

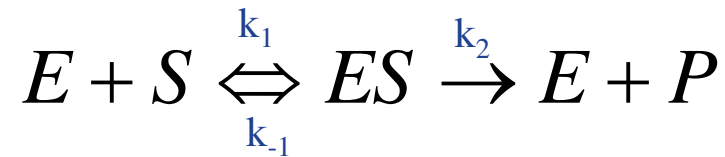
ITC in Enzyme Characterization



imagination at work

Enzyme Kinetics

K_M is analogous to K_d



$$K_M = \frac{k_2 + k_{-1}}{k_1}$$

When $k_{-1} \gg k_2$

$$K_M = K_d$$



imagination at work

Enzyme Kinetics

V_{\max} is proportional to [enzyme]

$$V_{\max} = k_{\text{cat}}[E]$$

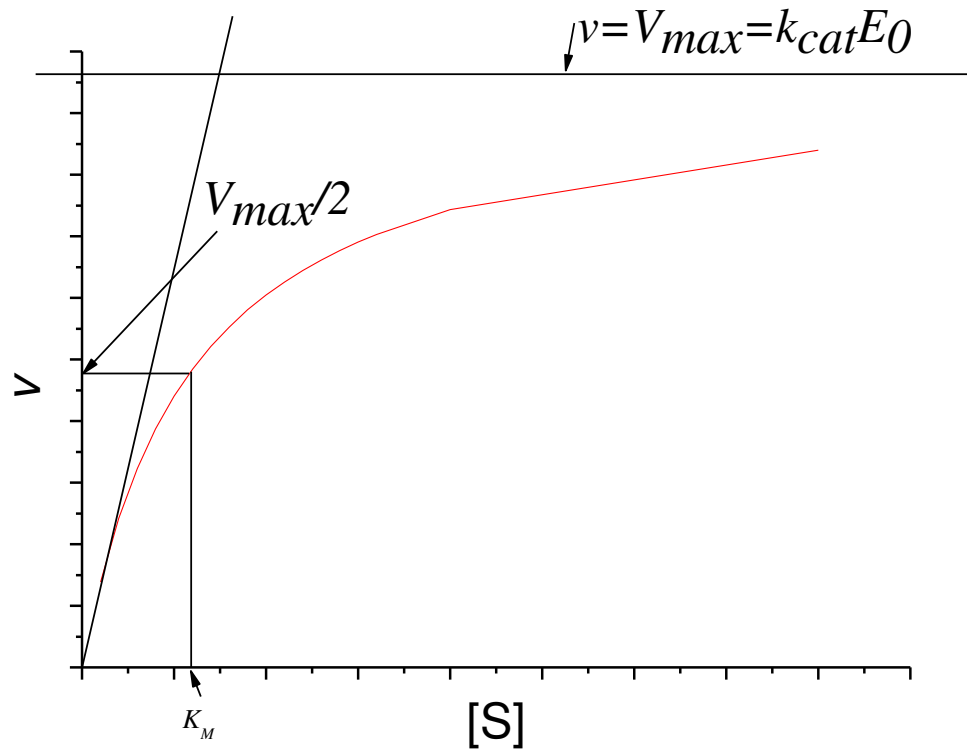
K_m is independent of [E]

Velocity of reaction is given by Michaelis-Menten equation

$$v = \frac{[S]}{K_M + [S]} k_{\text{cat}}[E]$$



Classical Enzyme Kinetics



Two ITC methods

Multiple substrate injections

- Low enzyme concentration
- Steady state conditions

Continuous assay

- Higher enzyme concentration
- Single injection of substrate



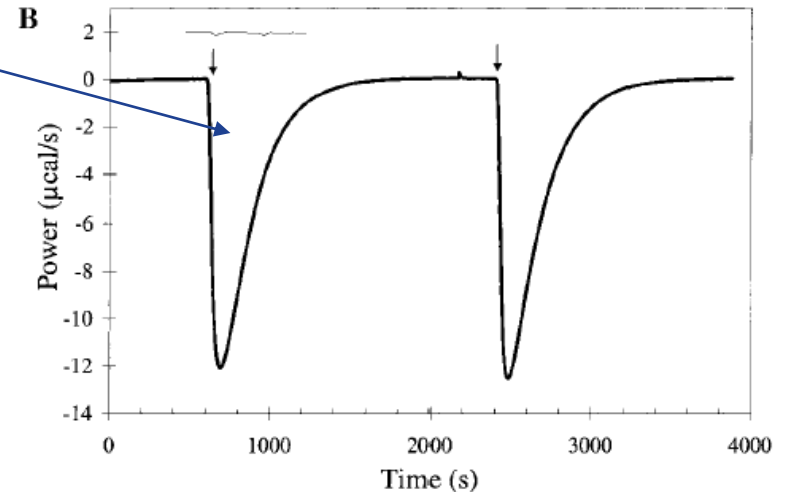
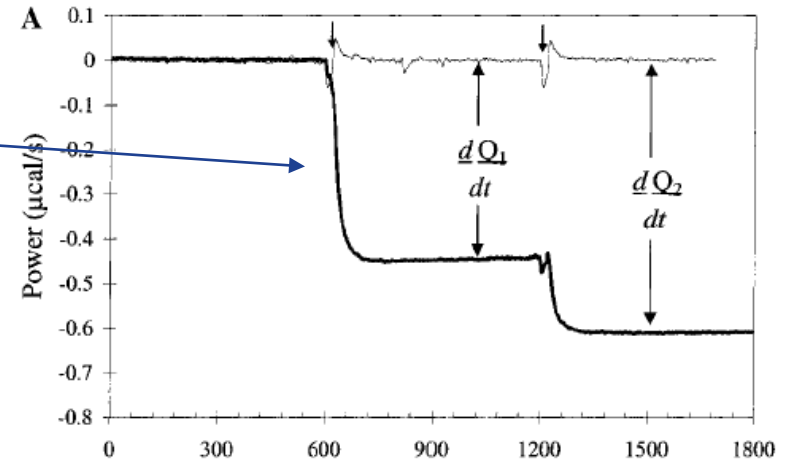
Enzyme Kinetics By ITC

