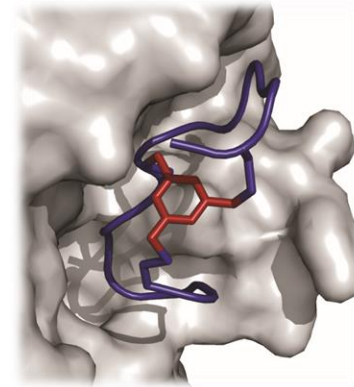
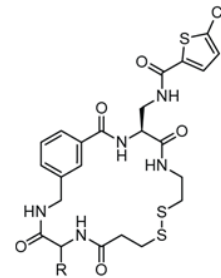
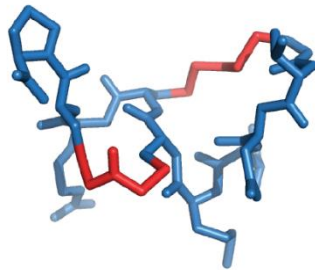


Cyclic peptides and high-throughput technologies for addressing unmet medical needs



My background

drug development technologies

(bi)cyclic peptides macrocycle drugs

phage display ligand development combinatorial chemistry

high-throughput screening

My background

drug development technologies

(bi)cyclic peptides macrocycle drugs

phage display ligand development combinatorial chemistry

high-throughput screening

diseases & targets

protease targets

plasma kallikrein, FXII,
FXI, thrombin, ...

thrombotic
diseases

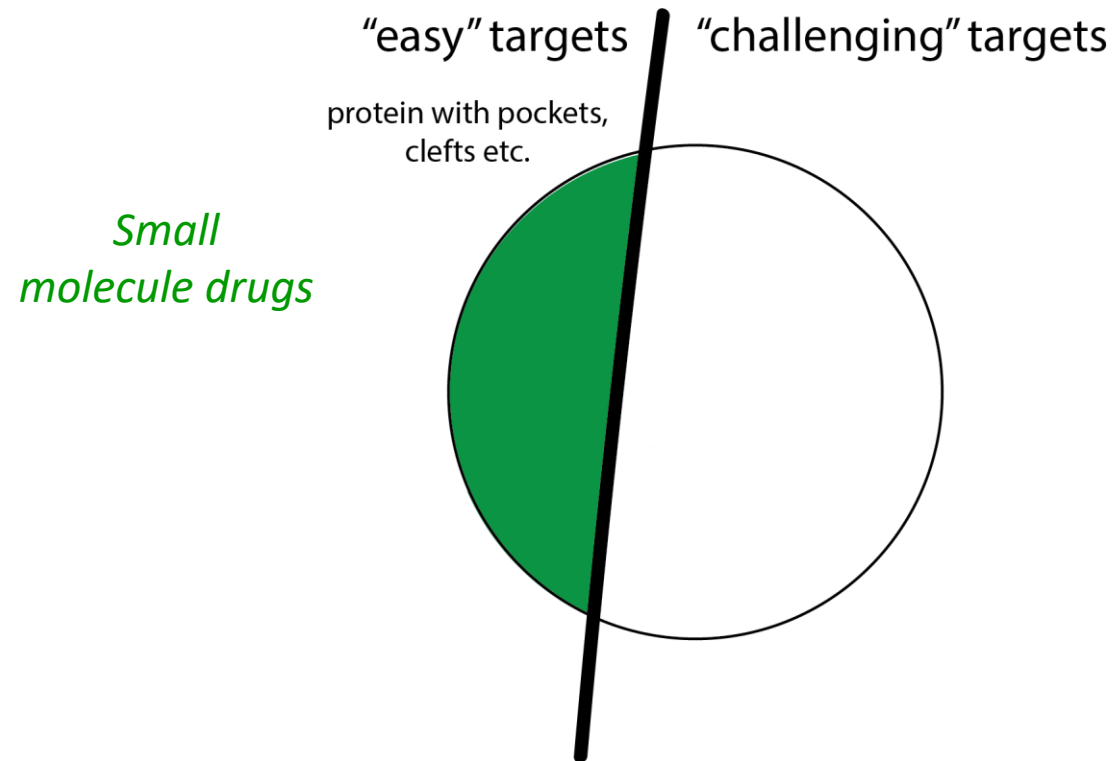
protein-protein
interactions

IL-23R, KRAS,
 β -catenin, Keap1, ...

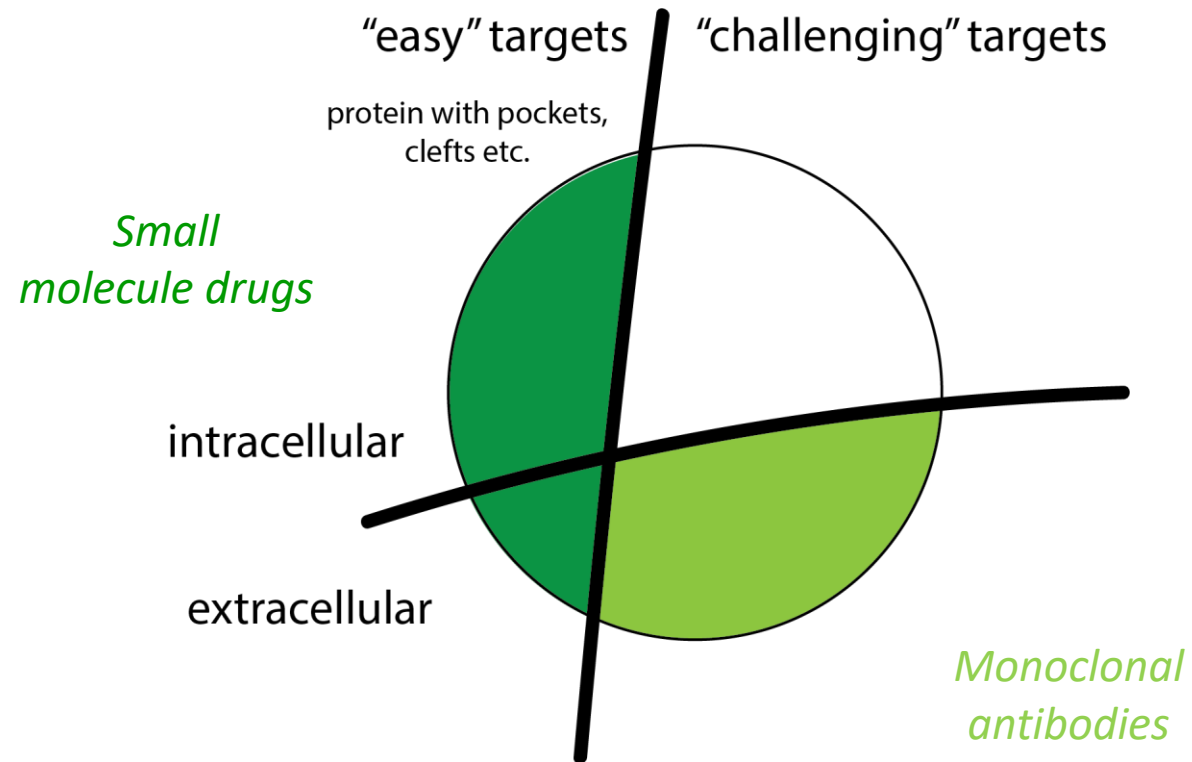
cancer



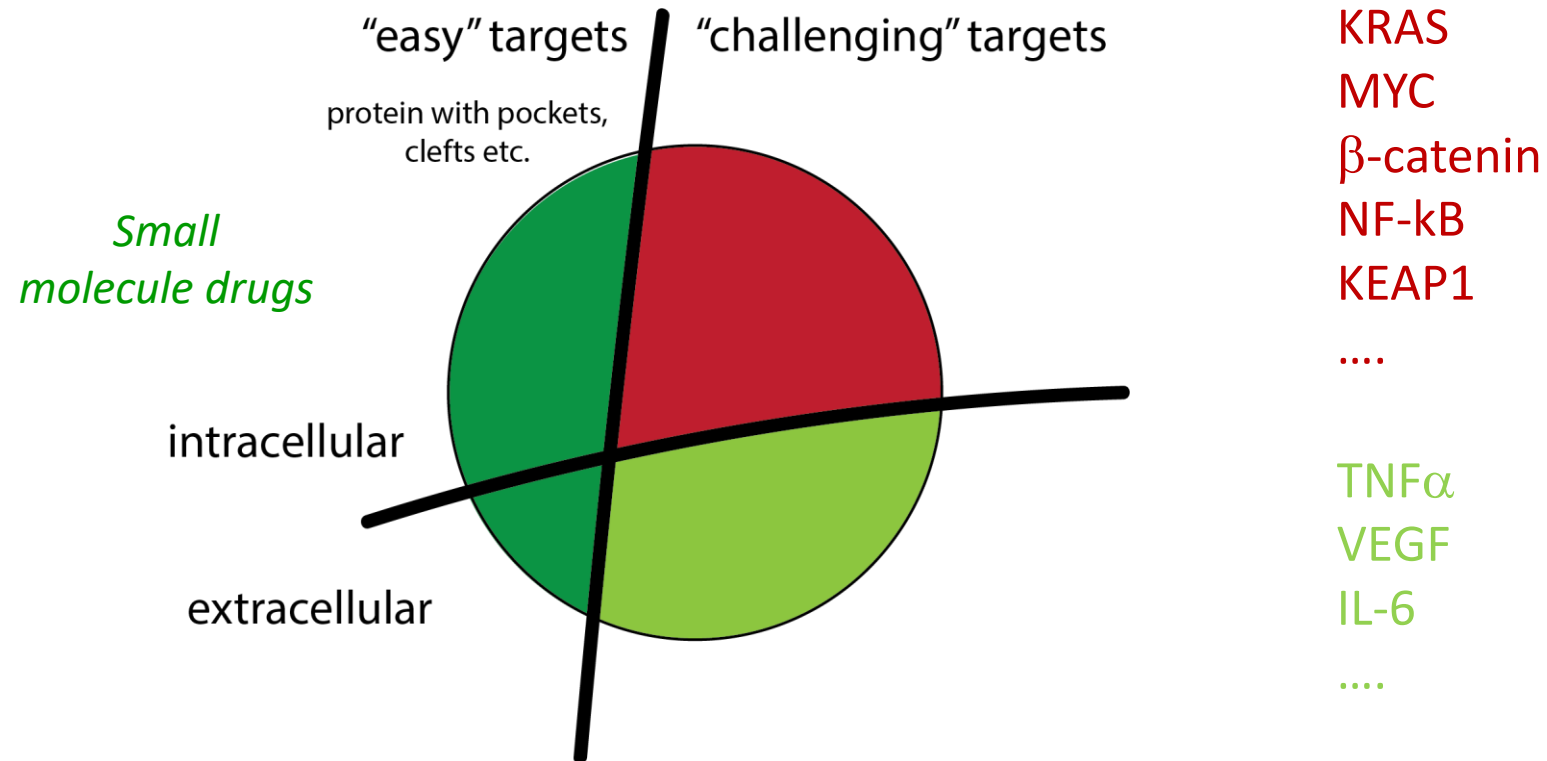
Why new drug modalities / technologies?



Why new drug modalities / technologies?

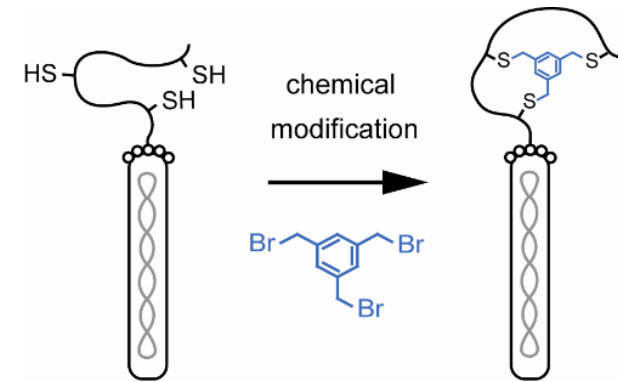
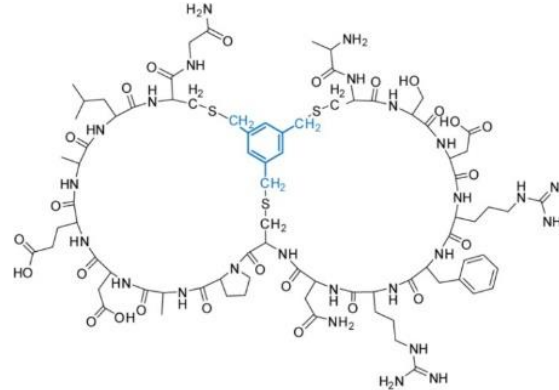


Why new drug modalities / technologies?

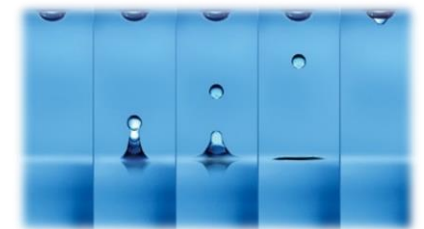
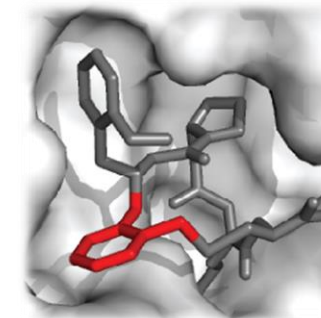
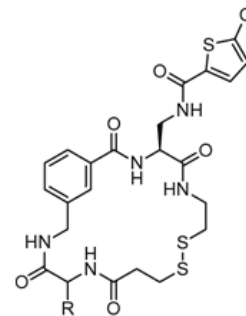


Talk overview

- # 1. Bicyclic peptides developed by phage display

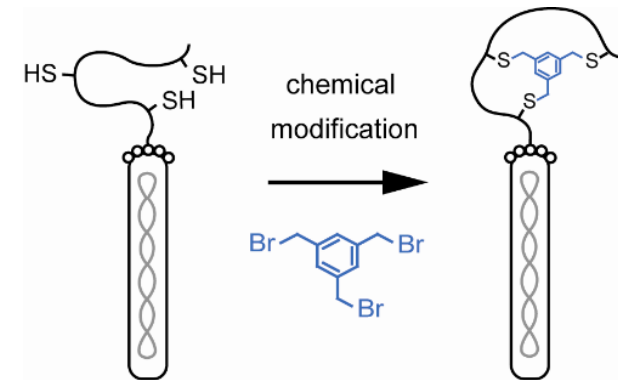
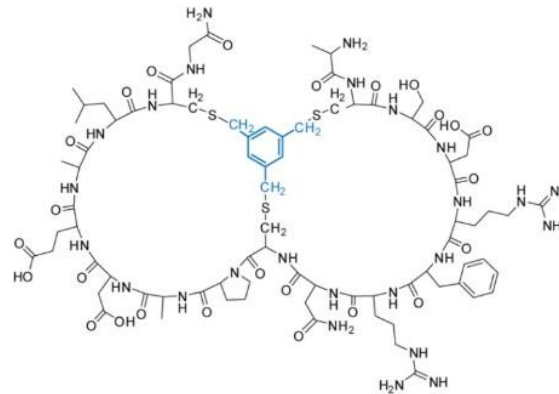


2. Peptide macrocycles developed by high-throughput synthesis at a nanomole scale

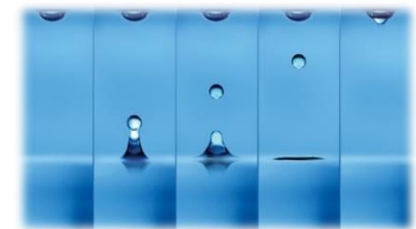
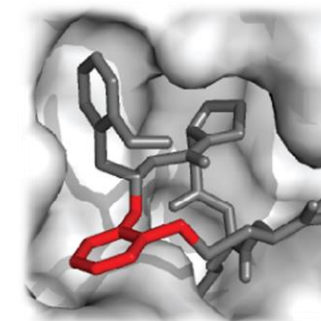
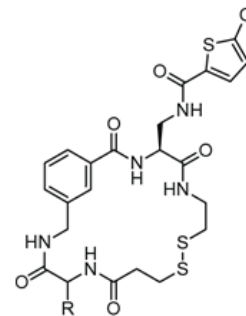


Talk overview

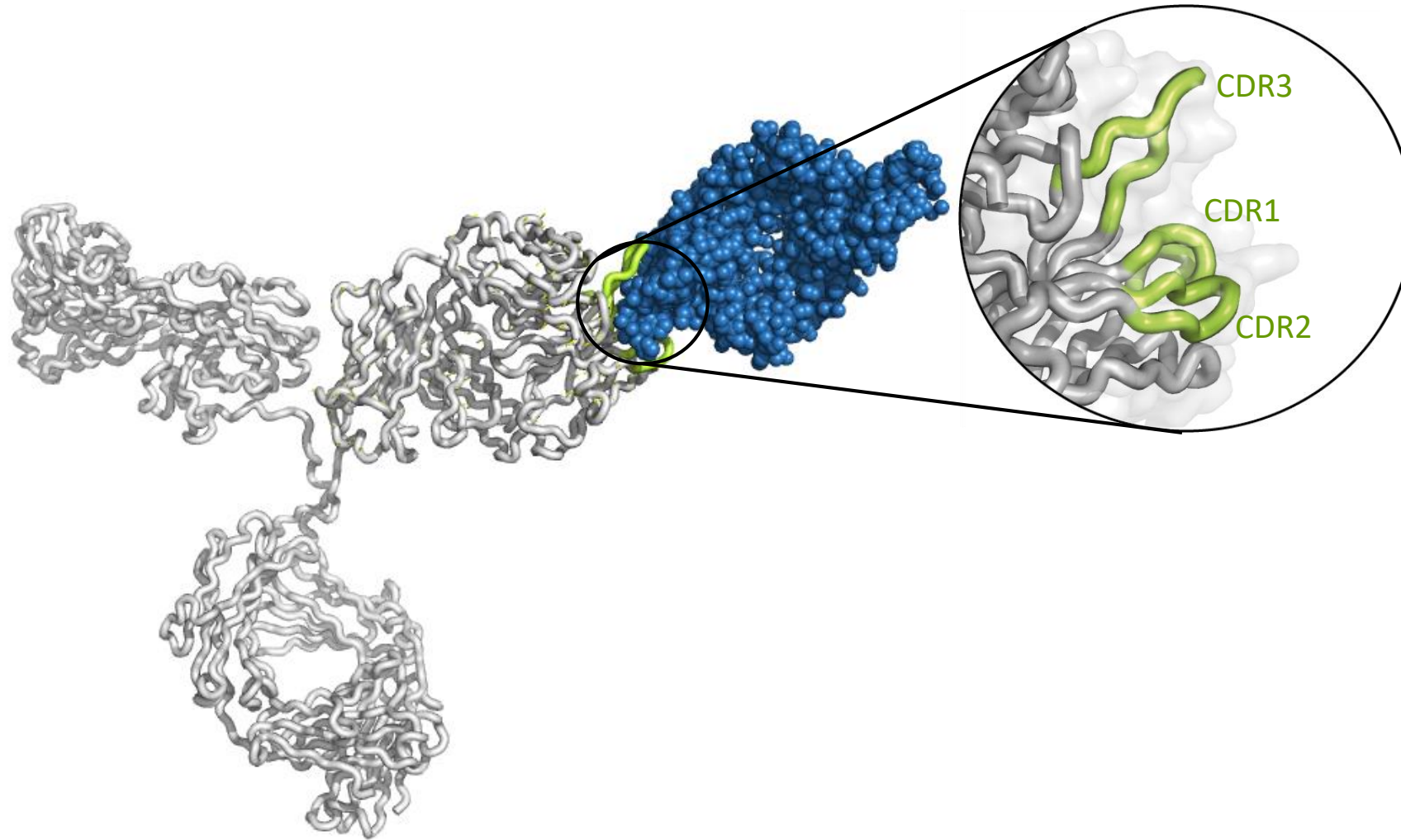
1. Bicyclic peptides developed by phage display



2. Peptide macrocycles developed by high-throughput synthesis at a nanomole scale



Miniaturization of antibodies



Miniaturization of antibodies

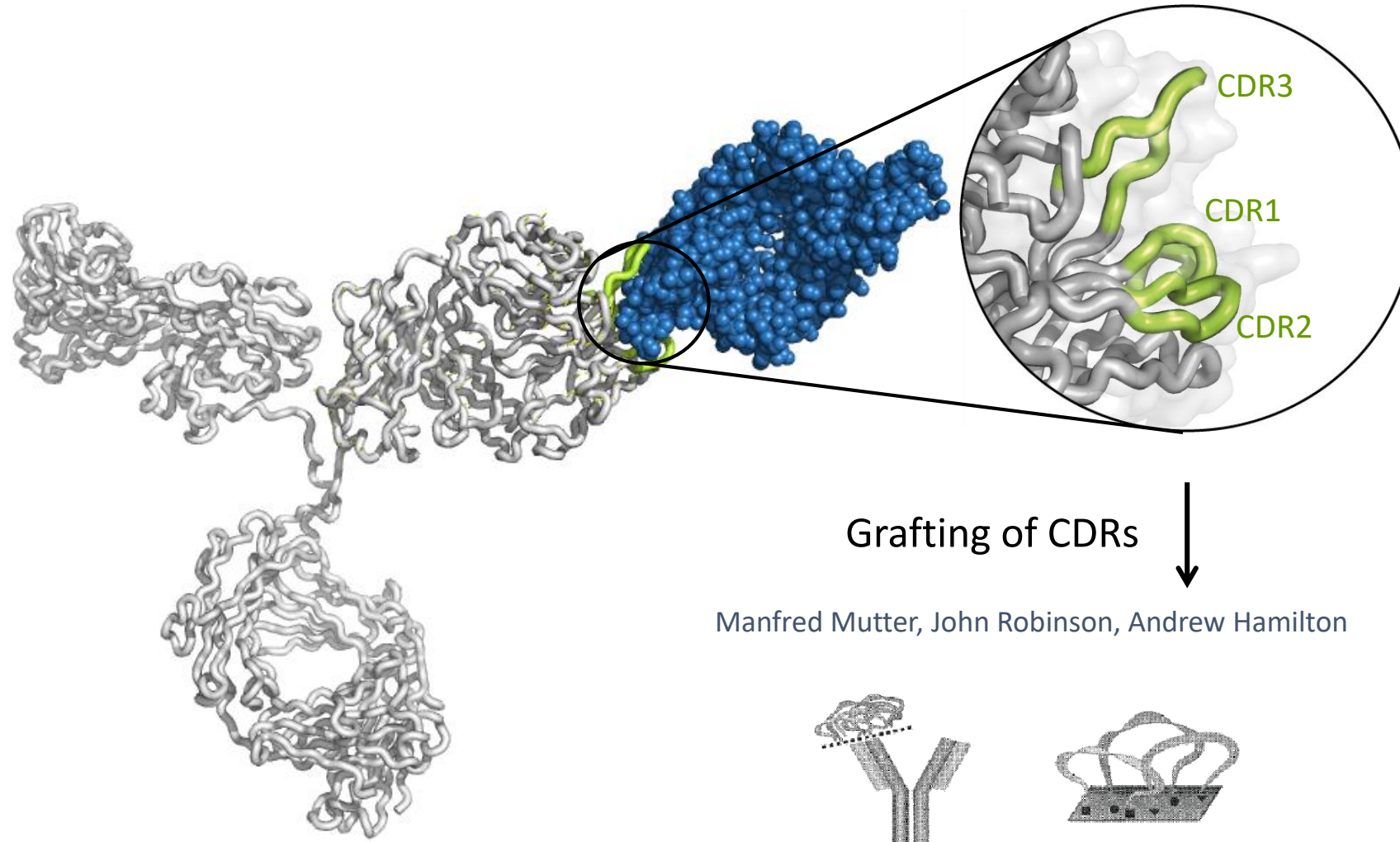
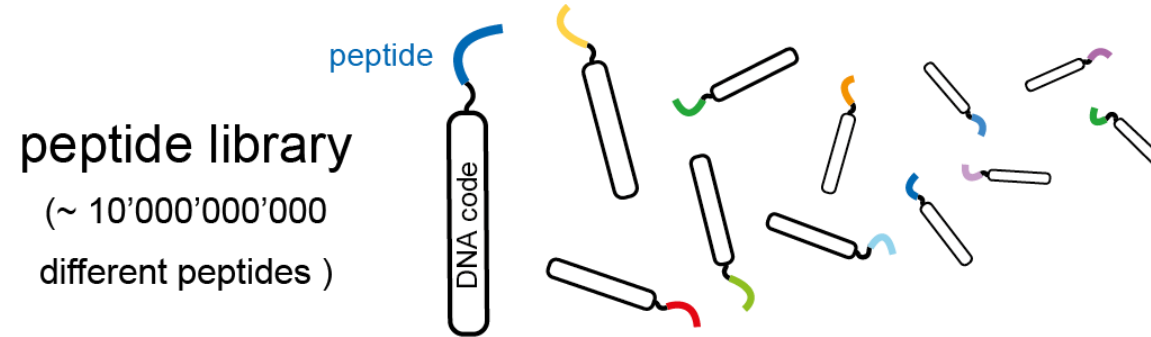


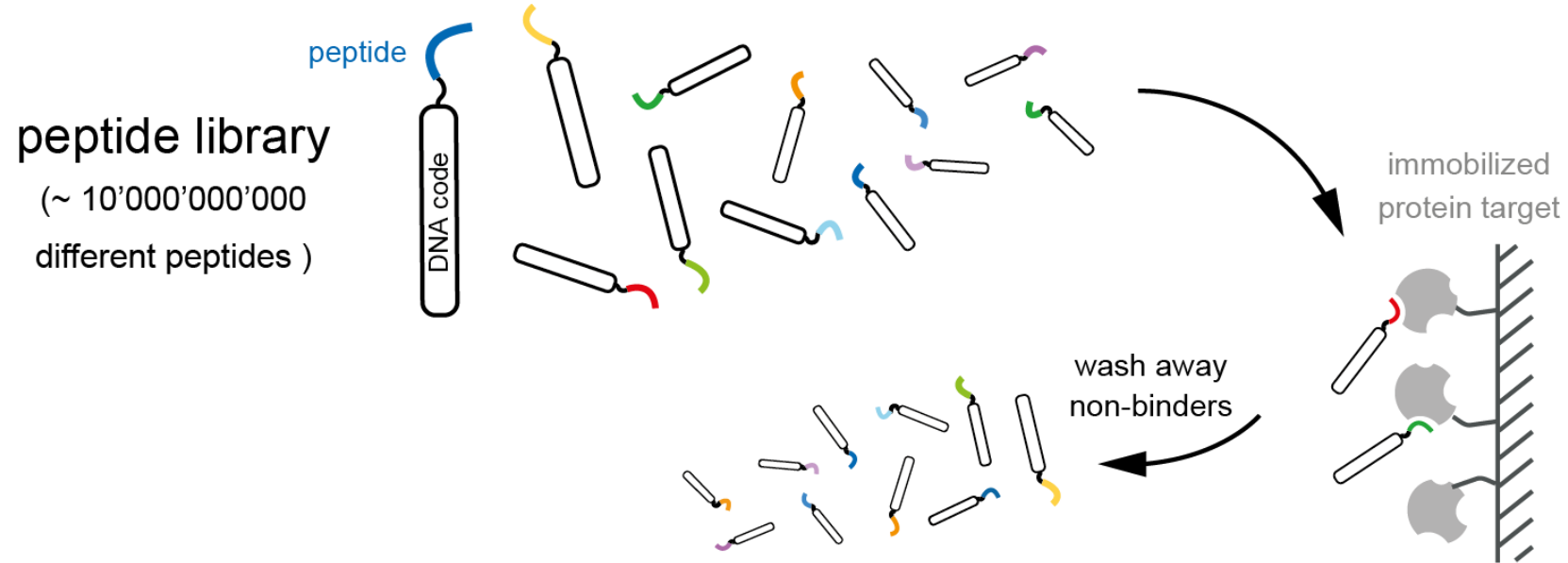
Figure from Mutter, *et al.*, *Angewandte*, 1996)

