

Identification of Covalent JNK Inhibitor

2024/11/20

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Axcelead Drug Discovery Partners, Inc.

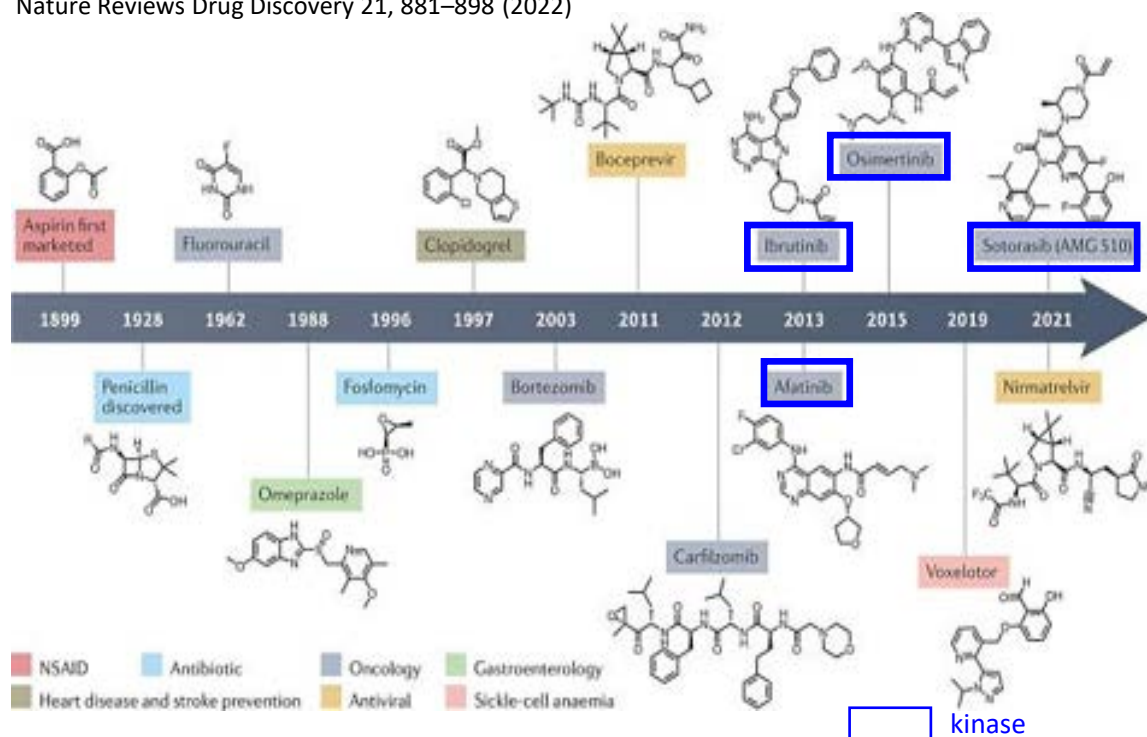
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Background

- ◆ Rising of covalent inhibitor: targeted EGFR, BTK, JAK etc. covalent kinase inhibitor
- ◆ Cysteinome: Ligandable cysteine, relationship with disease
- ◆ Screening trend: PROTAC > Degradar > **Covalent inh.** > Reversible inh. > RNA binder

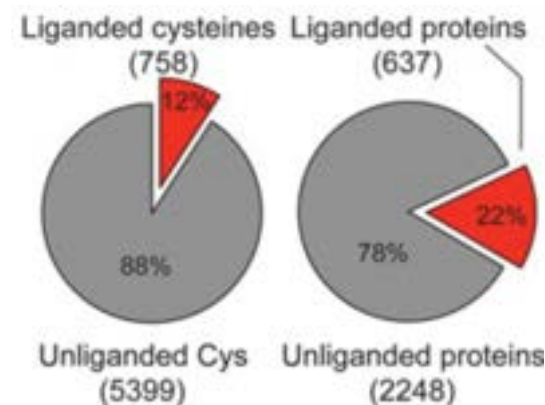
Covalent inhibitor history

Nature Reviews Drug Discovery 21, 881–898 (2022)



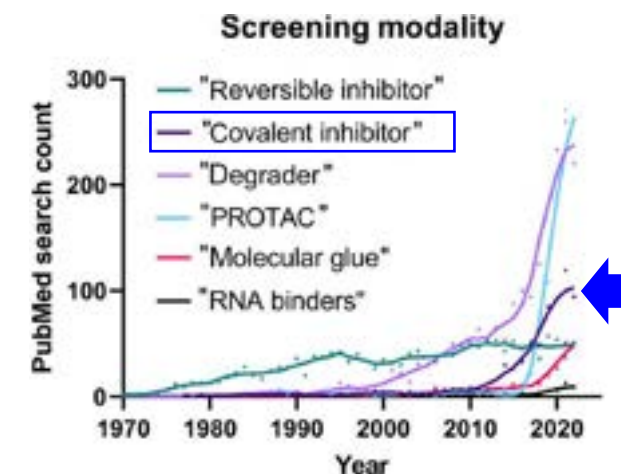
Ligandable cysteine

Nature. 2016, 534, 570–574.



HTS trend

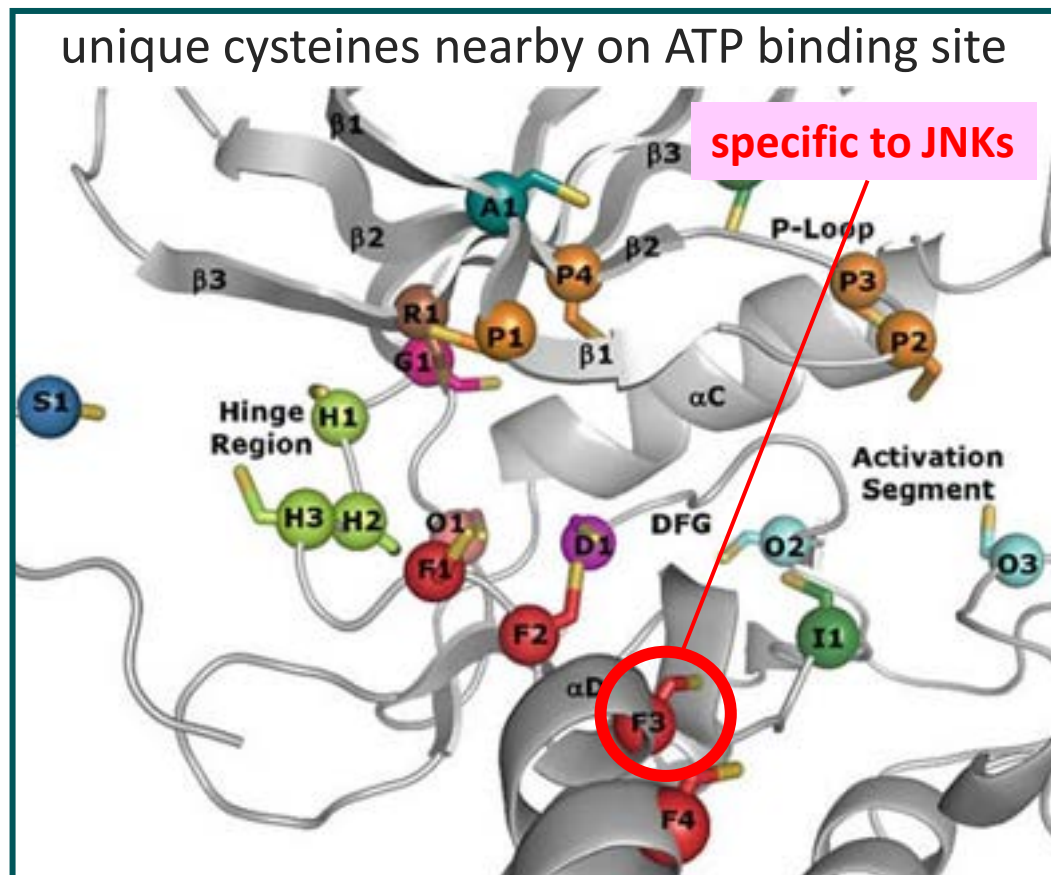
Drug Discovery Today, 28, 2023



Research on covalent drug is gaining attention.

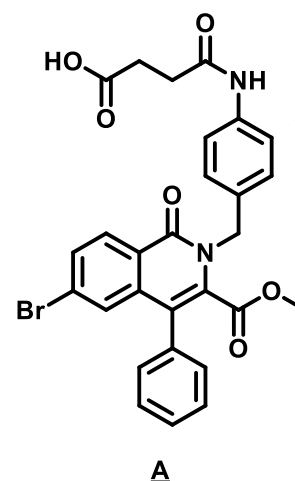
Concept of Research

- ✓ c-Jun N-terminal kinases (JNKs) are key signal transducers in the MAPK signaling pathways, involved in inflammation, cancer, and neurodegeneration.