$$Power = \frac{dQ}{dt}$$

$$Q = n\Delta H_{app} = [P]_{total} V\Delta H_{app}$$

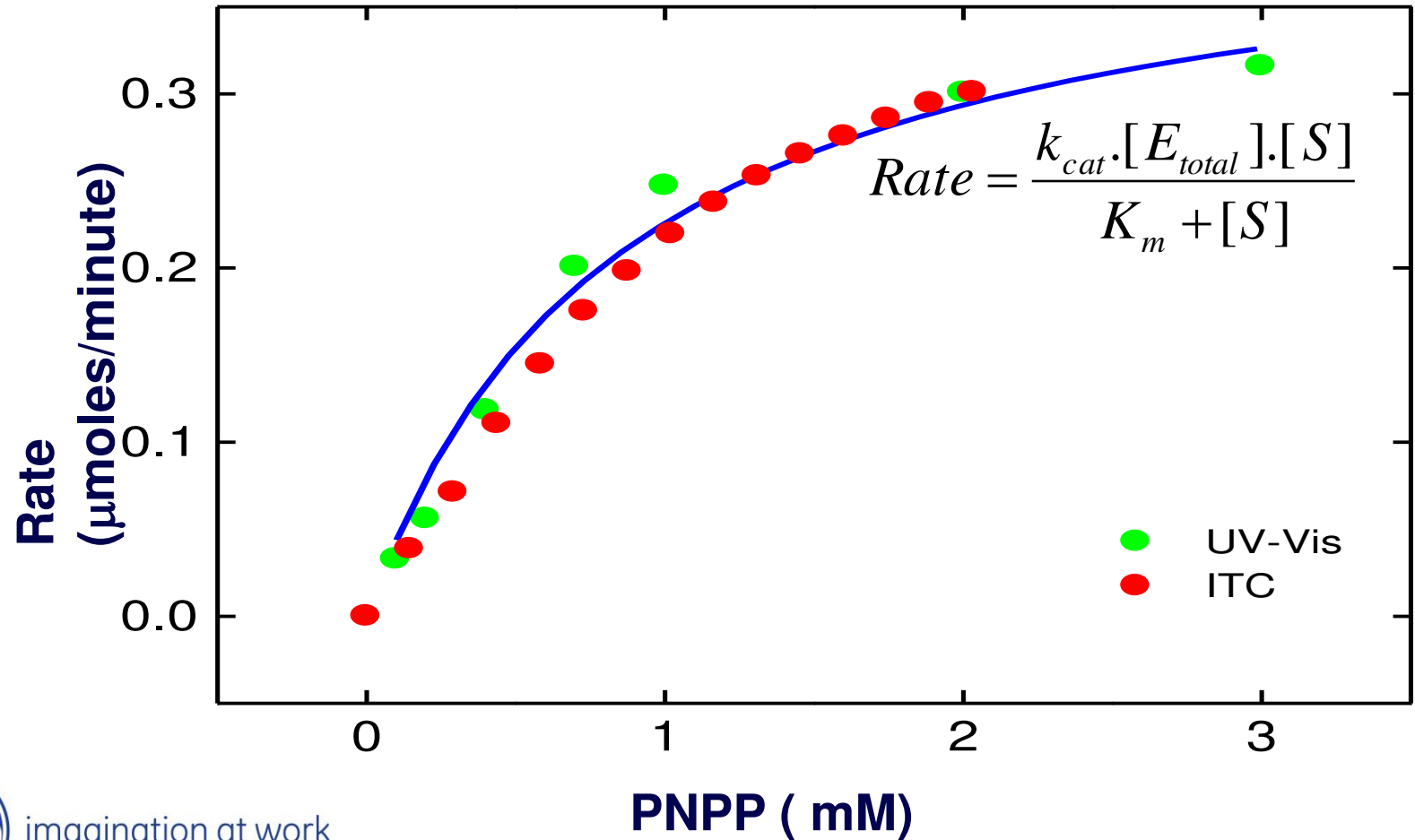
$$\frac{dQ}{dt} = \frac{d[P]}{dt} V\Delta H_{app}$$

$$Rate = \frac{d[P]}{dt} = \frac{1}{V\Delta H_{app}} \frac{dQ}{dt}$$



imagination at work

Enzyme Kinetics



[Enzyme] and [Substrate]

Following each injection of substrate there should be no appreciable depletion of substrate (<5%) prior to the next injection.

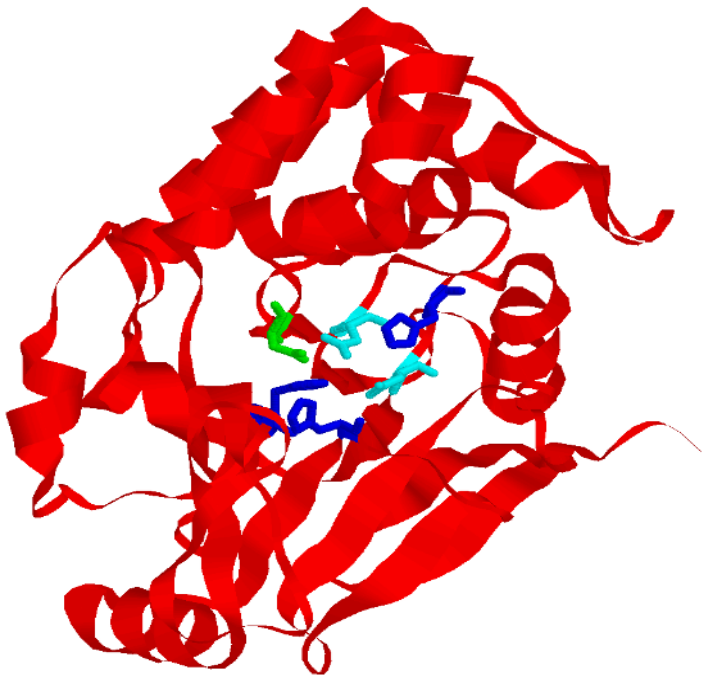
However substrate concentration cannot be too high otherwise $[S]$ will be above K_m after the first few injections.