Phage display technology



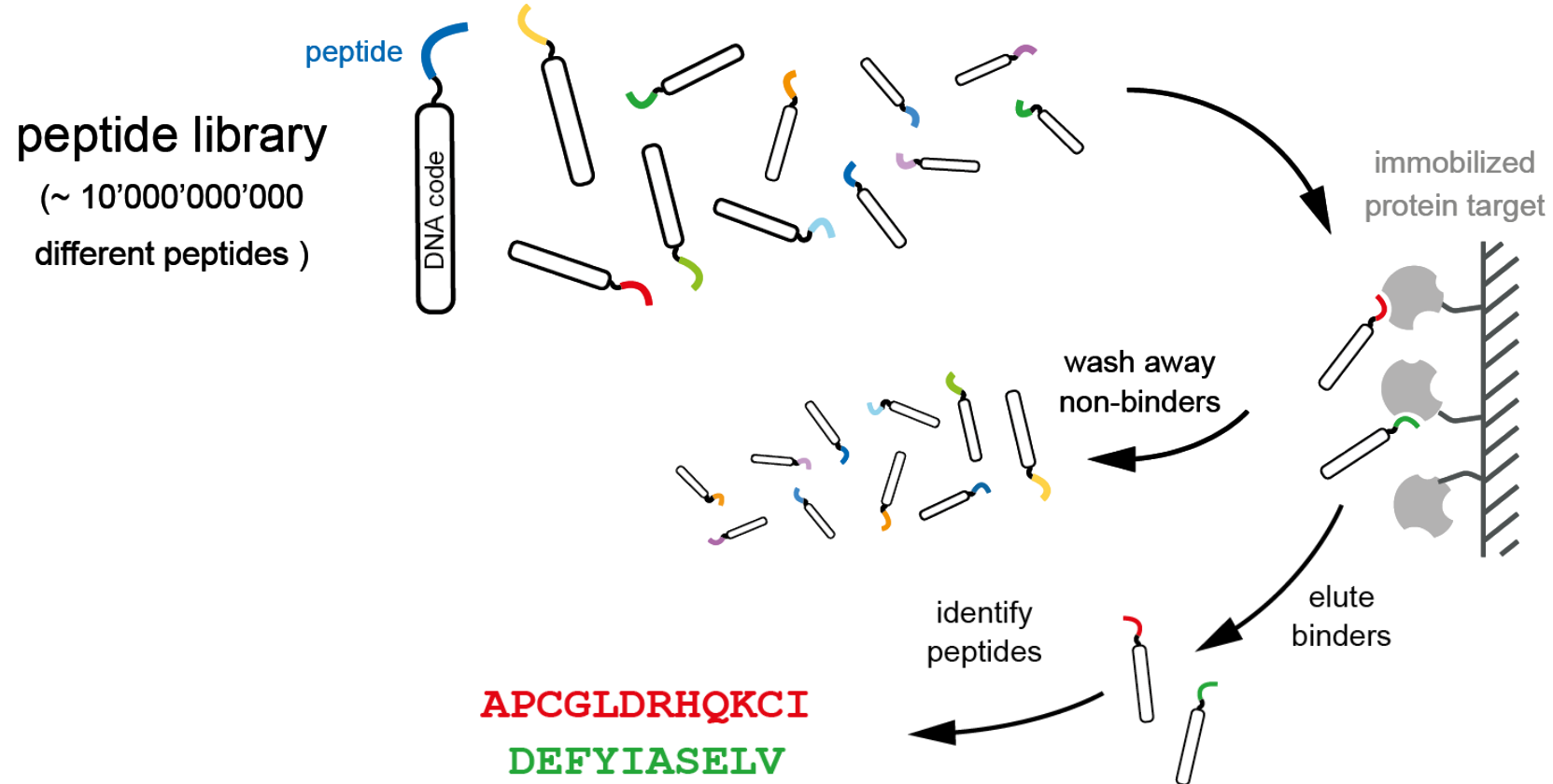
Smith, G.P., *Science*, 1985

Phage display technology



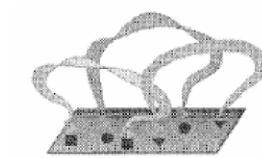
Smith, G.P., *Science*, 1985

Phage display technology

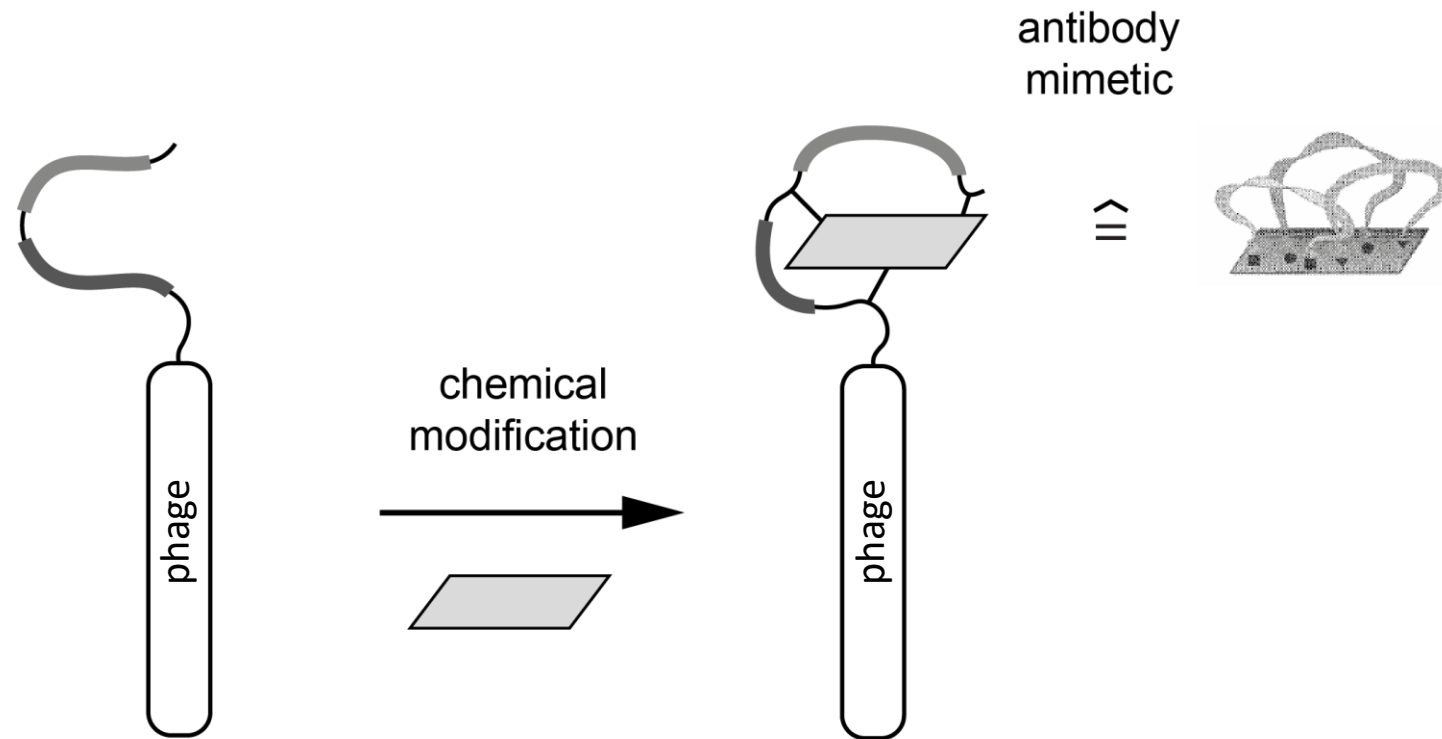


Smith, G.P., *Science*, 1985

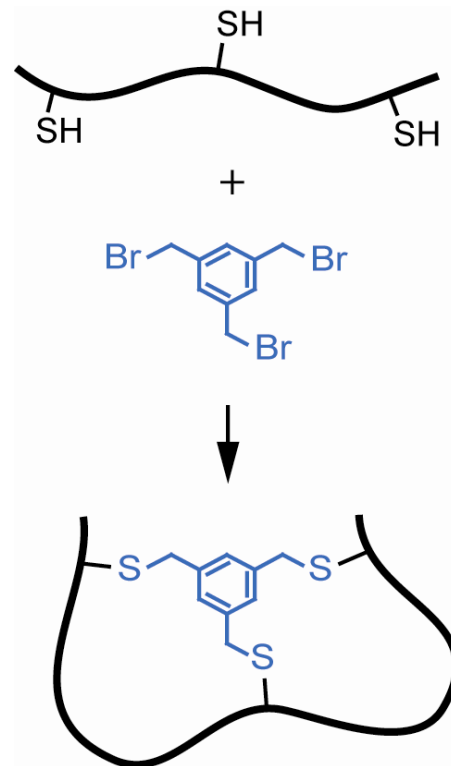
Phage-encoded antibody mimetics



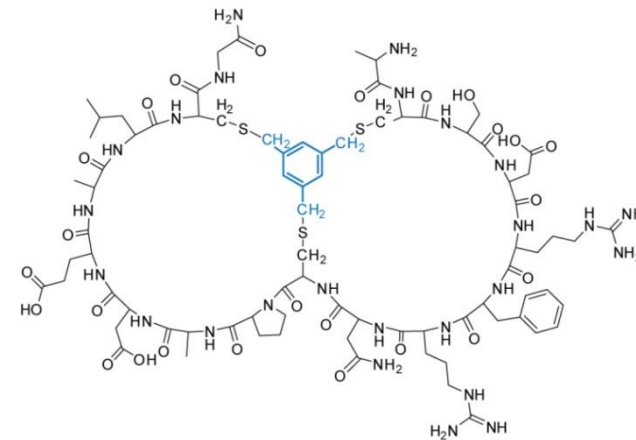
Phage-encoded antibody mimetics



Bicyclic peptides

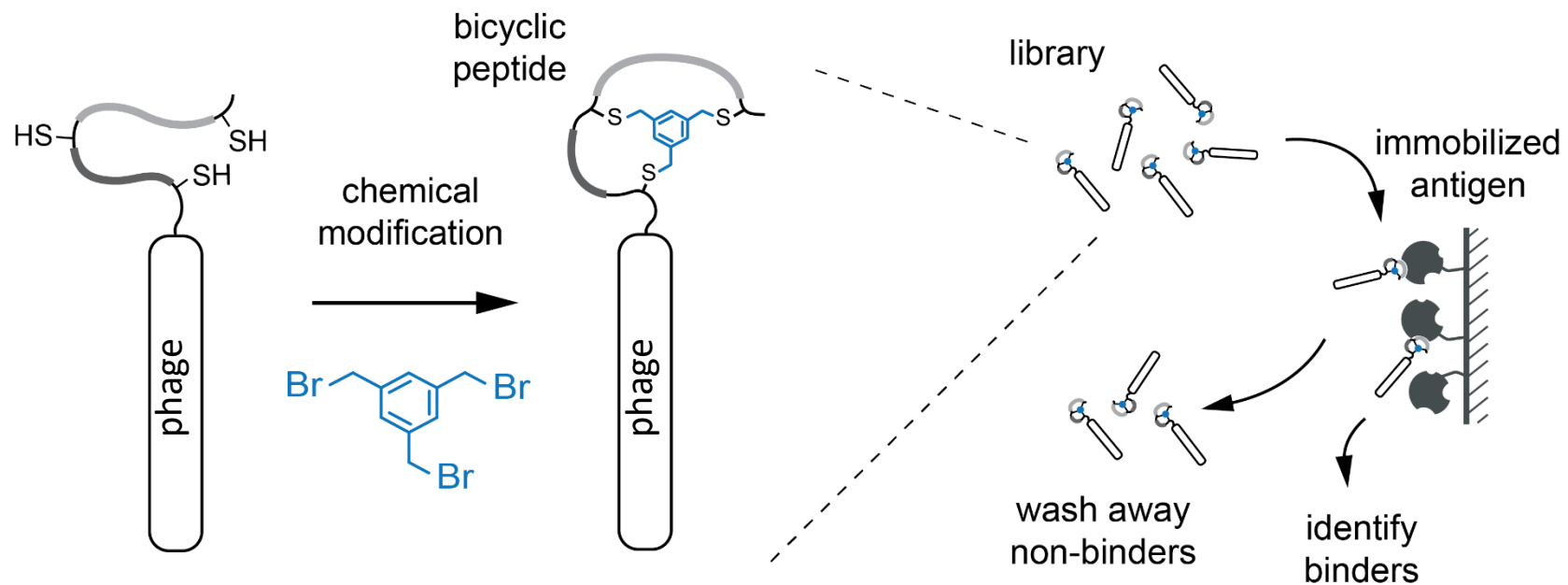


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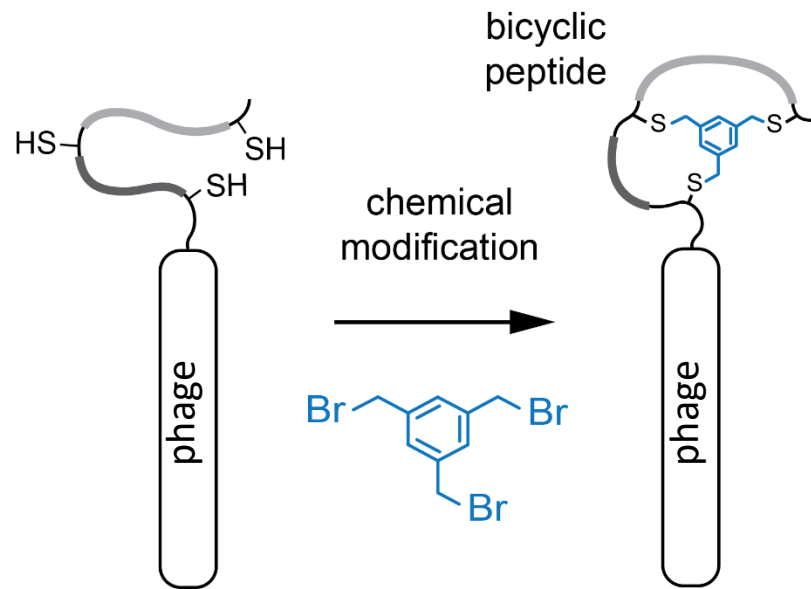


(Timmerman, P., Beld, J., Puijk, W.C. and
Meloan, R.H., *ChemBioChem*, 2005)

Phage selection of bicyclic peptides



Phage selection of bicyclic peptides

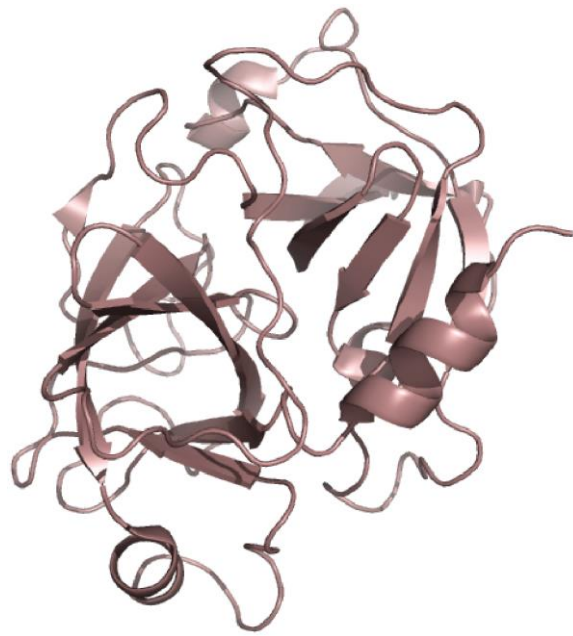


Laboratory of Molecular Biology (LMB),
Cambridge, UK




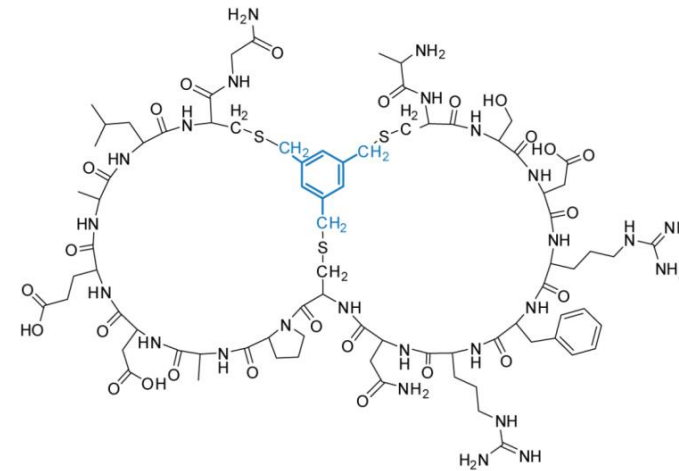
Sir Gregory
Winter

First phage-selected bicyclic peptide



plasma kallikrein


 $K_i = 2.9 \text{ nM}$



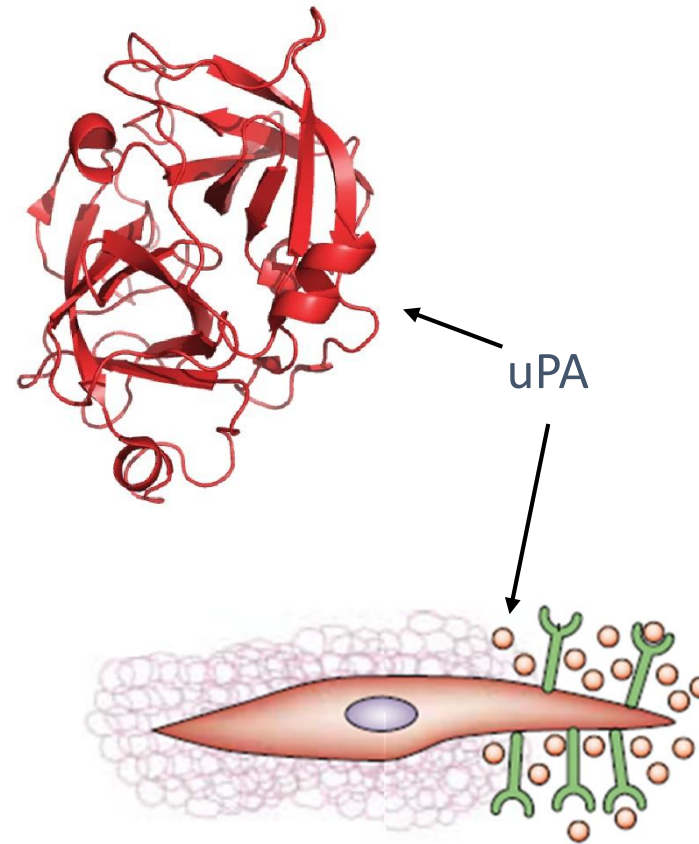
bicyclic peptide PK15

Heinis, C., Rutherford, T., Freund, S. and Winter, G., *Nat. Chem. Biol.*, 2009

Phage selection of bicyclic peptides

Example target: urokinase-type plasminogen activator (uPA)

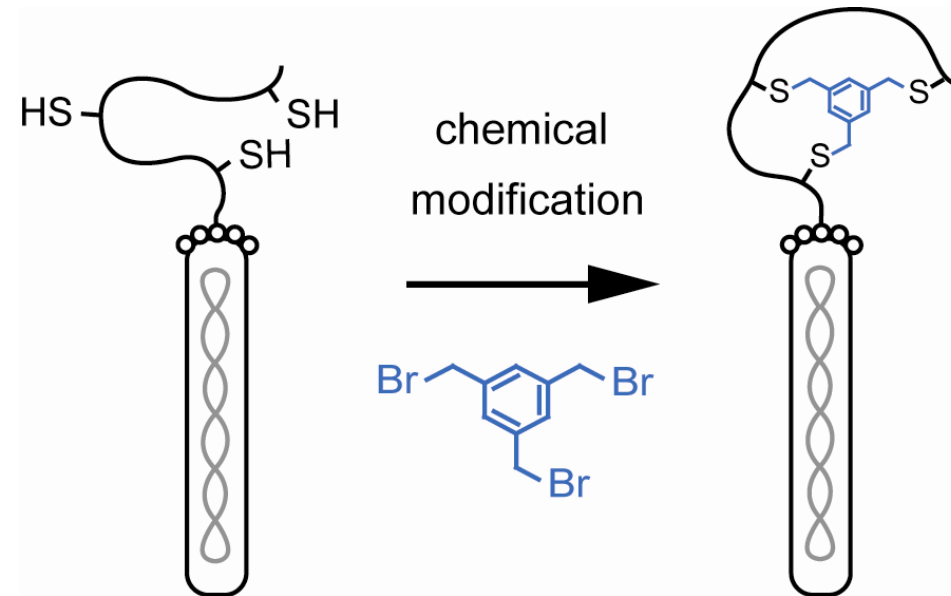
- Trypsin-like serine protease
- Overexpressed in some tumor cells (secreted)
- Proteolytic activity was reported to promote tumor growth and metastasis formation



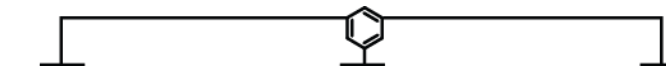
Phage library



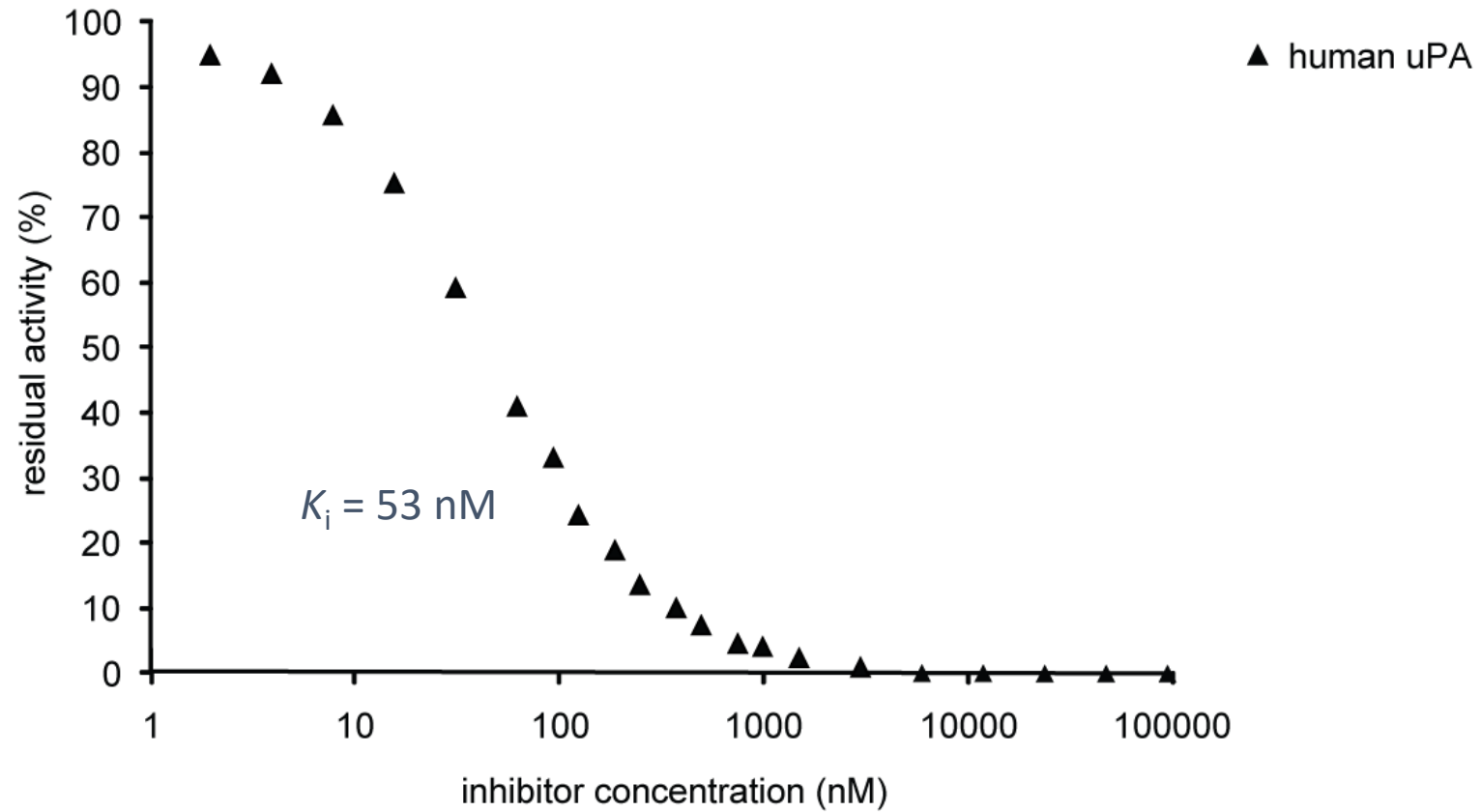
library size:
 $> 4 \times 10^9$
transformants



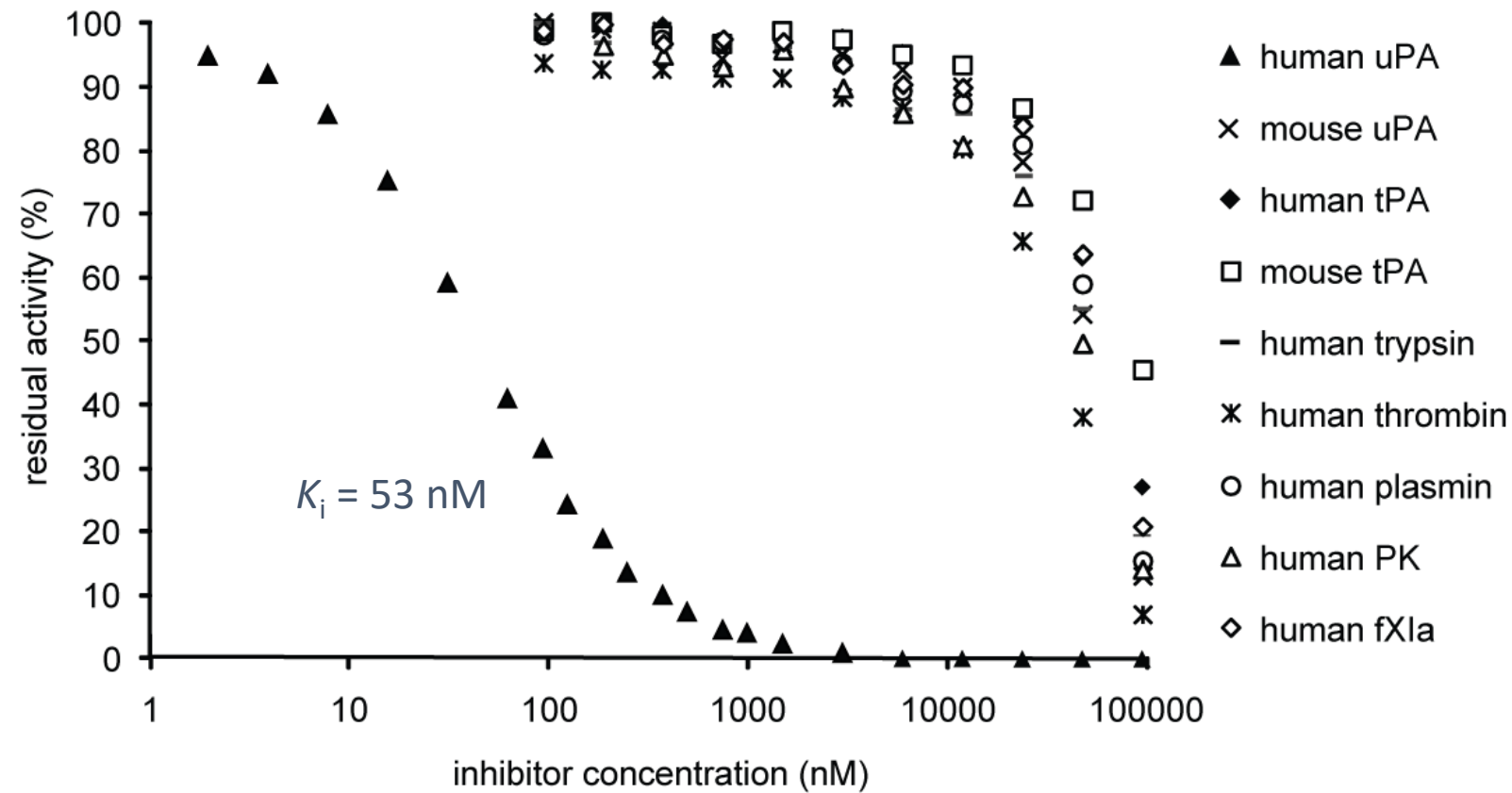
Isolated peptides

library																		K_i (μ M)
UK1	A	C	N	A	K	F	S	G	C	V	G	R	G	G	H	C	G	3.6
UK2	A	C	N	A	K	F	S	L	C	G	N	S	V	F	G	C	G	
UK3	A	C	N	A	K	F	S	G	C	R	Y	S	A	L	V	C	G	
UK4	A	C	N	A	Y	F	S	G	C	N	S	L	T	G	G	C	G	2.6
UK5	A	C	N	A	Y	F	S	G	C	D	W	A	V	Q	H	C	G	
UK6	A	C	N	A	R	F	S	G	C	Q	P	S	P	A	A	C	G	
UK7	A	C	N	E	R	F	S	M	C	G	Q	M	G	L	R	C	G	11.1
UK8	A	C	N	N	K	F	T	L	C	G	S	R	Q	I	I	C	G	
UK9	A	C	N	W	K	F	S	L	C	E	T	Q	R	N	Q	C	G	
UK10	A	C	N	S	R	F	S	G	C	Q	I	D	L	L	M	C	G	12.3
UK11	A	C	N	S	K	F	T	G	C	R	A	G	T	G	L	C	G	
UK12	A	C	N	S	K	F	T	Q	C	G	A	P	R	G	T	C	G	
UK13	A	C	N	S	R	F	A	L	C	S	P	S	S	Q	M	C	G	0.071
UK14	A	C	N	S	K	Y	S	G	C	Q	R	S	P	A	S	C	G	
UK15	A	C	N	S	K	Y	S	G	C	G	G	Q	R	Y	D	C	G	
UK16	A	C	N	S	R	Y	S	L	C	T	G	A	I	L	T	C	G	0.071
UK17	A	C	N	E	R	Y	A	L	C	G	I	L	G	F	Q	C	G	
UK18	A	C	S	R	Y	E	V	D	C	R	G	R	G	S	A	C	G	0.071
UK19	A	C	A	E	Q	V	I	D	C	R	G	R	G	G	L	C	G	

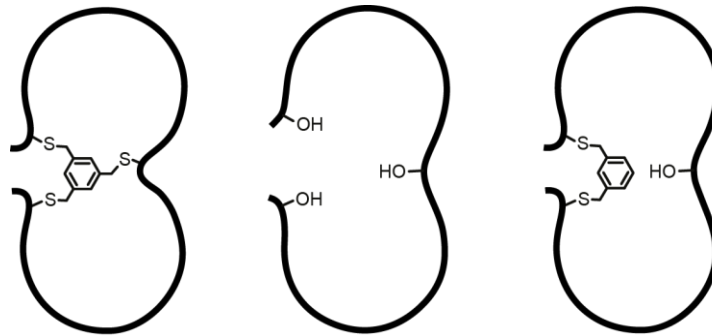
Inhibitory activity of UK18



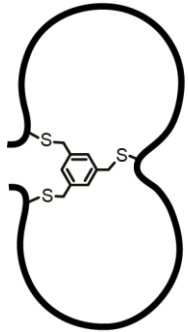
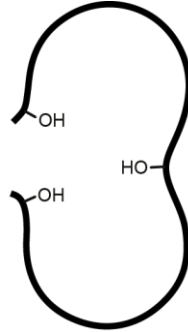
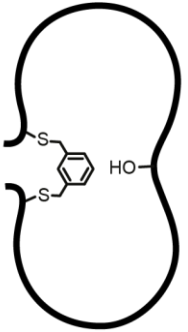
Inhibitory activity of UK18



Is bicyclic configuration important?

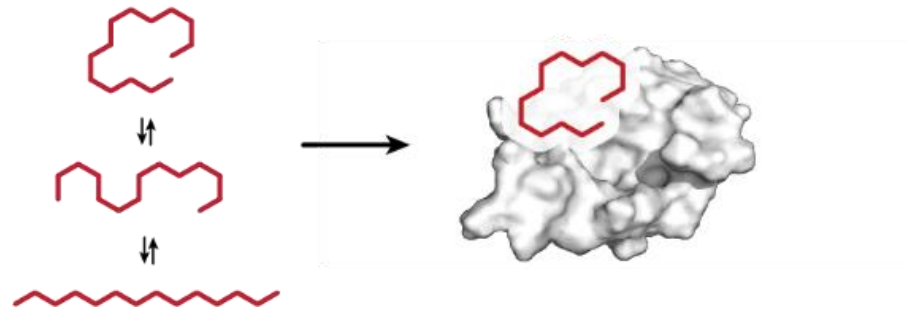


Is bicyclic configuration important?