Gehring, M., Covalent kinase inhibitors: an overview. In: Topics in Medicinal Chemistry. Springer, Berlin/Heidelberg, Germany, pp. 1–52, 2020.

previously reported
JNK inhibitor*



JNK inhibition $IC_{50} = 9.2 \text{ nM}$

potential warhead site search

compound design from SAR

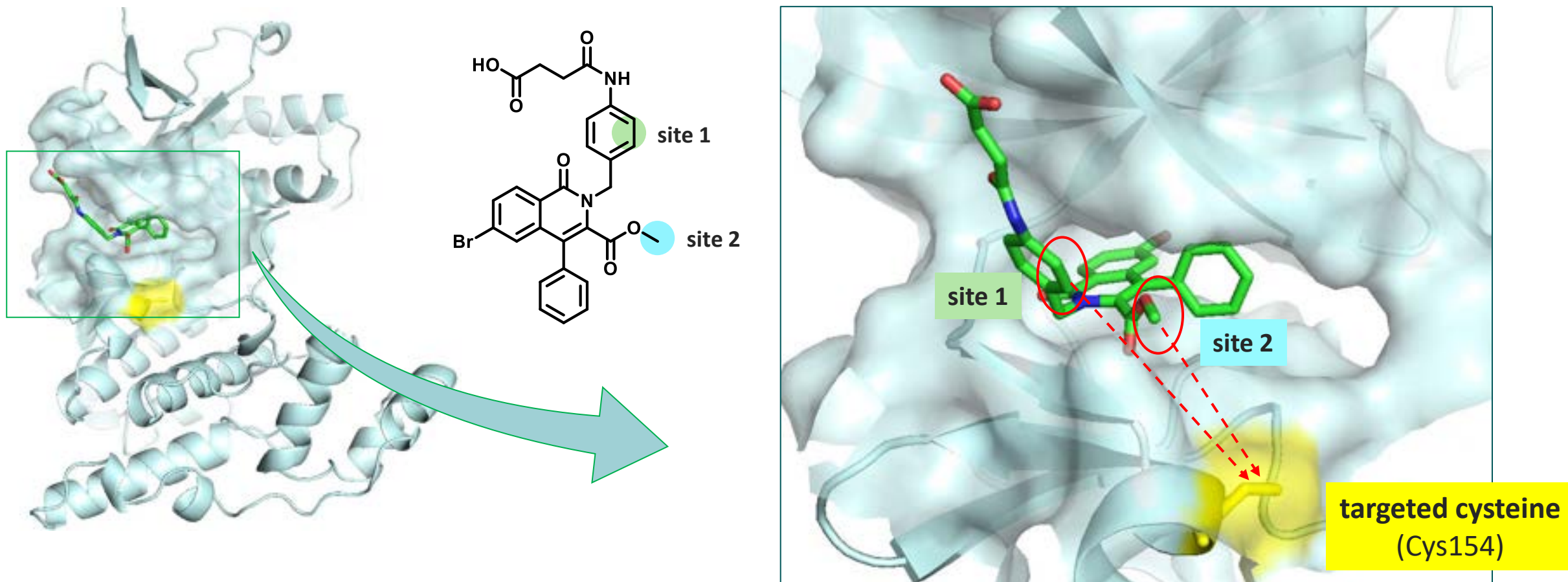
prioritization by docking score

Novel covalent JNK inhibitor

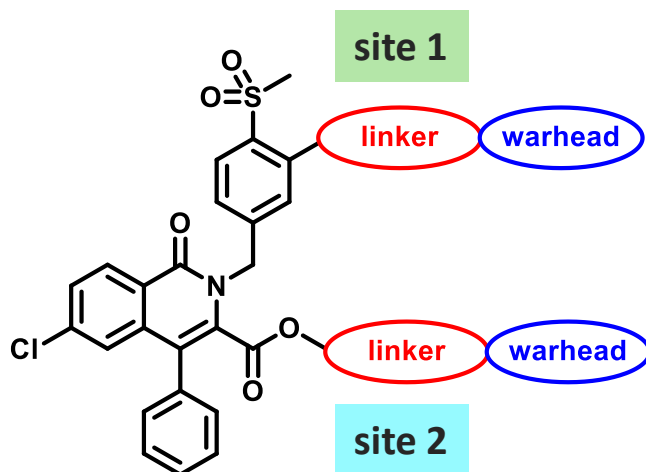
- inhibition activity ↑
- selectivity ↑

Potential warhead site search

Docking pose of compound A to JNK3 protein (PDBID : 2ZDT)

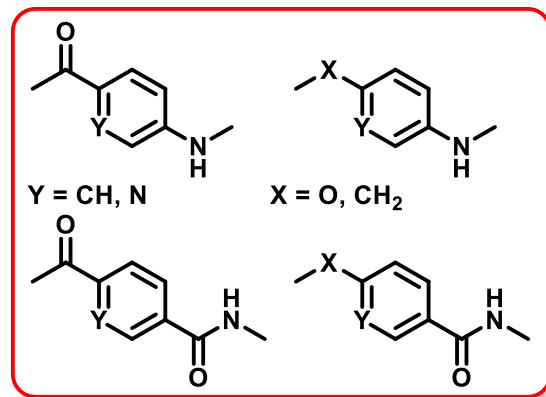


Site 1 and 2 were preferred to introduce warheads.