If enzyme-substrate affinity is high then enzyme concentration needs to be low

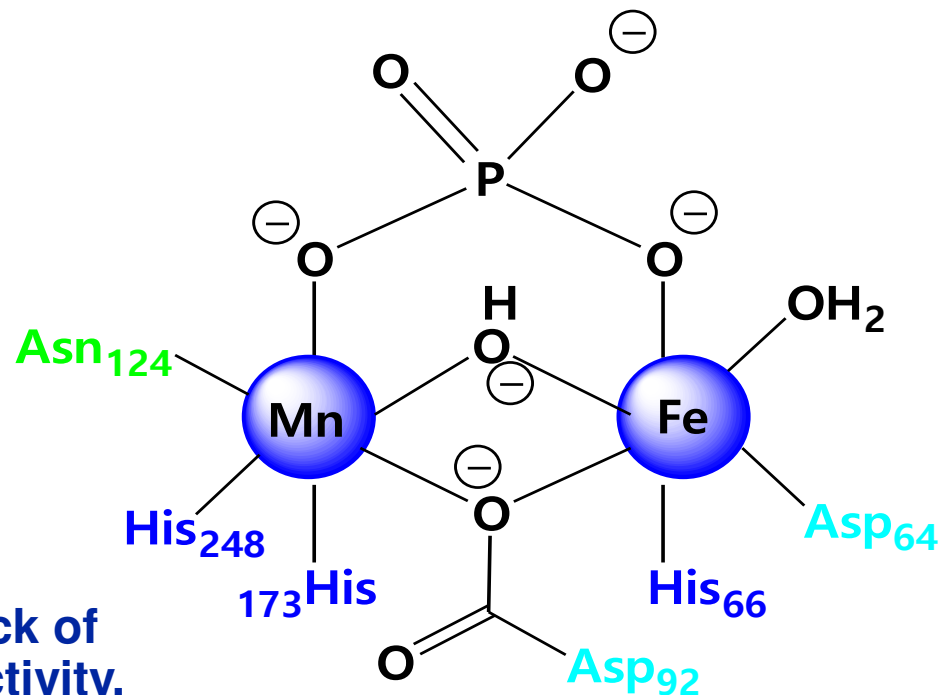
However this relies upon there being a sufficiently large enthalpy to detect.



PP1- γ Phosphatase



Phosphatases catalyse the hydrolytic removal of a phosphate group. In this study we examine a **serine/threonine phosphatase (PP1- γ)**.



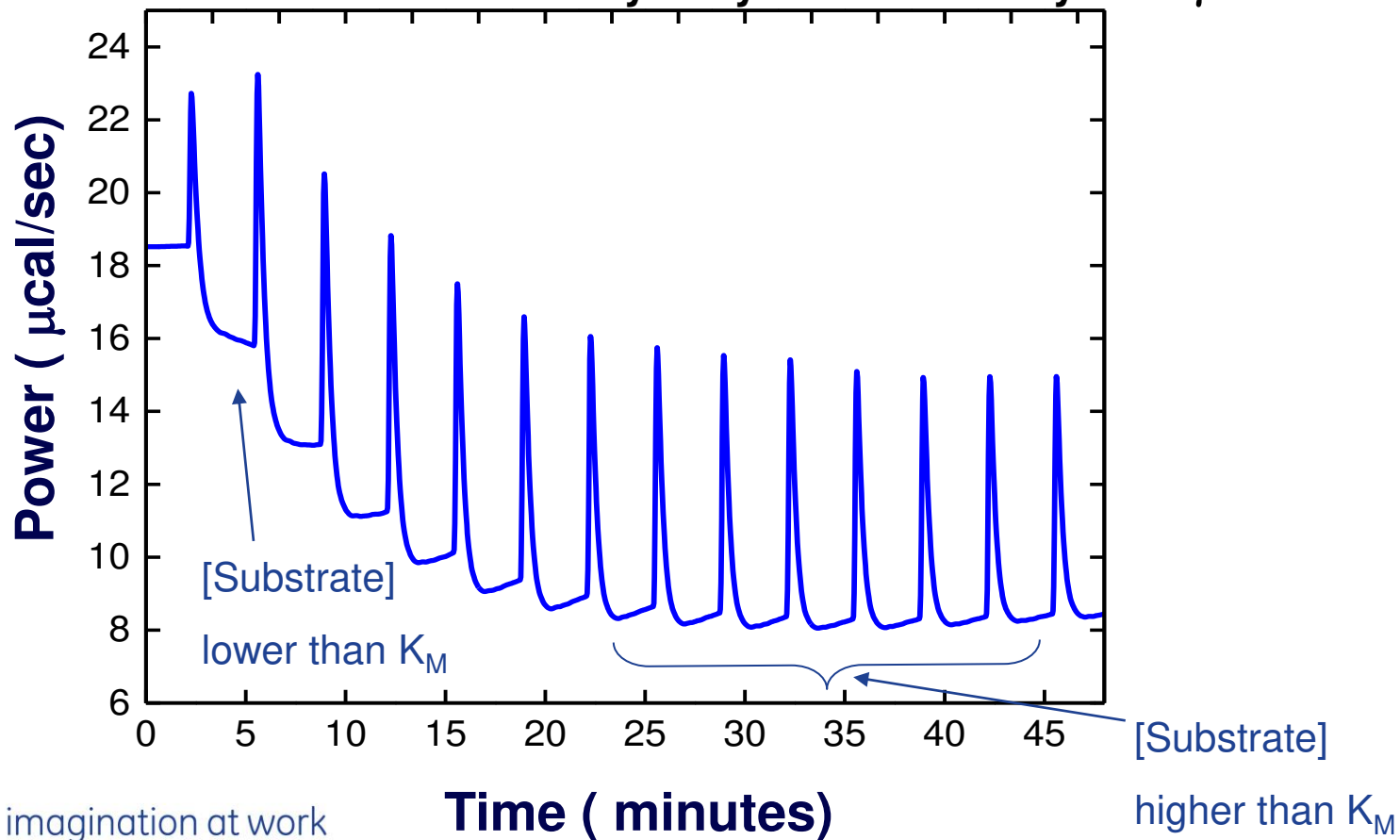
This enzyme catalyses the direct attack of water and it requires metal ions for activity.



imagination at work

Enzyme Kinetics

Raw calorimetric data for the measurement of the reaction rate for the hydrolysis of PNPP by PP1- γ .



imagination at work

Time (minutes)

