& Chem Informatics
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Kevin Parkes
Jose Gallego
Katarina Jansson
Peter Lind

Vice President Discovery Research

Bertil Samuelsson

Professor Tim Chambers

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