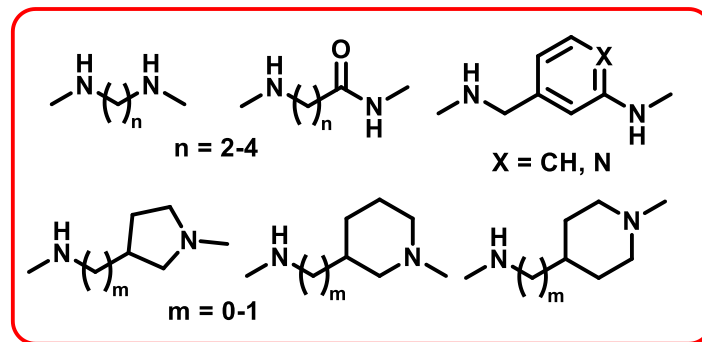
site 1

linkers



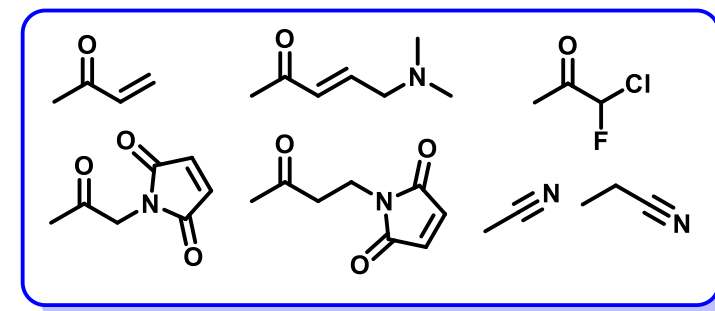
site 2

linkers



previously reported^{*)}
warheads

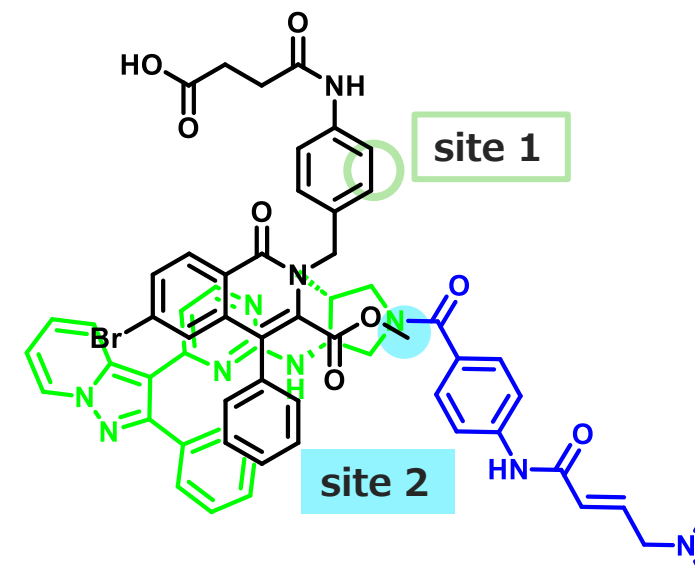
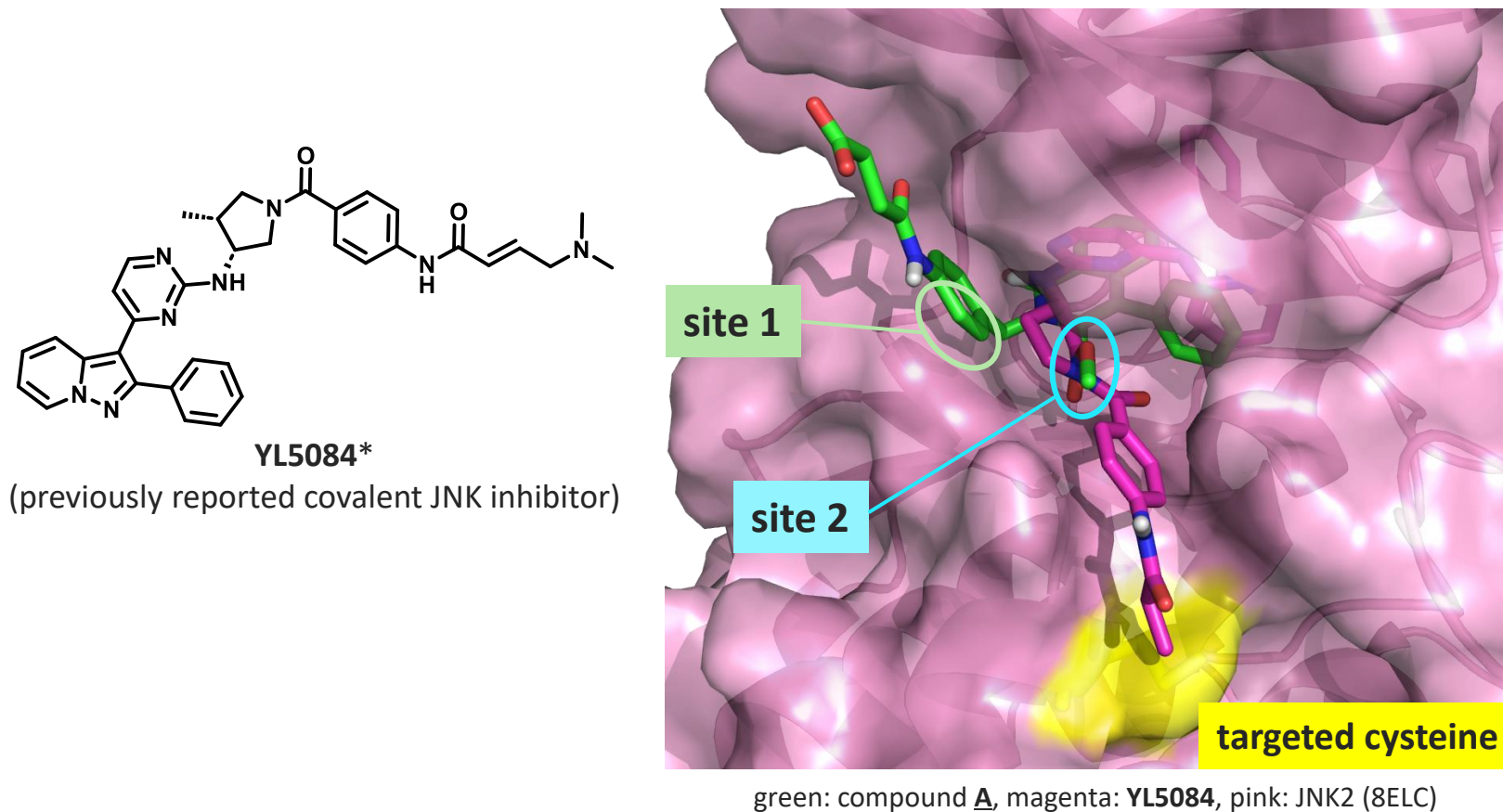
warheads



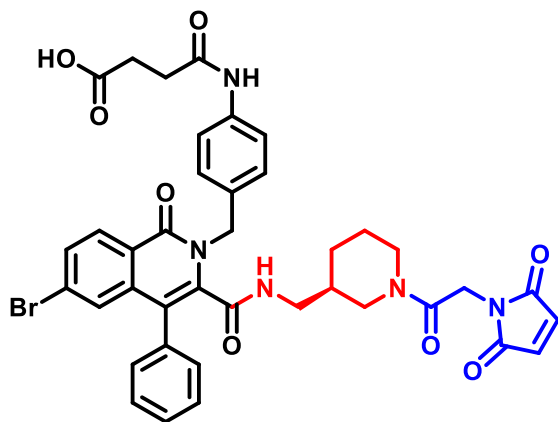
Narrow down the linker candidates
(size, length, stretching direction)
from SAR information.

Prioritization of compounds to be synthesized

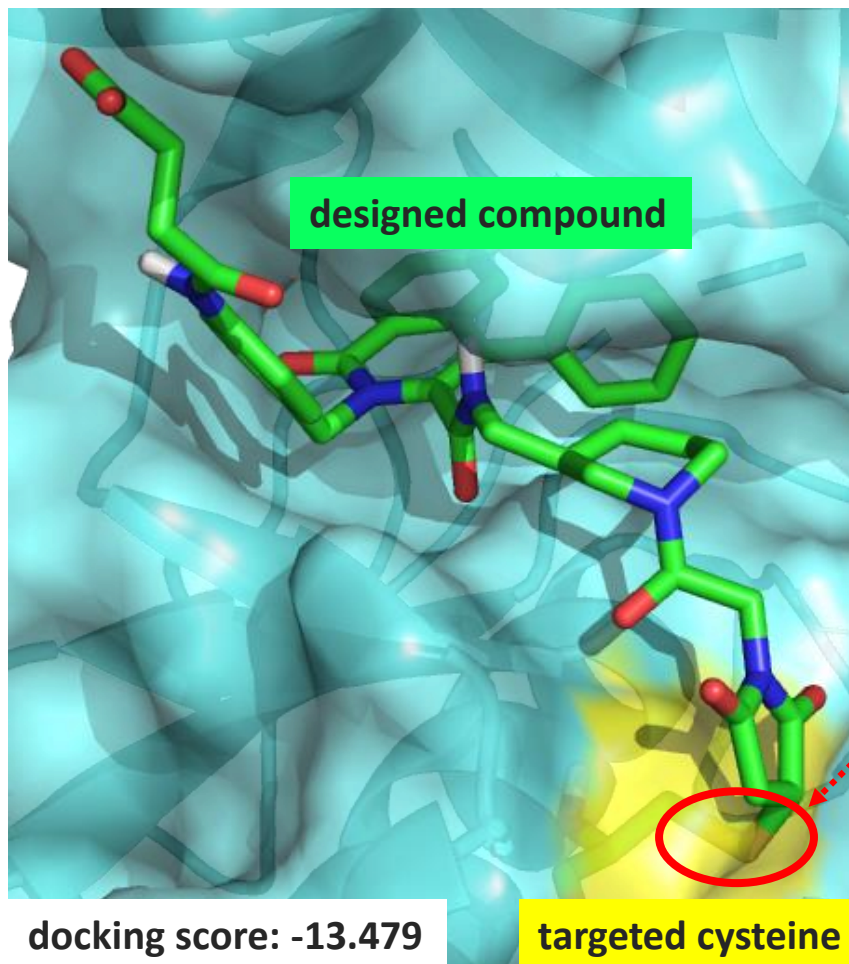
Fig. superposition of **A** and **YL5084**



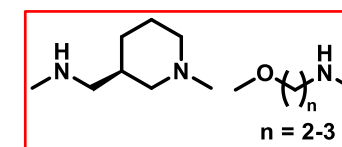
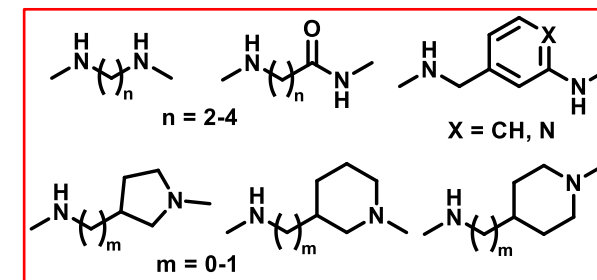
Based on previously reported JNK2 covalent inhibitor, the priority was raised for compounds that extend the linker from site 2.



designed compound

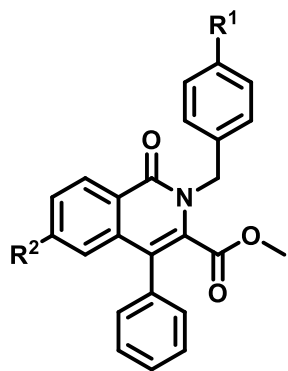


covalent bond
formation

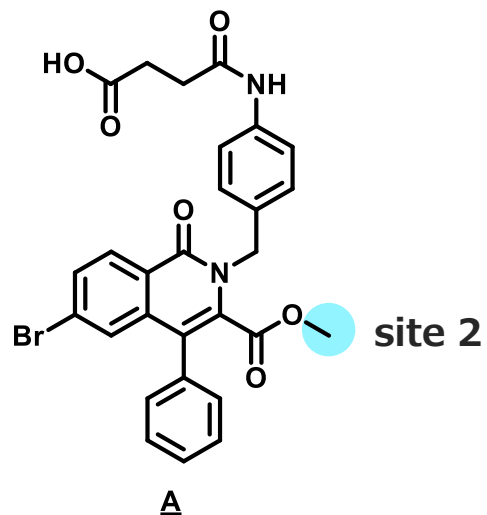


Linkers were prioritized based on **docking score** and **synthetic feasibility**.