[Substrate] higher than K_M

Enzyme Kinetics

ITC

$$K_m = 1.2(\pm 0.2) \text{ mM}$$

$$k_{\text{cat}} = 0.6 (\pm 0.1) \text{ s}^{-1}$$

$$V_{\text{max}} = 0.43 \text{ } \mu\text{M min}^{-1}$$

UV

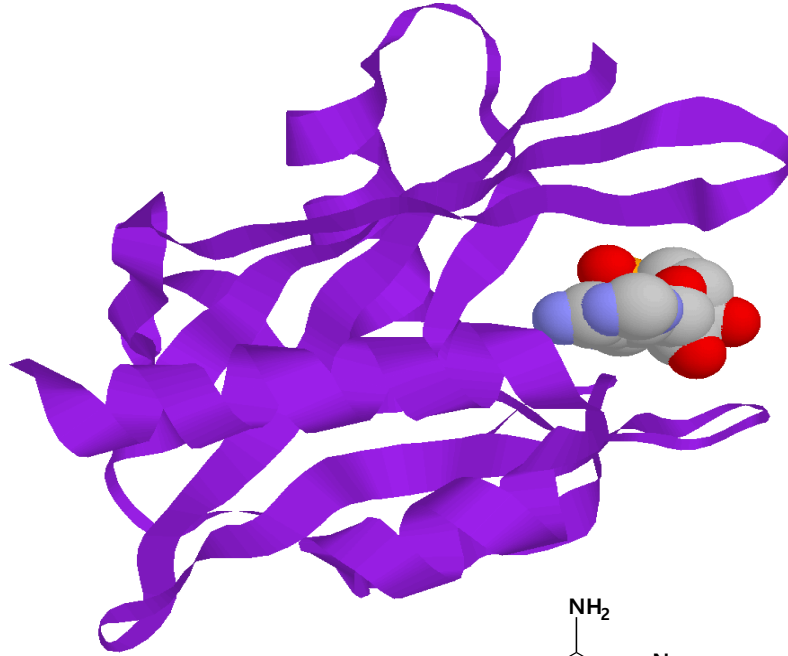
$$K_m = 0.9(\pm 0.2) \text{ mM}$$

$$k_{\text{cat}} = 0.5 (\pm 0.1) \text{ s}^{-1}$$

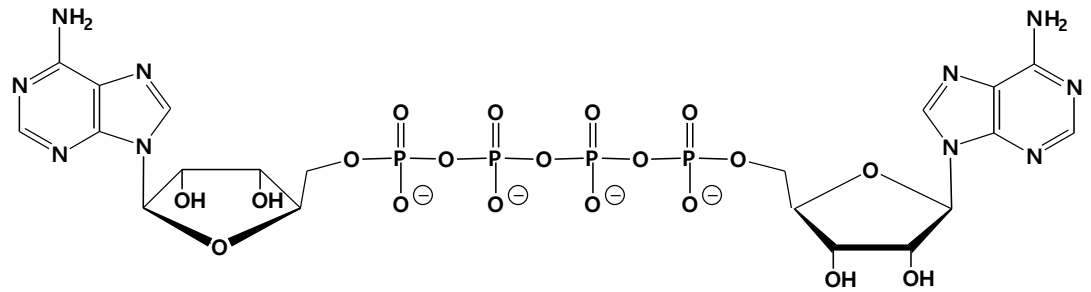
$$V_{\text{max}} = 0.39 \text{ } \mu\text{M min}^{-1}$$



