# Development of a kinetic ATPase assay for mechanism of inhibition studies

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#### INTRODUCTION

ATPases are involved in multiple aspects of cellular function and are attractive targets for novel cancer therapeutics. Key to designing new drugs targeting ATPases is the ability to measure inhibition of ATP turnover. We routinely use the endpoint ADP Glo assay for ATPase activity that is well suited to screening, however a kinetic assay would be better suited to complex mechanism of inhibition (MoI) studies.

#### ASSAY DEVELOPMENT WITH THE PHOSPHATE SENSOR DETECTION REAGENT

Phosphate sensor concentration was optimised to maximise signal to background and reduce cost.

#### MECHANISM OF INHIBITION RESULTS

Both compounds were non-competitive with respect to ATP so are able to bind equally well to the apo enzyme or enzyme-ATP complex. The compounds differed in their Mol with respect to the DNA substrate. Compound A was uncompetitive with respect to DNA so is able to only bind to the enzyme-DNA complex. Compound B showed mixed inhibition with respect to DNA so is able to bind to both the apo enzyme and enzyme-DNA complex with a preference for the apo enzyme.



Figure 1. ATP hydrolysis schematic. ATPases catalyse the hydrolysis of adenosine triphosphate (ATP) to adenosine diphosphate (ADP) and inorganic phosphate (P<sub>i</sub>). The energy from ATP hydrolysis is utilised by the enzyme to achieve a cellular function.

#### **INITIAL ASSAY DEVELOPMENT**

Initially four kinetic ATPase assay formats were investigated; three Transcreener assays from Bellbrooks to measure ADP levels and the phosphate sensor reagent from Thermo Fisher to measure P<sub>i</sub> levels.



Figure 4. Substrate and detection reagent concentration optimisation. Each substrate was titrated in turn against a range of phosphate sensor concentrations. The second substrate was included in excess to approximate a single substrate system.

Excellent Z' values were achieved at low enzyme concentrations of 1-4 nM highlighting the sensitivity of this detection reagent.





#### **Inhibitor A**



Figure 2. Schematics for A) the three Bellbrooks Transcreener kinetic ADP detection assays, B) the Thermo Fisher phosphate sensor reagent and C) the endpoint ADP Glo assay.

As part of the initial assay development a compound interference check was performed in each assay format. 21%, 14% and 36% (FI, TR-FRET, FP respectively) of the 14 compounds tested showed compound interference in the Transcreener assays compared to 0% in the phosphate sensor assay.

Transcreener	Transcreener	Transcreener ADP <sup>2</sup>	<sup>2</sup> Phosphate
ADP <sup>2</sup> FI assay	ADP <sup>2</sup> FP assay	TR-FRET Red assay	sensor assay
Inhibition	Inhibition	Inhibition	Inhibition

Figure 5. Linearity check and confirmation of compound inhibition under final assay conditions. Compound  $IC_{50}$  curves were used to select an appropriate concentration range to use in the Mol assays.

### MECHANISM OF INHIBITION DETERMINATION

Detailed mechanism of inhibition (MoI) studies were performed to determine whether compounds were competitive, non-competitive or uncompetitive with respect to the substrates. Kinetic characterisation assays were performed in which  $K_{\rm M}$ and  $V_{\rm max}$  kinetic parameters were assessed at different inhibitor concentrations.



Figure 7. Detailed MoI studies with two compounds of interest. Initial rates were calculated in the presence of 10  $\mu$ M phosphate sensor reagent between 5-16 minutes. Initial rates in RFU/second were plotted against substrate concentration. Using GraphPad Prism this data was fitted to equations for competitive, non-competitive, uncompetitive and mixed inhibition and the best fit determined using AIC. This data was also fitted to the Michaelis Menten equation to determine the  $K_{\rm M}$  and  $V_{\rm max}$  values plotted here to show the trend with increasing compound concentration which can indicate the type of inhibition. The second substrate was included in excess to approximate a single substrate system.

#### SUMMARY

Several detection technologies for continuous measurement of ATPase activity were assessed here. The phosphate sensor reagent was selected as the most promising as it was not prone to interference with our compounds and was very sensitive. We successfully used this assay to perform detailed MoI studies and were able to demonstrate differences in the MoI of representative compounds from two series of interest. Using the kinetic assay format for these studies significantly reduced both resource and reagent consumption compared to using an endpoint assay format.



Figure 3. Compound interference with each assay format. Interference assays were run in the absence of enzyme by spiking in a set amount of product, any variation from the expected signal in the presence of compound is the result of interference with the detection technology.

The phosphate sensor assay was selected to take forwards.

Figure 6. Compound inhibition schematics. A competitive compound can only bind to the apo enzyme. A noncompetitive compound can bind equally well to the apo enzyme or enzyme complex. An uncompetitive compound can only bind to the enzyme substrate complex. Often the mechanism lies in between one of these cases and is referred to as mixed inhibition.