Table. SAR information of compound A^{*)}

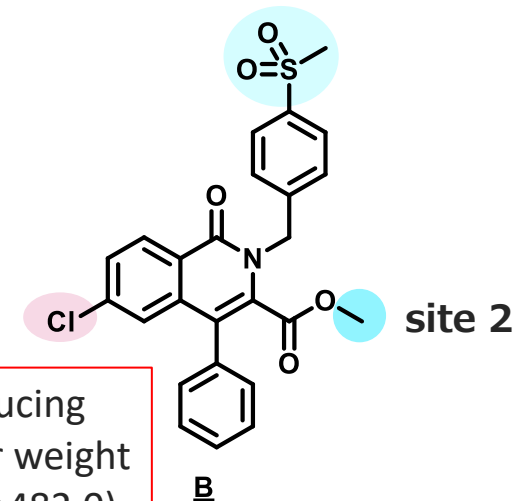


R ¹	R ²	JNK1 IC ₅₀ (nM)
NHCO(CH ₂) ₂ CO ₂ H	Br	9.2
SO ₂ Me	Br	30
SO ₂ Me	Cl	28



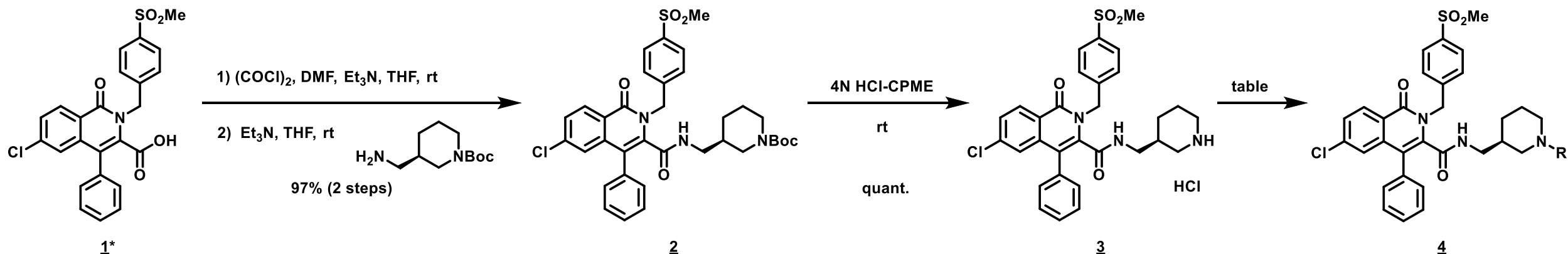
for avoiding β oxidation

for reducing
molecular weight
(563.4 \Rightarrow 482.0)



Functional groups which are not related introducing warheads were modified based on SAR information.

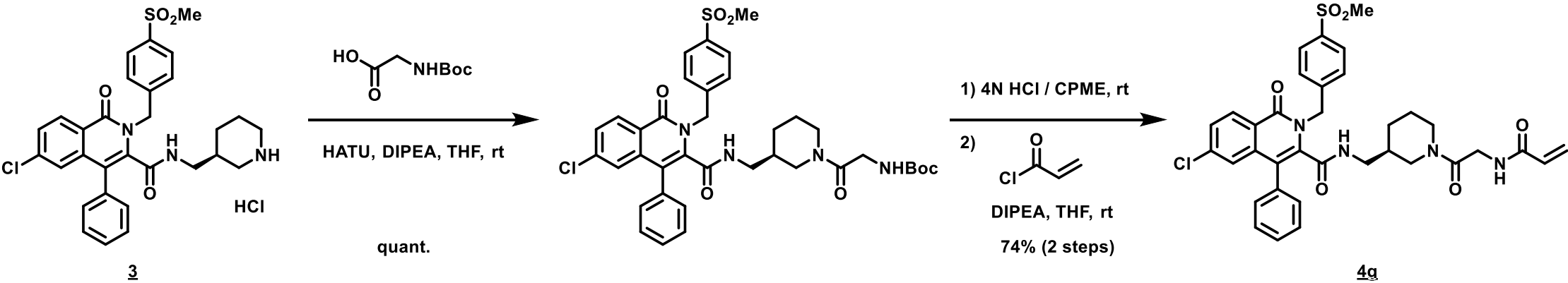
Synthesis of designed compounds



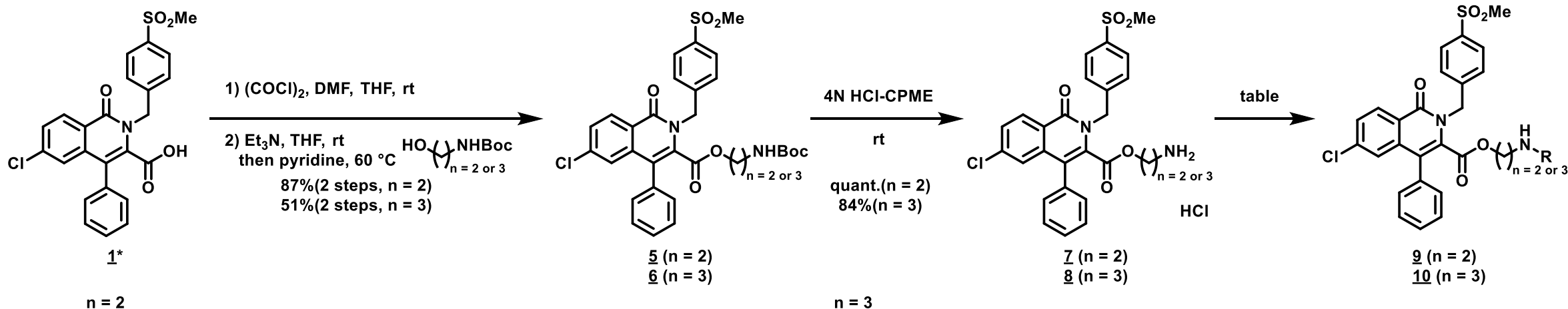
compound	R	Reaction conditions	Yield
4a		DIPEA, THF	77%
4b		DIPEA, THF	72%
4c		K ₂ CO ₃ , MeCN	72%

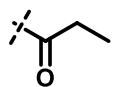
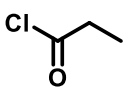
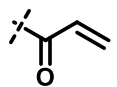
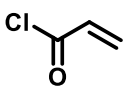
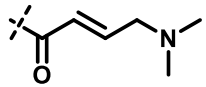
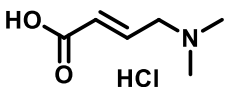
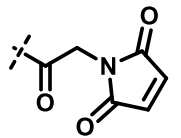
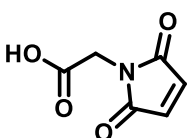
compound	R	Reaction conditions	Yield
4d		HATU, DIPEA, THF	76%
4e		HATU, DIPEA, THF	82%
4f		HATU, DIPEA, THF	93%

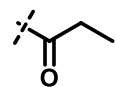
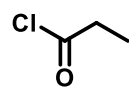
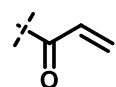
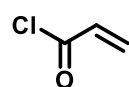
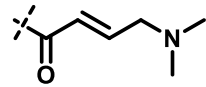
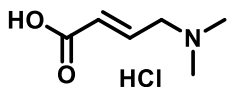
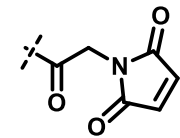
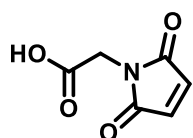
Synthesis of designed compounds



