



Inhibiting CK2 α from outside the active site

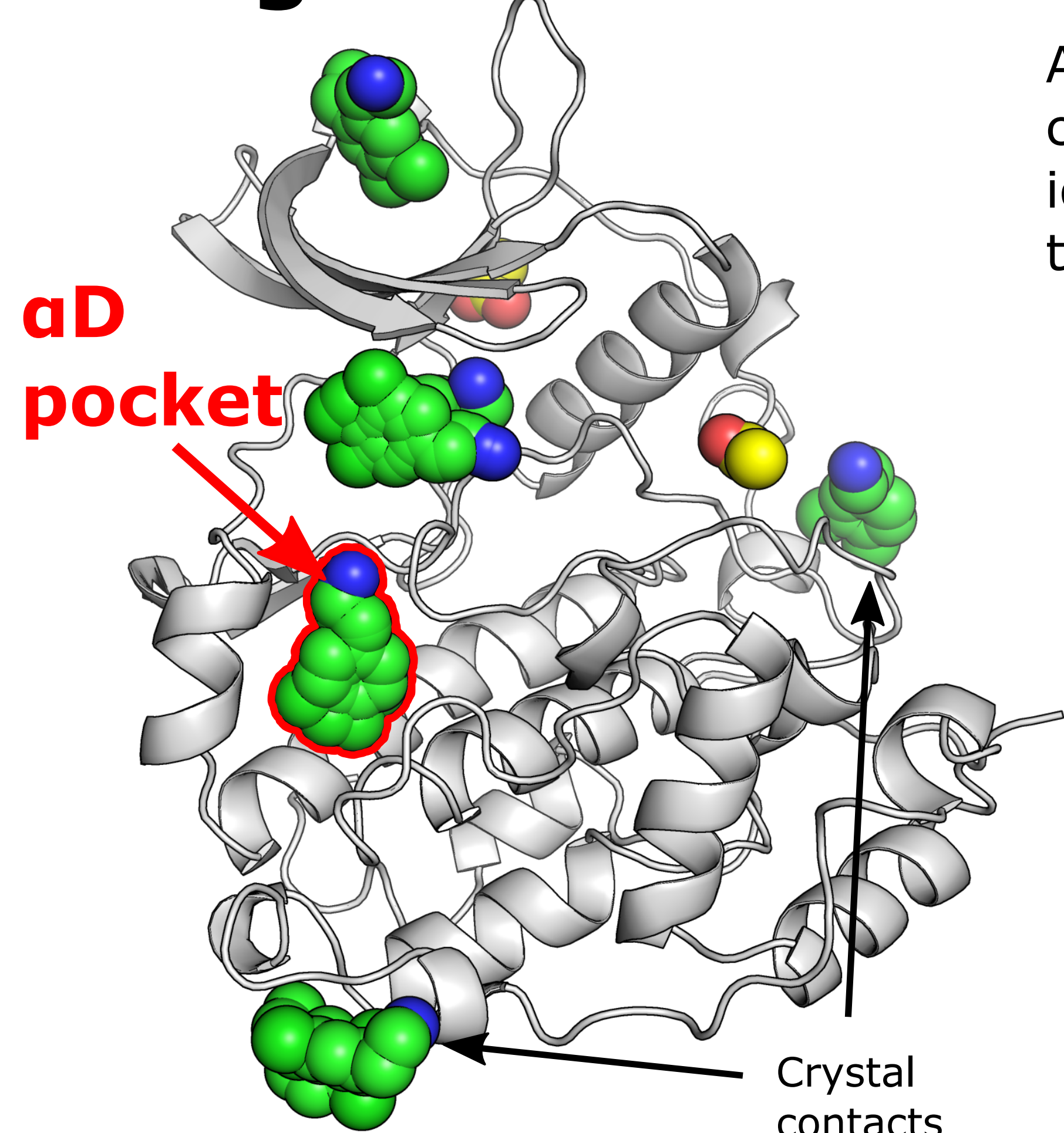


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Introduction

CK2 is a highly conserved kinase with pro survival and anti-apoptotic effects on cells. It is often over expressed in cancer cells in which it promotes their proliferation by multiple mechanisms.¹⁻² A number of potent CK2 α inhibitors, that target only the ATP site, have been shown to inhibit the growth of a range of cancer cell lines and one of these, CX-4945 has progressed to phase II clinical trials.³ Although described as highly selective CX-4945 inhibits at least 12 other kinases with nanomolar IC₅₀s. Strategies to inhibit CK2 α without targeting the ATP binding site offer the promise of enhanced selectivity as well as new mechanisms of action. Here we report the creation and characterisation of a unique inhibitor of CK2 α that targets a novel cryptic pocket and was developed using fragment based methods.