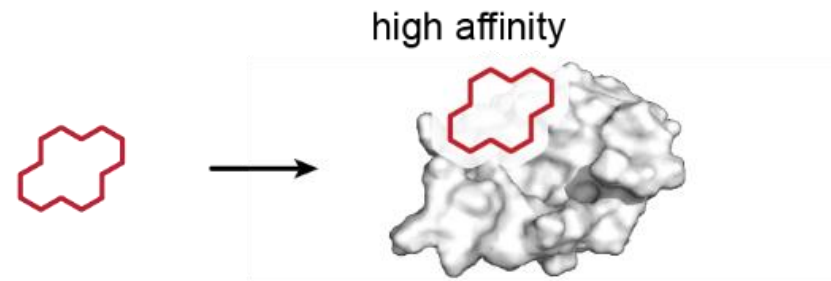
			
Inhibitory constant (K_i)	53 nM	17'500 nM	383 nM

Smaller entropic penalty \rightarrow Higher affinity

Linear

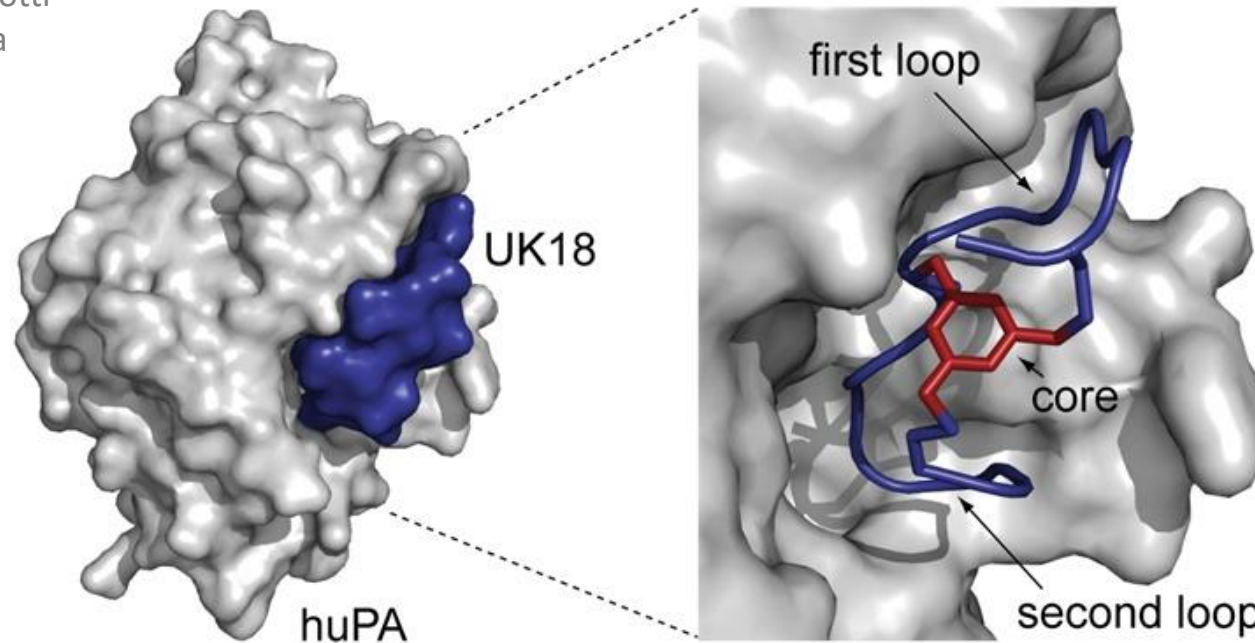


Cyclic



Structure of uPA-UK18 complex

Collaboration with
Prof. Giuseppe Zanotti
University of Padua



1.9 Å resolution

> 700 Å² binding interface

Bicyclic peptides to a broad range of targets

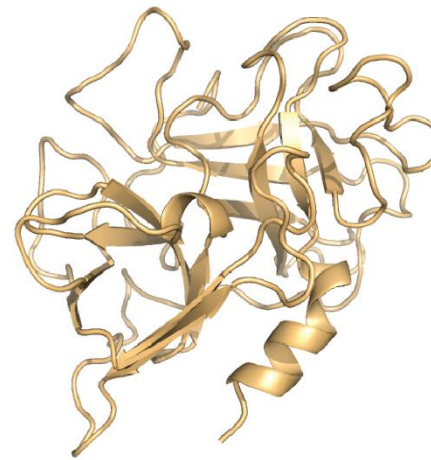
Plasma kallikrein	$K_i = 300 \text{ pM}$
uPA	$K_i = 28 \text{ nM}$
MMP-2	$K_i = 1.5 \text{ nM}$
PSA	$K_i = 47 \text{ nM}$
Her2	$K_d = 270 \text{ nM}$
Notch1 receptor	$K_d = 170 \text{ nM}$
β -catenin	$K_d = 6 \text{ }\mu\text{M}$
CAIX	$K_i = 50 \text{ nM}$
Sortase A	$K_i = 1.2 \text{ }\mu\text{M}$
Factor XIIa	$K_i = 370 \text{ pM}$
G-actin	$K_d = 5 \text{ nM}$
TNF α	$K_d = 300 \text{ nM}$
FKBP12	$K_d = 0.5 \text{ }\mu\text{M}$
IL-17	$K_d = 300 \text{ nM}$
Factor XIa	$K_i = 10 \text{ nM}$

....

Bicyclic peptides to a broad range of targets

Plasma kallikrein	$K_i = 300 \text{ pM}$
uPA	$K_i = 28 \text{ nM}$
MMP-2	$K_i = 1.5 \text{ nM}$
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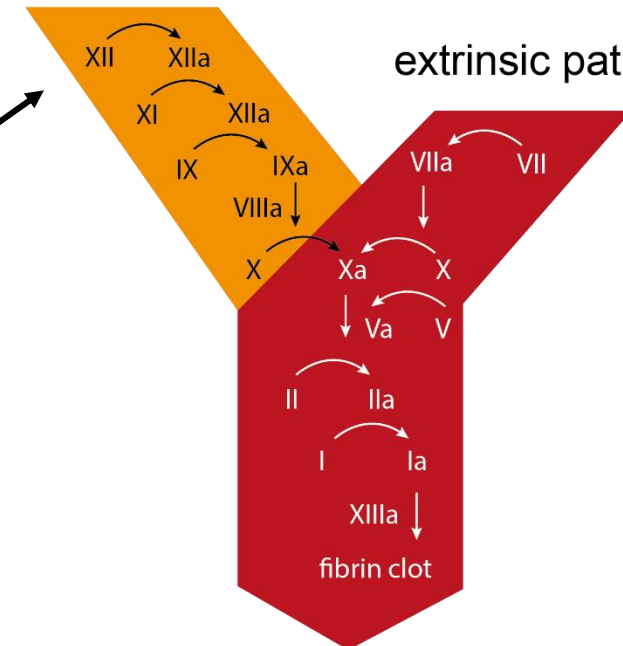
....



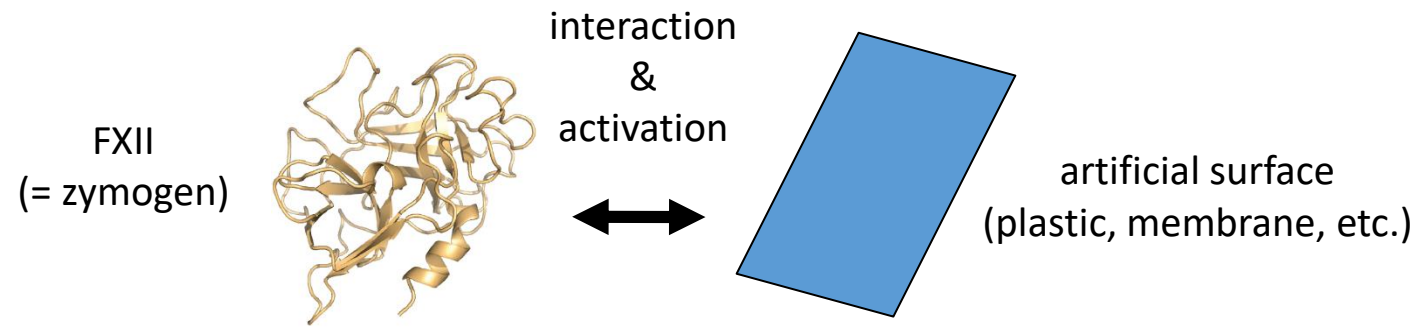
FXIIa

intrinsic pathway

extrinsic pathway

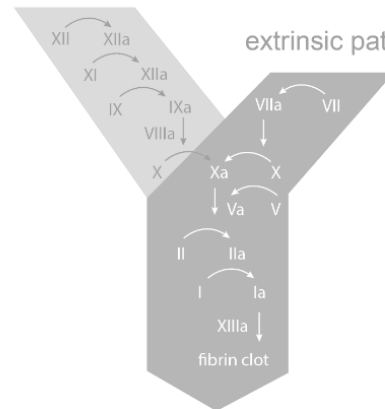


Contact activation



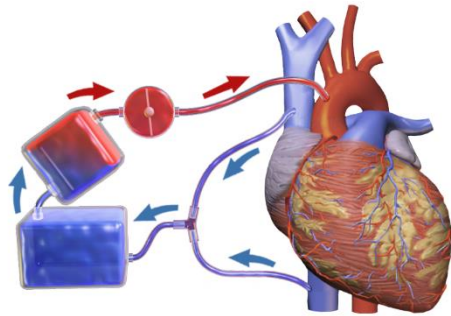
intrinsic pathway

extrinsic pathway

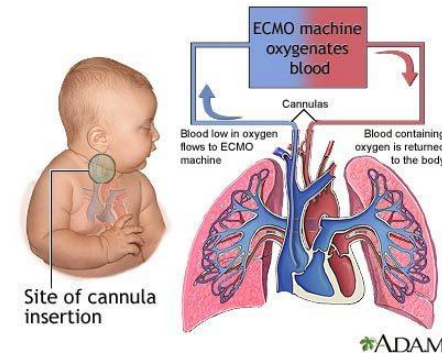


Contact activation in CPB and ECMO

Cardiopulmonary bypass surgery
(CPB)

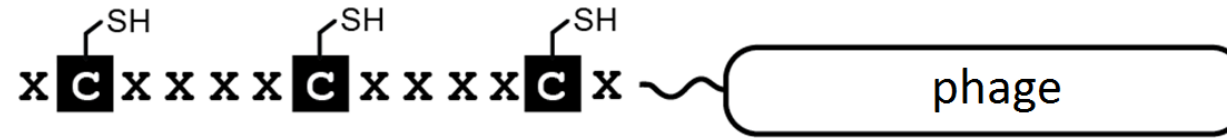


Extracorporeal membrane oxygenation
(ECMO)

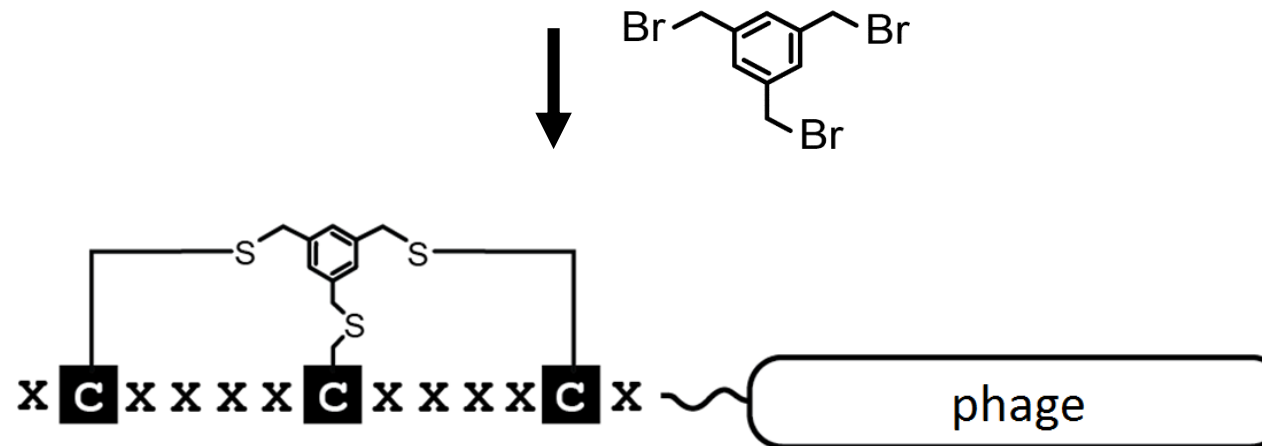


- Strong anticoagulation required (heparin) → bleeding risks
- Activation of kinnin-kininogen pathway → inflammation, organ damage

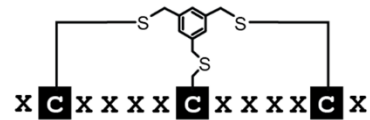
Phage selection of bicyclic peptide FXIIa inhibitor



X = random amino acid, $> 3 \times 10^8$ different peptides

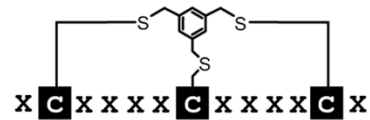


Isolated peptides



peptide:	sequence:	abundance:
FXII301	A C D A R P C P Q T Y C L	28
FXII302	Q C N A R P C P S S Y C R	2
FXII303	G C M G R P C P V S Y C F	2
FXII304	S C G G R P C P P A Y C K	22
FXII305	G C L G R P C P M A Y C S	13
FXII306	G C W A R P C P L A L C Q	1
FXII307	G C A A R P C P L T A C W	1
FXII308	G C H G R P C P L Q Y C K	1
FXII309	R C Y A N P C P I S Y C R	1
FXII310	S C S G R R C P P S Y C K	1
FXII311	V C V Q K F C W R G W C P	4
FXII312	A C Q Q Q F C W R G W C P	2
FXII313	L C E Y T L C W R G W C P	2
FXII314	H C R Y V F C W R G W C P	2
FXII315	N C V N R Y C W R G W C S	1
FXII316	W C V M Q K C L M Q Y C E	2
FXII317	Y C V K D K C L Q A M C S	2
FXII318	S C V V G K C L V Q Y C A	1
FXII319	H C V Q G K C L V Y M C G	1
FXII320	Q C V P F K C L L H L C A	1
FXII321	G C V L G K C L Q D F C Q	1
FXII322	Y C V P L K C L W D R C E	1
FXII323	R C S A G G C C E R V C E	1
FXII324	T C T T Q G C A F R L C S	1
FXII325	W C T H Q R C V K A R C C	1
FXII326	Y C Q G T K C V K E R C C	1
FXII327	L C Q P R K C V K A L C C	1
FXII328	S C A N Q R C V K S L C C	1
FXII329	T C L C K R C I K E L C C	1
FXII330	V C T Y R K C V K E L C C	1
FXII331	G C Q A R A C A K F L C C	1
FXII332	G C G R N I C V K N L C C	1

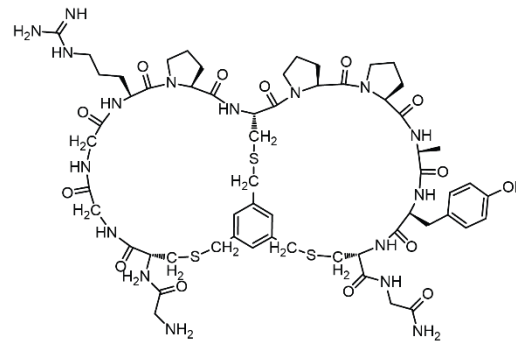
Isolated peptides



peptide:	sequence:	abundance:	K_i for FXIIa (μM):
FXII301	A C D A R P C P Q T Y C L	28	20.5 \pm 5.2
FXII302	Q C N A R P C P S S Y C R	2	4.7 \pm 1.5
FXII303	G C M G R P C P V S Y C F	2	5.0 \pm 1.3
FXII304	S C G G R P C P P A Y C K	22	3.1 \pm 0.5
FXII305	G C L G R P C P M A Y C S	13	5.0 \pm 1.5
FXII306	G C W A R P C P L A L C Q	1	10.2 \pm 4.6
FXII307	G C A A R P C P L T A C W	1	33.5 \pm 5.9
FXII308	G C H G R P C P L Q Y C K	1	11.2 \pm 4.4
FXII309	R C Y A N P C P I S Y C R	1	
FXII310	S C S G R C P P S Y C K	1	7.8 \pm 3.2
FXII311	V C V Q K F C W R G W C P	4	> 75
FXII312	A C Q Q Q F C W R G W C P	2	
FXII313	L C E Y T L C W R G W C P	2	
FXII314	H C R Y V F C W R G W C P	2	
FXII315	N C V N R Y C W R G W C S	1	
FXII316	W C V M Q K C L M Q Y C E	2	
FXII317	Y C V K D K C L Q A M C S	2	
FXII318	S C V V G K C L V Q Y C A	1	
FXII319	H C V Q G K C L V Y M C G	1	
FXII320	Q C V P F K C L L H L C A	1	
FXII321	G C V L G K C L Q D F C Q	1	
FXII322	Y C V P L K C L W D R C E	1	
FXII323	R C S A G G C C E R V C E	1	
FXII324	T C T T Q G C A F R L C S	1	> 75
FXII325	W C T H Q R C V K A R C C	1	
FXII326	Y C Q G T K C V K E R C C	1	
FXII327	L C Q P R K C V K A L C C	1	
FXII328	S C A N Q R C V K S L C C	1	
FXII329	T C L C K R C I K E L C C	1	
FXII330	V C T Y R K C V K E L C C	1	
FXII331	G C Q A R A C A K F L C C	1	
FXII332	G C G R N I C V K N L C C	1	

Bicyclic peptide FXIIa inhibitor

1st generation



$$K_i = 1.2 \mu\text{M}$$

Baeriswyl, V., Calzavarini, S., Gertschheimer, C., Diderich, P.,
Angelillo-Scherrer, A. and Heinis, C., *J. Med. Chem.*, 2012

Screening larger structural diversities

3x3 library

x **C** x x x **C** x x x **C** x ~ phage

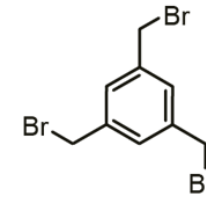
4x4 library

x **C** x x x x **C** x x x x **C** x ~ phage

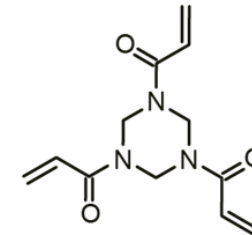
6x6 library

A **C** x x x x x x **C** x x x x x x **C** G ~ phage

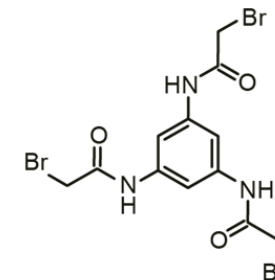
TBMB



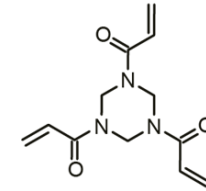
TATA



TBAB



Bicyclic peptide isolated with TATA



3x3 library

peptide:	sequence:	abundance:	K_i (μM):
FXII501	K C W G A C L N T C P	4	1.2
FXII502	R C W G A C L N V C S	2	
FXII503	K C W G A C L N A C P	1	
FXII504	V C W G A C L N V C P	1	1.9
FXII505	Q C W G A C L N V C N	4	6.3
FXII506	A C W G A C Q N V C R	5	
FXII507	F C N P Y C M M K C A	1	
FXII508	D C N P Y C Q M K C M	1	
FXII509	R C W Y G C V P Q C Y	6	1.0
FXII510	F C T K E C L Q S C Y	3	
FXII511	R C L Q K C F P P C L	1	
FXII512	G C C Y L R L S C R	2	0.16
FXII513	G C C Y L R Y L C S	1	0.35



4x4 library

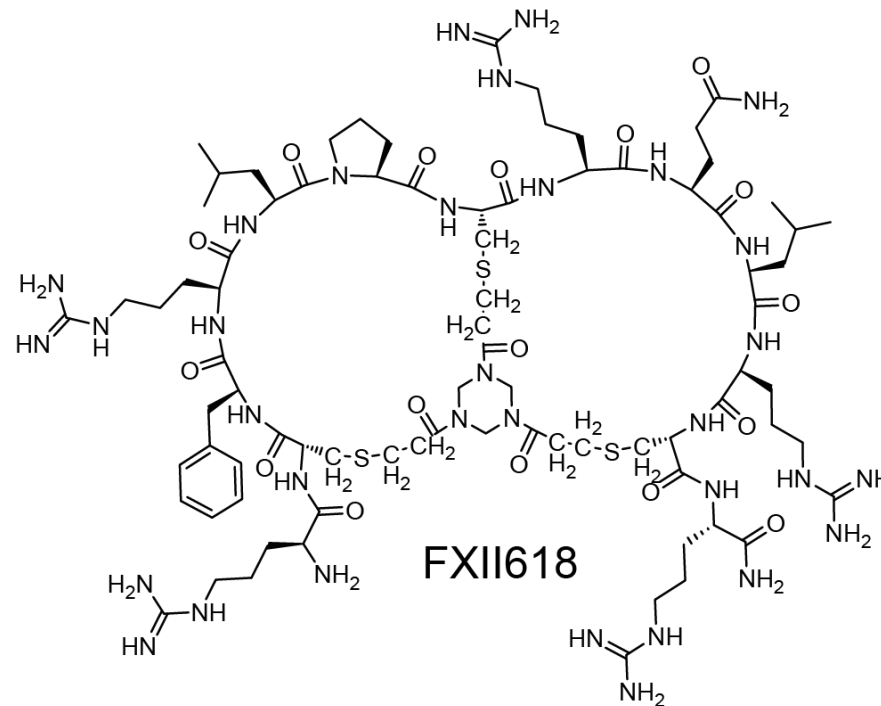
peptide:	sequence:	abundance:	K_i (μM):
FXII514	L C S R L P C E Q L P C N	11	10.2
FXII515	L C P R L S C H Q L G C S	4	2.5
FXII516	R C F R L P C A Q L S C E	1	0.16
FXII517	N C Q Y M R C P P M S C L	3	
FXII518	S C M A T Q C L N Q L Y G	4	



6x6 library

peptide:	sequence:	abundance:	K_i (μM):
FXII519	A C W G A A A G C V Q G A S N C G	1	
FXII520	A C W G A A A G C V S V H Q S C G	1	
FXII521	A C W G A A A K C S Q T W S G C G	1	
FXII522	A C W G A A A K C P Q T A A Q C G	1	
FXII523	A C W G A A A T C N Q Q Q L V C G	1	
FXII524	A C W G A A A H C Q L P K D R C G	1	
FXII525	A C W G A A V G C N S P R G P C G	1	
FXII526	A C W G A A V S C E T Q S S L C G	1	
FXII527	A C W G A A L S C N P Q G S R C G	1	
FXII528	A C W G A A Q R C K V M Q G E C G	1	8.5
FXII529	A C W G A A Q K C K M N G Q N C G	1	3.7
FXII530	A C W G A A Q L C G R P Q M P C G	1	
FXII531	A C W G A A Q A C I P Q M A A C G	1	
FXII532	A C W G A A Q H C Q A S N N S C G	1	
FXII533	A C W G A A S G C N Q T G M D C G	1	
FXII534	A C W G A A S A C Q Q G G A K C G	1	
FXII535	A C W G A A S R C T V I S G R C G	1	
FXII536	A C W G A A R G C P D Q H G V C G	1	
FXII537	A C W G A A K L C G A A R D G C G	1	
FXII538	A C W G A A E R C K Q A D N S C G	1	

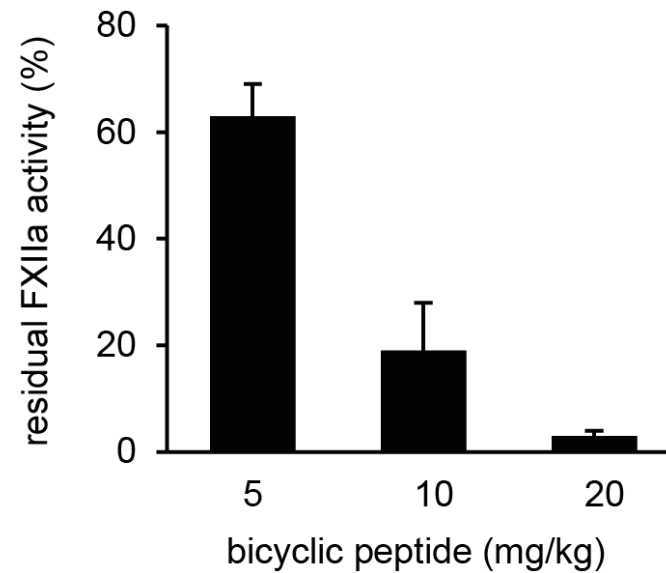
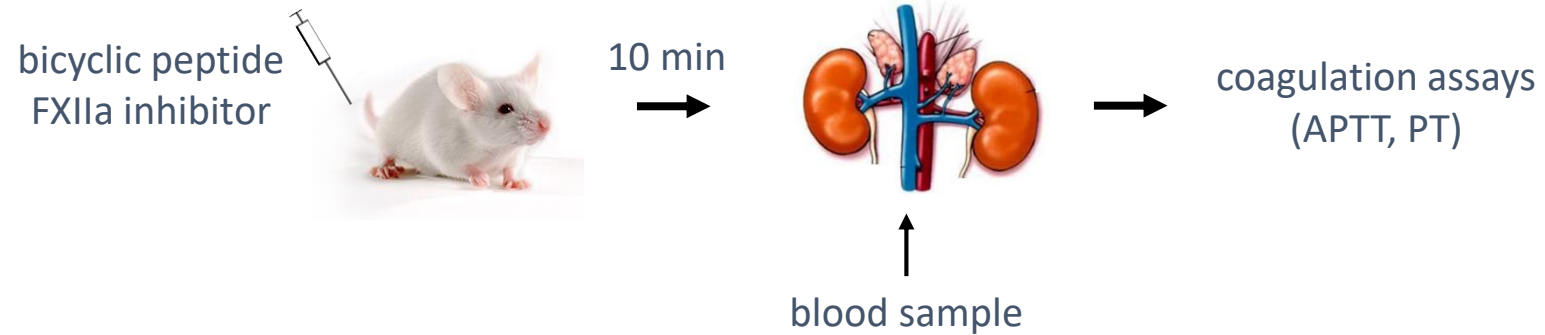
Affinity maturation



Protease	K_i (nM)
hFXIIa	8 ± 0.7
mFXIIa	93 ± 4
htPA	$> 50,000$
huPA	$> 50,000$
hFVIIa	$> 50,000$
hFXa	$> 50,000$
hFXIa	$> 50,000$
hPK	$> 50,000$
hthrombin	$> 50,000$
hplasmin	$> 50,000$

Baeriswyl, V., Calzavarini, S., Chen, S., Zorzi, A., Bologna, L.,
 Angelillo-Scherrer, A. and Heinis, C., *ACS Chem. Biol.*, 2015

Inhibition of intrinsic coagulation pathway (in vivo)



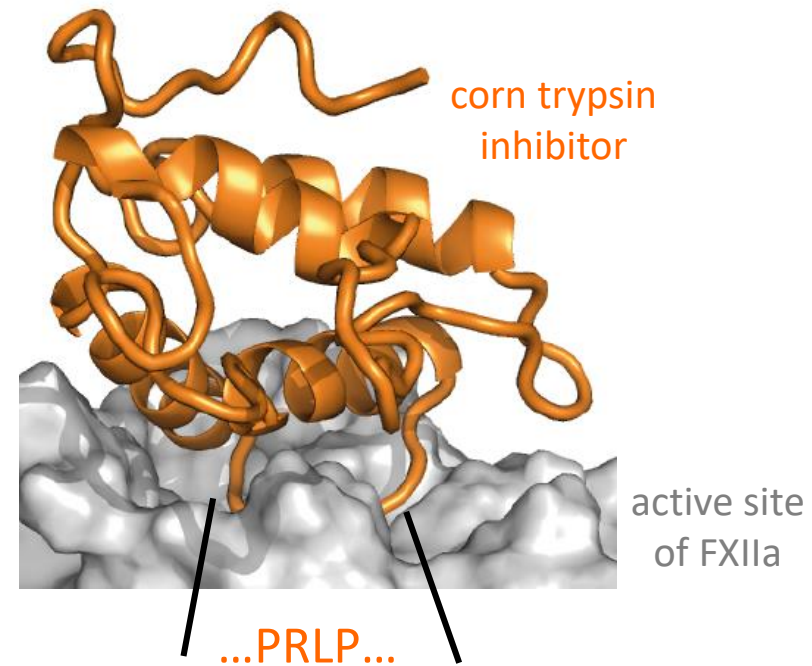
Collaboration with group
Prof. Anne Angelillo-Scherrer
(Inselspital Bern)

Sequence similarity to natural FXIIa inhibitor

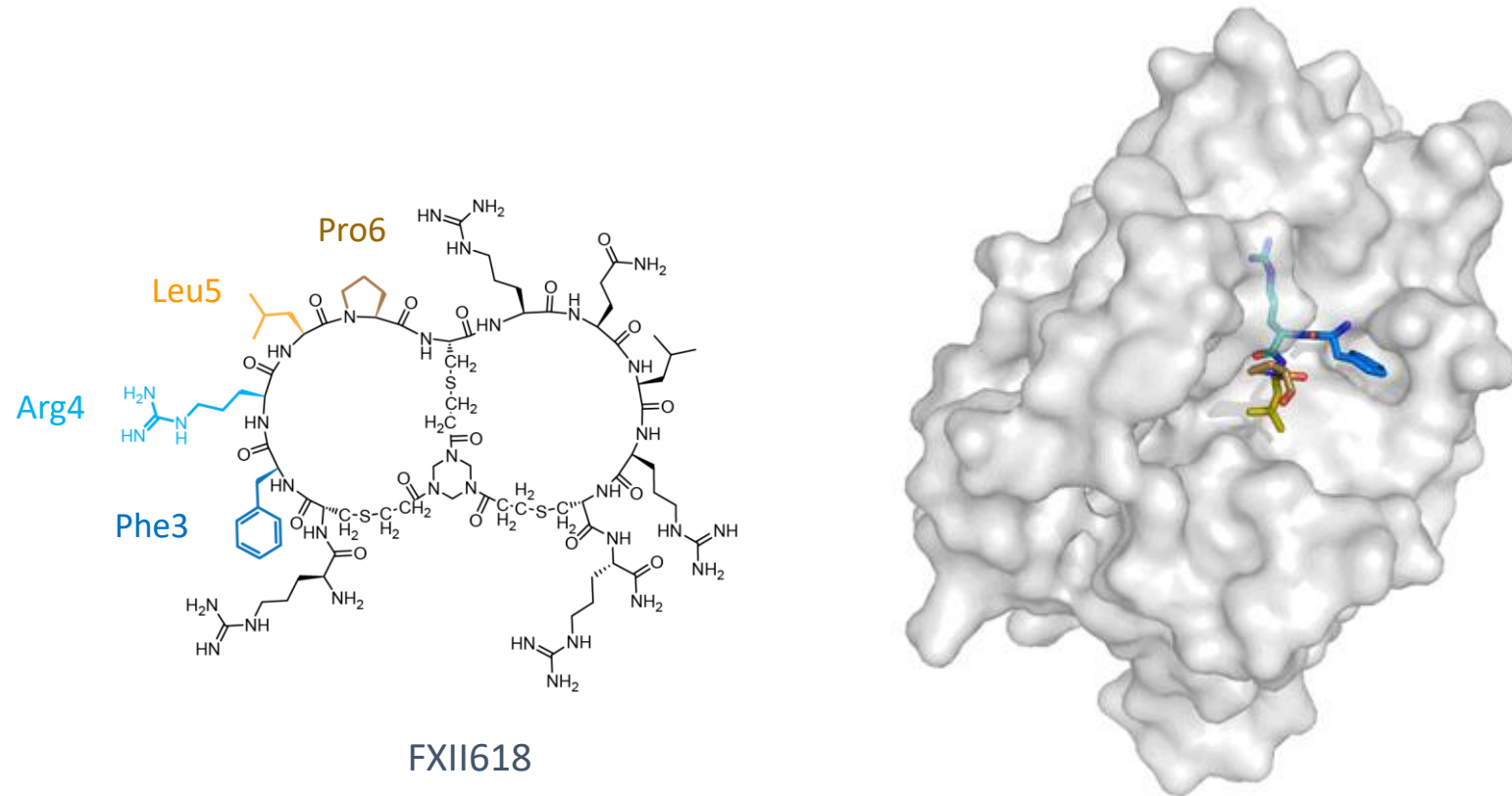
peptides selected against FXIIa:

FXII572	R	C	P	R	L	P	C	A	Q	L	Q	C	P
FXII573	I	C	P	R	R	L	P	C	A	Q	L	R	C
FXII574	V	C	P	R	R	L	P	C	G	Q	L	R	C
FXII575	I	C	P	R	R	L	P	C	R	Q	L	S	C
FXII576	V	C	P	R	R	L	P	C	H	Q	L	R	C
FXII577	R	C	P	R	R	L	P	C	E	Q	L	R	C
FXII578	I	C	P	R	R	L	P	C	A	Q	L	V	C
FXII579	L	C	P	R	R	L	P	C	G	Q	L	R	C
FXII580	I	C	P	R	R	L	P	C	G	Q	L	S	C
FXII581	R	C	P	R	R	L	P	C	E	Q	L	A	C
FXII582	G	C	P	R	R	L	P	C	A	Q	L	R	C
FXII583	I	C	P	R	R	L	P	C	S	Q	L	R	C
FXII584	K	C	P	R	R	L	P	C	G	Q	L	R	C
FXII585	A	C	P	R	R	L	P	C	A	Q	L	P	C
FXII586	R	C	P	R	R	L	P	C	S	Q	L	V	C
FXII587	R	C	P	R	R	L	P	C	A	Q	L	E	C
FXII588	I	C	P	R	R	L	P	C	A	Q	L	K	C
FXII589	L	C	P	R	R	L	P	C	A	Q	L	R	C
FXII590	A	C	P	R	R	L	P	C	S	Q	L	K	C
FXII591	F	C	P	R	R	L	P	C	A	Q	L	K	C
FXII592	V	C	P	R	R	L	P	C	G	Q	L	R	C
FXII593	R	C	P	R	R	L	P	C	A	Q	L	H	C
FXII594	G	C	P	R	R	L	P	C	A	Q	L	Q	C
FXII595	R	C	P	R	R	L	P	C	E	Q	L	H	C
FXII596	P	C	P	R	R	L	P	C	G	Q	L	R	C
FXII597	V	C	P	R	R	L	P	C	H	Q	L	R	C
FXII598	V	C	P	R	R	L	P	C	A	Q	L	R	C
FXII599	V	C	A	R	R	L	P	C	A	Q	L	K	C
FXII600	A	C	F	R	R	L	P	C	L	Q	L	A	C
FXII601	R	C	F	R	R	L	P	C	S	Q	L	T	C
FXII602	R	C	A	R	R	L	P	C	H	Q	L	H	C
FXII603	R	C	A	R	R	L	P	C	H	Q	L	K	C
FXII604	K	C	L	R	R	L	P	C	D	Q	L	P	C
FXII605	G	C	F	R	R	L	P	C	S	Q	L	K	C
FXII606	W	C	L	R	R	L	P	C	E	Q	L	M	C

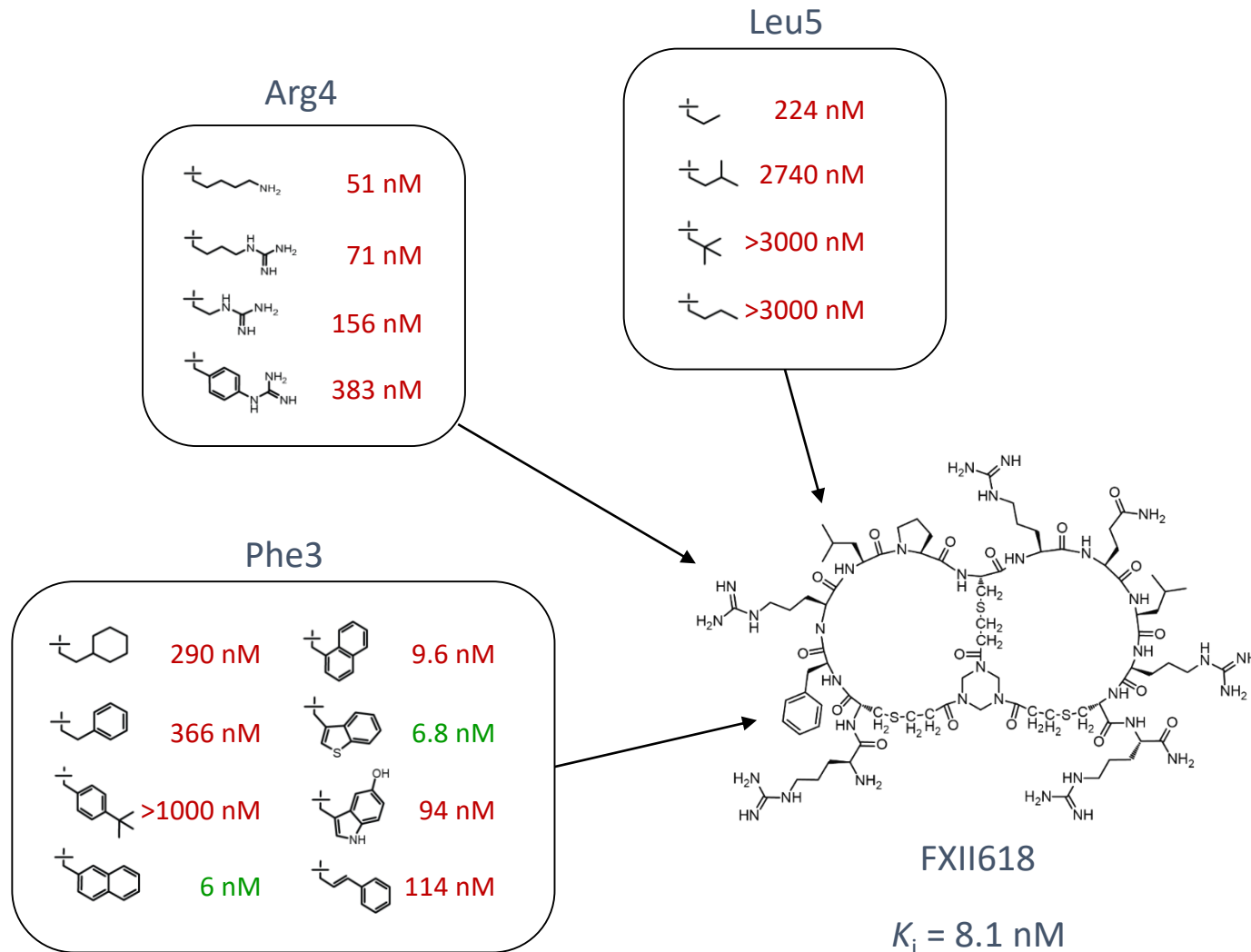
Corn trypsin inhibitor (CTI):



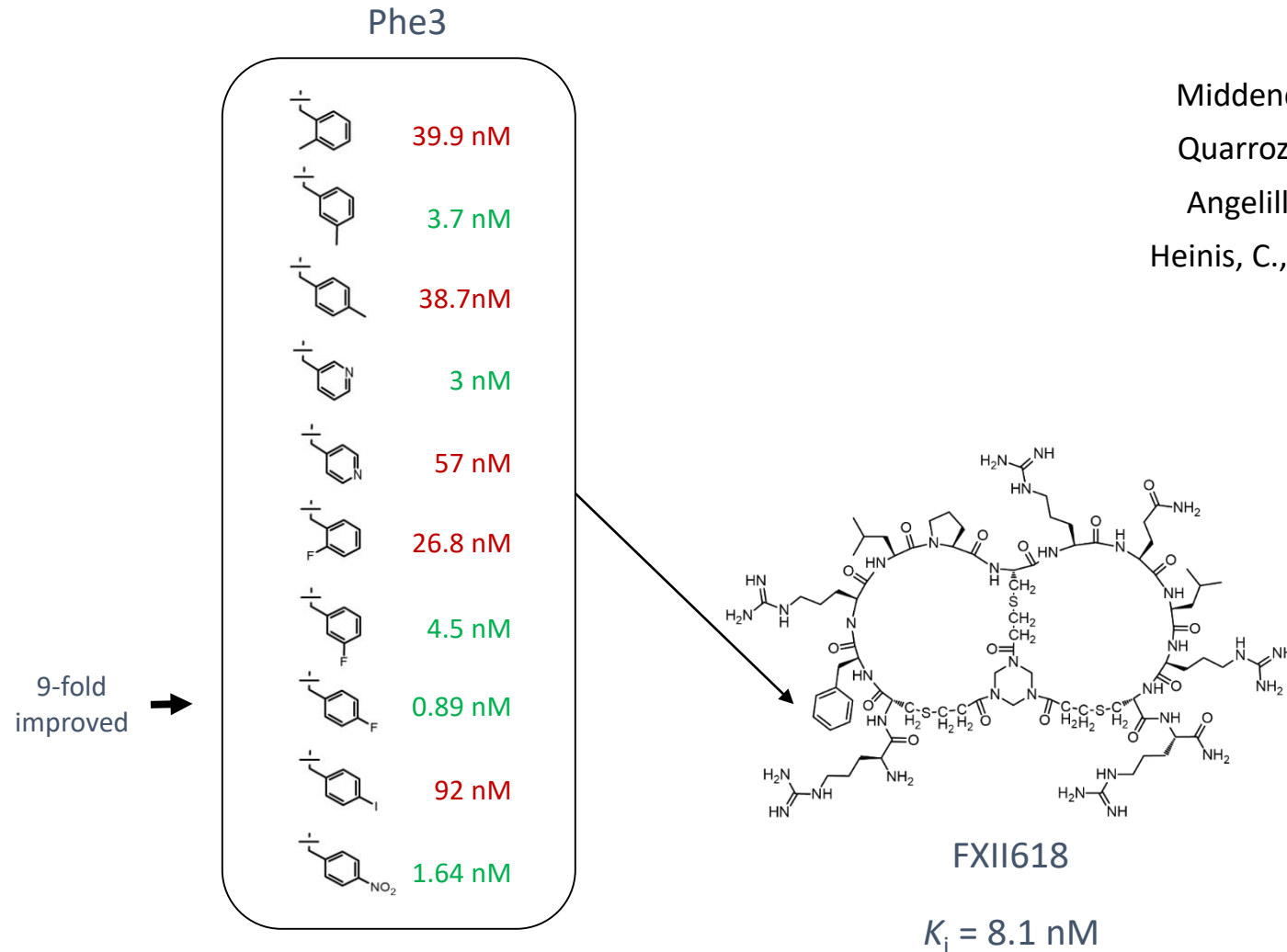
Structure model



Affinity maturation with non-natural amino acids



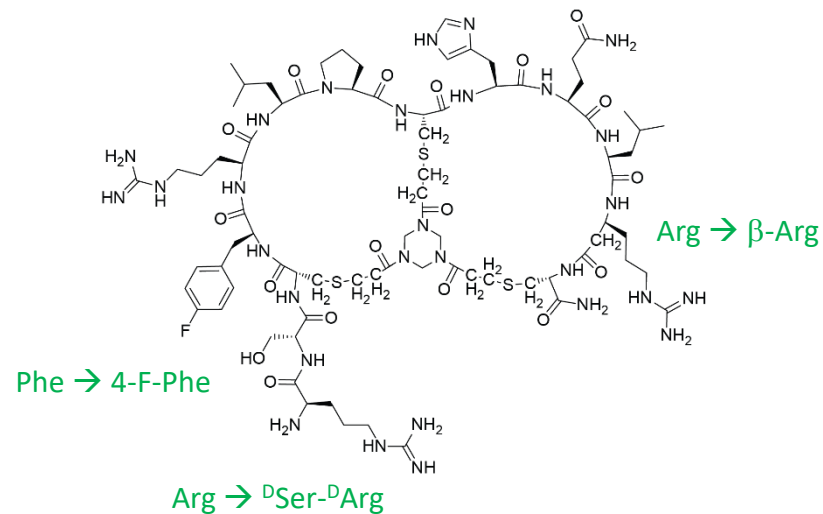
Affinity maturation with non-natural amino acids



Middendorp, S.J., Wilbs, J.,
 Quarroz, C., Calzavarini, S.,
 Angelillo-Scherrer, A. and
 Heinis, C., *J. Med. Chem.*, 2016

Combination of all mutations

FXII900



Affinity: $K_i = 370 \text{ pM}$

Selectivity:
(over homologous
plasma proteases) $> 50'000\text{-fold}$

Stability:
(human plasma) $t_{1/2} > 4 \text{ days}$

Activity
in plasma:
(APTT, human, ex vivo) $EC_{2x} = 1.3 \text{ }\mu\text{M}$

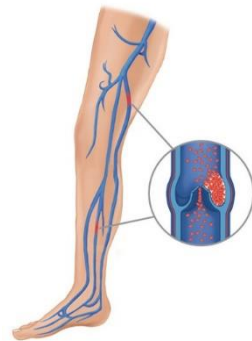
Pharmacokinetics:
elimination half-life $t_{1/2} (\beta)$
20 min (mouse)
15 min (rabbit)
40 min (pig)

Collaboration with
Prof. Robert Rieben, University of Bern



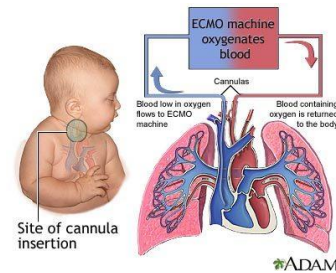
Evaluation of inhibitor in vivo

1. Thrombosis



Ferric chloride-
induced thrombosis
mouse model

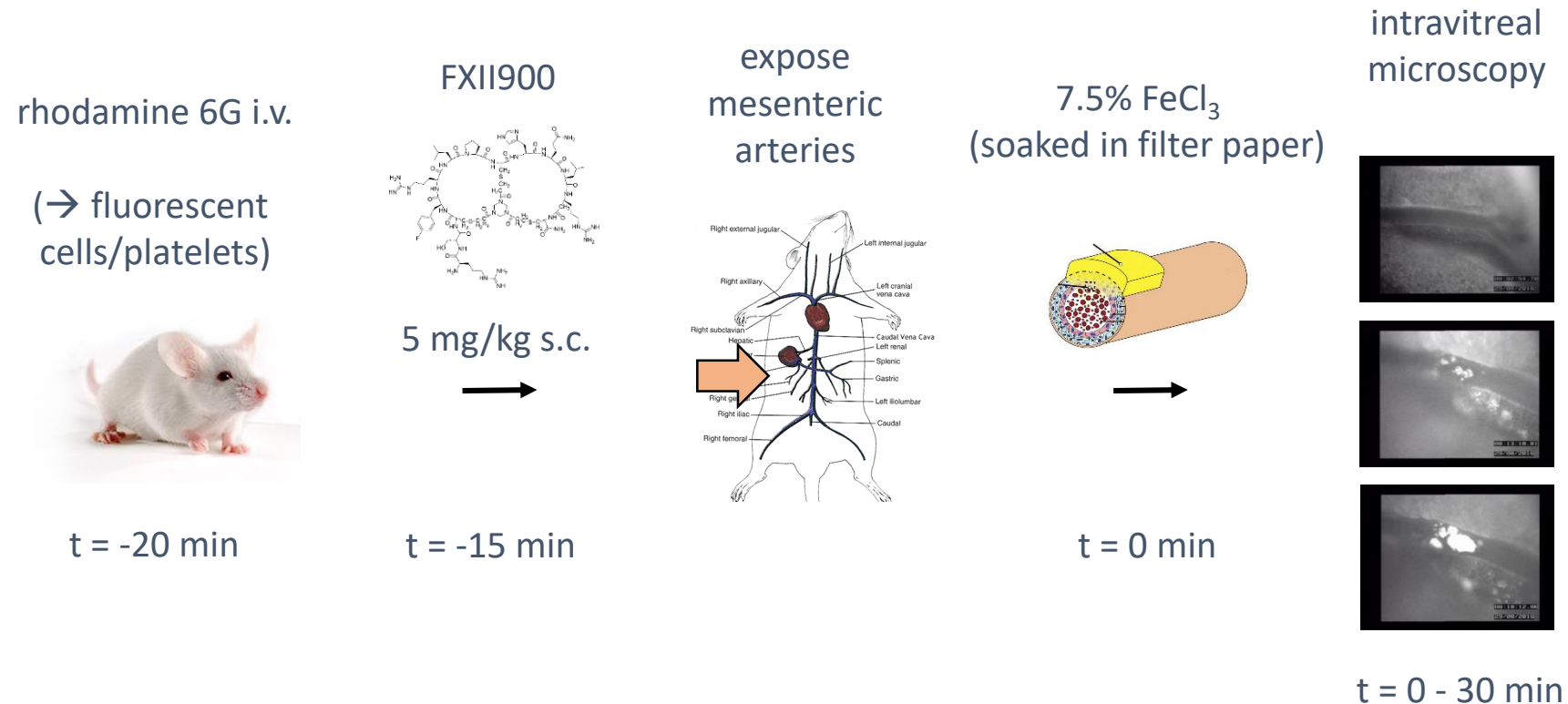
2. Contact activation



Extracorporeal
membrane oxygenation
(ECMO) in rabbits

Prevention of thrombosis

Ferric chloride-induced thrombosis mouse model

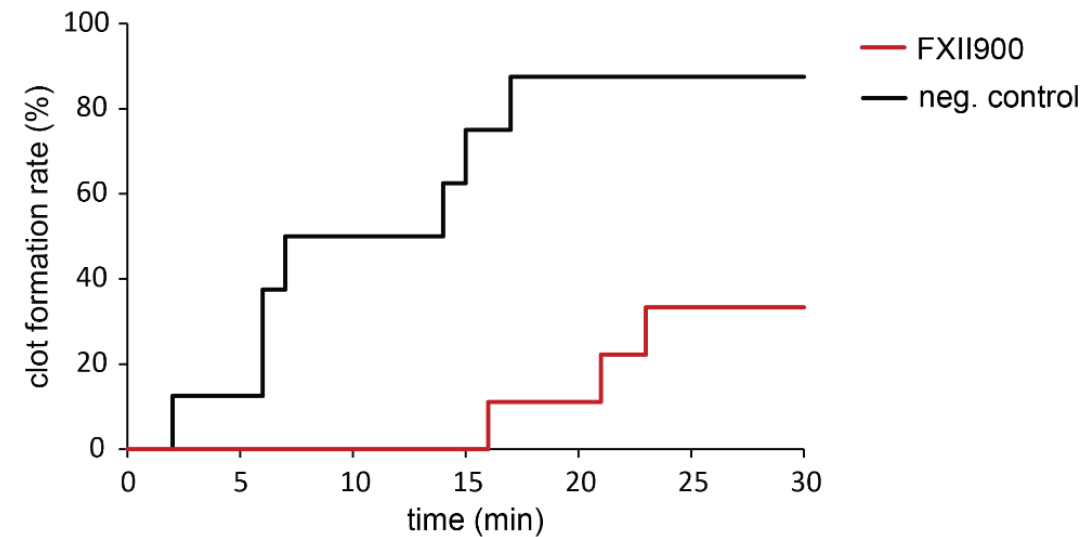


Study performed in collaboration with
Prof. Anne Angelillo-Scherrer and Ms Raja Price (University of Bern)

Prevention of thrombosis

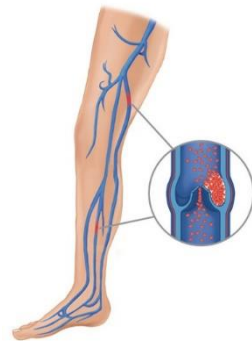
Ferric chloride-induced thrombosis mouse model

Time to clot formation:



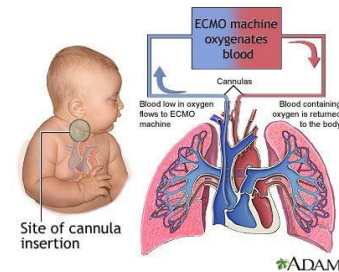
Evaluation of inhibitor in vivo

1. Thrombosis



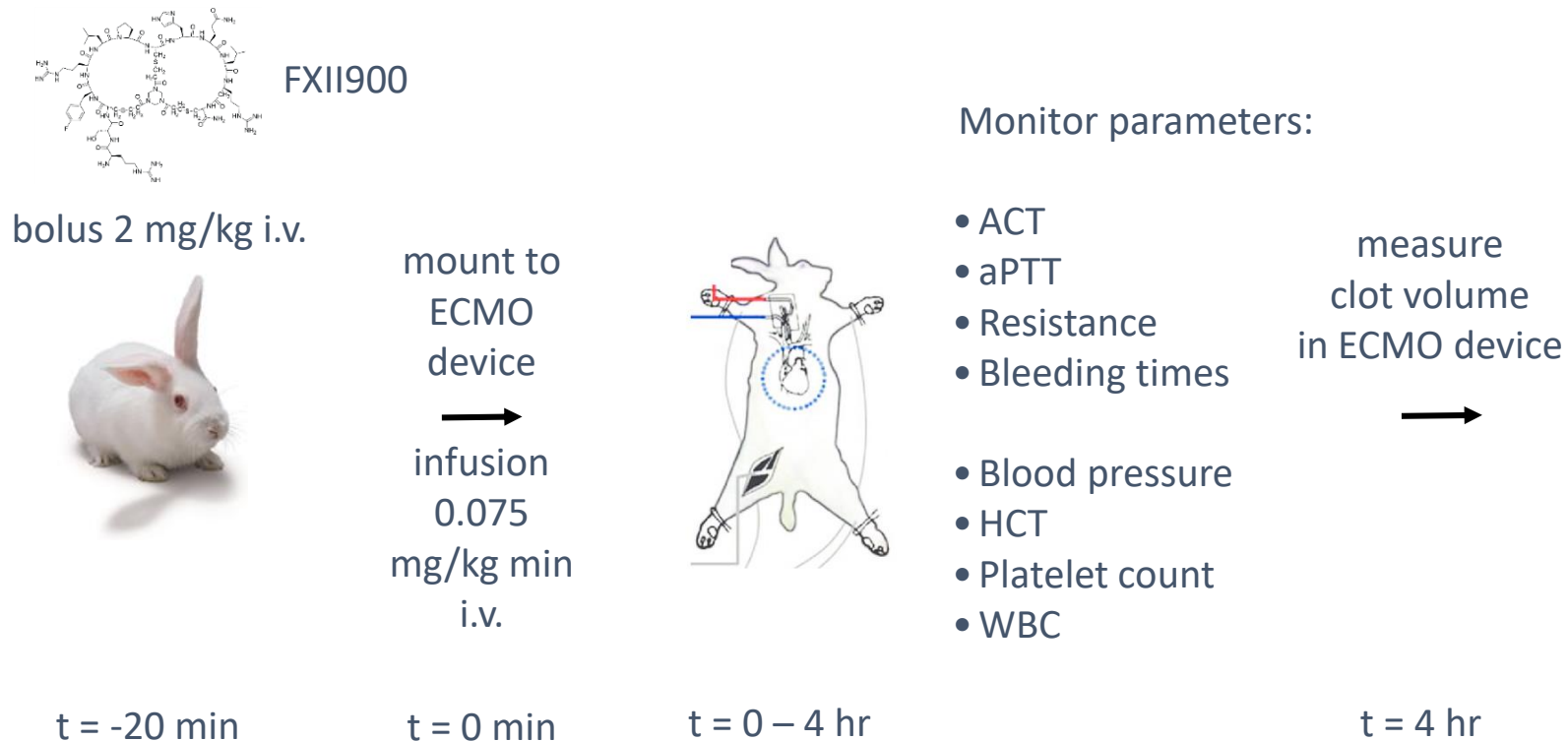
Ferric chloride-
induced thrombosis
mouse model

2. Contact activation



Extracorporeal
membrane oxygenation
(ECMO) in rabbits

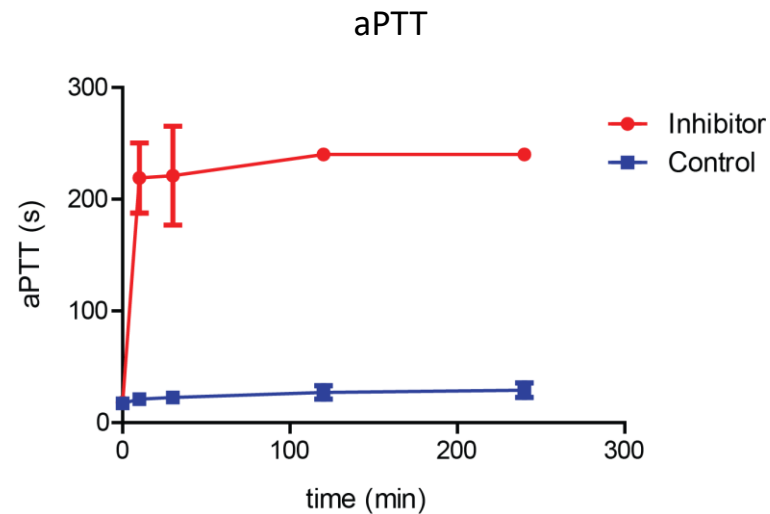
Inhibition of coagulation during ECMO in rabbit model



Study performed in collaboration with
Prof. Keith Cook and Ms Alida Cooke (Carnegie Mellon University)

Inhibition of coagulation during ECMO in rabbit model

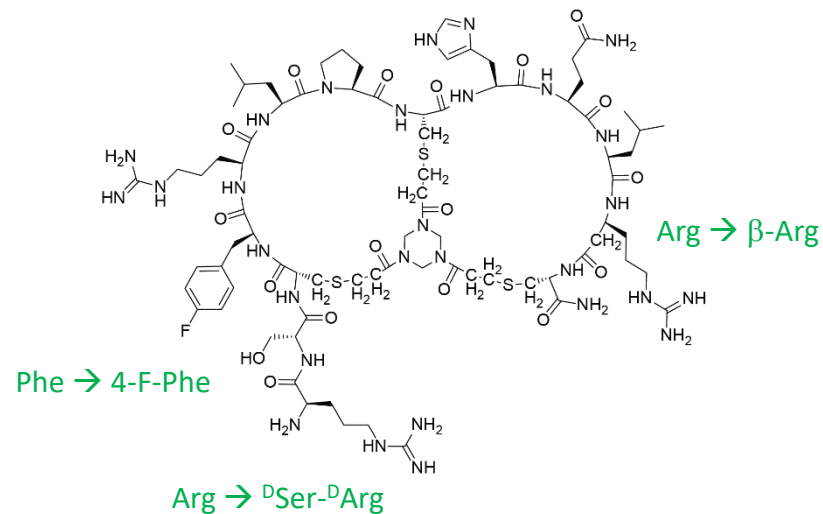
- Maximal prolongation of aPTT
- Clot volume in ECMO oxygenator was reduced > 3-fold
- Normal bleeding time



Study performed in collaboration with
Prof. Keith Cook and Ms Alida Cooke (Carnegie Mellon University)

Combination of all mutations

FXII900



Affinity: $K_i = 370 \text{ pM}$

Selectivity:
(over homologous
plasma proteases) $> 50'000\text{-fold}$

Stability:
(human plasma) $t_{1/2} > 4 \text{ days}$

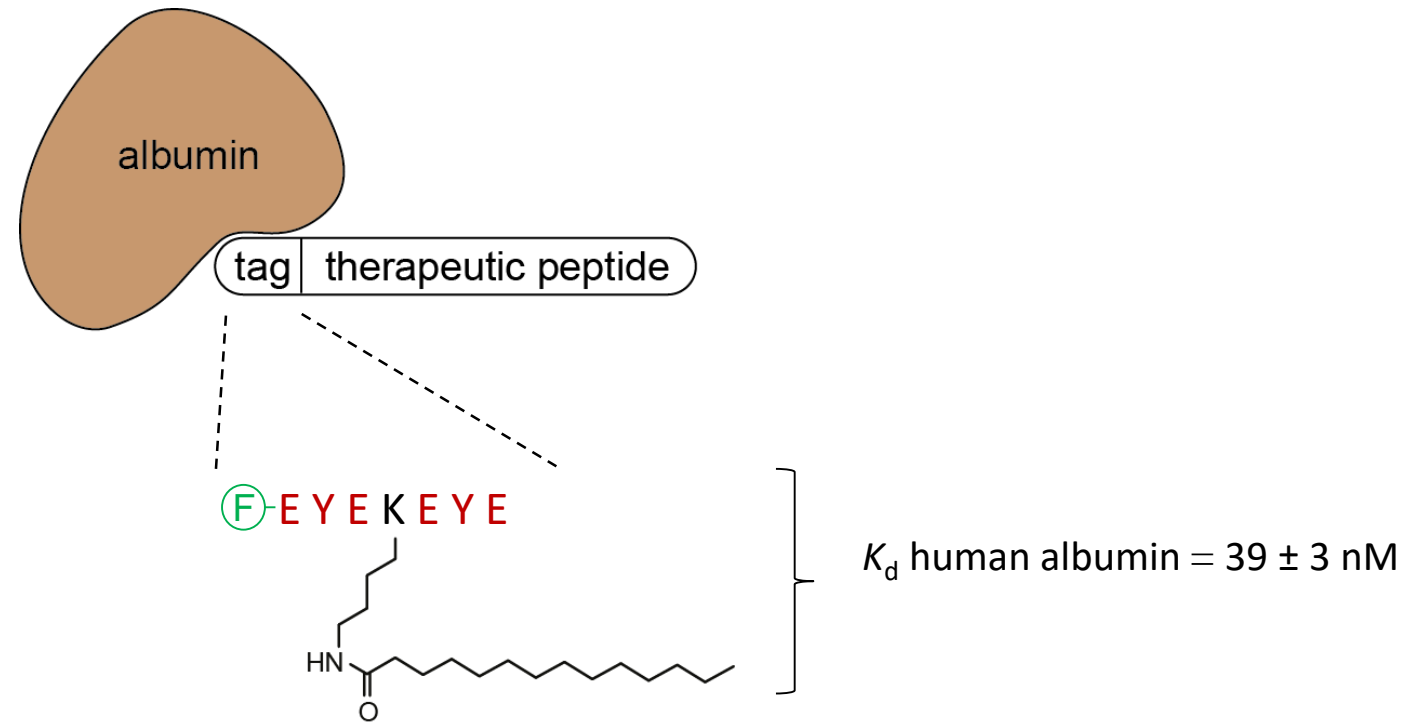
Activity
in plasma: $EC_{2x} = 1.3 \text{ }\mu\text{M}$
(APTT, human, ex vivo)

Pharmacokinetics:

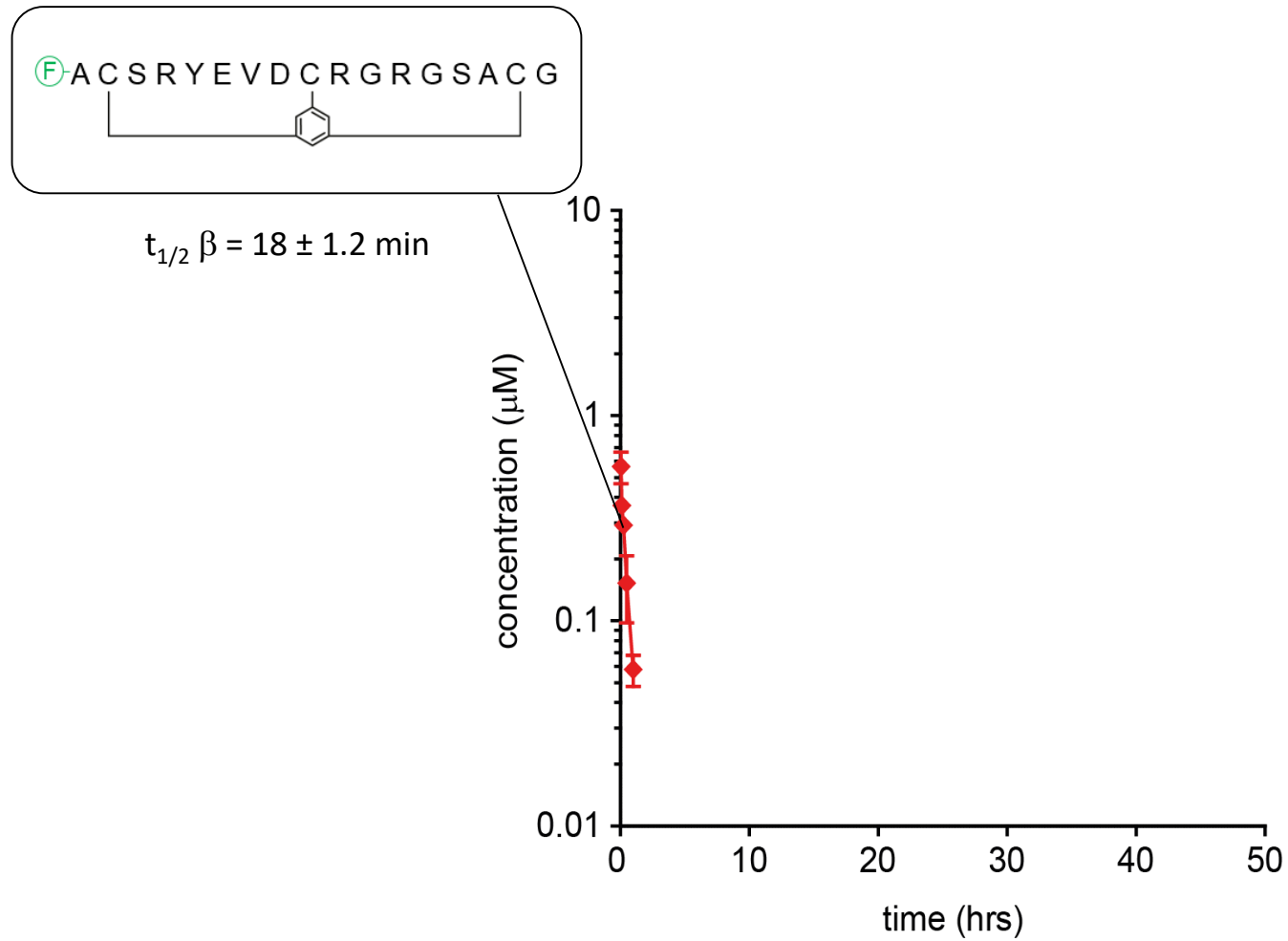
20 min (mouse)
elimination half-life $t_{1/2}$ (β)
15 min (rabbit)
40 min (pig)

Collaboration with
Prof. Robert Rieben, University of Bern

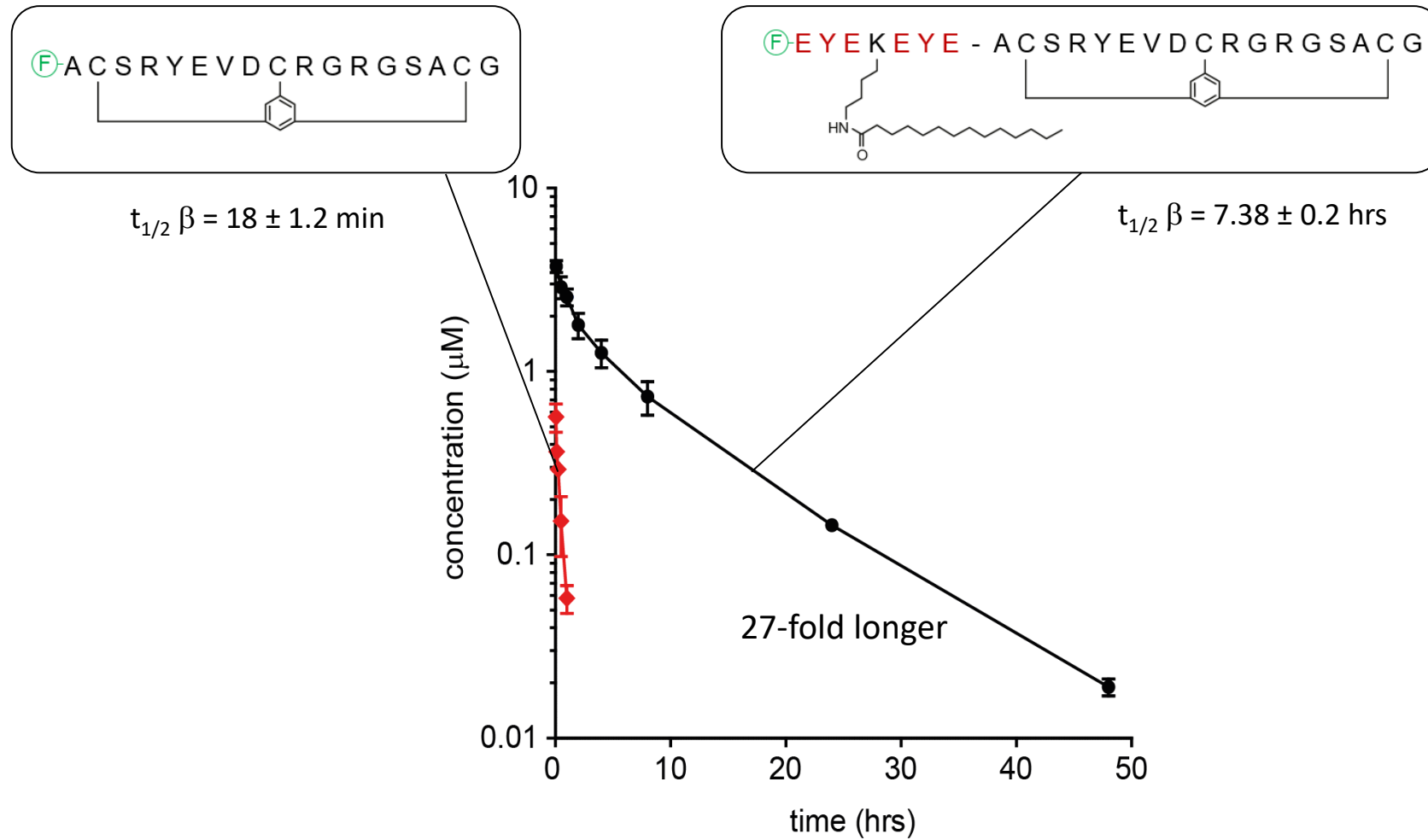
Short peptidic albumin-binding tag



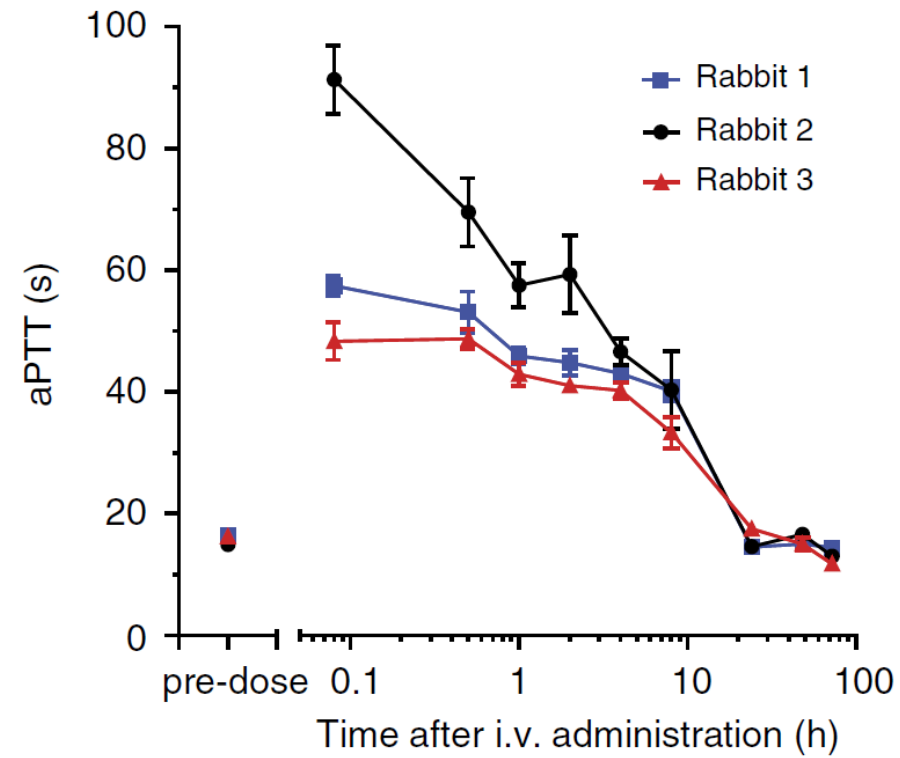
Pharmacokinetics in rats



Pharmacokinetics in rats



Application to FXII inhibitor

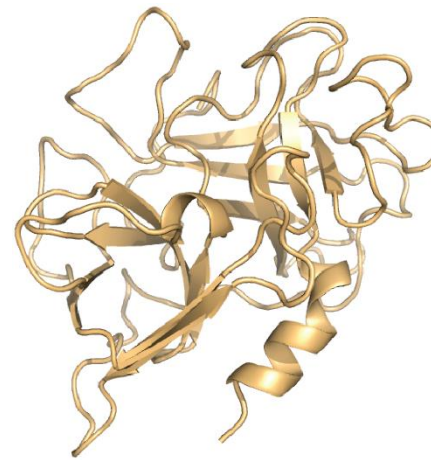


Zorzi, A., Wilbs, J., Middendorp, S., Deyle, K. and Heinis, C., *Nature Communications*, 2017

Bicyclic peptides to a broad range of targets

Plasma kallikrein	$K_i = 300 \text{ pM}$
uPA	$K_i = 28 \text{ nM}$
MMP-2	$K_i = 1.5 \text{ nM}$
PSA	$K_i = 47 \text{ nM}$
Her2	$K_d = 270 \text{ nM}$
Notch1 receptor	$K_d = 170 \text{ nM}$
β -catenin	$K_d = 6 \text{ }\mu\text{M}$
CAIX	$K_i = 50 \text{ nM}$
Sortase A	$K_i = 1.2 \text{ }\mu\text{M}$
Factor XIIa	$K_i = 370 \text{ pM}$
G-actin	$K_d = 5 \text{ nM}$
TNF α	$K_d = 300 \text{ nM}$
FKBP12	$K_d = 0.5 \text{ }\mu\text{M}$
IL-17	$K_d = 300 \text{ nM}$
Factor XIa	$K_i = 10 \text{ nM}$

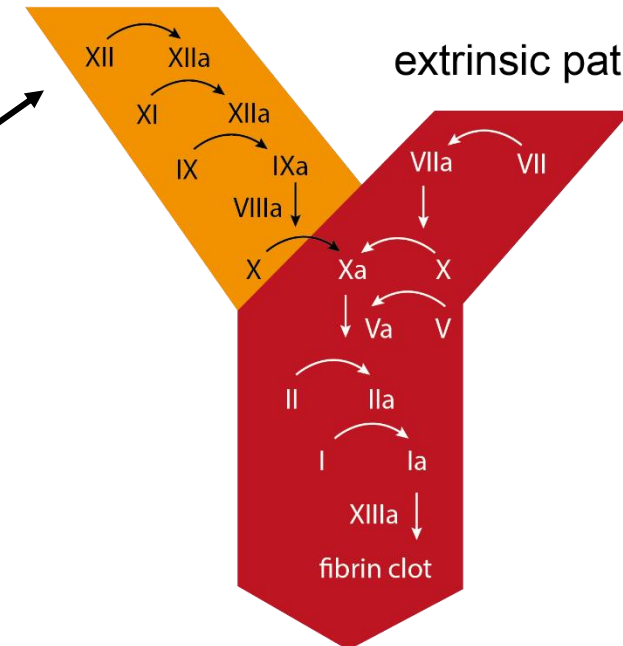
....



FXIIa

intrinsic pathway

extrinsic pathway



Spin-off company Bicycle Therapeutics

Spin-off

- Founded in 2009, based in Cambridge, UK
- > 120 employees
- Listed at Nasdaq (BCYC)

Three bicyclic peptides in clinical trials

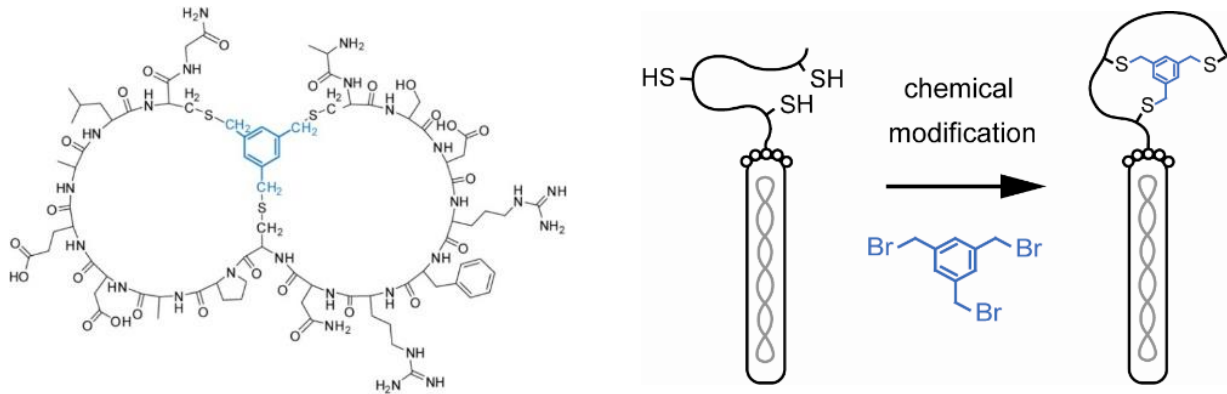
- BT5528 (anti-EphA2 peptide drug conjugate, Phase I/II)
- BT8009 (anti-Nectin-4 peptide drug conjugate, Phase I/II)
- BT5528 (anti-MT1-MMP peptide drug conjugate, Phase I/IIa)

Bicycle

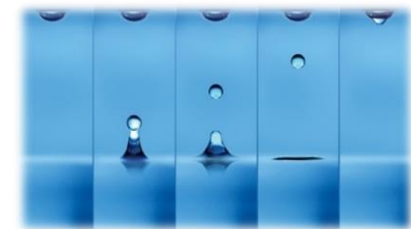
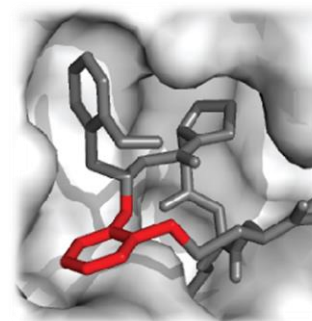
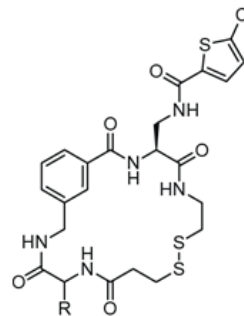


Talk overview

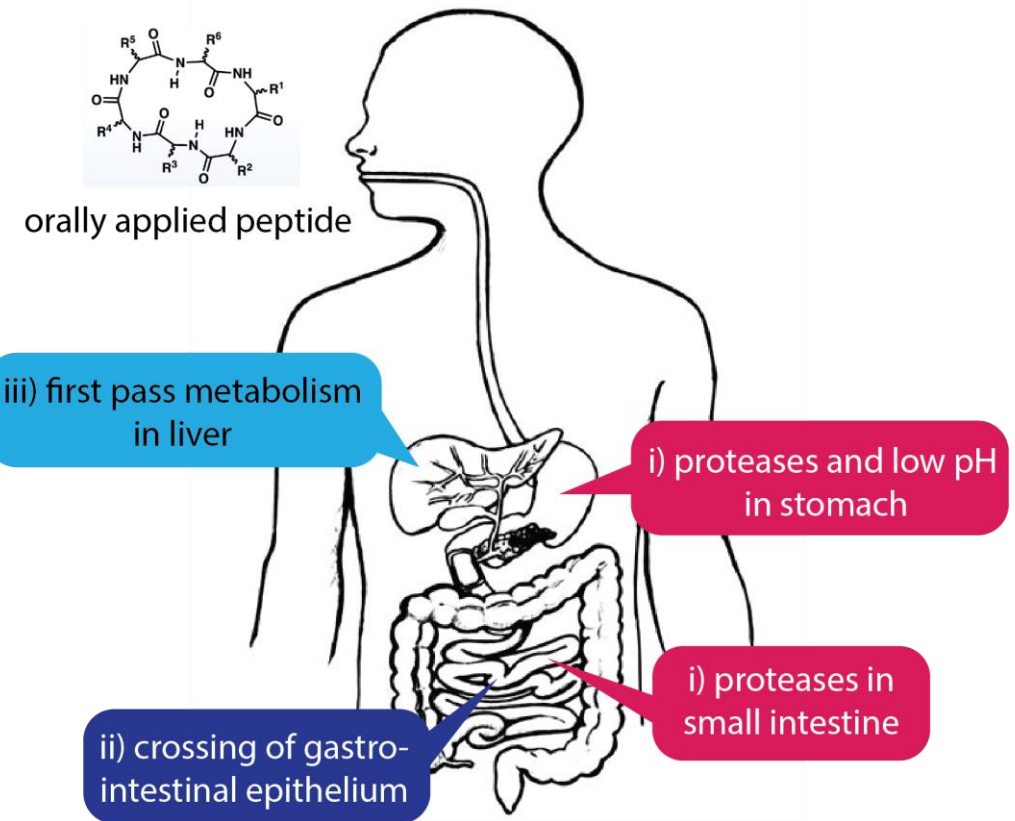
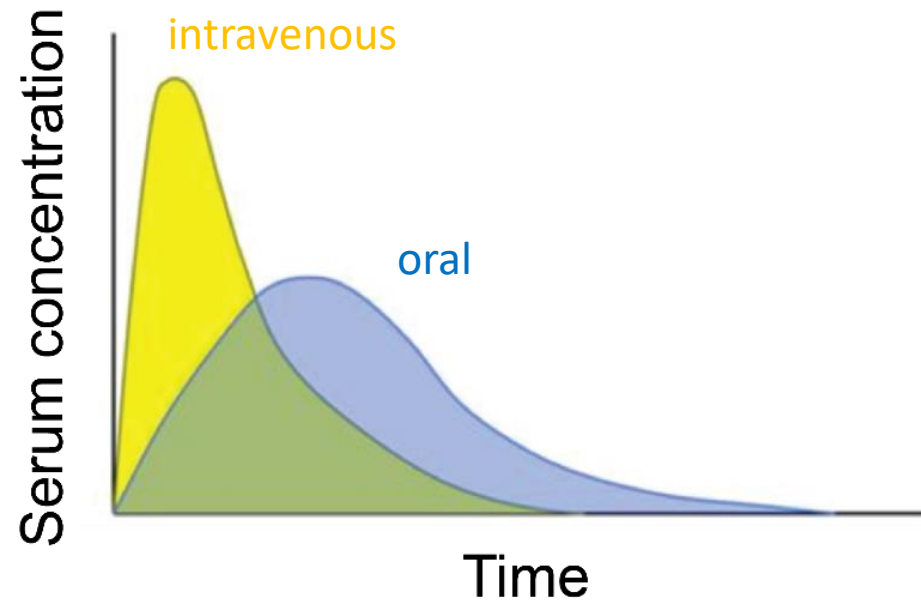
1. Bicyclic peptides developed by phage display



2. Peptide macrocycles developed by high-throughput synthesis at a nanomole scale



Can peptides be applied orally?



Which cyclic peptides can cross membranes?

Oral vs. not oral macrocyclic drugs

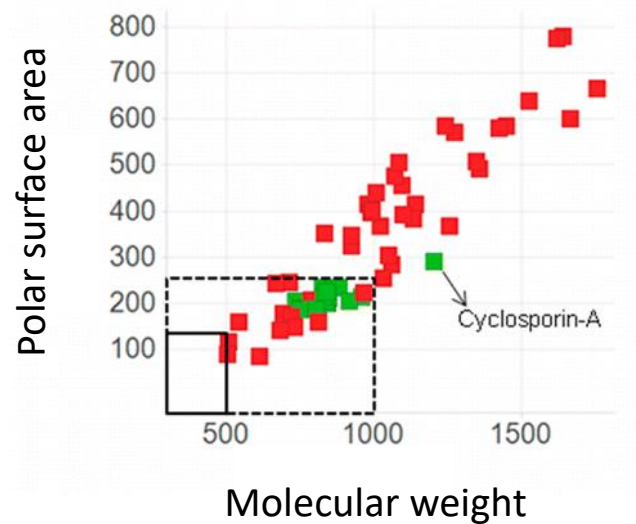


Figure from Giordanetto and Kihlberg,
J. Med. Chem., 2014

Journal of
**Medicinal
Chemistry**

How Big Is Too Big for Cell Permeability?

Pär Matsson[†] and Jan Kihlberg^{*†}

Which cyclic peptides can cross membranes?

Oral vs. not oral macrocyclic drugs

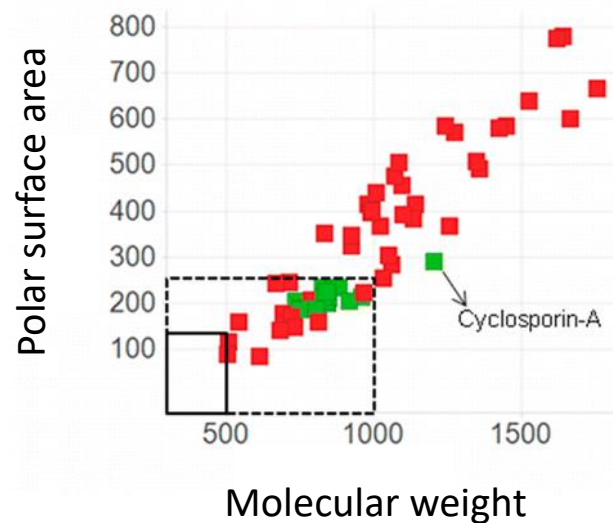


Figure from Giordanetto and Kihlberg,
J. Med. Chem., 2014

Journal of
**Medicinal
Chemistry**

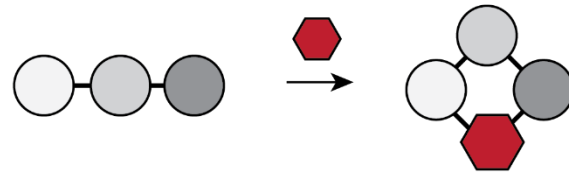
How Big Is Too Big for Cell Permeability?

Pär Matsson[†] and Jan Kihlberg^{*,†}

Key factors:

- Molecular weight: < 750 Da
- Polar surface area: < 250 Å²
- Number of H-bond donors: < 6

Combinatorial synthesis of peptide-based macrocycles



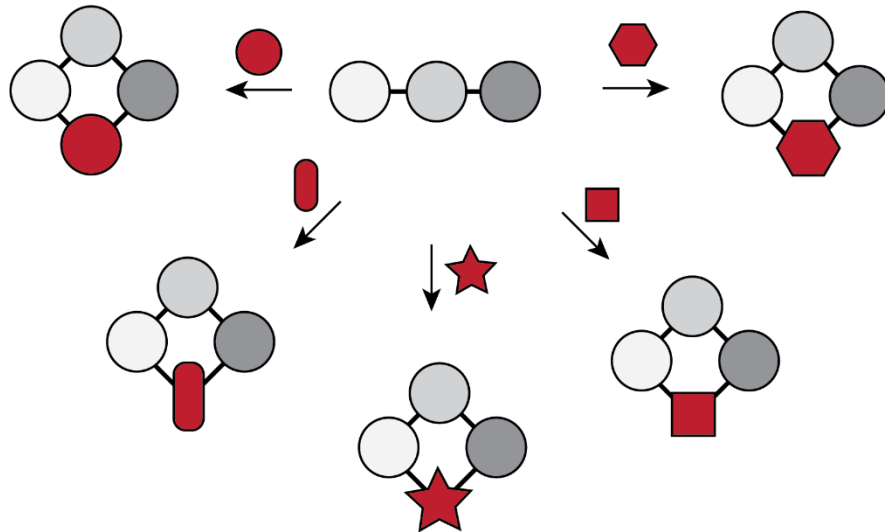
Combinatorial synthesis of peptide-based macrocycles

Number example:

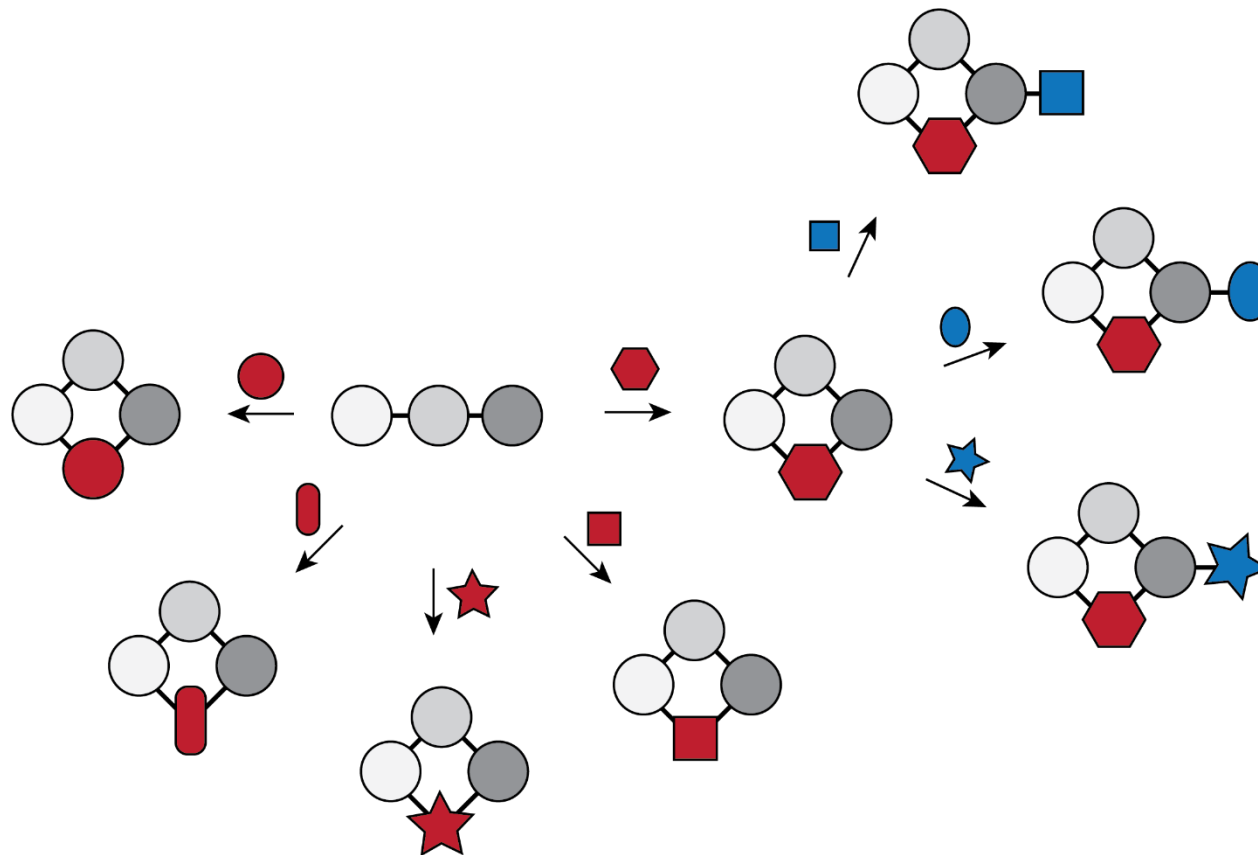
1,000 peptides

x

10 linkers



Combinatorial synthesis of peptide-based macrocycles



Number example:

1,000 peptides

x

10 linkers

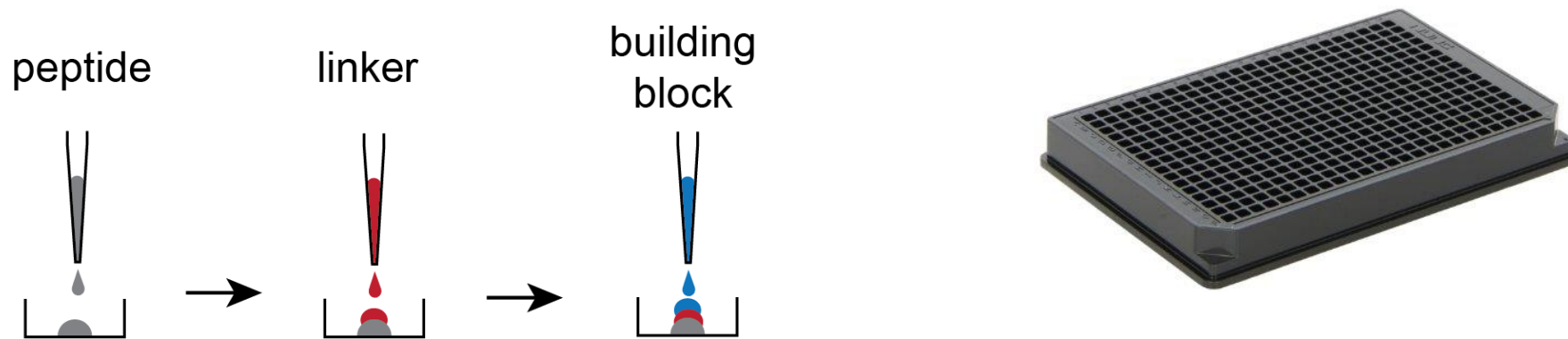
x

10 building blocks

= 100,000 macrocycles

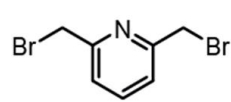
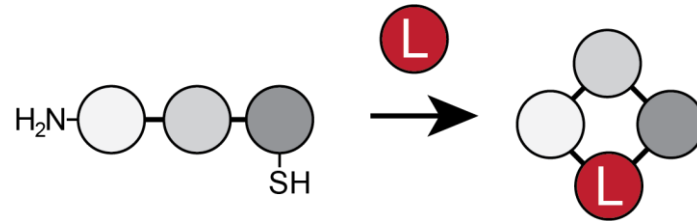
Combinatorial synthesis of peptide-based macrocycles