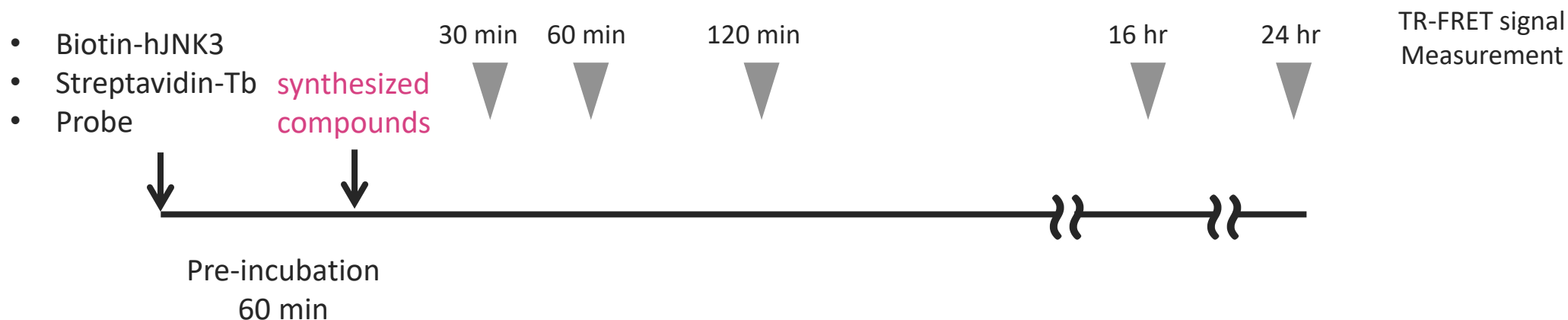
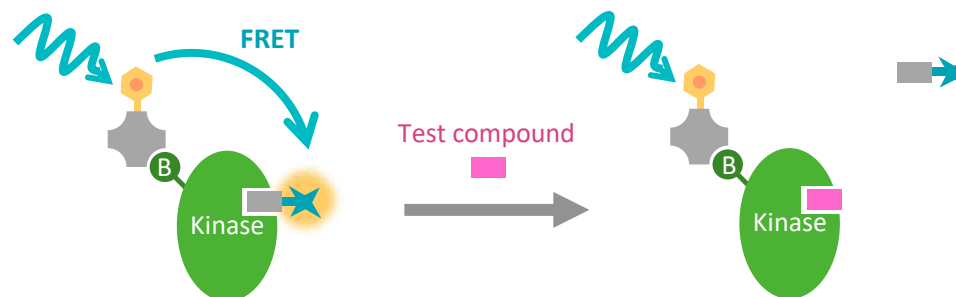
Synthesis of designed compounds



compound	R	Reaction conditions	Yield
9a		 DIPEA, THF	84%
9b		 DIPEA, THF	81%
9c		 HCl HATU, DIPEA, THF	61%
9d		 HATU, DIPEA, THF	99%

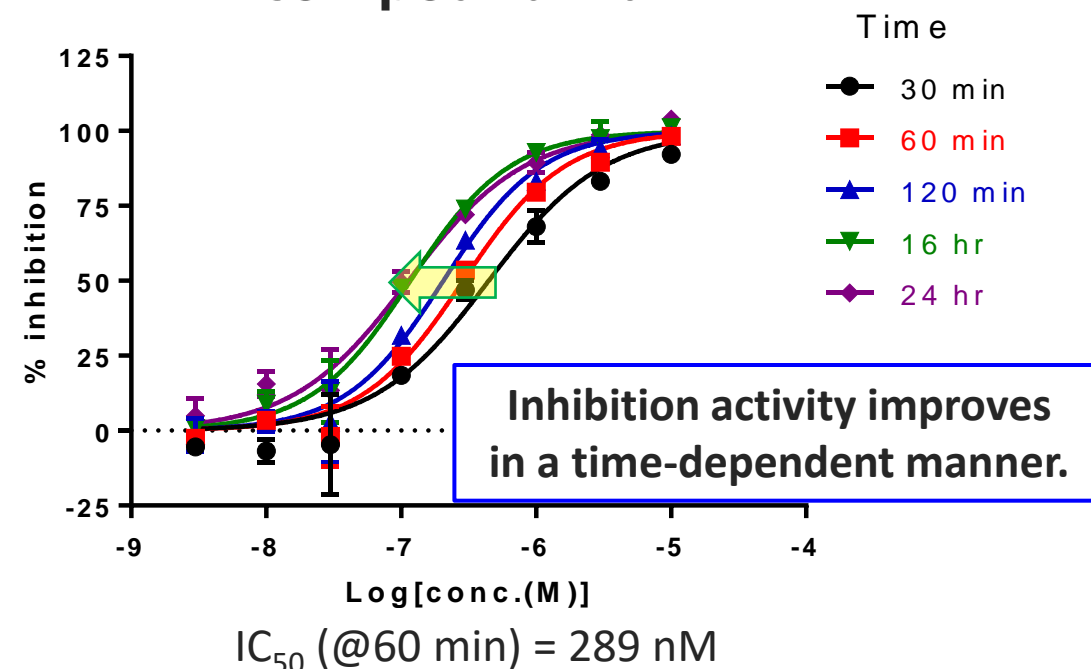
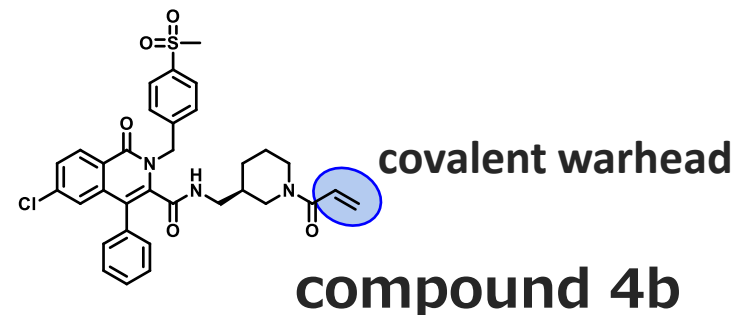
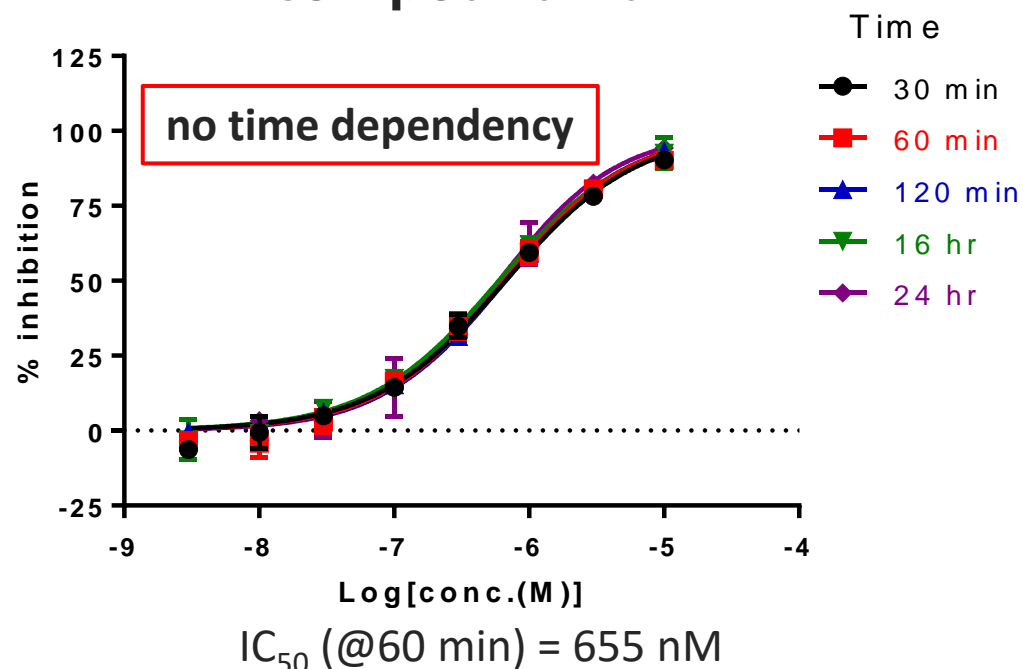
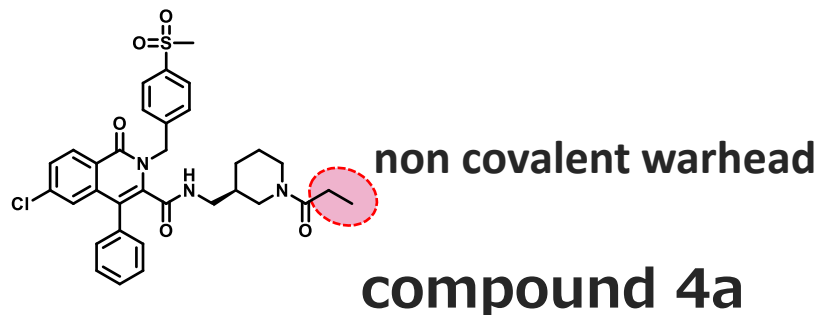
compound	R	Reaction conditions	Yield
10a		 DIPEA, THF	87%
10b		 DIPEA, THF	82%
10c		 HCl HATU, DIPEA, THF	92%
10d		 HATU, DIPEA, THF	58%

Probe displacement TR-FRET assay



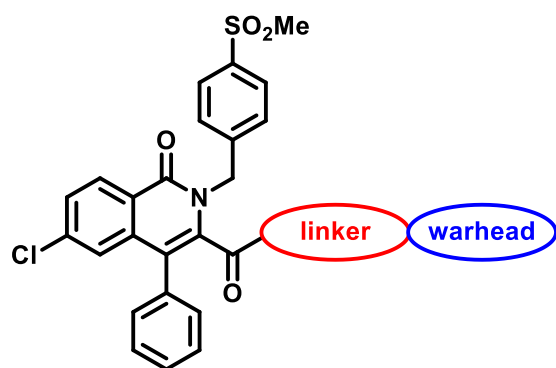
Time dependency of the compound is determined by the increase in inhibition during the incubation time.