1. Fragment screen



A high concentration X-ray crystallographic screen identified a novel site behind the α D loop.

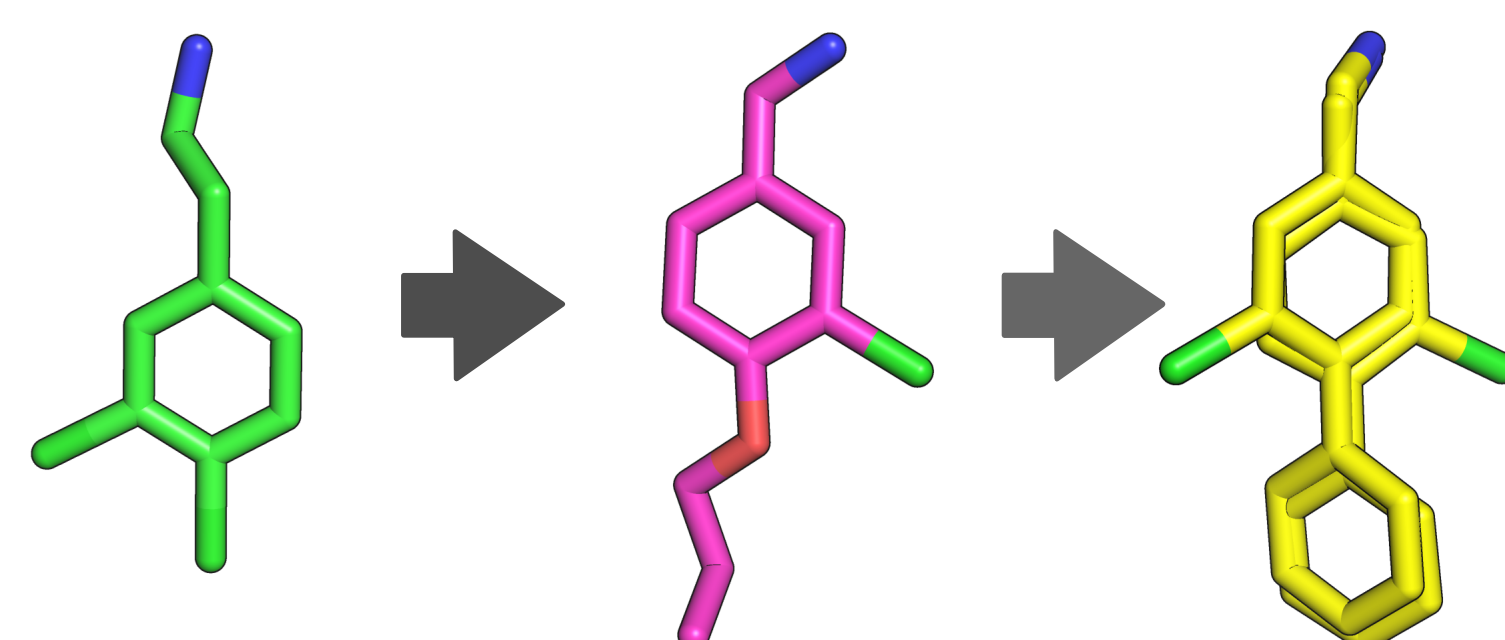
Scan the QR code to see videos.



1. Fragment binding

2. Fragment growth

Aim: Increase selectivity and affinity for the α D pocket



Final fragment has
-significantly increased affinity
-increased selectivity for the α D site