Enzyme Kinetics

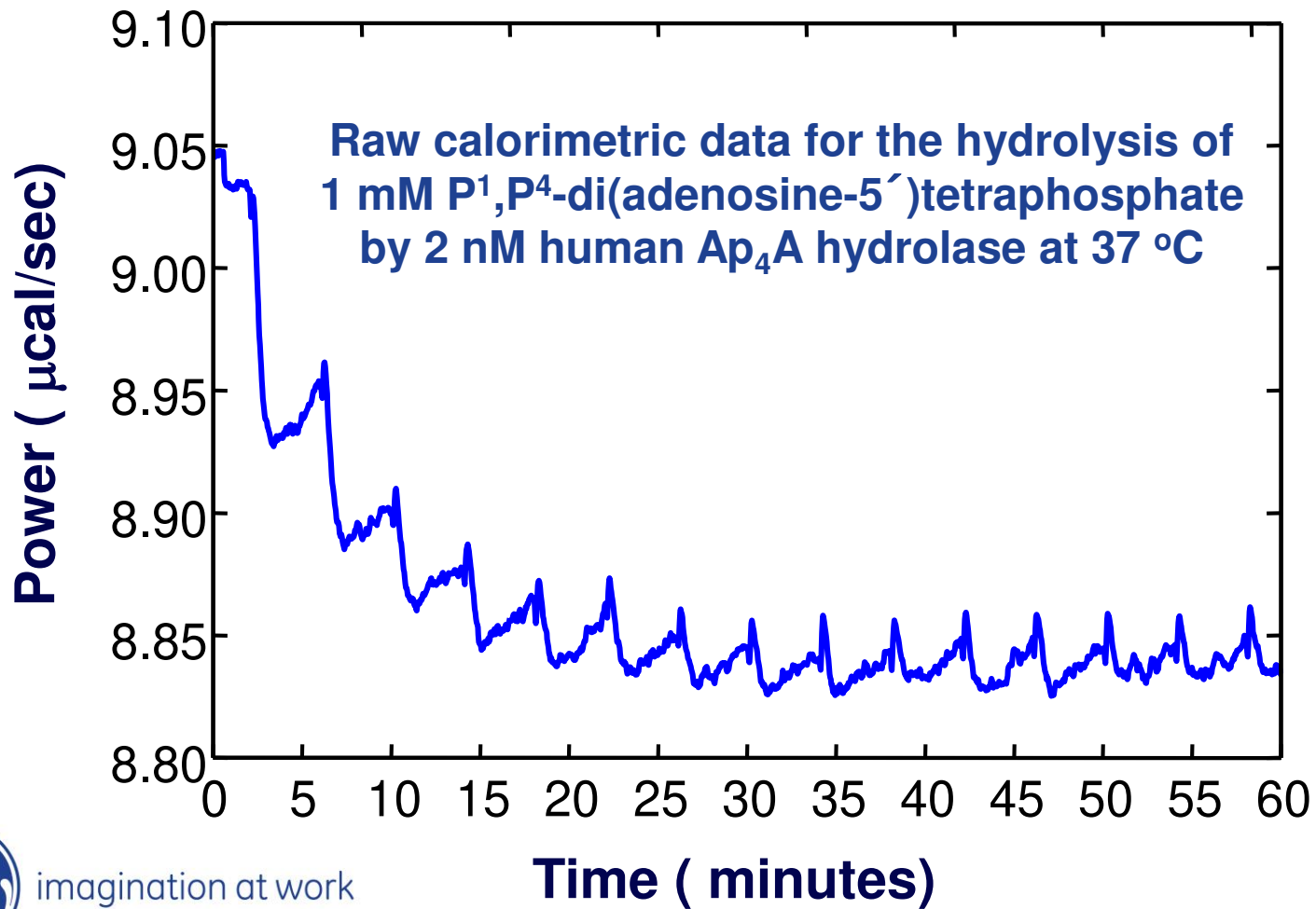


Ap4A Hydrolase

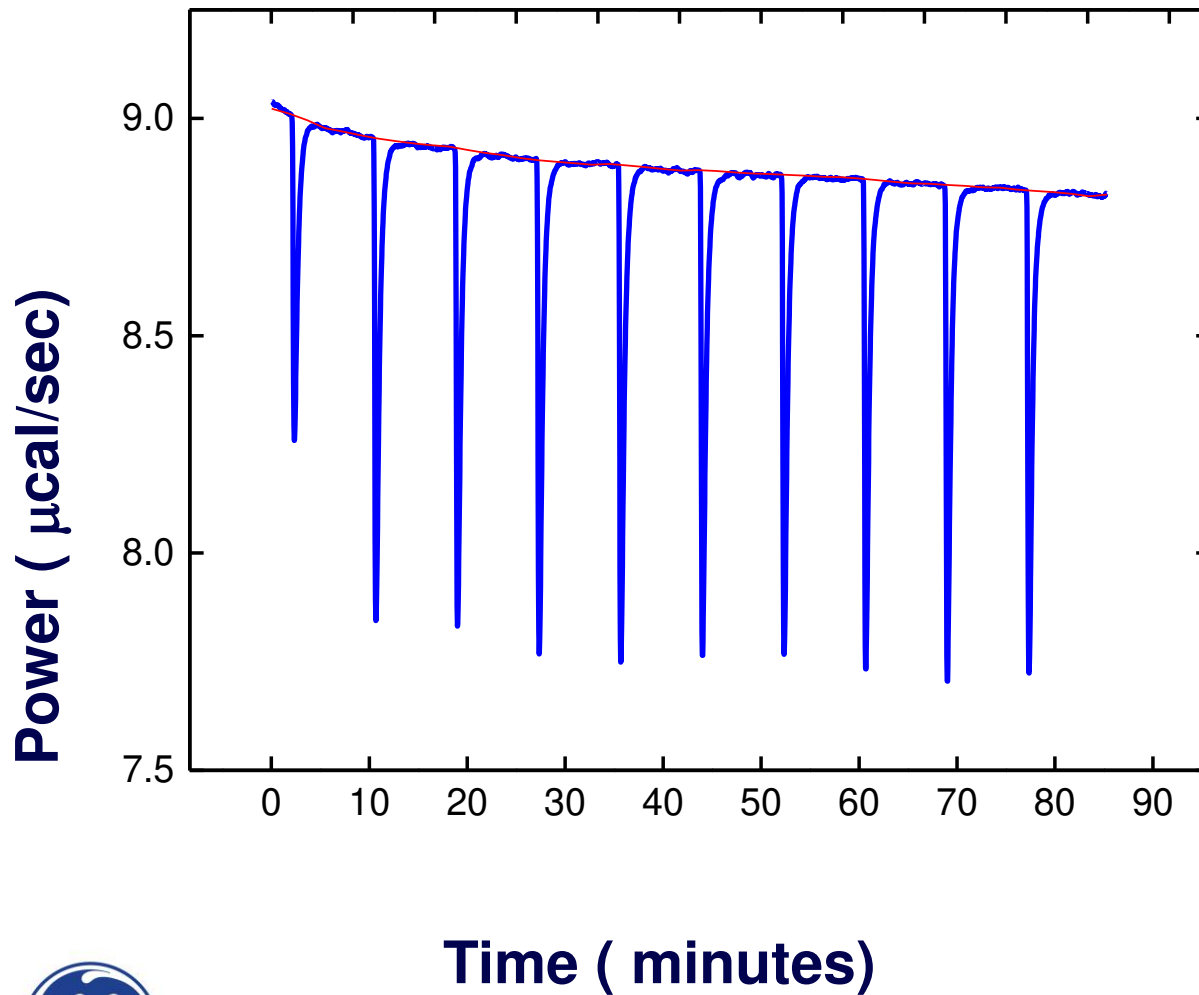


imagination at work

Enzyme Kinetics



Enzyme Kinetics

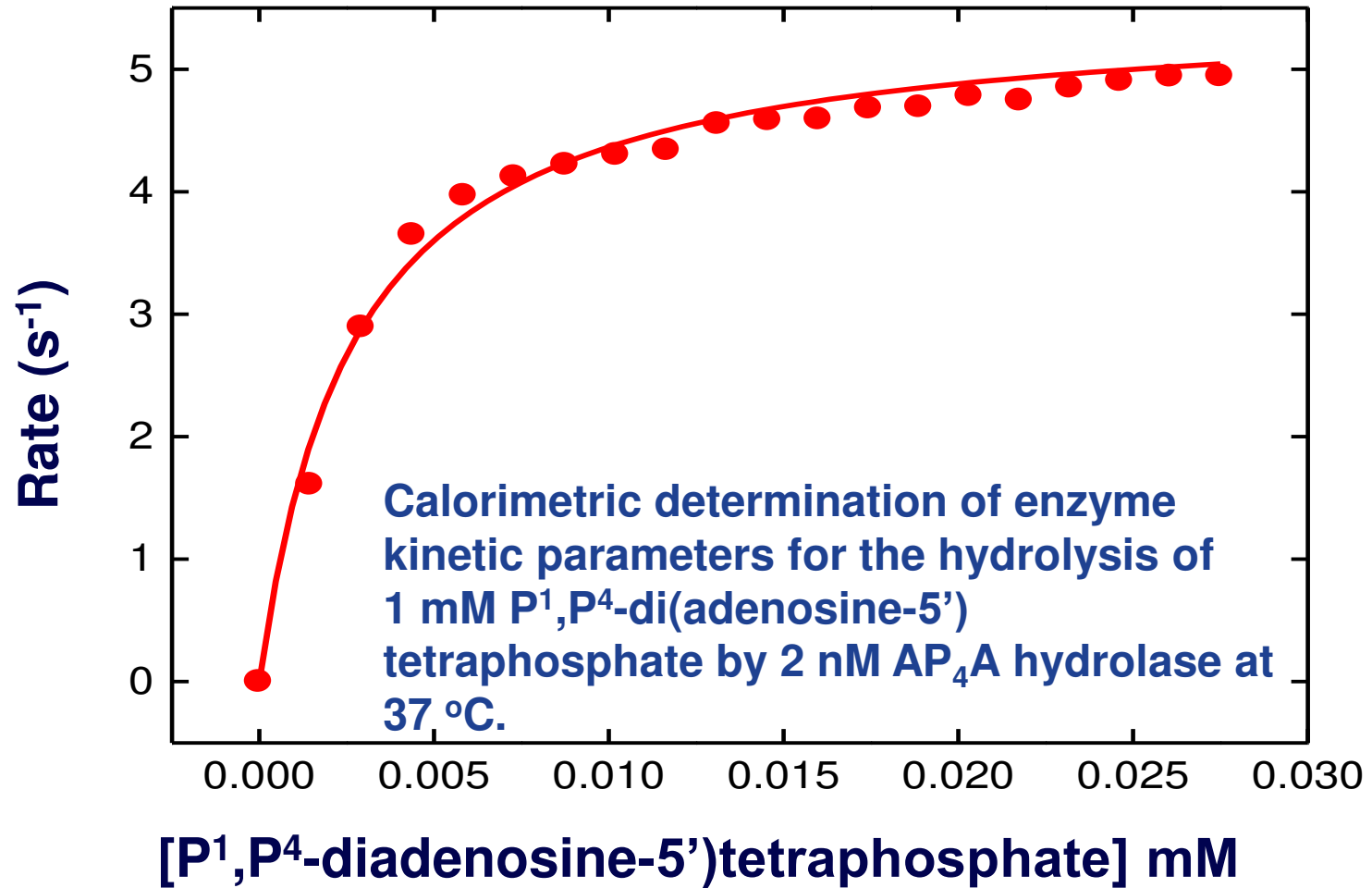


Determination of ΔH_{app}
for the hydrolysis
of 1 mM P₁,P₄-
di(adenosine-5')
tetraphosphate