Assessing Time Dependency in Binding Assay : Representative Results



- ✓ Retained inhibition of JNK3.
- ✓ Compound **4b** inhibited JNK3 in a time-dependent manner.

Assessing Time Dependency in Binding Assay : Overall Results



linker	warhead	compound	time dependency	IC ₅₀ (nM) @ 60 min.
–	–	<u>B</u>	–	28
<hr/>				
		4a	–	655
		4b	○	289
		4c	–	553
		4d	–	347
		4e	–	408
		4f	○	0.34
		4g	○	116

linker	warhead	compound	time dependency	IC ₅₀ (nM) @ 60 min.
		9a	–	196
		9b	–	251
		9c	–	177
		9d	○	0.23
<hr/>				
		10a	–	102
		10b	–	102
		10c	○	72
		10d	○	0.14

- ✓ 6 compounds (4b, 4f, 4g, 9d, 10c, 10d) inhibited JNK3 in a time-dependent manner.
- ✓ Inhibition activity of 3 compounds (4f, 9d, 10d) were improved from compound A.

Whole Protein MS of JNK3 + synthesized compounds

Condition of LC-MS

UPLC: Waters Acquity UPLC I-class

Column: Acquity BEH C4, 300 Å, 1.7µm, 2.1 x 50mmL

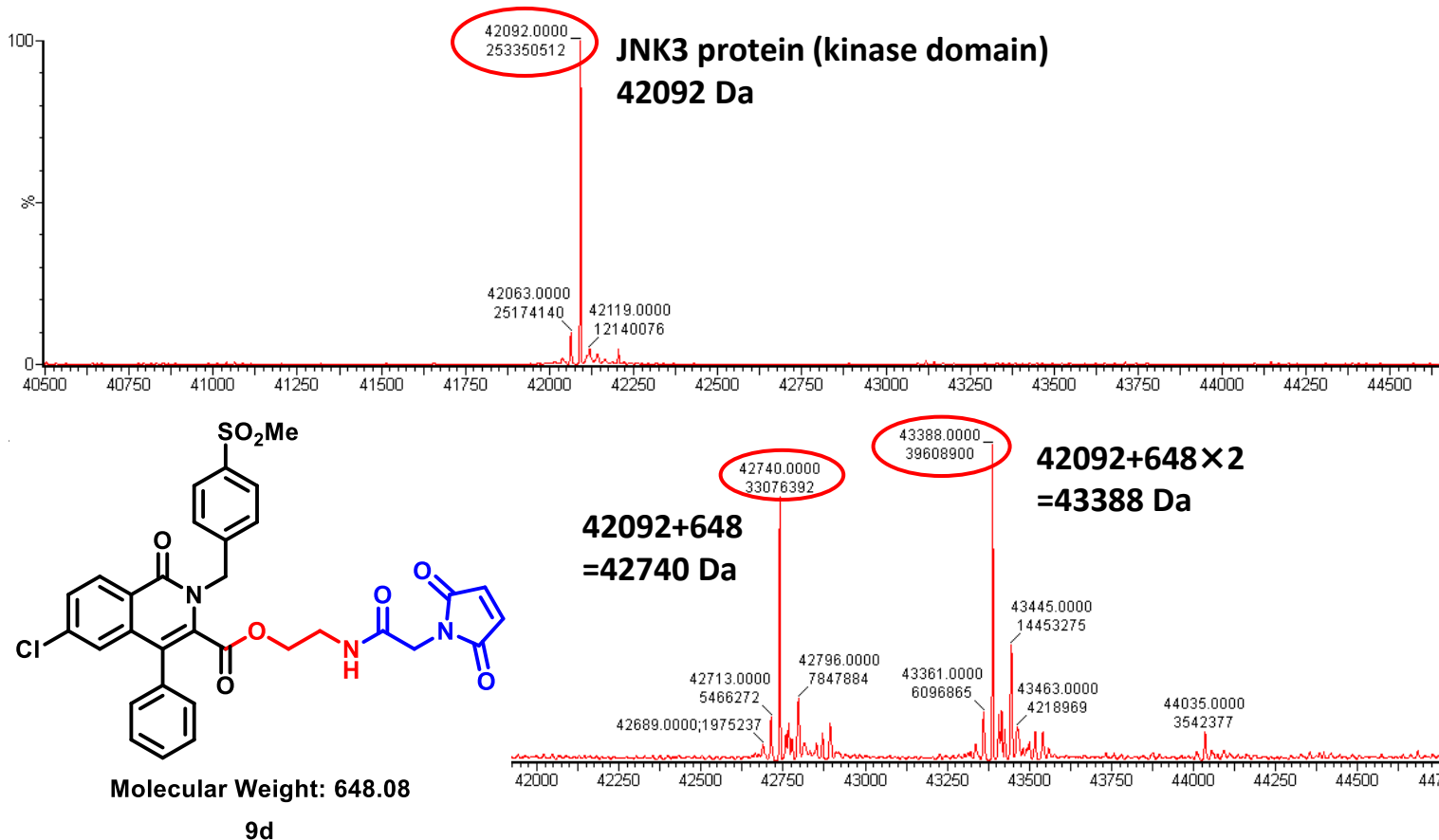
Mobile phase A: TFA/ water (0.25/1000)

Mobile phase B: TFA / MeCN (0.25/1000)

MS: Waters Xevo G2-S ToF

Electrospray ionization, Positive ion mode

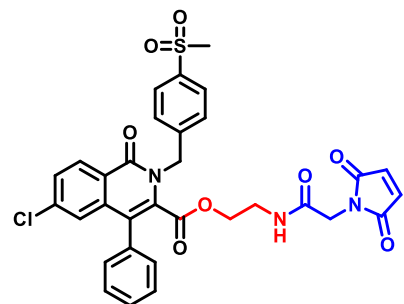
Data analysis: MassLynx, MaxEnt I



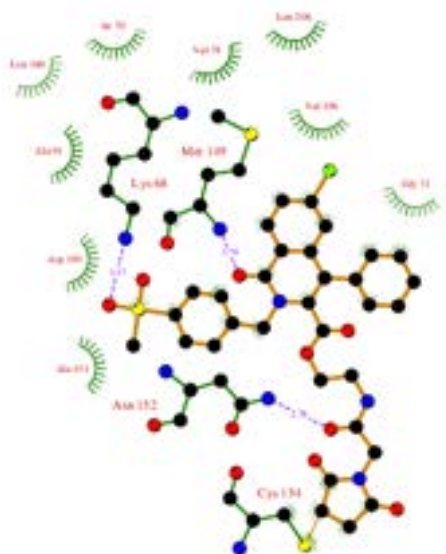
comp. ID	Number of adduct	covalent binding
4b	1	single
4f	1,2,3,4	multi
4g	1	single
9d	1,2	multi
10d	1,2	multi

Covalent adduct of **9d** with JNK3 protein was confirmed by protein-MS.

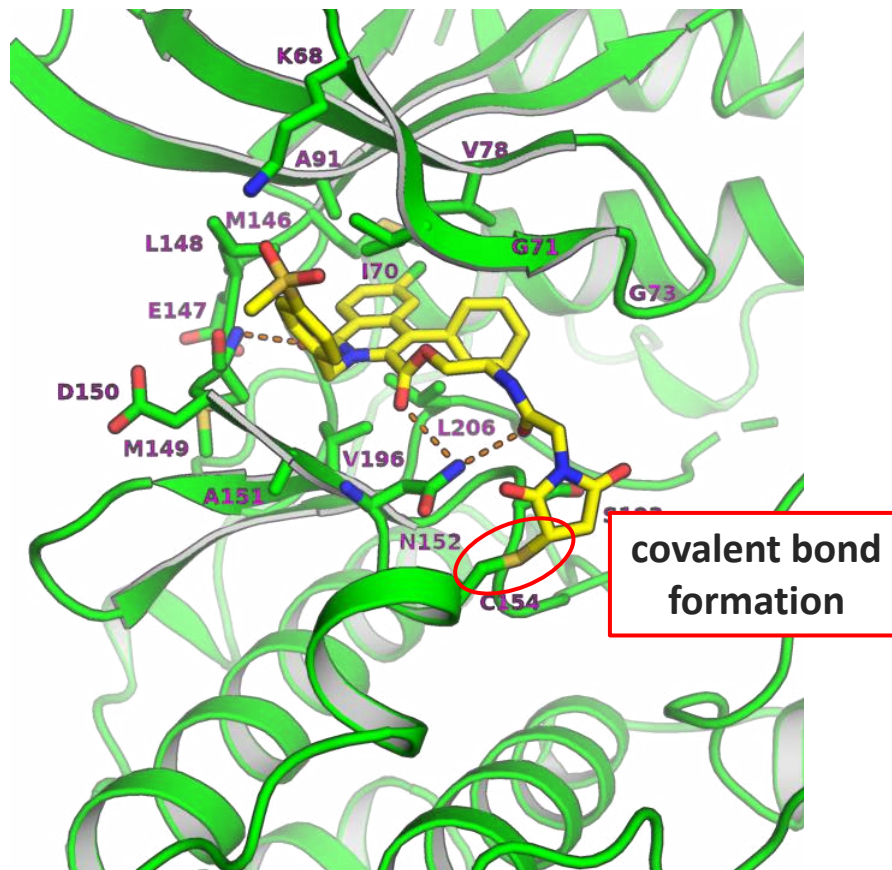
Several compounds confirmed covalent adducts.