$K_d = 630\mu\text{M}$

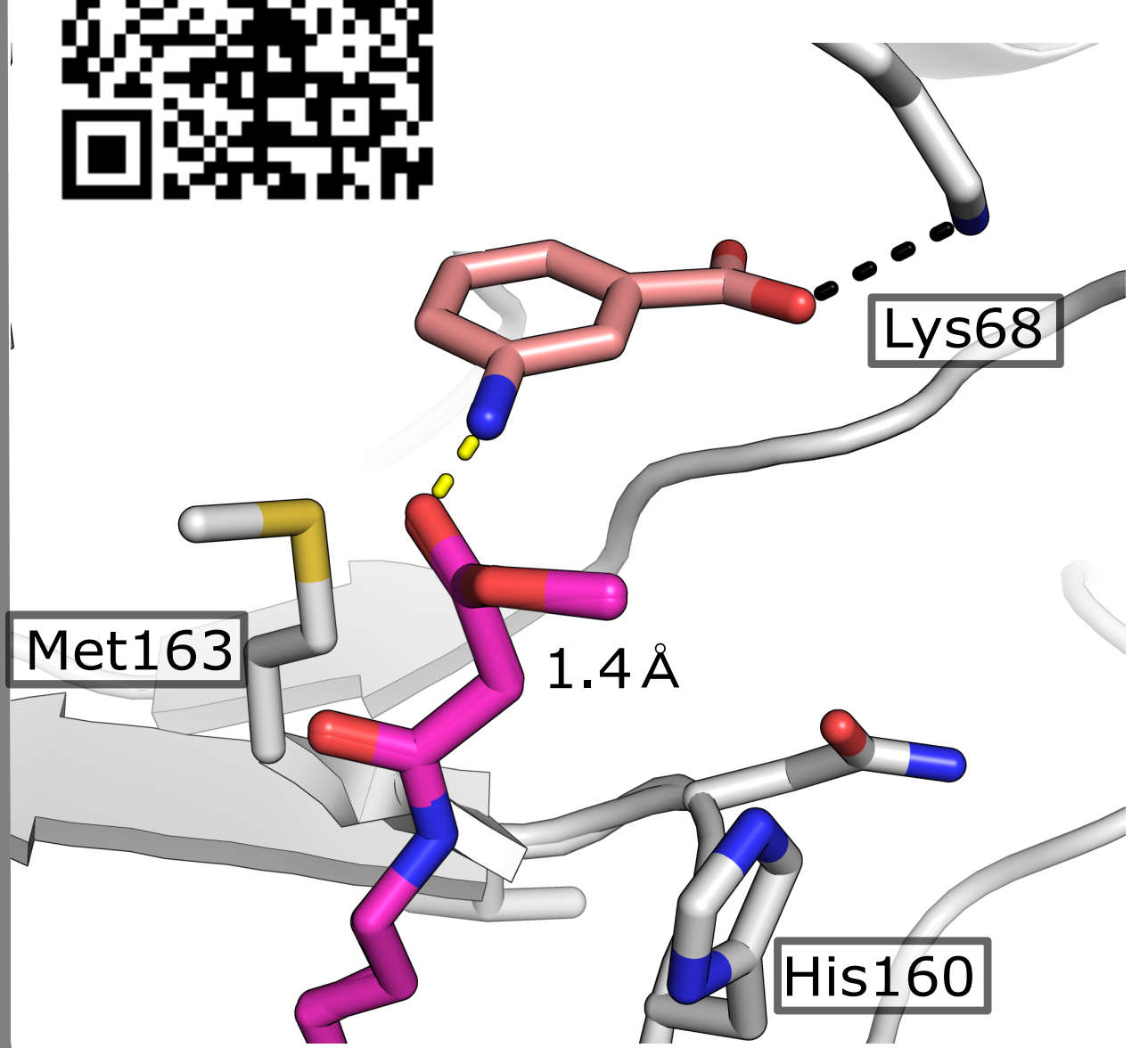
$250\mu\text{M}$

3. Linker design

Aim: link the ATP and α D site fragments

A library of linkers was designed and tested.

A channel that connects the 2 sites was discovered, allowing the use of shorter, more efficient linkers.