Combine reagents, react & screen

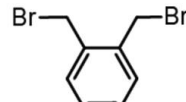


No purification → large libraries

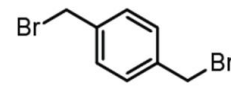
Efficient cyclization reaction: Thiol-to-amine ligation



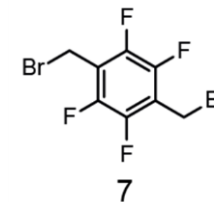
1



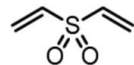
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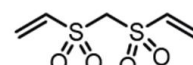
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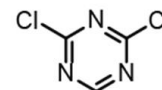
7



4

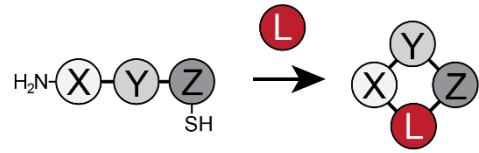


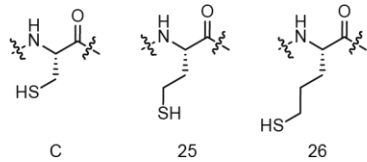
5



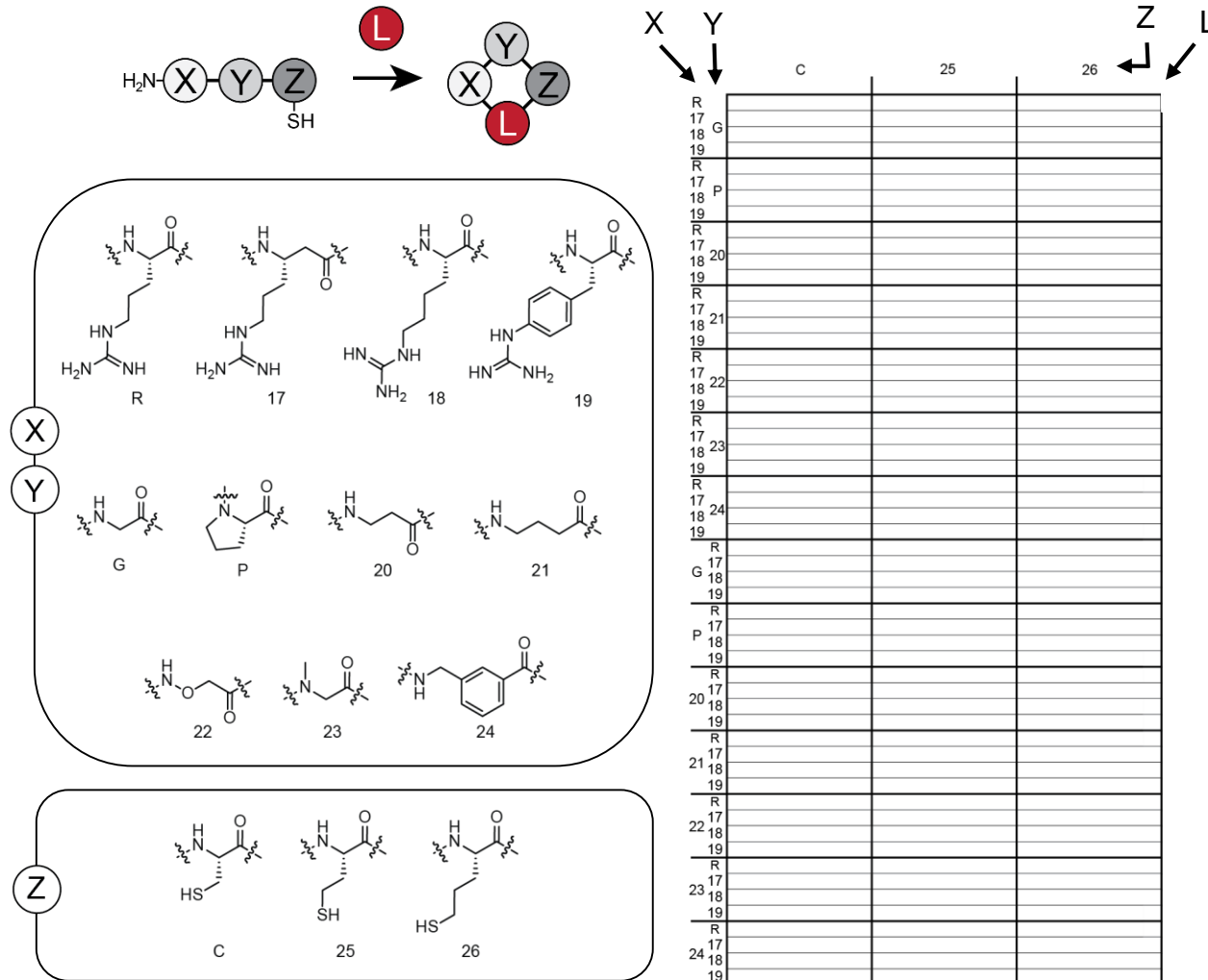
6

Pilot-scale library



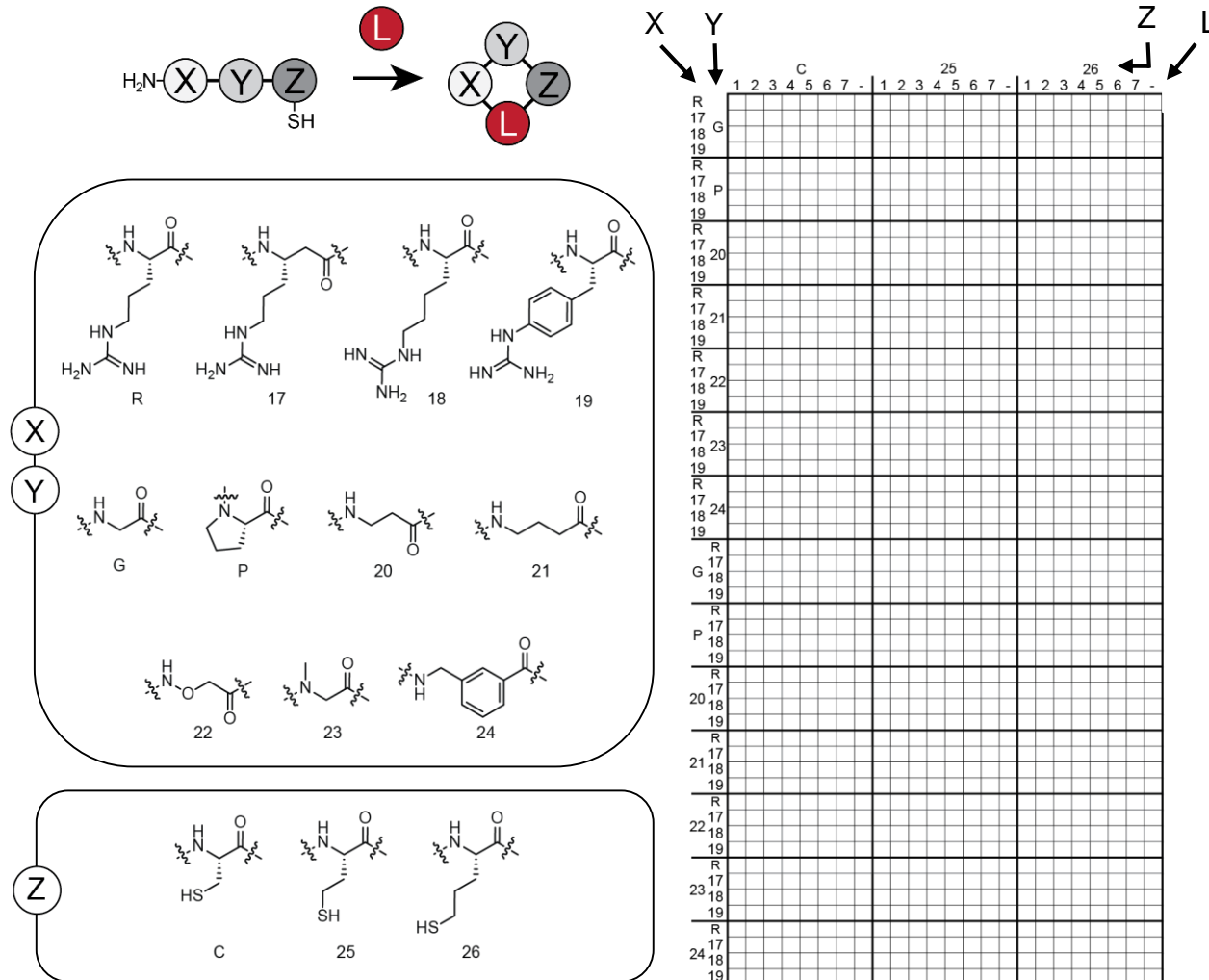


Pilot-scale library

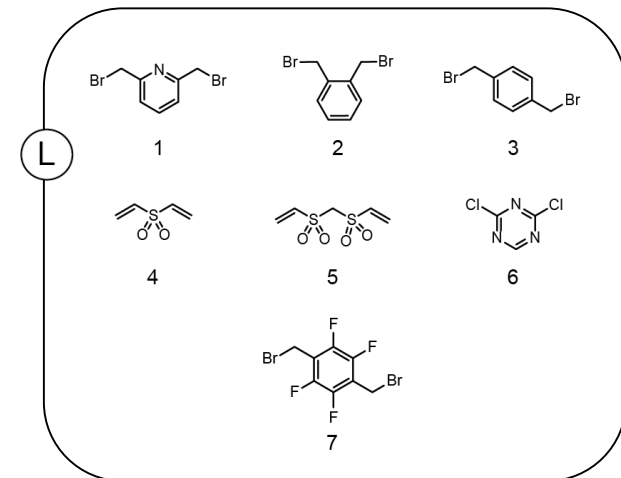


Solid-phase peptide
synthesis of 168
peptides

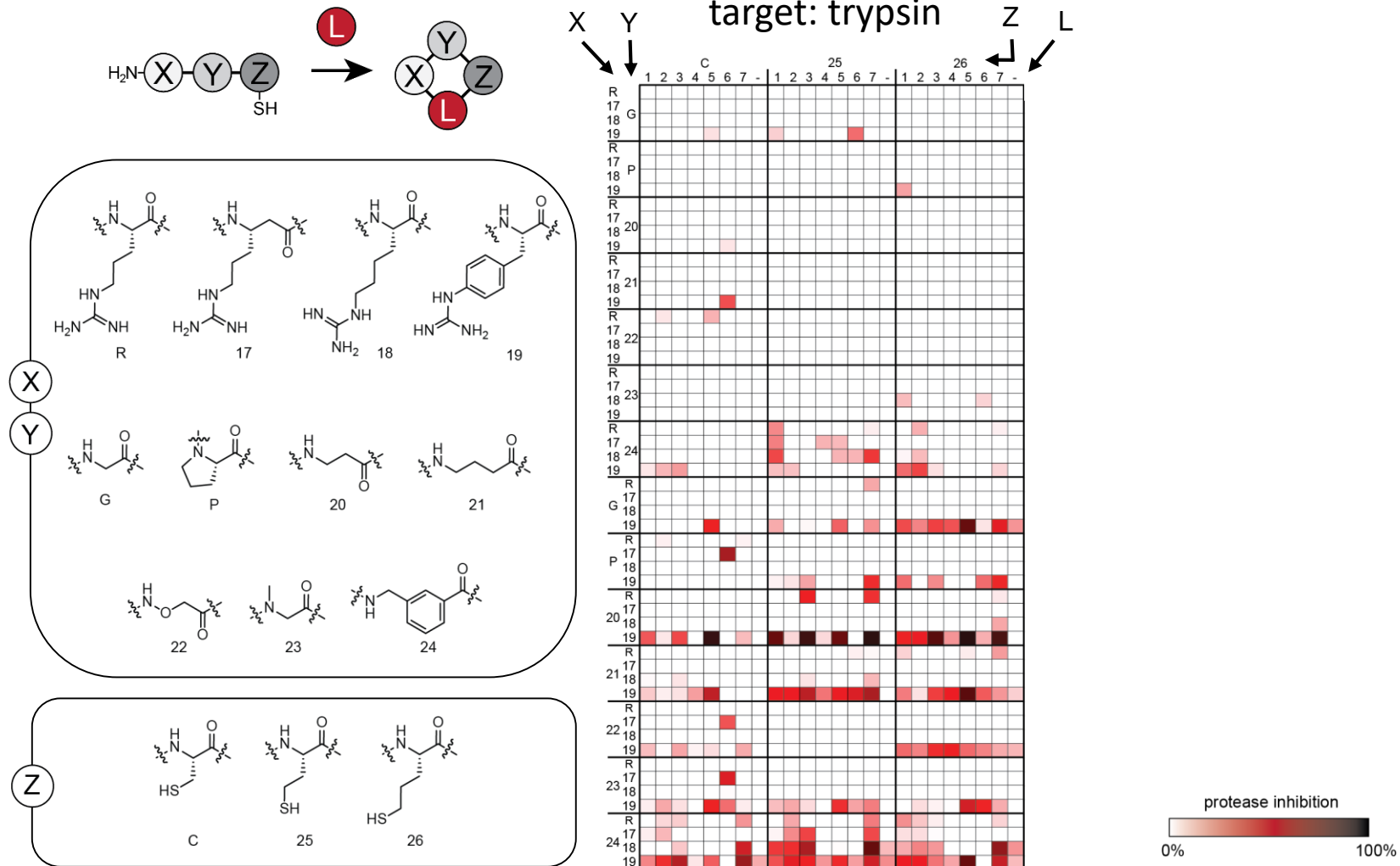
Pilot-scale library



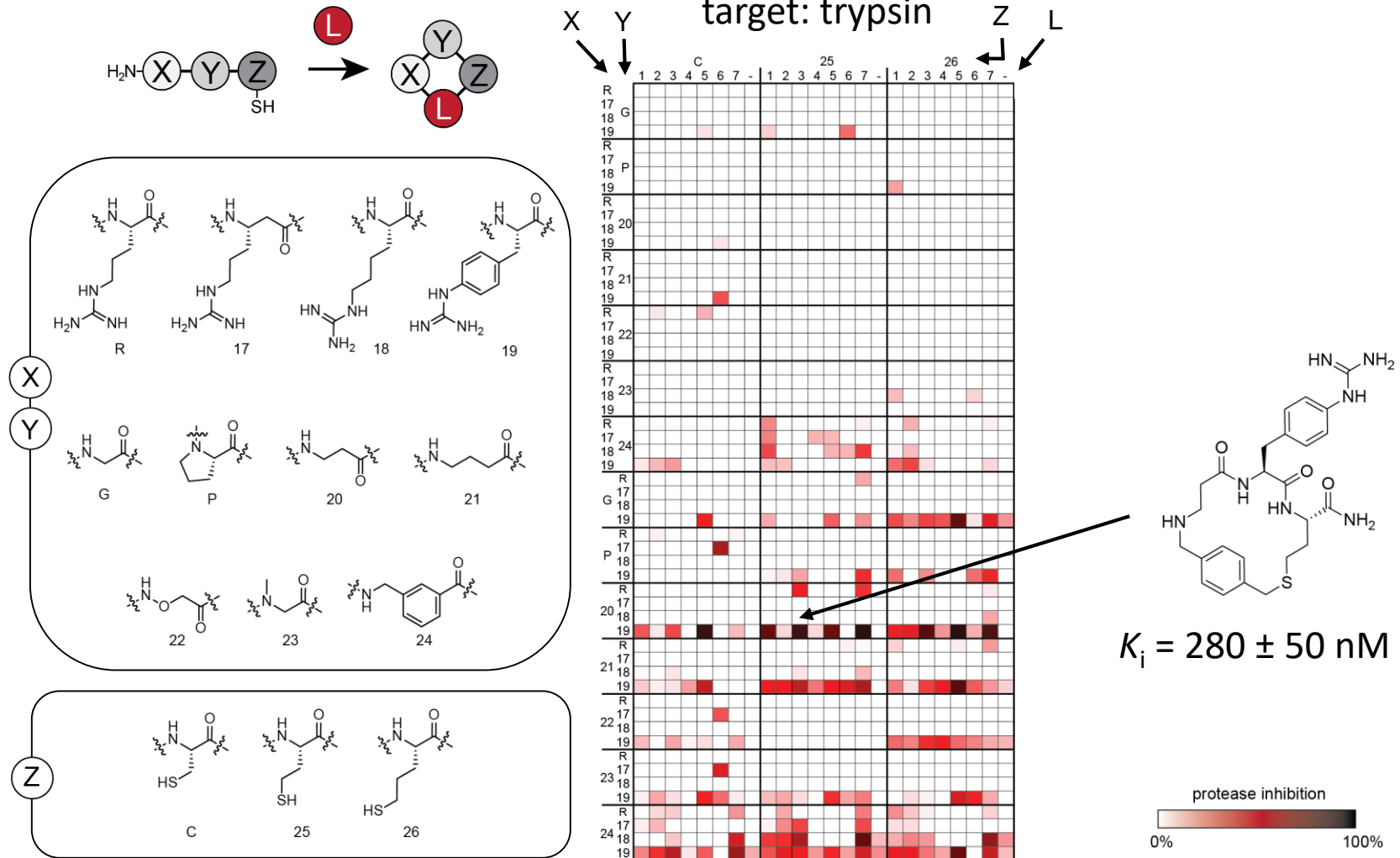
168 peptides × 7 linkers
= 1176 macrocycles