9d



JNK3/comp. 9d (1.7 Å)

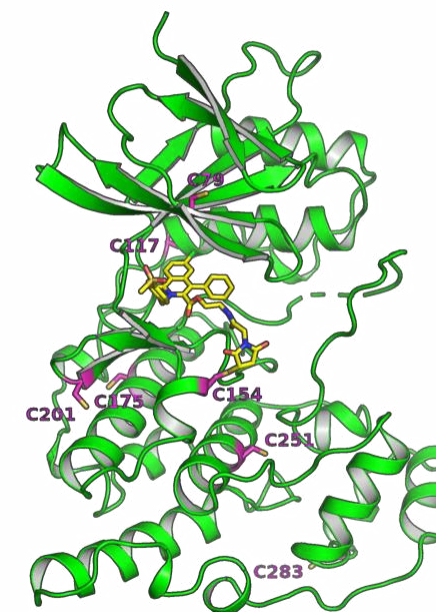


Only co-crystals to which **one** compound 9d is bound were obtained.

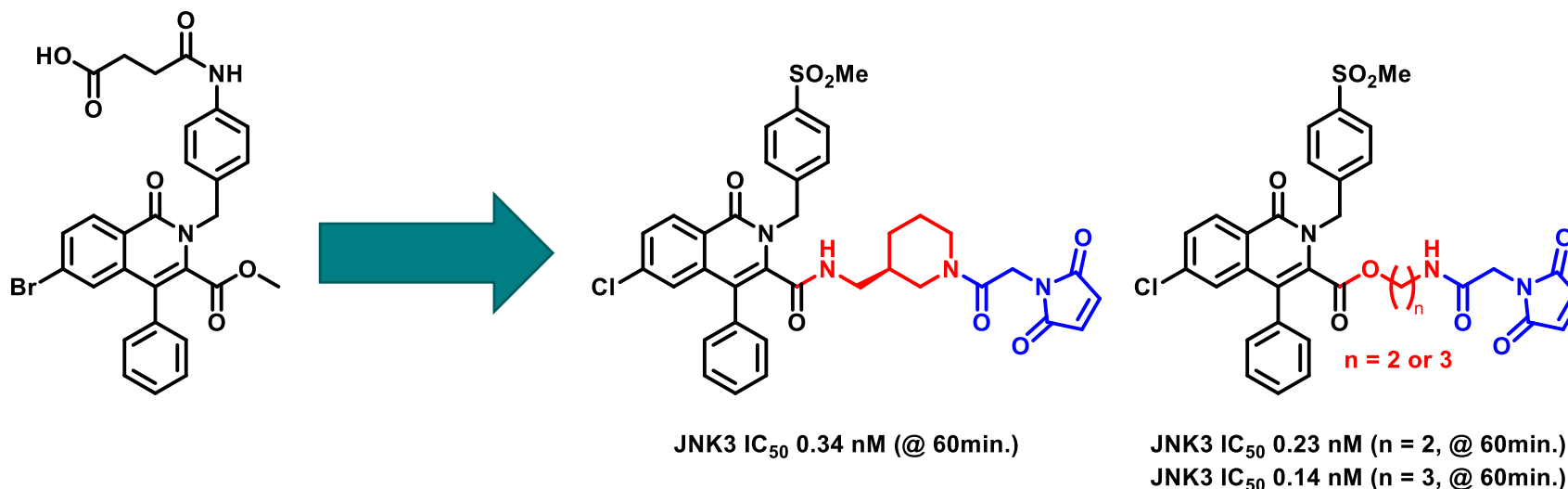
Covalent binding to the expected cysteine residue (C154) was confirmed, but the additional modification sites remain unidentified.

Mass spectroscopy indicated **9d** is bound to a few sites.

The potential covalent modification sites



The kinase domain contains **7** cysteines.



- ✓ Designed compounds **retained inhibition of JNK3.**
- ✓ Several compounds inhibited JNK3 **in a time-dependent manner.**
- ✓ **Adducts of designed compound with JNK3 protein** was confirmed by Protein-MS.
- ✓ **Covalent binding to the expected cysteine** was confirmed by X-ray crystallography.

Tasks to do in future

- Selectivity assay (JNK3/1 and 2)
- Site identification
- *In vivo* assay

Thank you !

Access to Accelerate Lead for Drug Discovery

Axcelead Drug Discovery Partners, Inc.

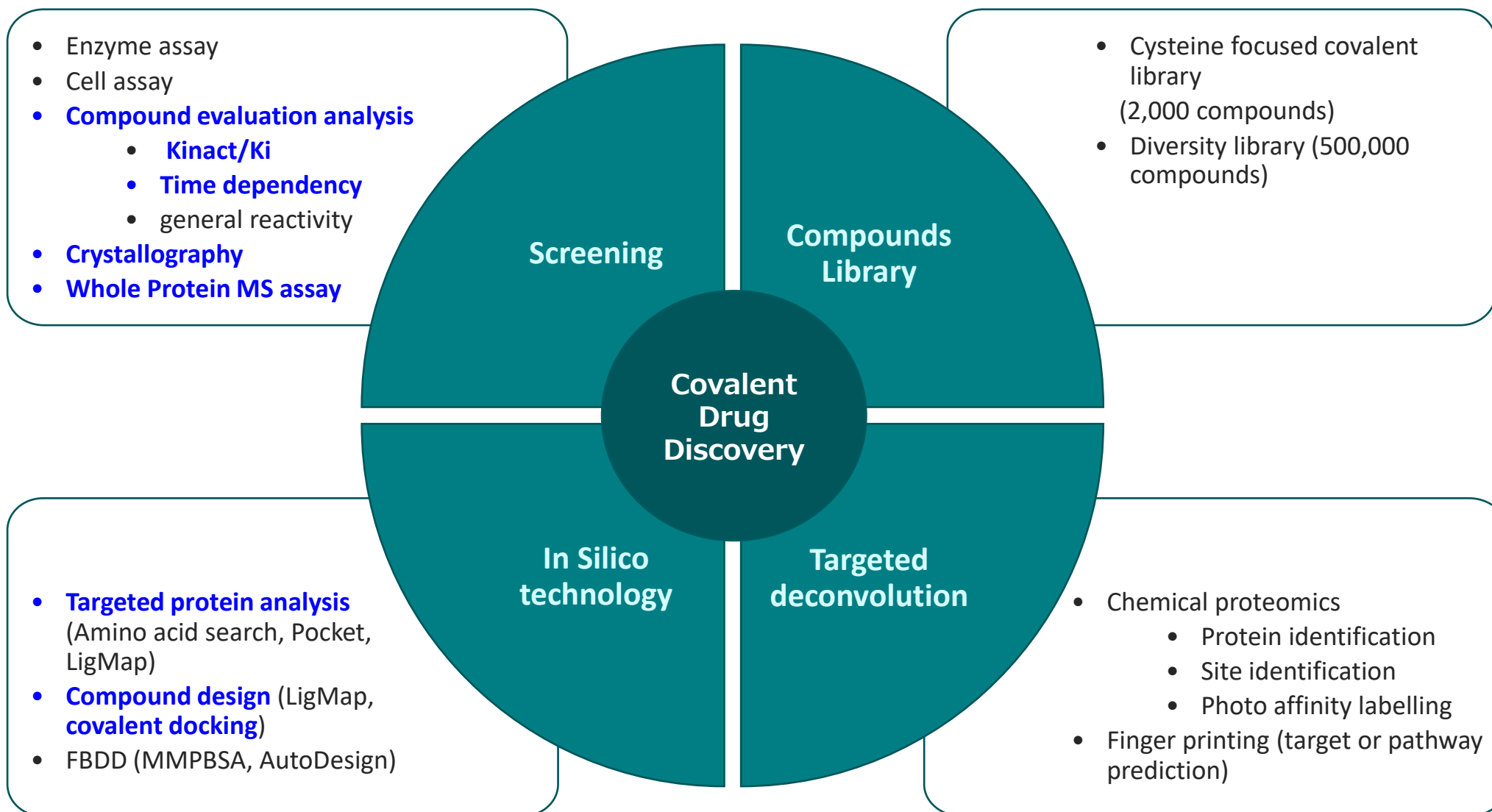
<https://axcelead-us.com/>

<https://www.axcelead.com/>

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appendix

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Reaction kinetic analysis of covalent inhibitor-1