Acknowledgements

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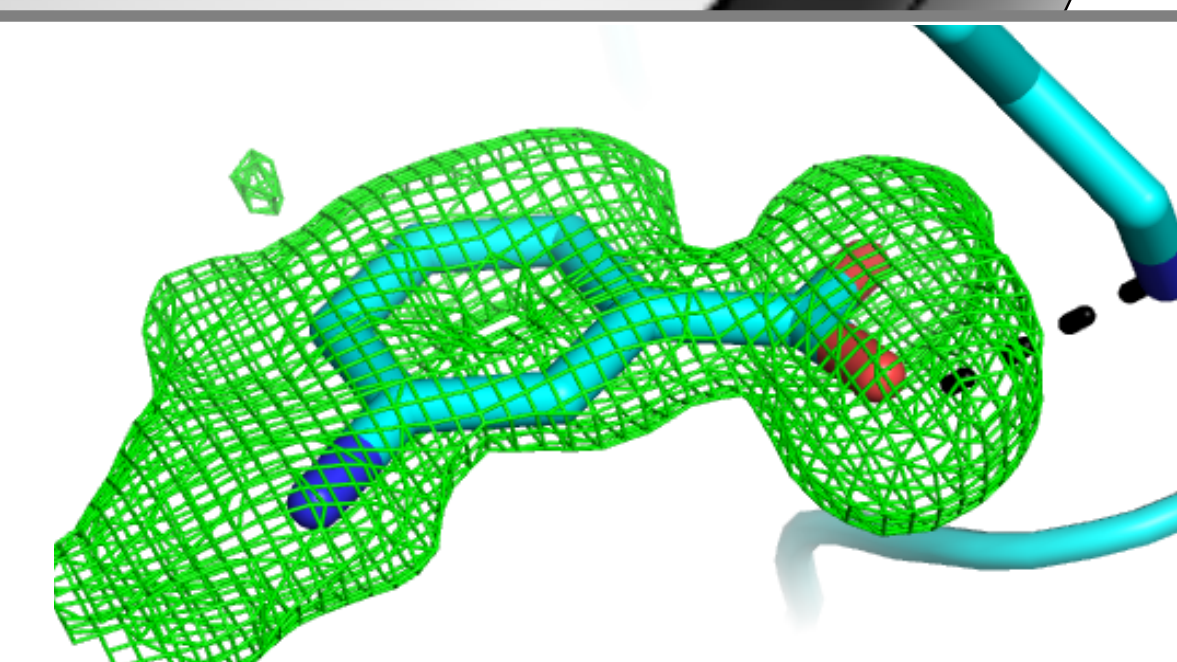
References

1. Ruzzene, M. & Pinna, L. a. Biochim. Biophys. Acta. 1804, 499–504 (2010).
2. Guerra, B. & Issinger, O. G. Electrophoresis 20, 391–408 (1999).
3. Martins, L. R. et al. Leukemia 28, 179–82 (2014)
4. J. Med. Chem. (2011) 54:2:635
5. PLOS ONE (2014) 9:4

4. ATP site fragment

Aim: Identify a fragment in the ATP site to link to the α D pocket.