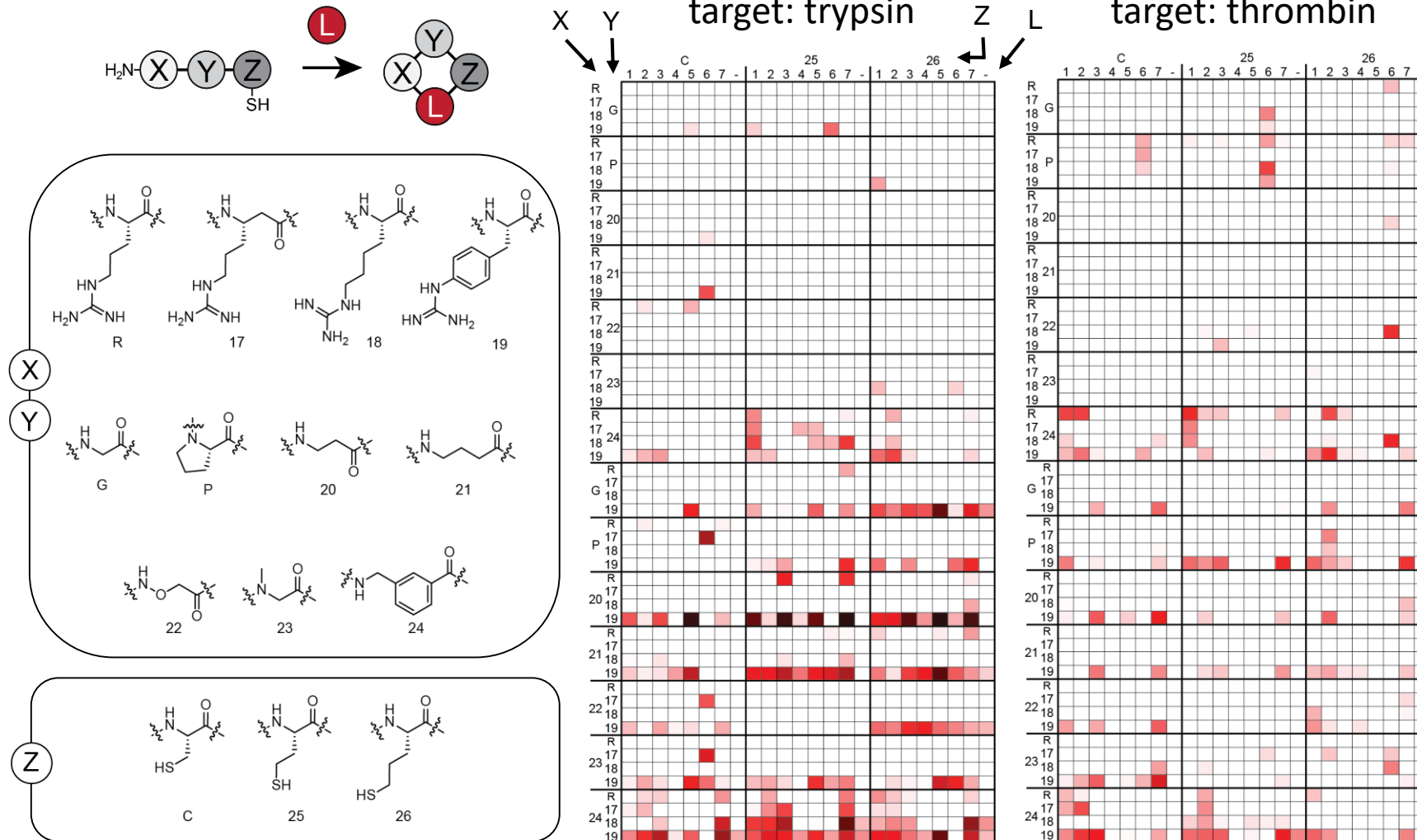
Pilot-scale library



Pilot-scale library



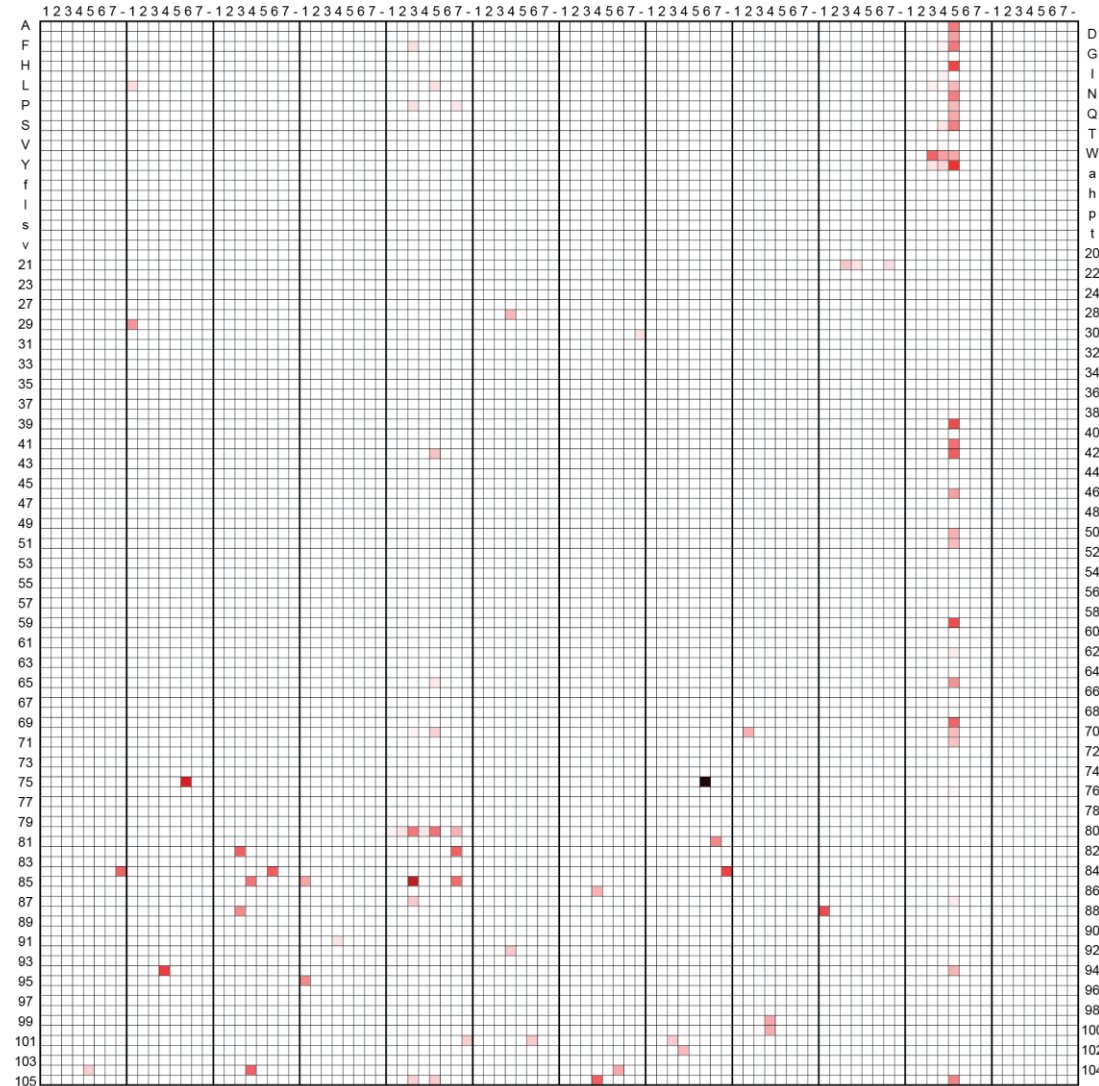
Pilot-scale library



8,988 macrocycle library



1,284 peptides × 7 linkers

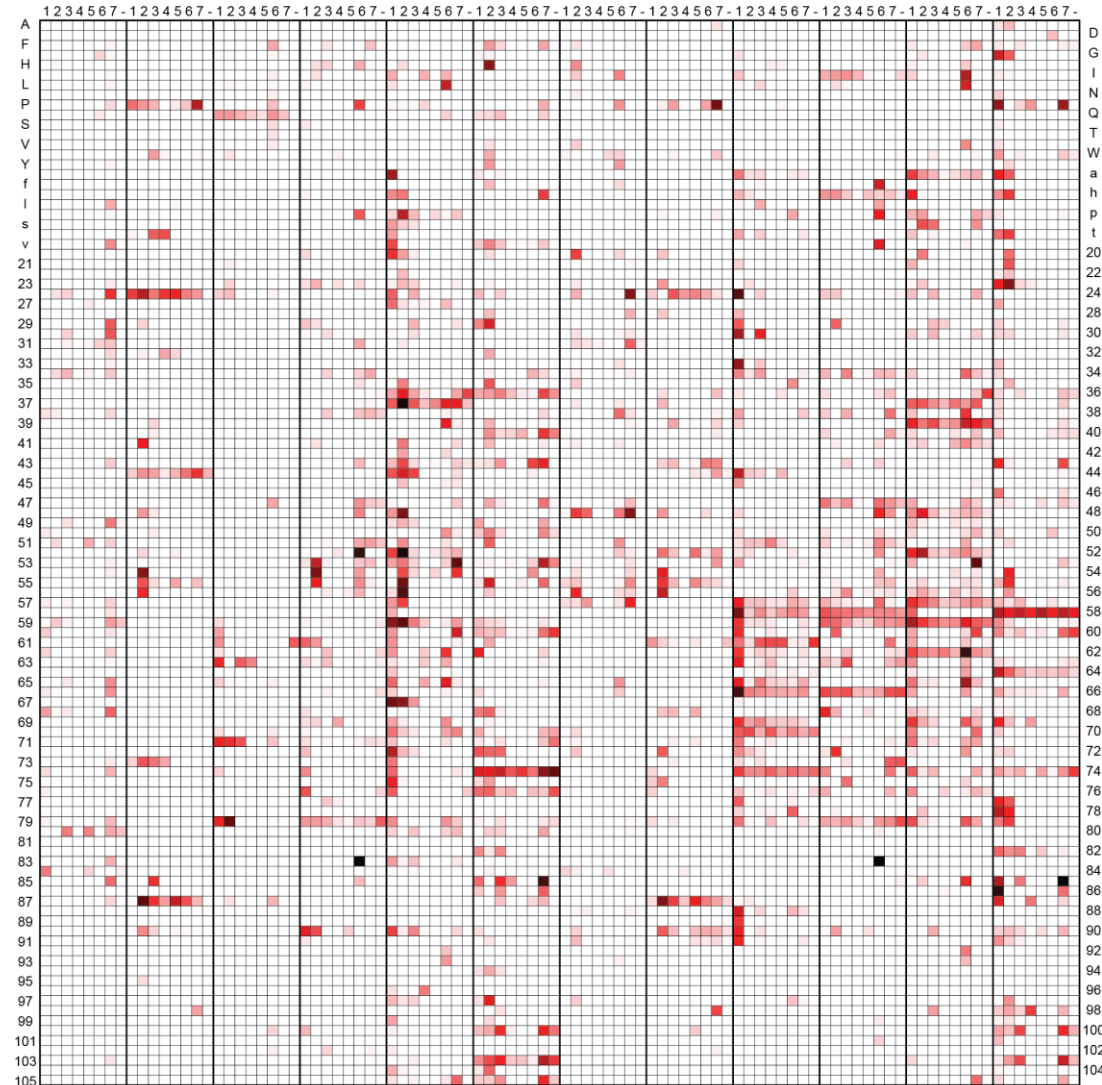


8,988 macrocycle library

thrombin inhibition

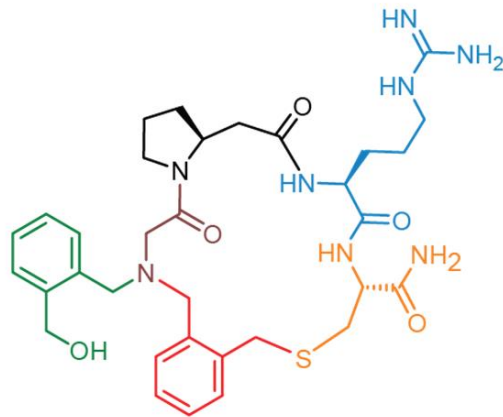


1,284 peptides × 7 linkers

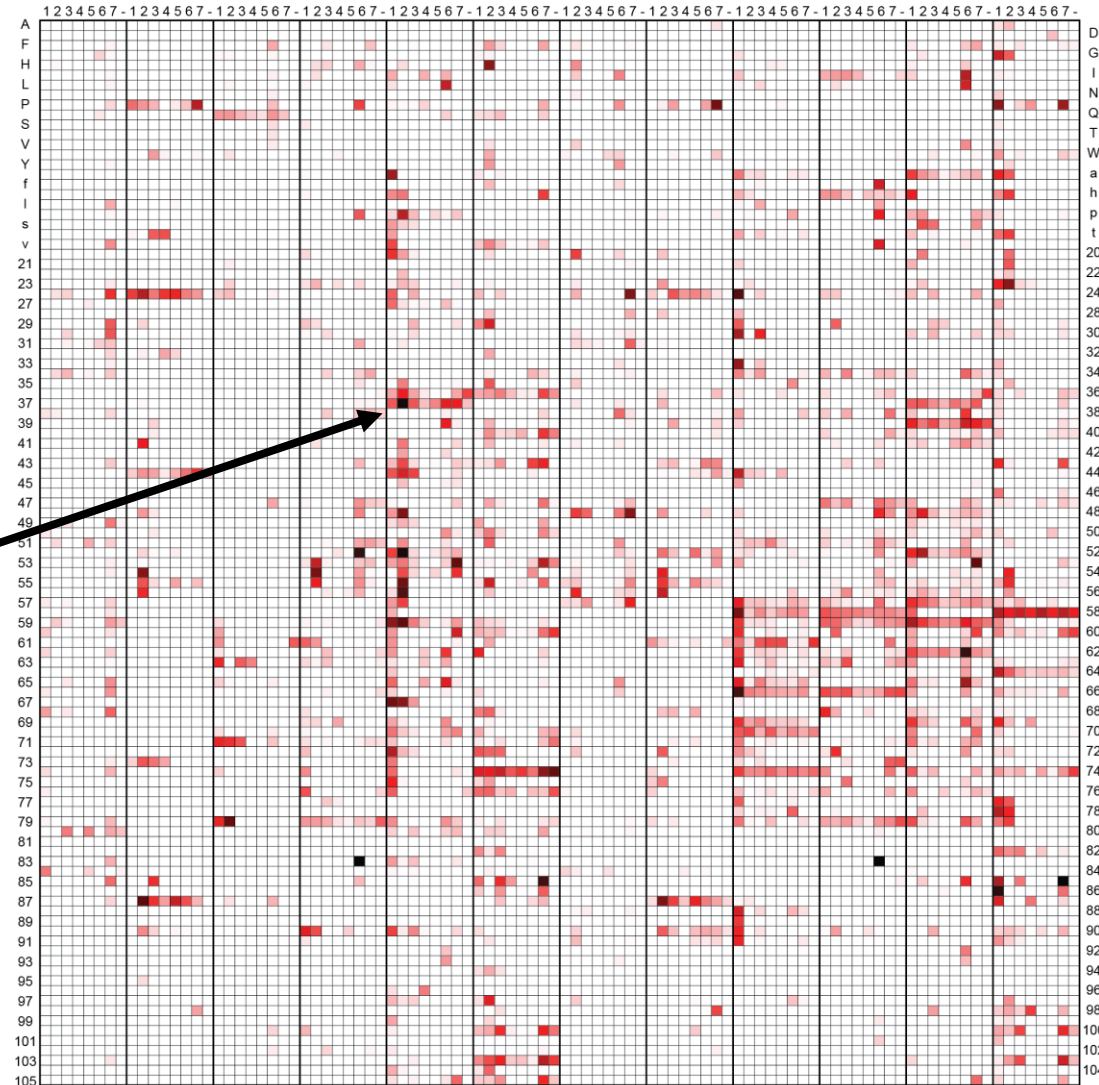


8,988 macrocycle library

thrombin inhibition

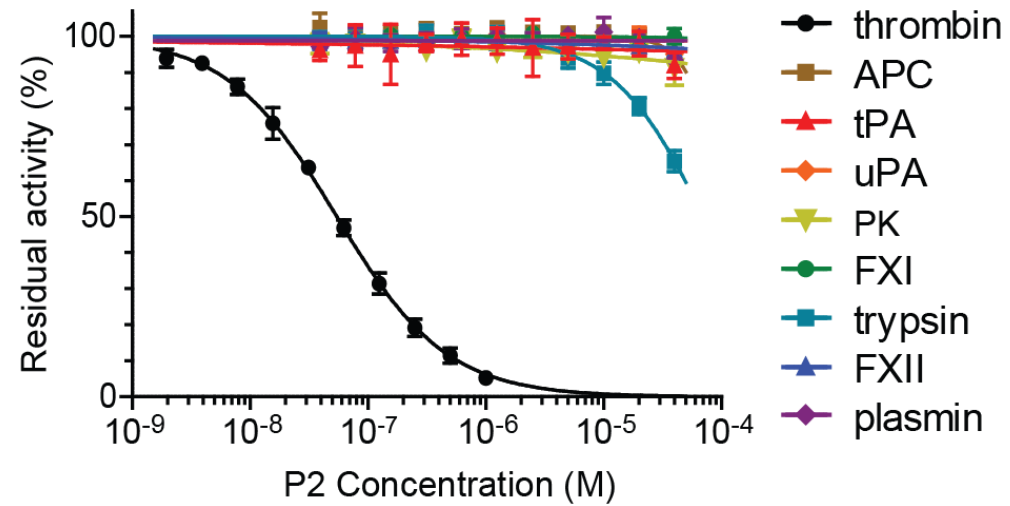


1,284 peptides × 7 linkers

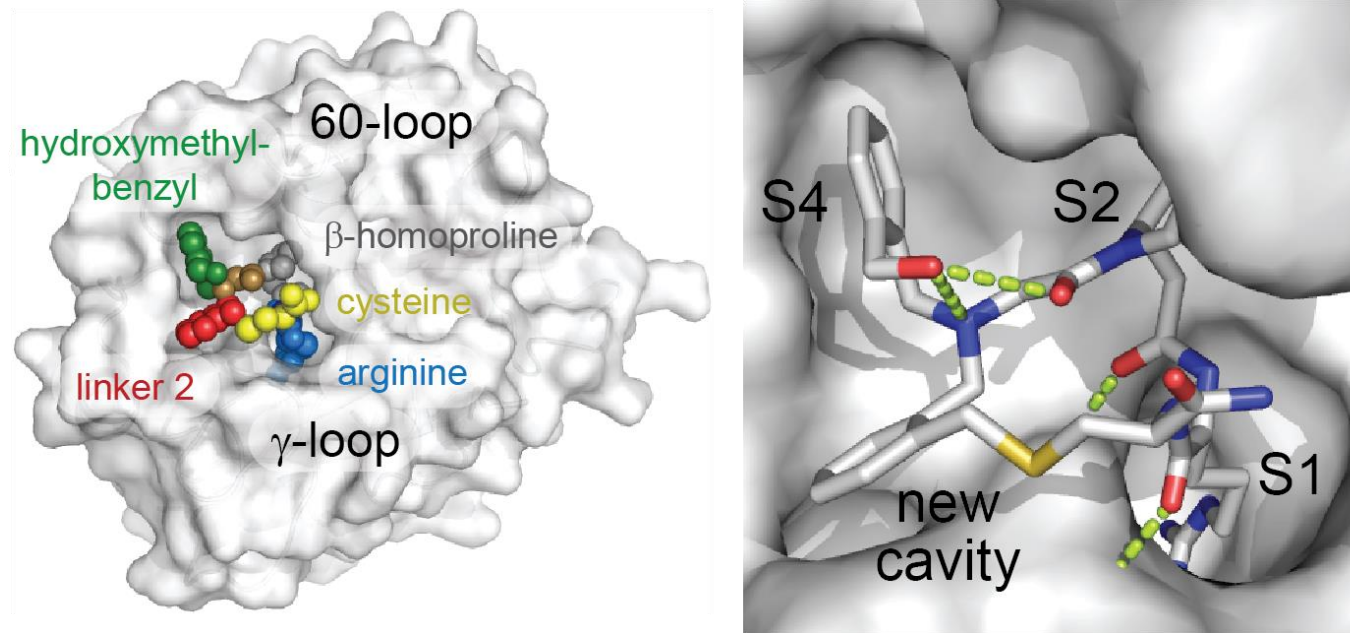


Thrombin inhibitor: specificity

$$K_i = 42 \text{ nM}$$

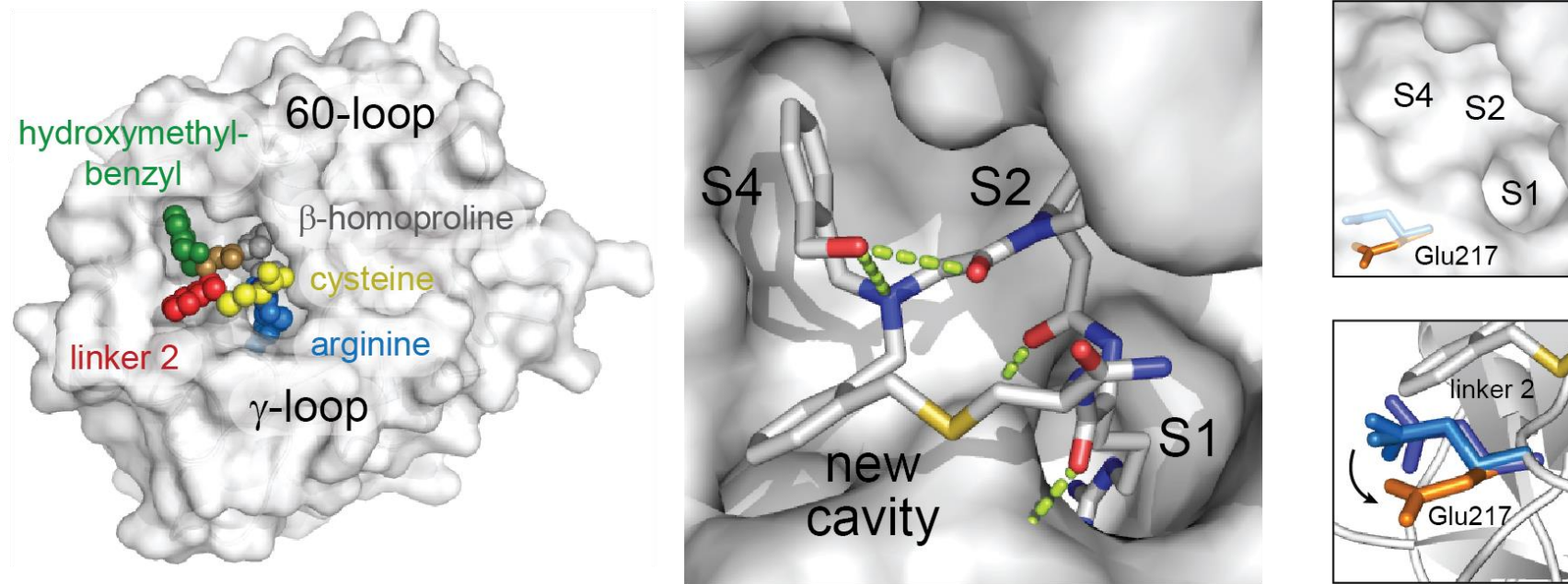


Thrombin inhibitor: structure



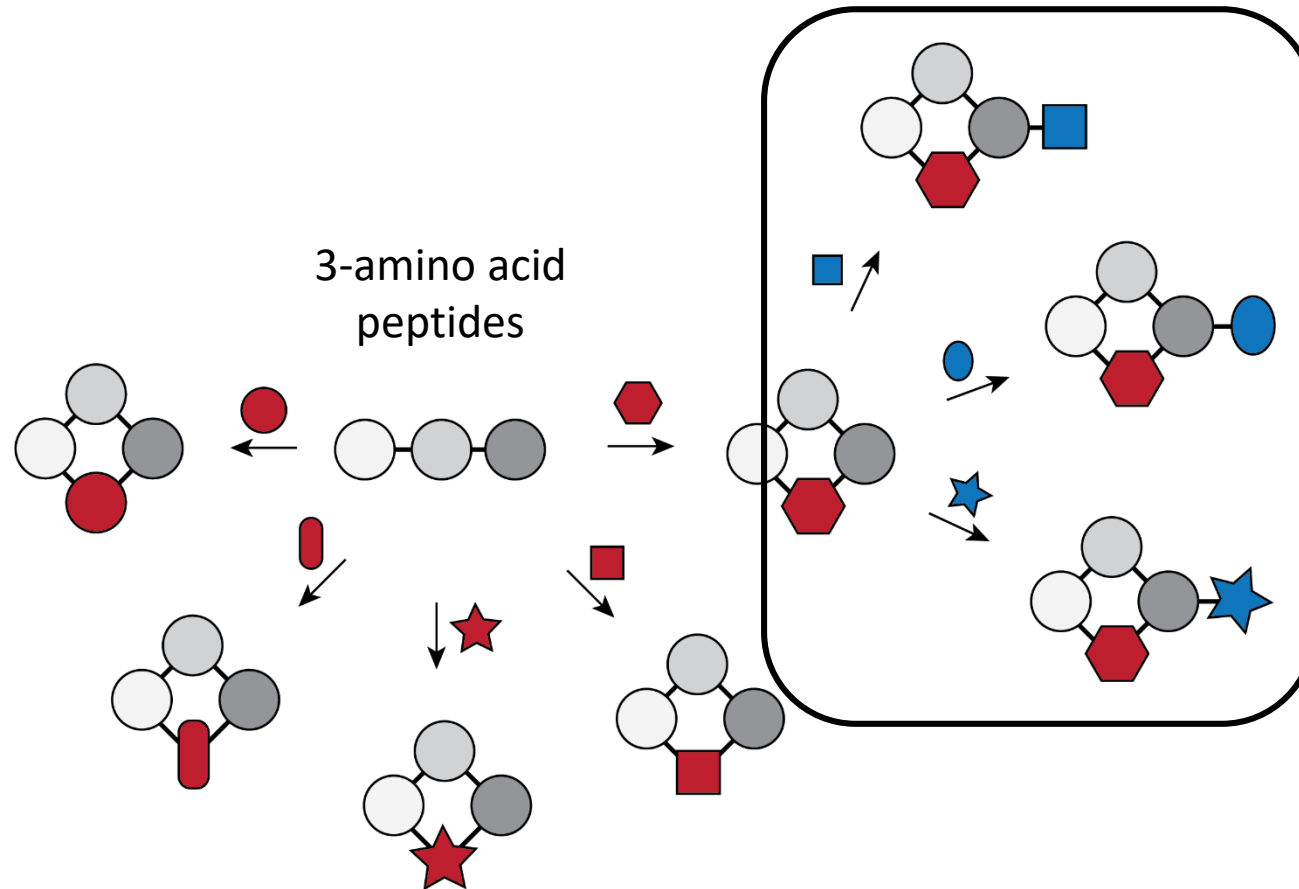
Structure solved by
Prof. Alessandro Angelini (Ca' Foscari University of Venice) and Prof. Laura Cendron (University of Padova)

Thrombin inhibitor: structure



Kale, S.S., Bergeron-Brlek, M., Wu, Y., Kumar, M.G., Pham, M.V., Bortoli, J., Vesin, J., Kong, X.-D., Franco Machado, J., Deyle, K., Gonschorek, P., Turcatti, G., Cendron, L., Angelini, A. and Heinis, C., *Science Advances*, 2019

Combinatorial synthesis of peptide-based macrocycles



Number example:

1,000 peptides

x

10 linkers

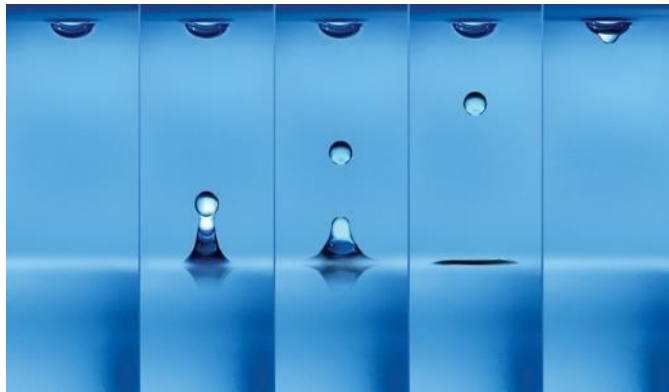
x

10 building blocks

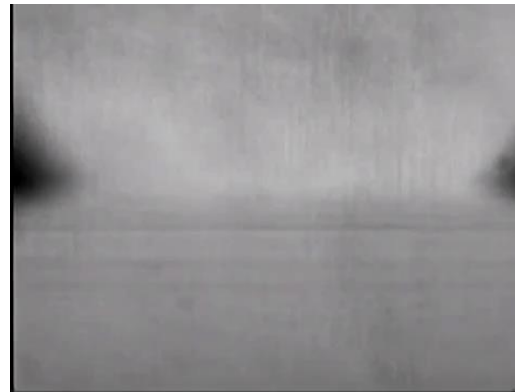
= 100,000 macrocycles

Macrocycles at a nanomole scale

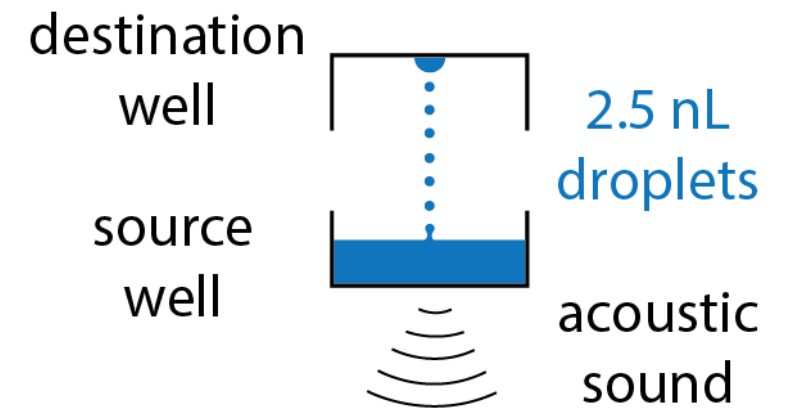
Acoustic droplet ejection (ADE)



Acoustic dispensing
(image from C&EN)



Acoustic dispensing
(film from Labcyte)



Macrocycles at a nanomole scale

Acoustic droplet ejection (ADE)



Our acoustic dispensing platform



1536 microwell plates

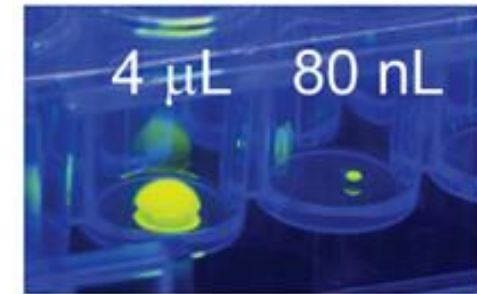


Photo of droplets (fluorescein)

Macrocycles at a nanomole scale

Acoustic droplet ejection (ADE)



Our acoustic dispensing platform



1536 microwell plates

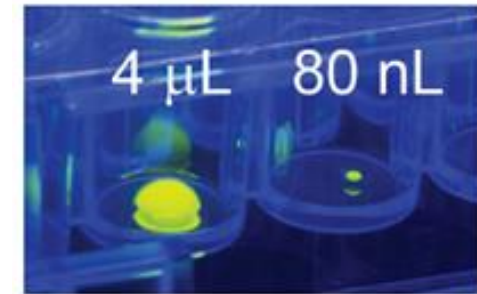


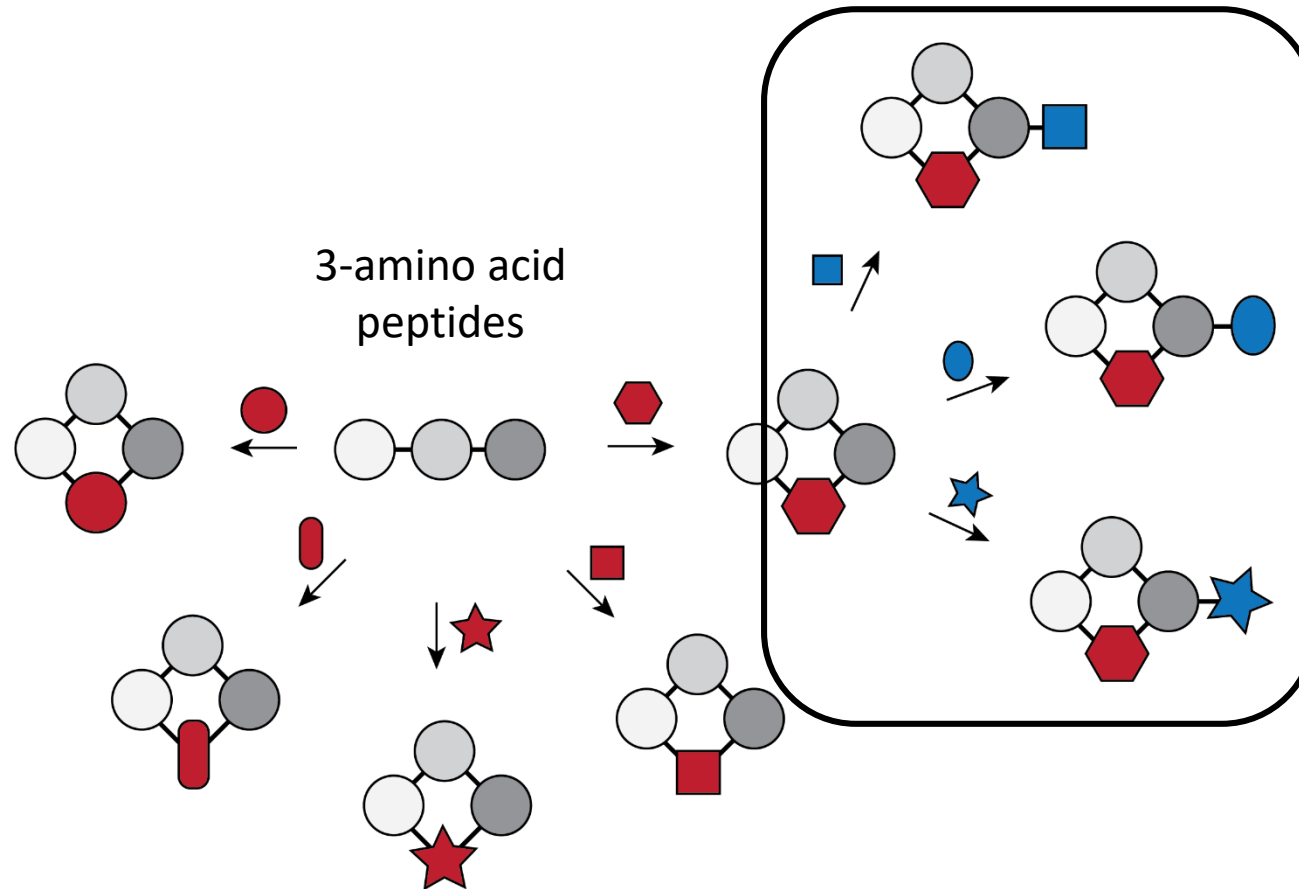
Photo of droplets (fluorescein)

synthesis and assay
in the same microwell plate

assay reagents



Combinatorial synthesis of peptide-based macrocycles



Number example:

1,000 peptides

x

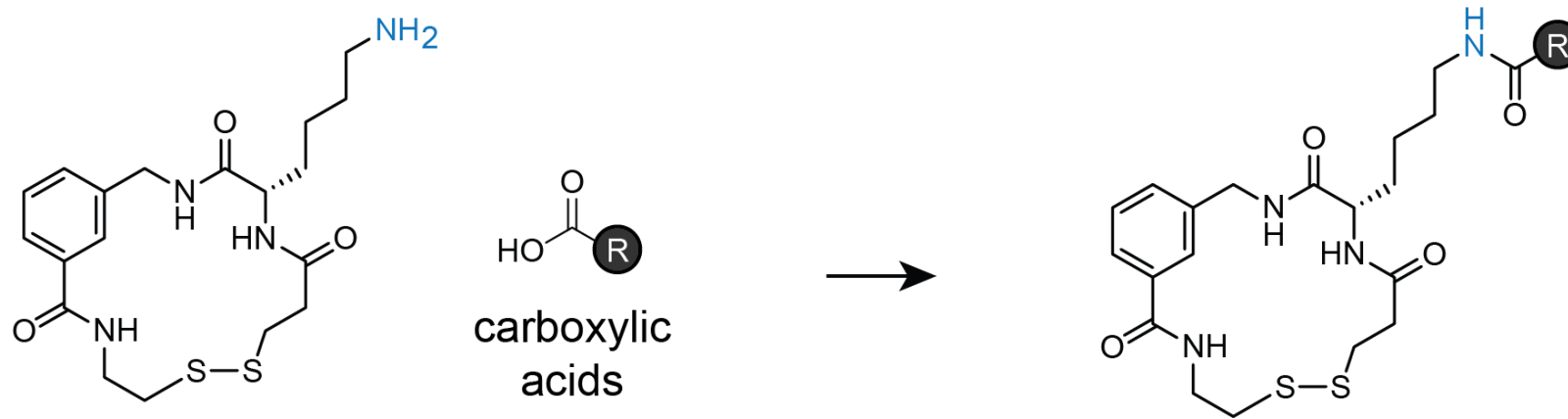
10 linkers

x

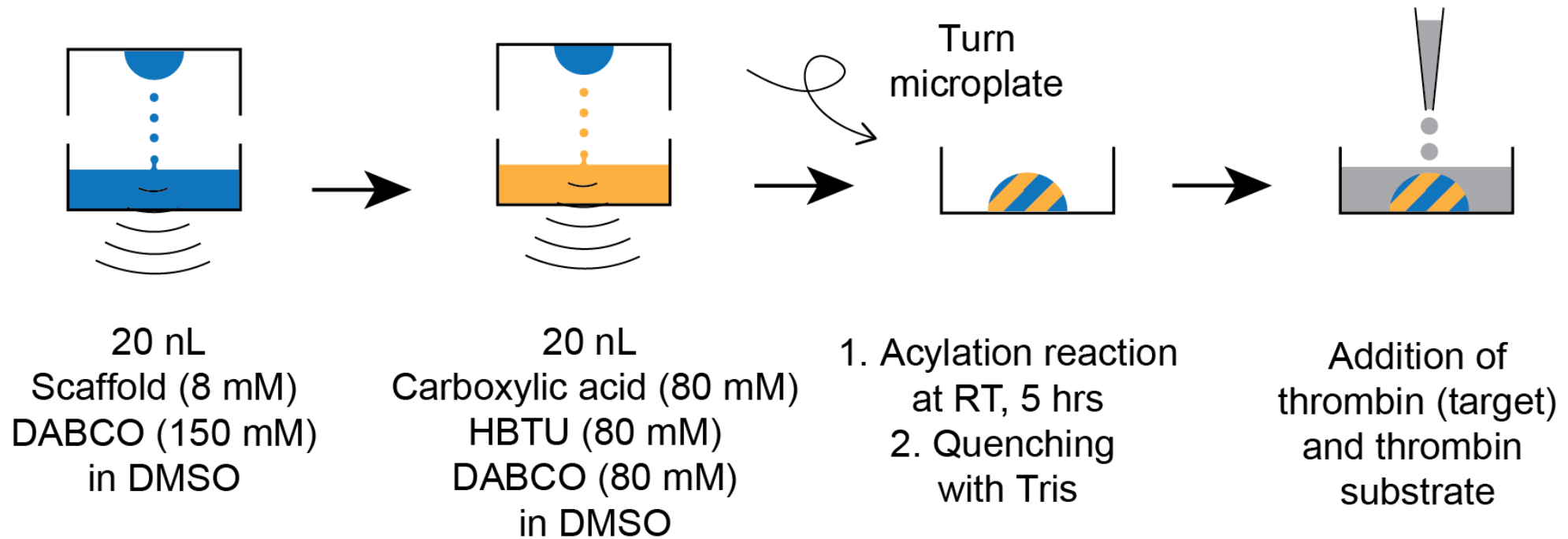
10 building blocks

= 100,000 macrocycles

Acylation of amino groups

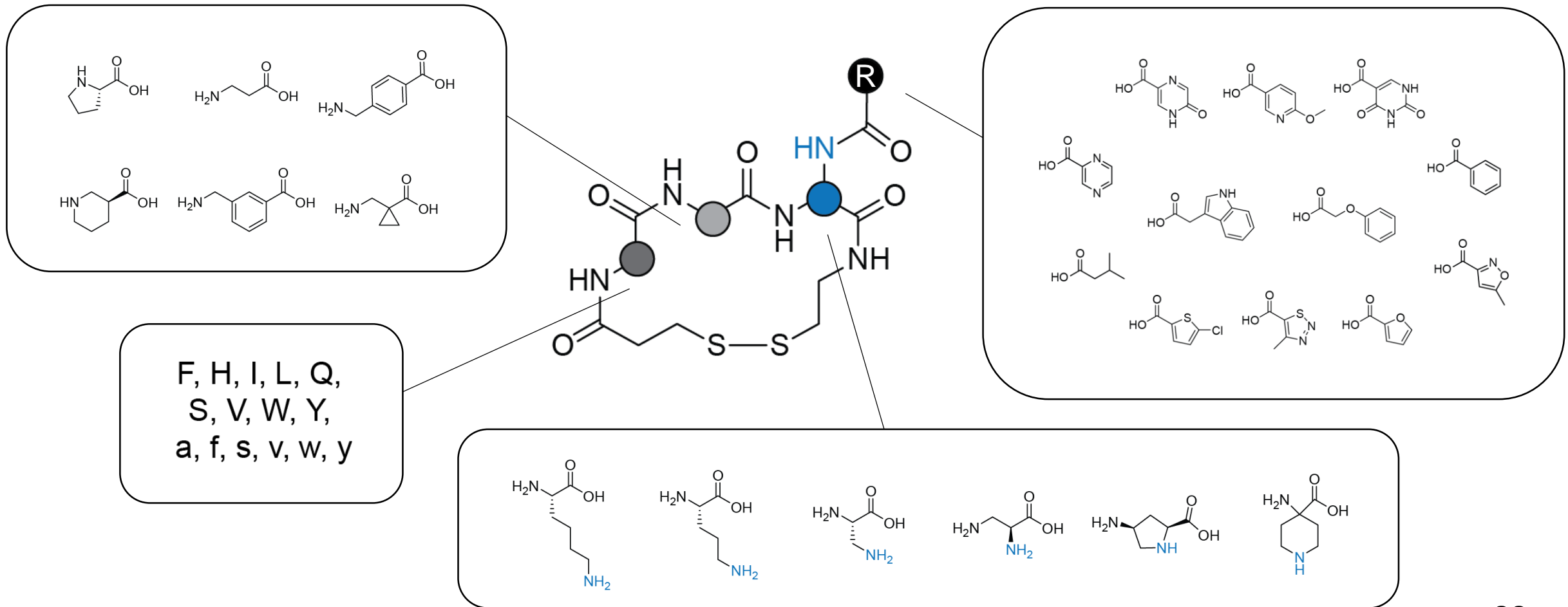


Synthesis of macrocycles at a nanomole scale

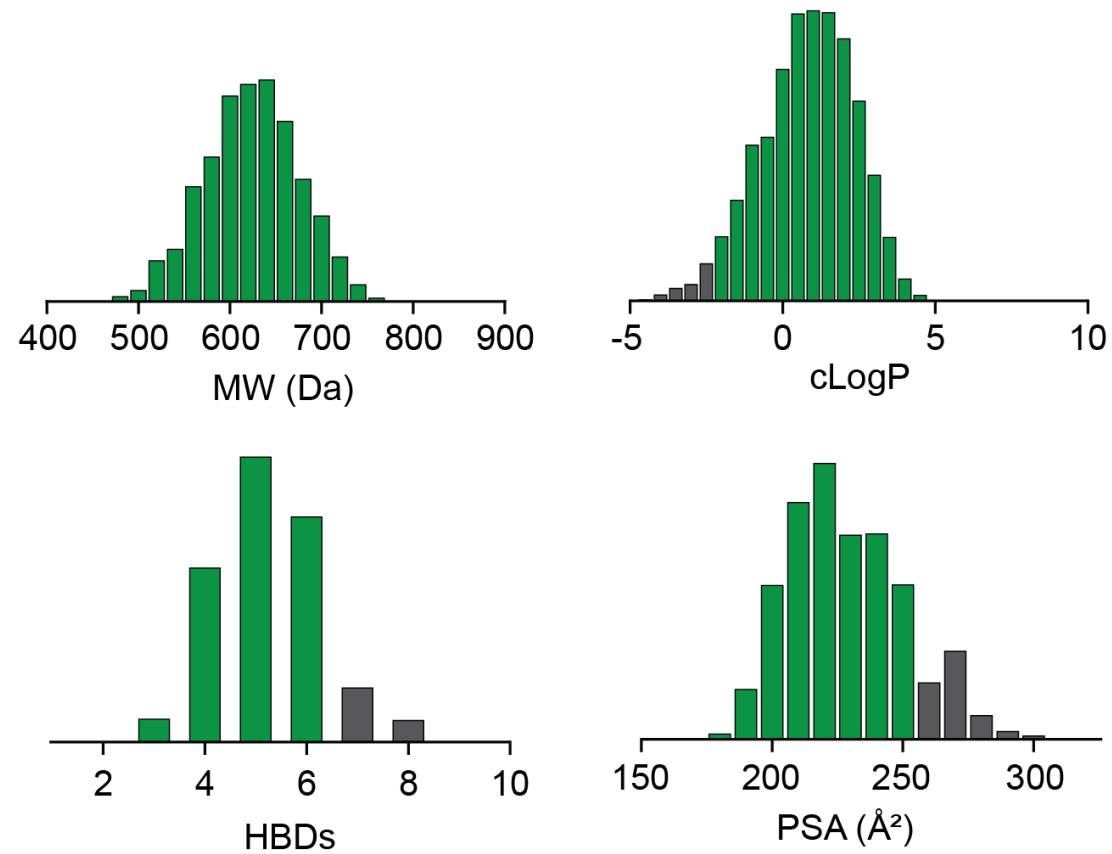


Synthesis of 3,840 macrocycles

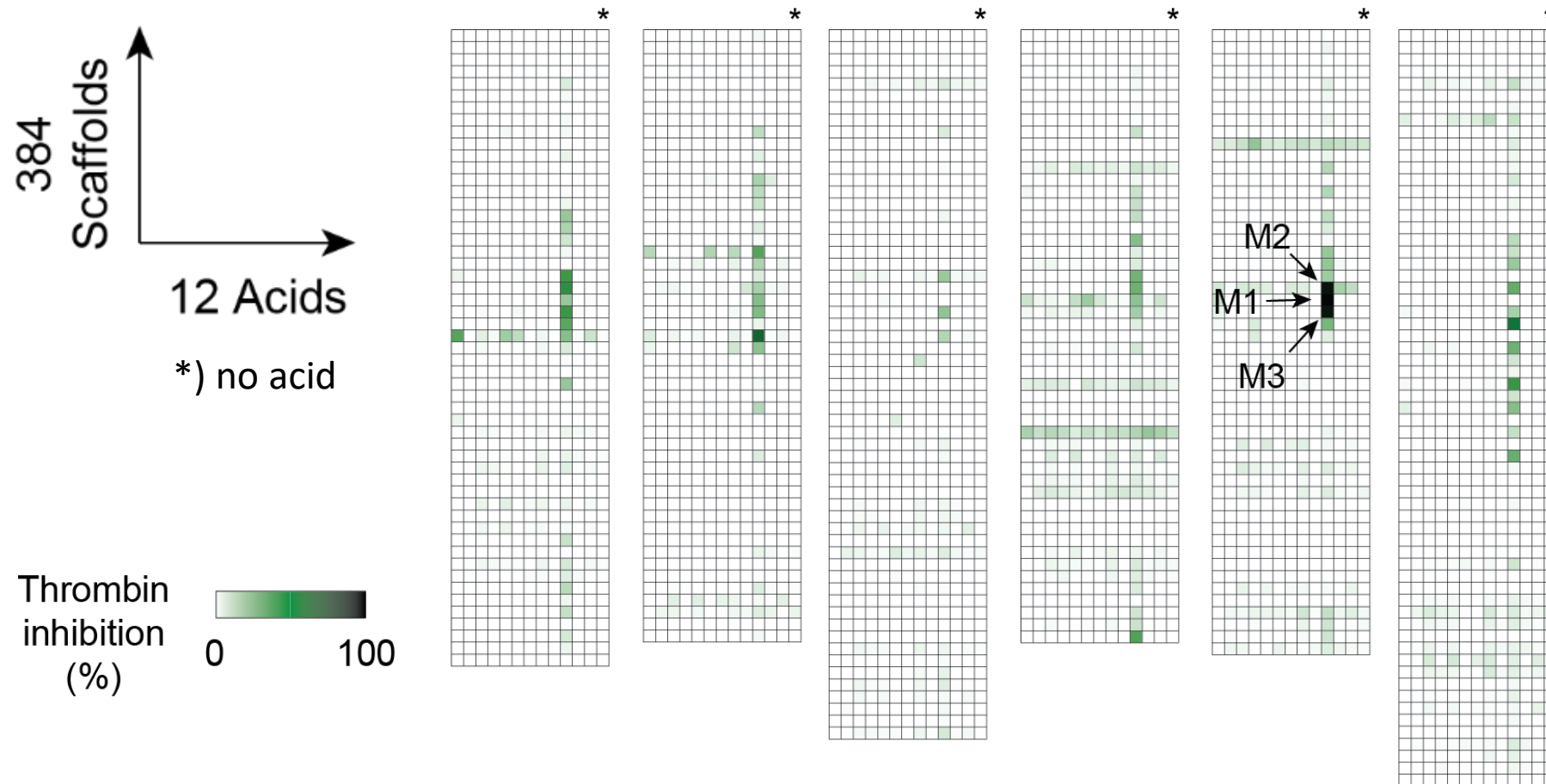
384 scaffolds \times 12 carboxylic acids



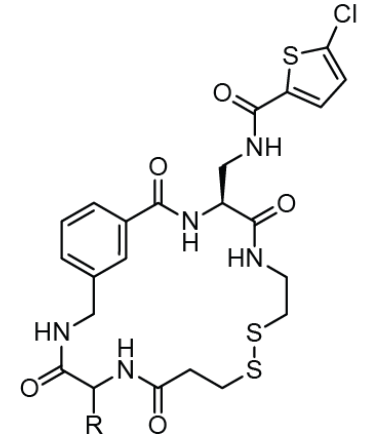
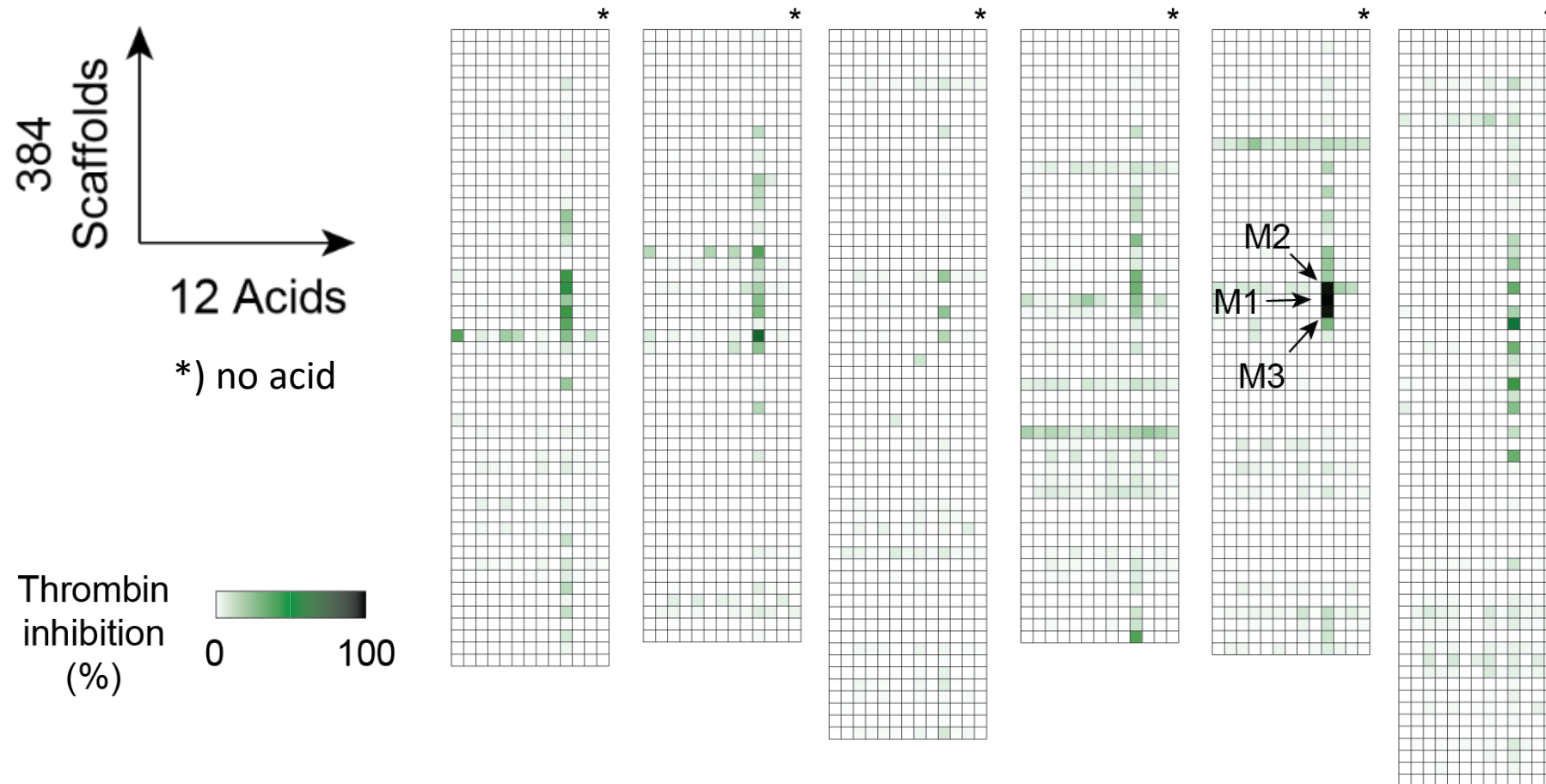
Synthesis of 3,840 macrocycles tailored for thrombin




Thrombin inhibition screen

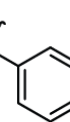



Thrombin inhibition screen



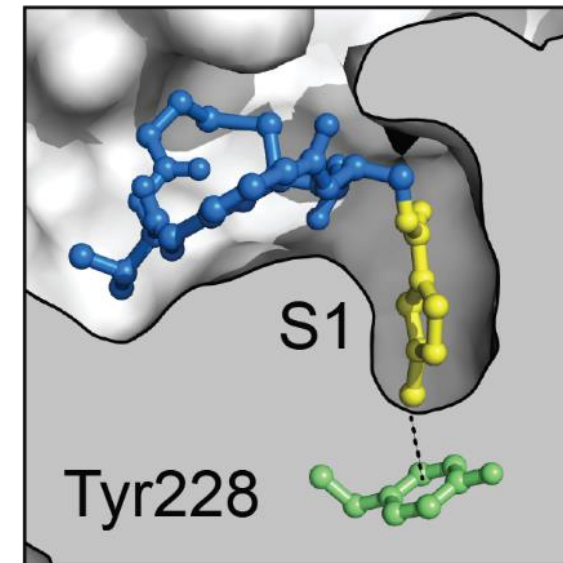
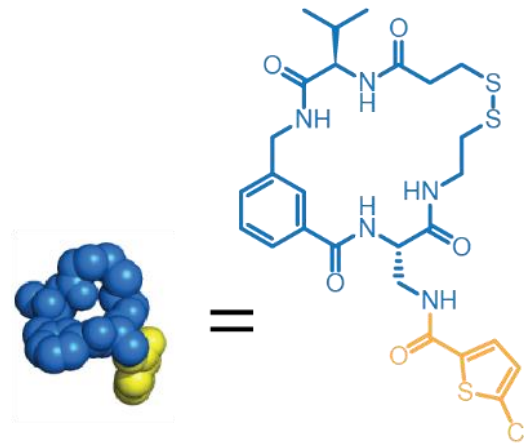
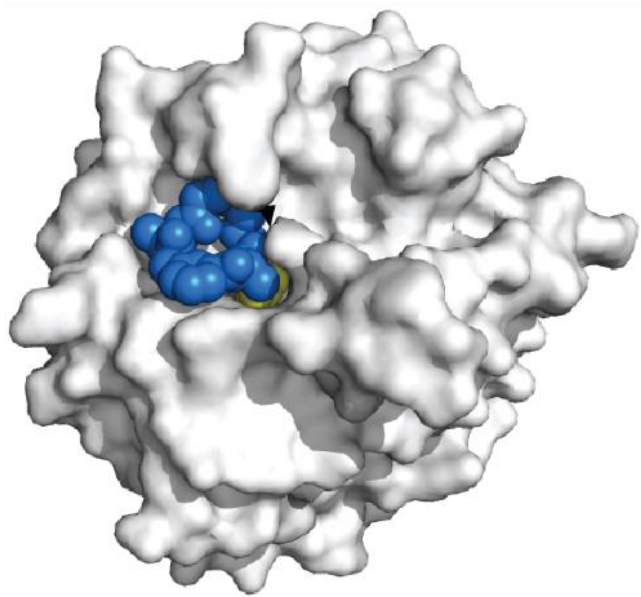
R = K_i (nM)

M1  44 ± 1

M2  165 ± 38

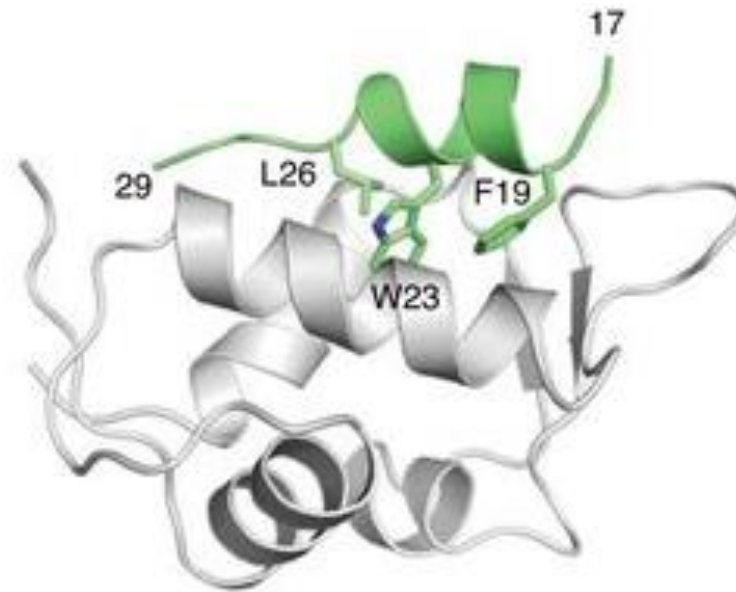
M3  125 ± 8

X-ray structure



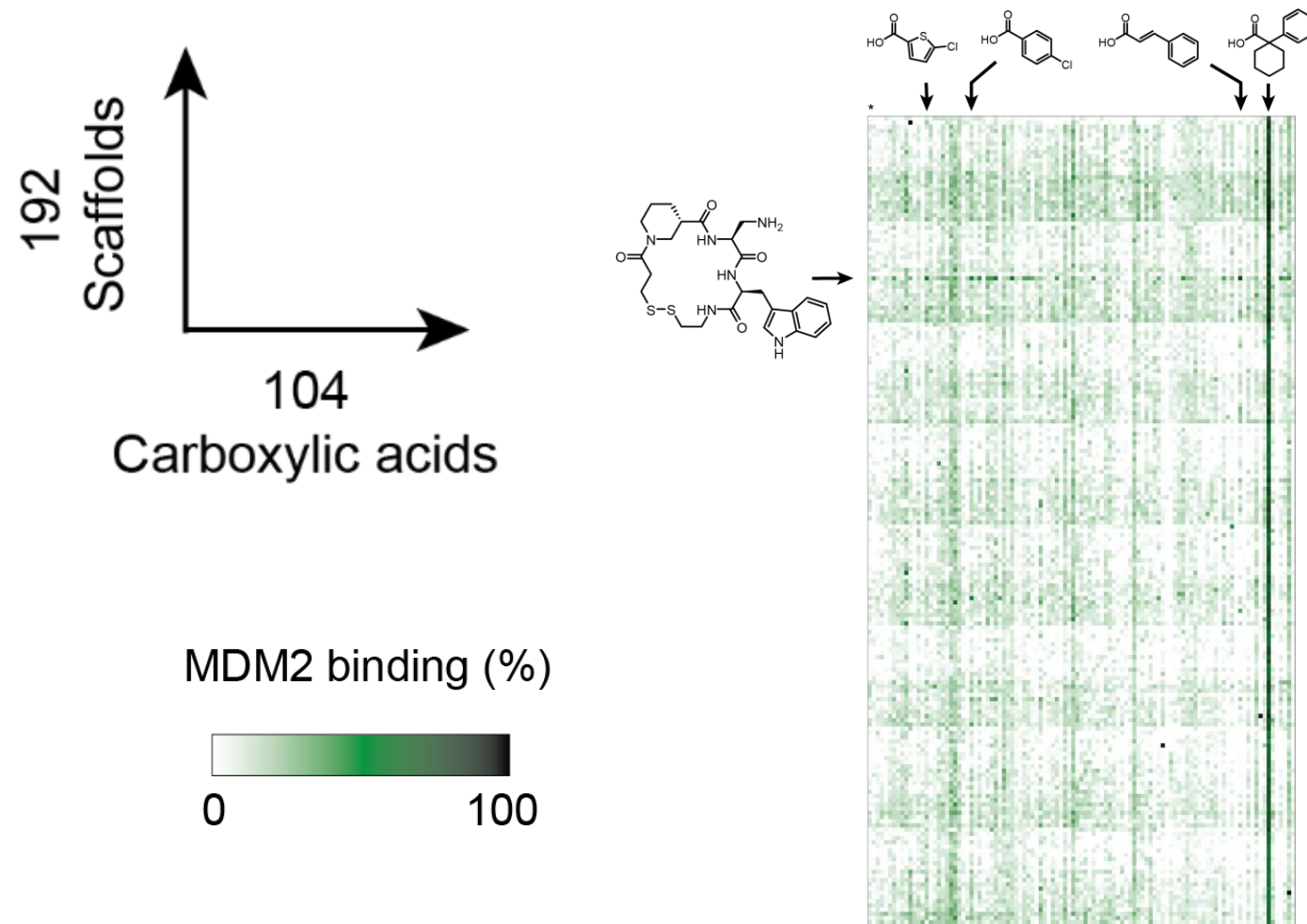
Structure solved by
Prof. Alessandro Angelini (Ca' Foscari University of Venice) and Prof. Laura Cendron (University of Padova)

Inhibition of protein-protein interactions (PPIs)

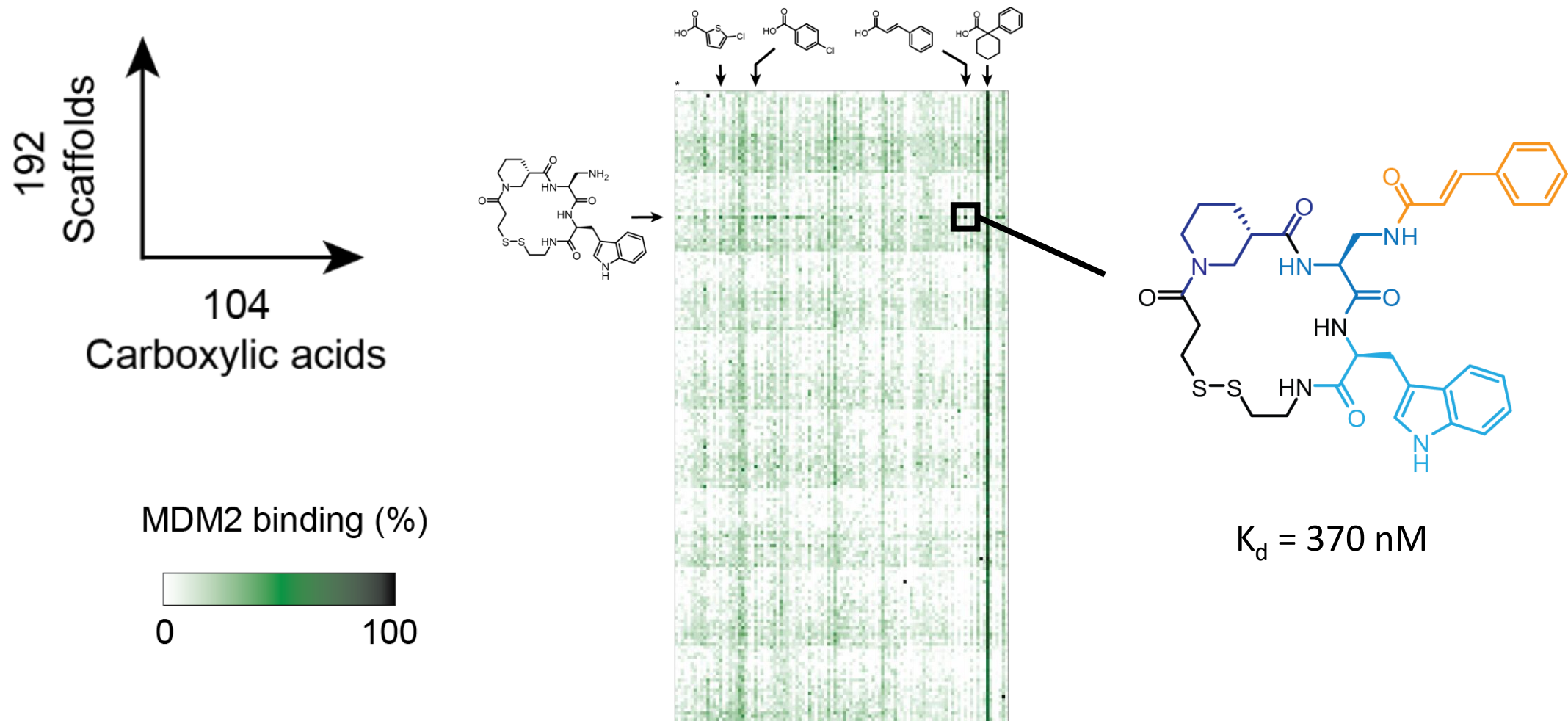


p53-MDM2

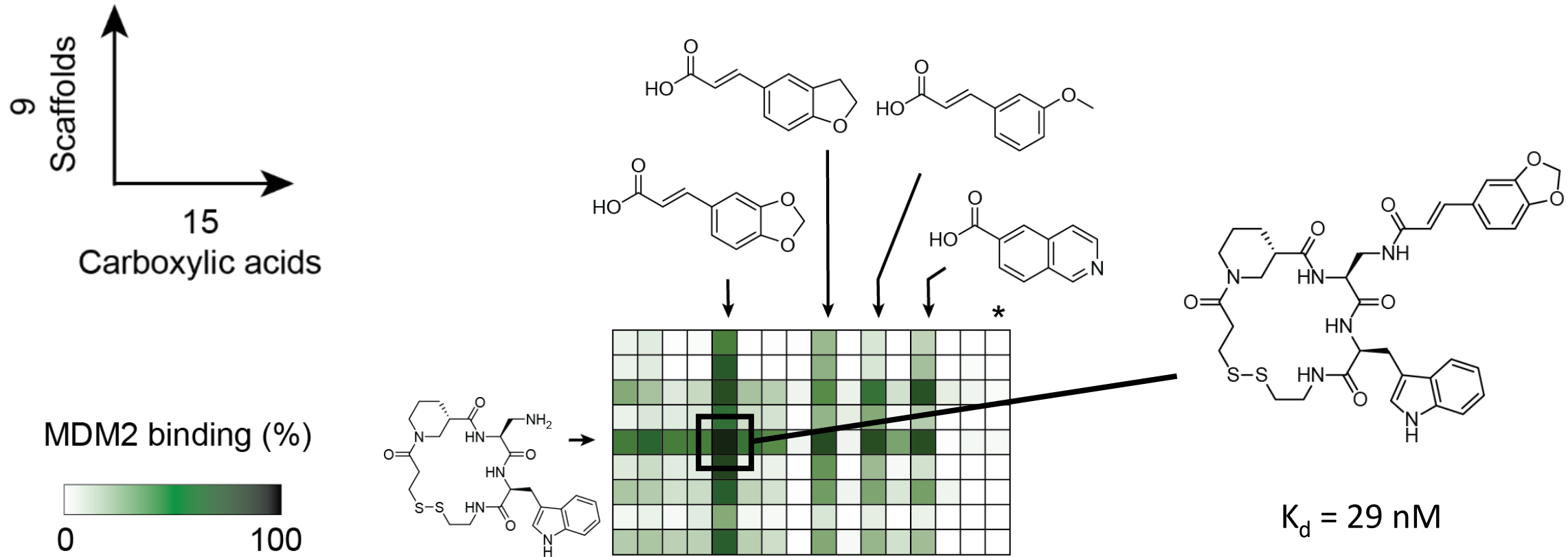
Synthesis of a 19,968 macrocycles tailored for MDM2/p53



Synthesis of a 19,968 macrocycles tailored for MDM2/p53



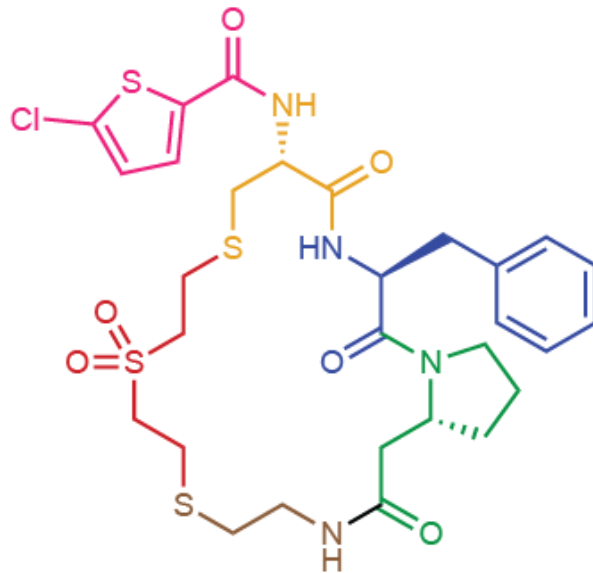
Rapid improvement of binding affinity



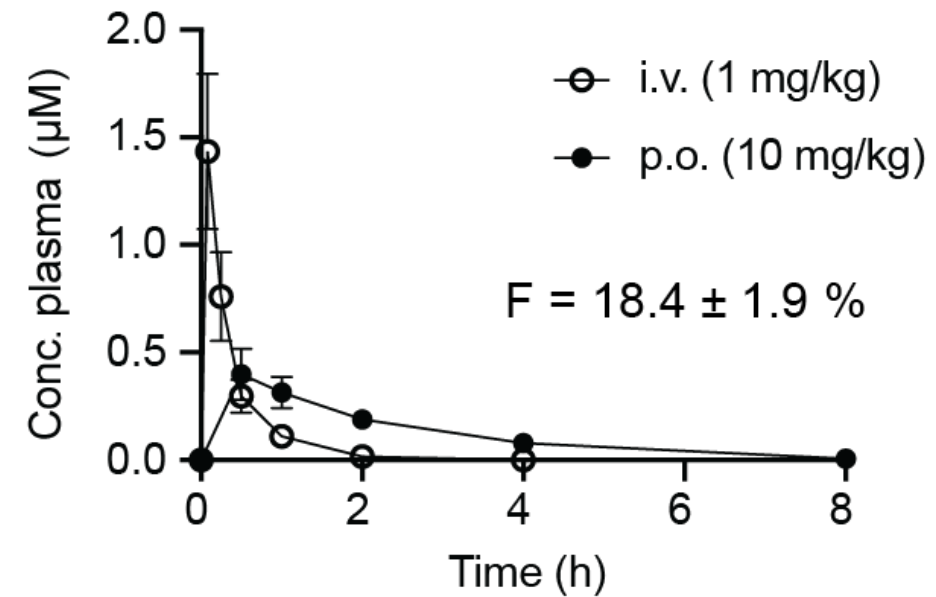
Habeshian, S., Merz, M.L., Mothukuri, G.K., Schüttel, M., Bognar, Z., Diaz-Perlas, C., Vesin, J., Bortoli Chapalay, J., Turcatti, G., Cendron, L., Angelini, A. and Heinis, C., *Nature Communications*, 2022

Are they membrane permeable / orally available?

Stable linkers

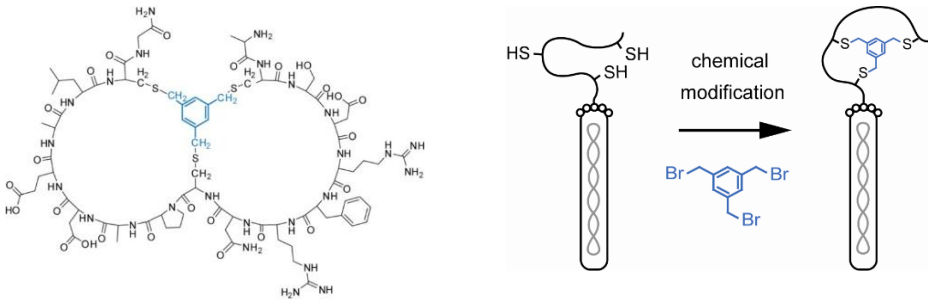


Oral availability (rat)



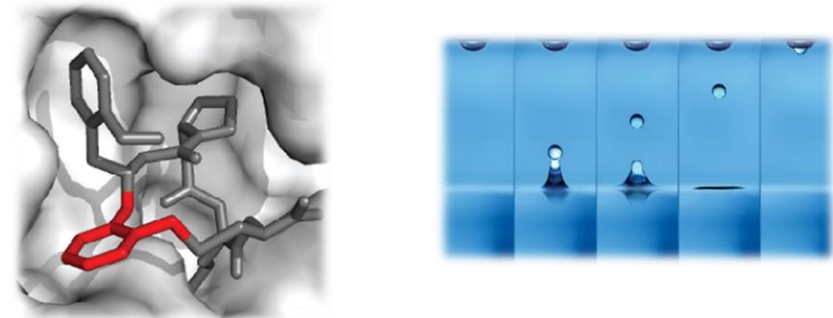
Summary

Bicyclic peptides & phage display



- High affinity, specificity
- Fast development
- Works with many targets
- Good for extracellular targets
- Limits: needs injection, short half-life

Macrocycles & screening



- High affinity, specificity
- Also intracellular targets
- Oral application
- Limits: harder to develop for challenging targets