by 100 nM AP₄A
hydrolase at 37 °C

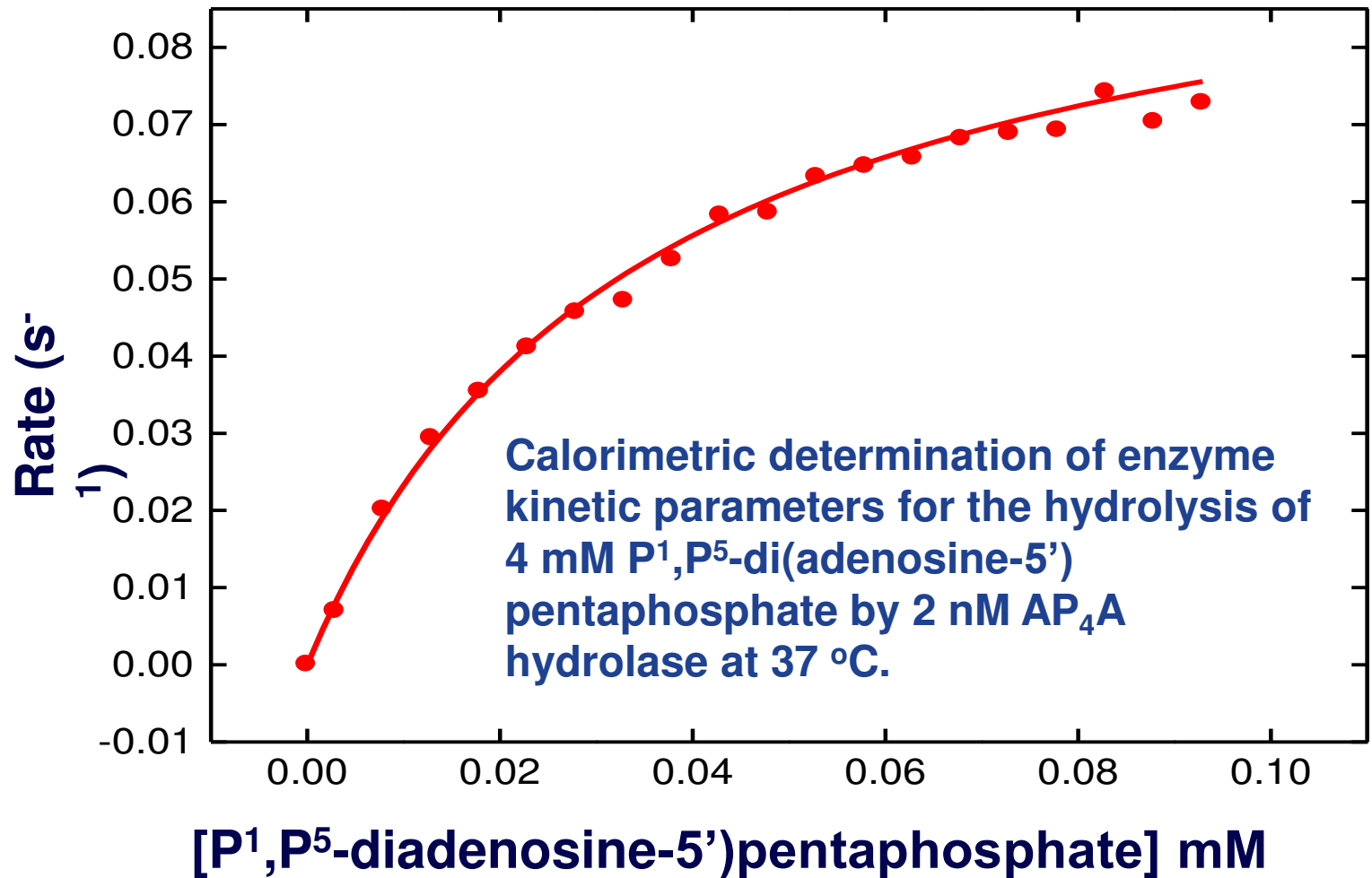


imagination at work

Enzyme Kinetics



Enzyme Kinetics



Enzyme Kinetics

Hydrolysis of diadenosine polyphosphates by Ap₄A Hydrolase

P¹,P⁴-di(adenosine-5') **P¹,P⁵-di(adenosine-5')** **ATP**
tetraphosphate **pentaphosphate**