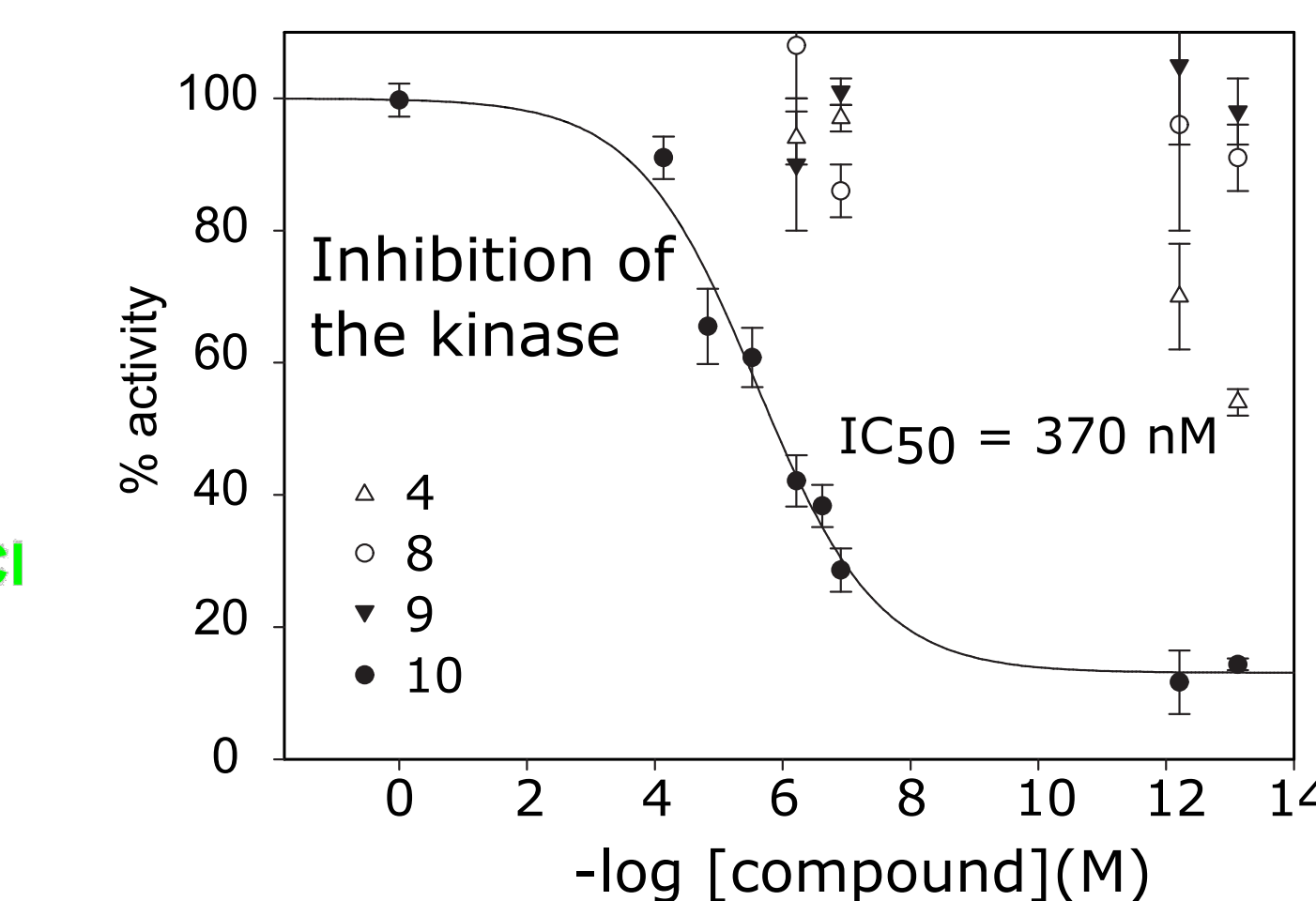
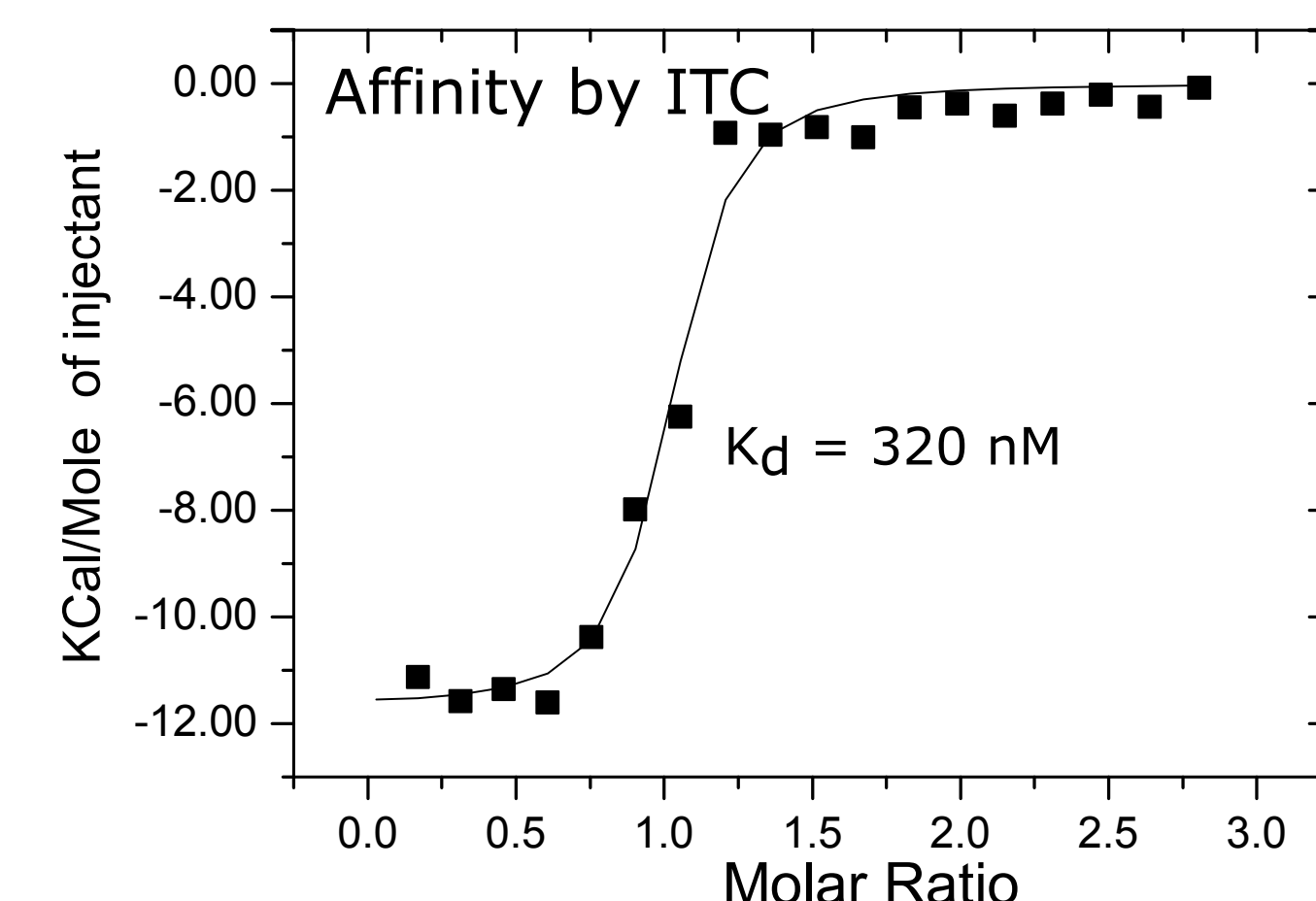
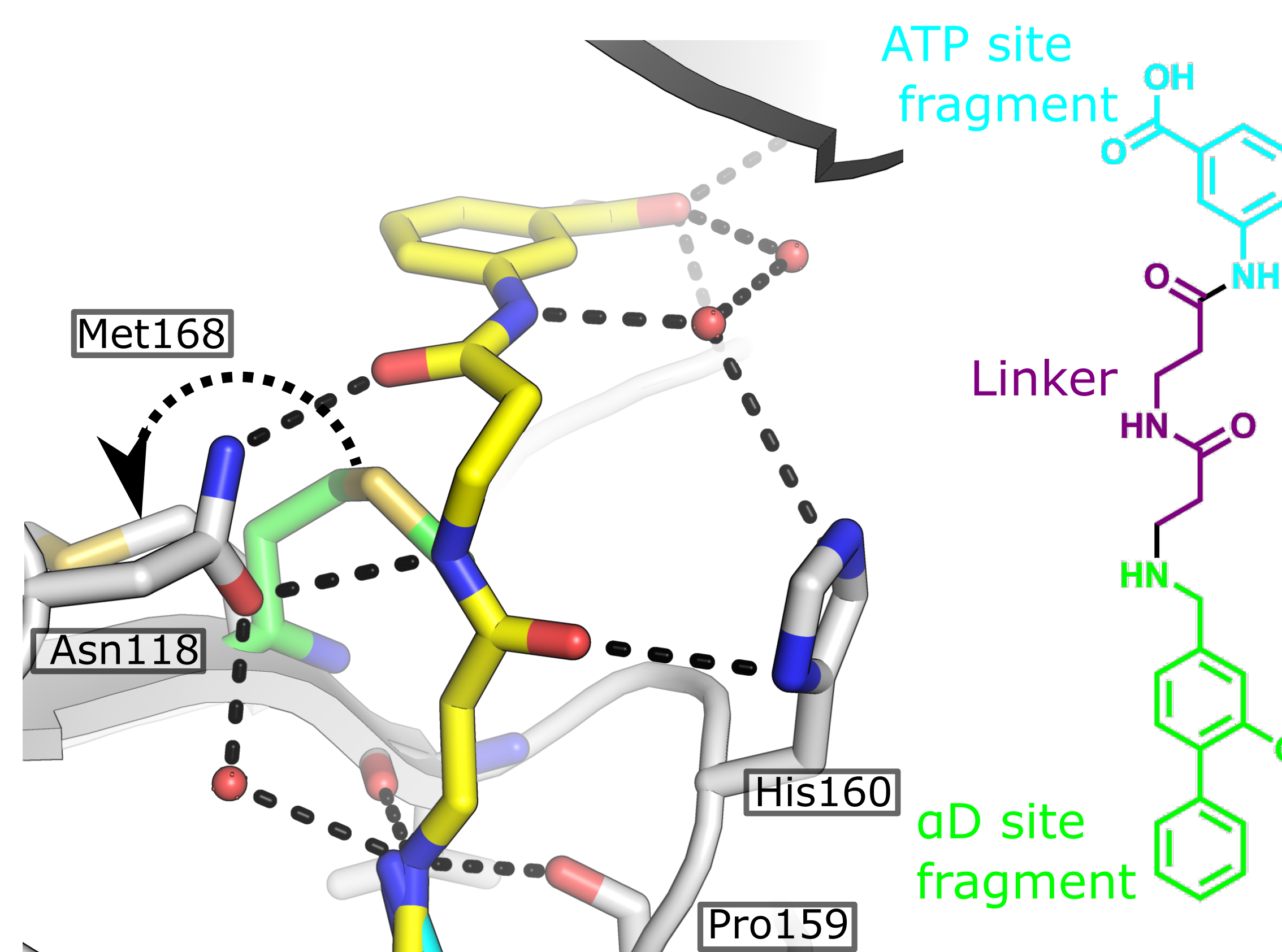
A weakly binding fragment was chosen so that the binding would be dominated by the none conserved α D site therefore giving an inhibitor specific for CK2 α .



