Proof of Concept for Discovering Allosteric Kinase Inhibitors by Characterising a MEK1 Type III Kinase Inhibitor Using a Fluorescent ATP Analogue

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INTRODUCTION

ATP binding pocket Activation Type IV Type I Type II Type III loop Allosteric DFG in DFG out Allosteric within **ATP binding** (inactive) (active) pocket

Small-molecule kinase inhibitors are an important cancer therapy approach, many targeting the

METHODS

Assay: GST-tagged MEK1 protein prepared in magnesium rich assay buffer was added to 384-well assay plates containing test inhibitor PD0325901 (mirdametinib), and pre-incubated for 30 min at room temperature (RT). Fluorescent non-hydrolysable ATP analogue was added, followed by a 30 min RT incubation. Finally, terbium-labelled anti-GST antibody was applied to the assay plate followed by a 2 hour RT incubation. The assay plate was read using the PHERAstar FSX plate reader on HTRF mode.

> Type III fluorescent ATP HTRF assay readout

Standard enzymatic assay readout

highly conserved ATP binding pocket to block ATP and kinase interactions. However, low selectivity and gatekeeper mutation development ultimately leads to acquired resistance, driving a new search for allosteric kinase inhibitors. Type III kinase inhibitors bind to target kinases allosterically within the catalytic domain, adjacent to the ATP binding-site. Once bound, these inhibitors allow and promote ATP binding, which we have leveraged to develop an assay to detect type III inhibitors using a fluorescent ATP analogue.



MEK1 HTRF binding assay schematic using nonhydrolysable fluorescent ATP analogue, which is advantageous for differentiating type III inhibitors. Schematics adapted from Cisbio.

Standard enzymatic kinase assay schematic with the limitation of similar assay readout for each kinase inhibitor type, prompting the development of more selective assays.

Data Analysis: GraphPad Prism version 9.4.1 was used to fit all dose-response curves to a four-parameter fit function and perform all statistical analysis.

RESULTS

Figure 1: Buffer optimisation enhances HTF	F assay signal over 10-fold when P	D0325901 is applied
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A) 25 ·	0.1 μM ATP	Buffer optimisation MEK1 + PD0325901 (10 μM) at 0.1 μM and 0.25 μM ATP
<u>p</u> 20 ·	0.25 μM ATP	T I I

B)

Fluorescent ATP titration comparing response with and without 10 μ M PD0325901

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Figure 1: (A) Buffer conditions were optimised to provide the highest inhibitor-dependent increase in assay signal, testing 24 different buffer compositions with various concentrations of buffer components (TCEP, Triton X-100, MgCl₂, NaCl & BSA) and buffering agents (phosphate, MOPS, Tris & HEPES). (B) In optimised buffer conditions, PD0325901 appeared to cause more than 10-fold increase in HTRF ratio when compared to MEK1 alone.

Figure 2: Dose-ratio titrations of fluorescent ATP and PD0325901 show pharmacological dependency impacting K_D and EC₅₀ respectively



0.0 0.3 1.0	0	
Fluorescent ATP	Fluorescent ATP	
$\mathbf{r} = \mathbf{r} + $	concentration (uM)	Log[PD0325901] (µM)
concentration (µW)		

Figure 2: (A1) Inhibitor EC₅₀ shows a leftward shift, therefore increasing in potency as ATP concentration rises. (A2) Fluorescent ATP concentration impact on PD0325901 EC₅₀. (B1) Leftward shift in K_D is observed as concentration of inhibitor increases. (B2) PD0325901 concentration effect on fluorescent ATP K_D.

CONCLUSION

An inhibitor-dependent increase in assay signal was observed, due to more fluorescent ATP binding to MEK1 causing an increased FRET signal. The leftward shift in ATP $K_{\rm D}$ in the doseratio experiment, confirms that PD0325901 is promoting the binding of ATP, indicative of a type III inhibitor.

This proof-of-concept assay will be used to detect further type III kinase inhibitors, particularly for targets lacking reference compounds, and to potentially identify and characterise new groups of allosteric inhibitors.