K_m (μM)
nd

2.68 ± 0.80

34.5 ± 1.8

k_{cat} (s^{-1})
nd

2.76 ± 0.03

0.05 ± 0.01

V_{max} (nM s^{-1})
nd

5.52

1.0

ΔH_{app} (kcal mol^{-1})
 8.6 ± 0.2

-19.24 ± 0.27

-16.38 ± 0.74

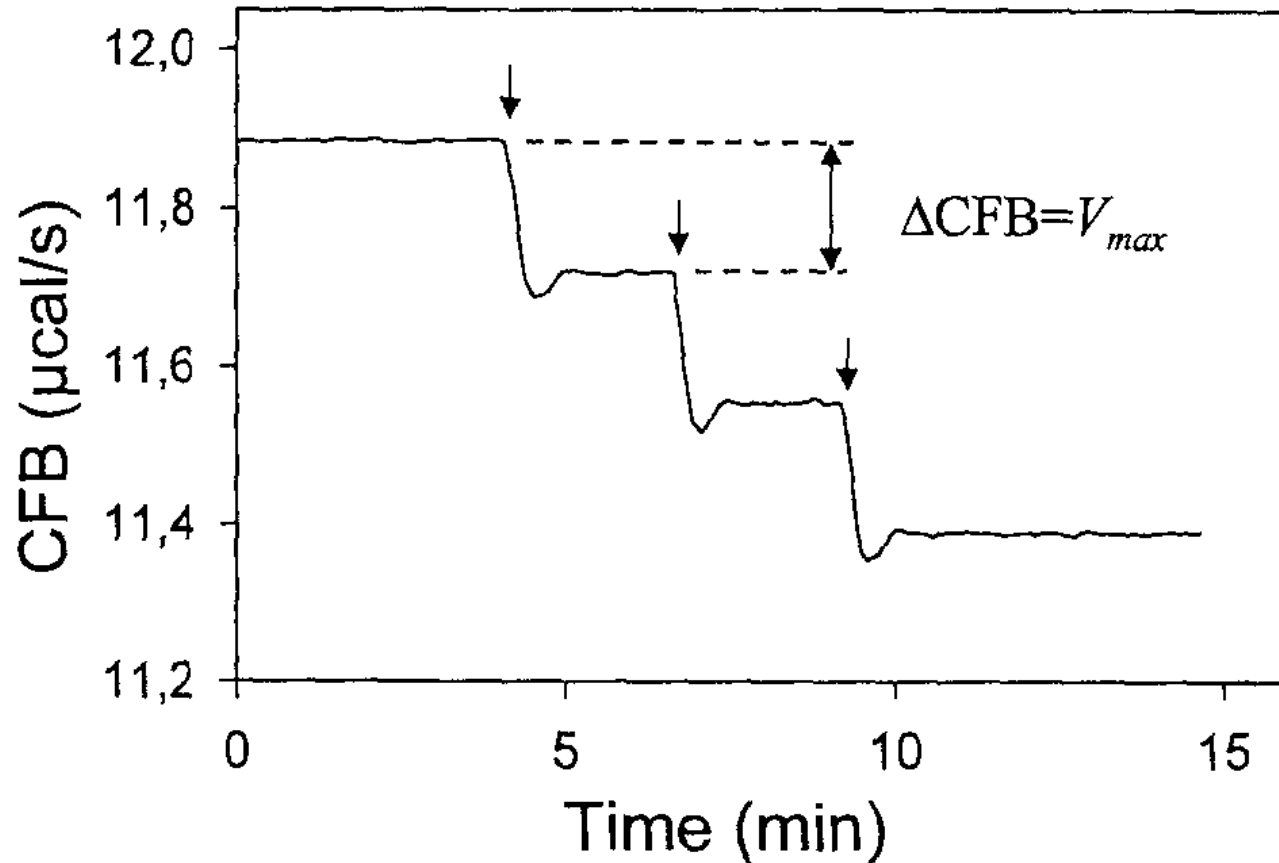
-



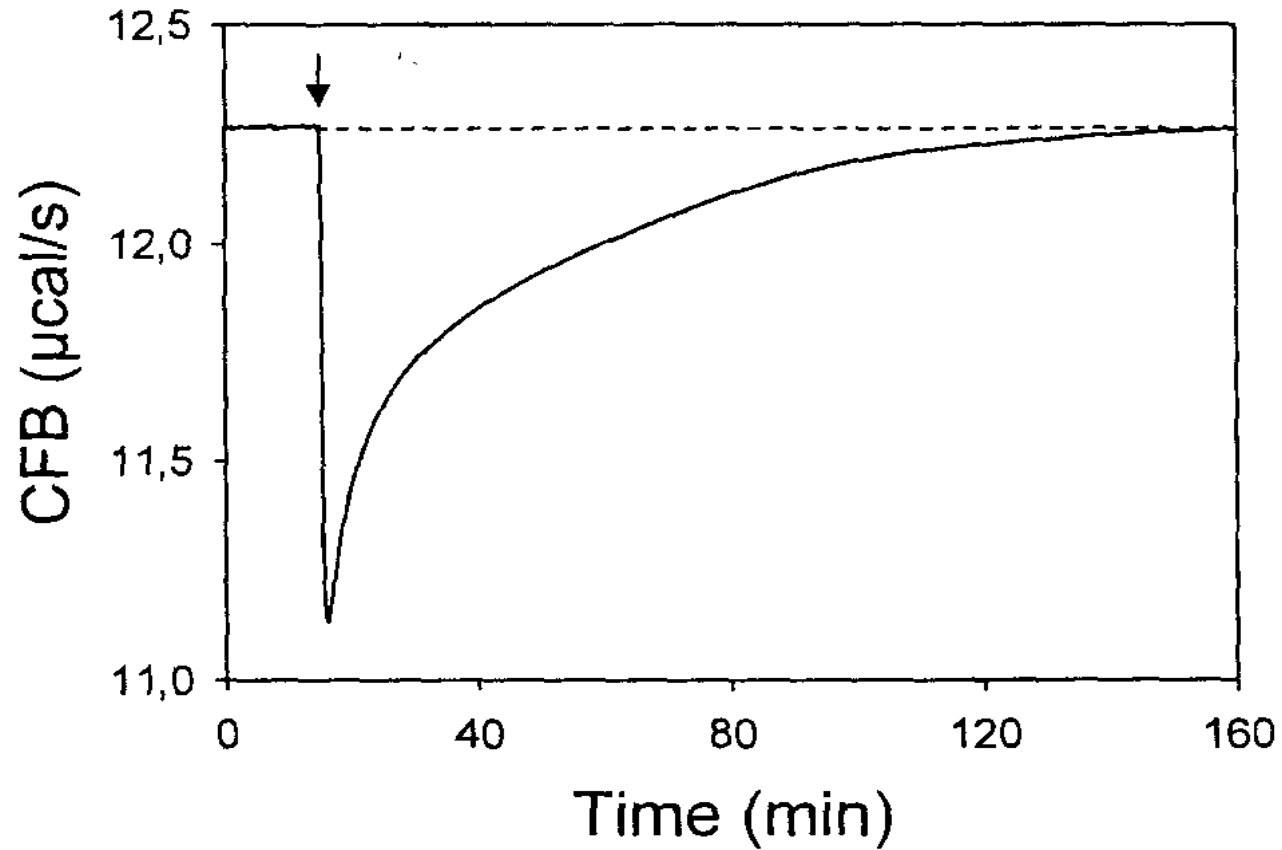
imagination at work

Enzyme Kinetics

Alternatively.....