5. Linked compound: CAM4066

CAM4066 successfully linked the α D pocket and the ATP site with a greater than 1000-fold improvement on the K_d of the isolated fragment

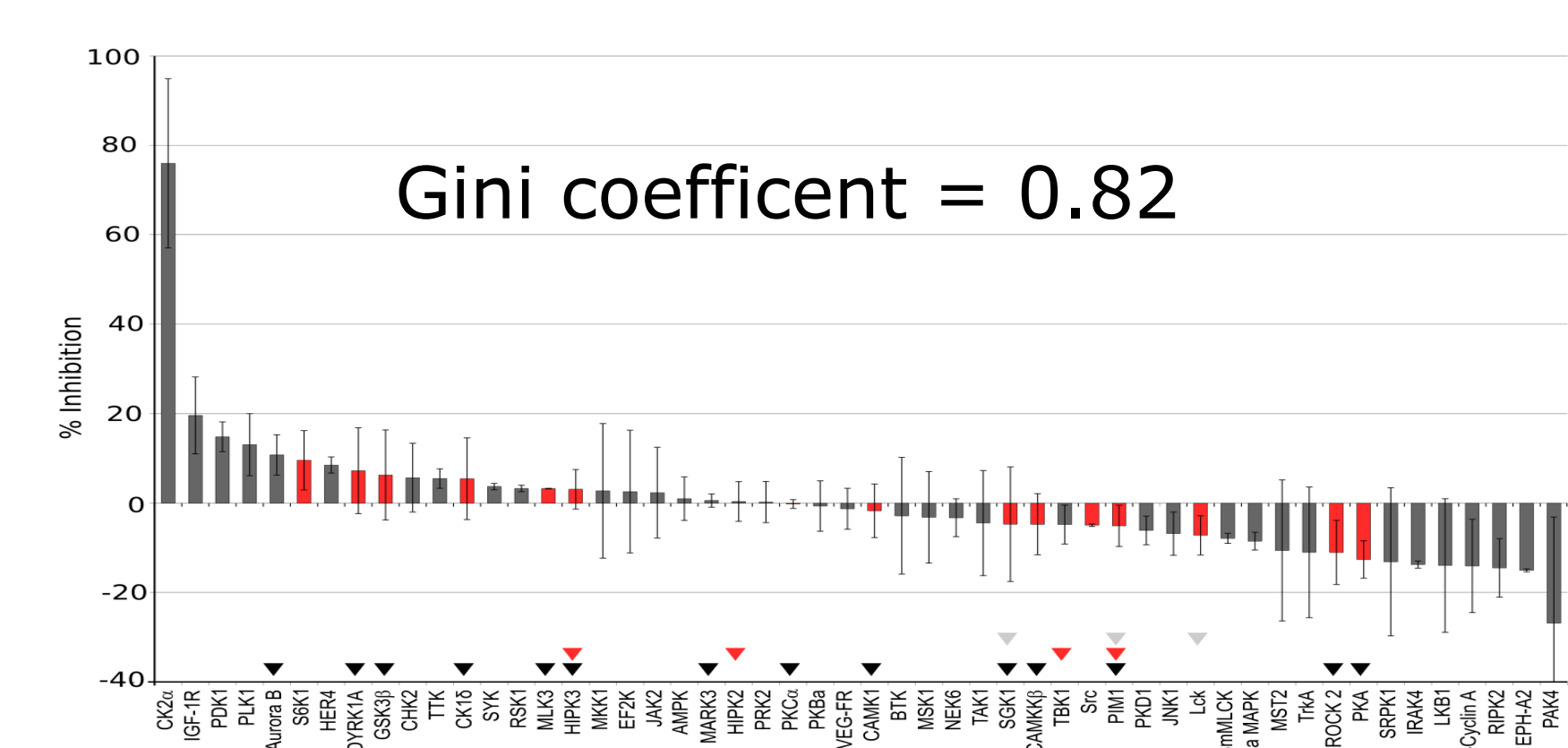
The linker forms an efficient hydrogen bonding next work with the channel linking to the ATP site