2-step IRREV



Inactivation Potency

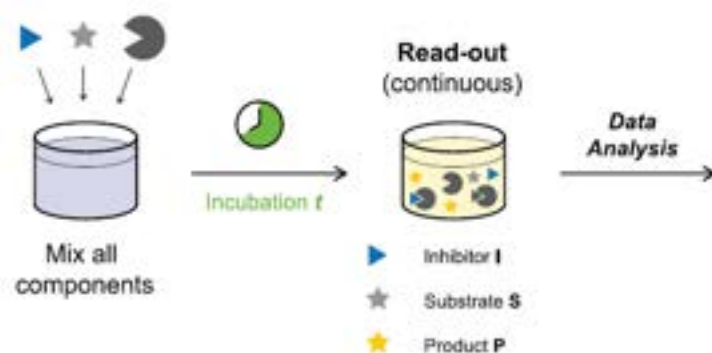
$$k_{\text{inact}}/K_I = \frac{k_3 k_{\text{inact}}}{k_4 + k_{\text{inact}}}$$

Inactivation Constant

$$K_I = \frac{[E][I]}{[EI]} = K_1 + \frac{k_{\text{inact}}}{k_3}$$

Covalent Inhibitor Binding Kinetics

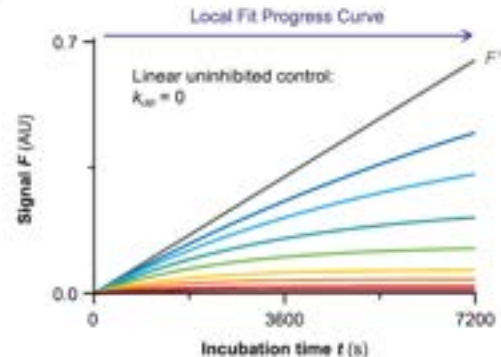
$$F_t = v_s t + \frac{v_i - v_s}{k_{\text{obs}}} [1 - e^{-k_{\text{obs}} t}] + F_0$$



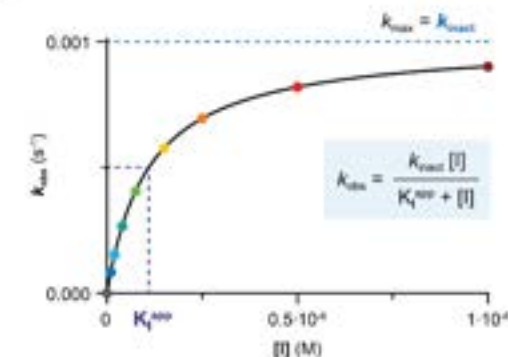
A



B Time-dependent Enzyme Activity Inhibition



C Correct K_I for substrate competition during incubation

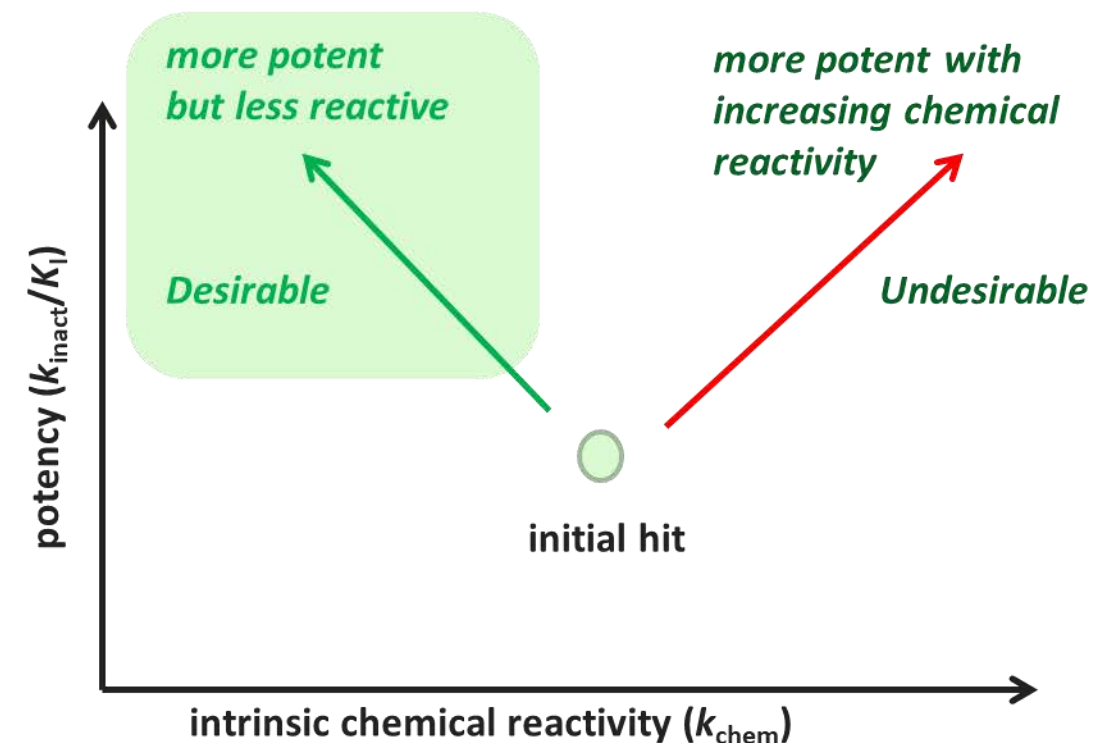
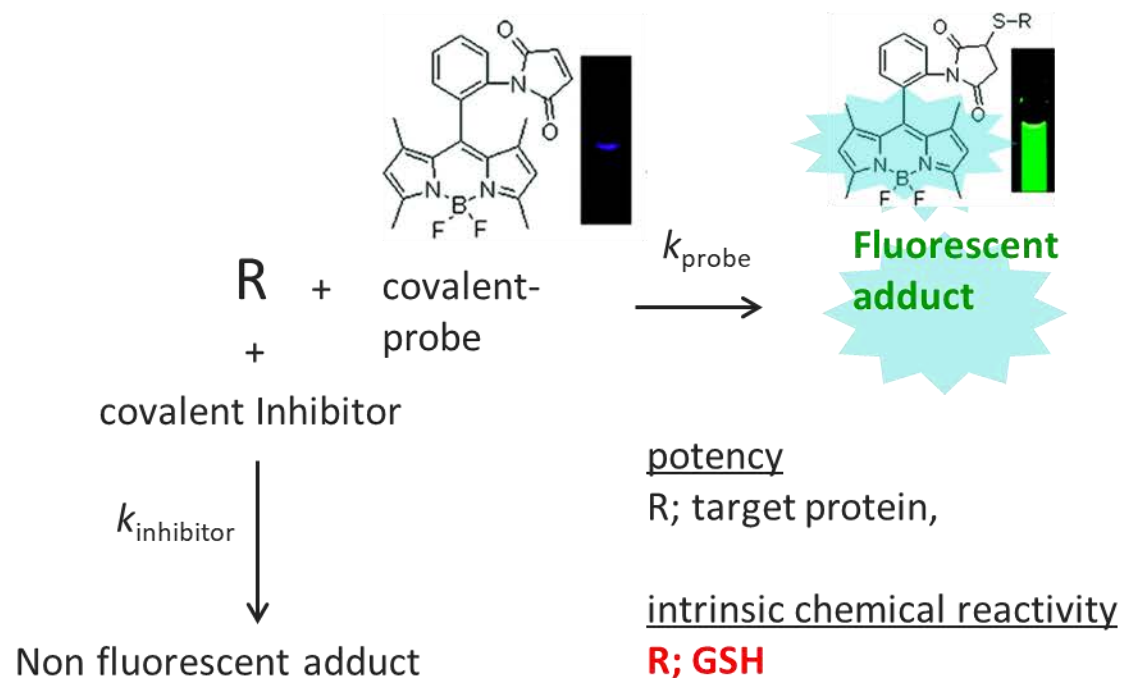


Progress curve analysis for two-step irreversible covalent inhibition.

Curr Protoc. 2022 Jun;2(6)

Inhibitor concentration-dependent K_{obs} reaches K_{inact} at saturating inhibitor concentration ($K_{\text{max}} = K_{\text{inact}}$).
Half-maximum $K_{\text{obs}} = \frac{1}{2}K_{\text{inact}}$ is reached when inhibitor concentration equals the apparent inactivation constant K_I^{app} .

簡易的一般反応性試験 (蛍光プローブ法による反応性試験)



SLAS Discov. 2017 Apr 1:2472555217704654. doi: 10.1177/2472555217704654.
 High-Throughput Quantitative Intrinsic Thiol Reactivity Evaluation Using a Fluorescence-Based Competitive Endpoint Assay.

- 一般反応性 K_{chem} が低く、標的への親和性 K_{inact}/K_i の高い化合物 が好ましい。