6. CAM4066-Validation

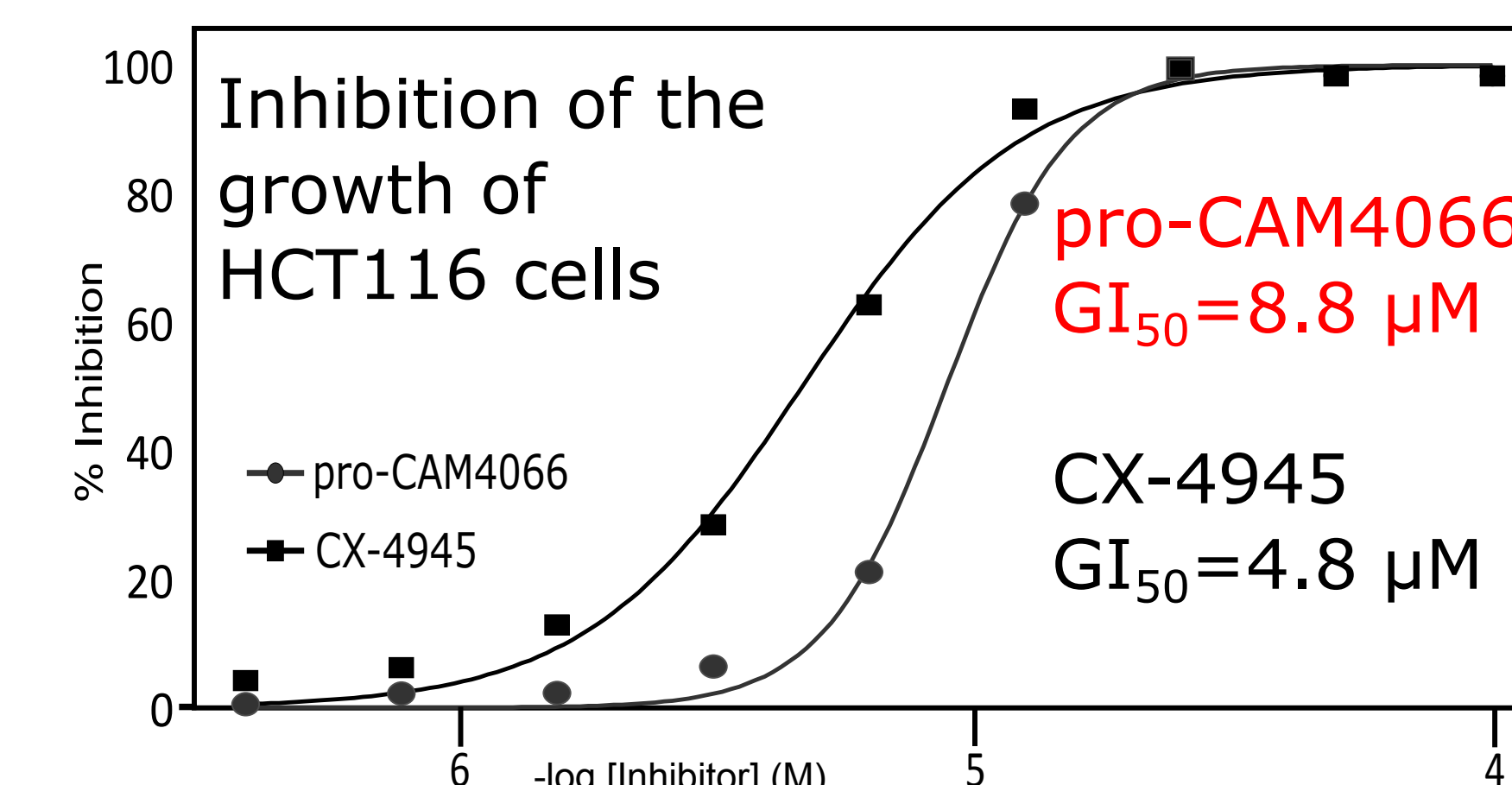
Selectivity screen

CAM4066 was screened against 52 diverse kinases
No significant off target inhibition was detected > Gini coefficient of 0.82
CAM4066 is the most selective CK2 α inhibitor to date.



ProCAM4066

An ester prodrug form of CAM4066 was active in cells:
- It inhibited cell growth at similar levels to the clinical trials candidate CX-4945.
- It showed good inhibition of specific CK2 α phosphorylation targets in HCT116 cells.



Hydrolysis of proCAM4066