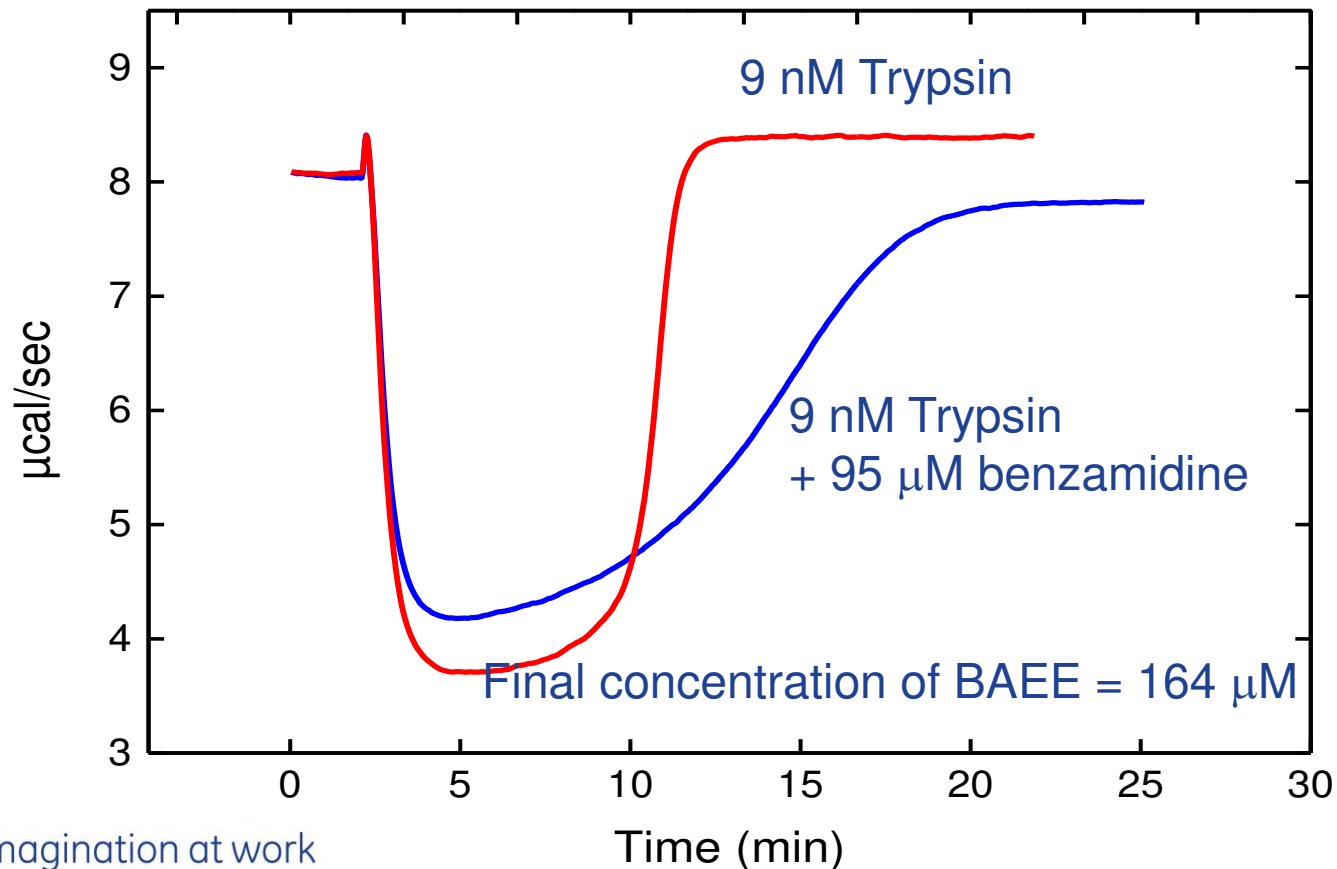
Enzyme Kinetics



imagination at work

Continuous Injection Method

Trypsin catalysed hydrolysis of BAEE and inhibition by benzamidine



Continuous Injection Method

From this raw data the **rate** at any time can be obtained using:

$$Rate = \frac{d[P]}{dt} = \frac{1}{V \cdot \Delta H_{app}} \frac{dQ}{dt}$$

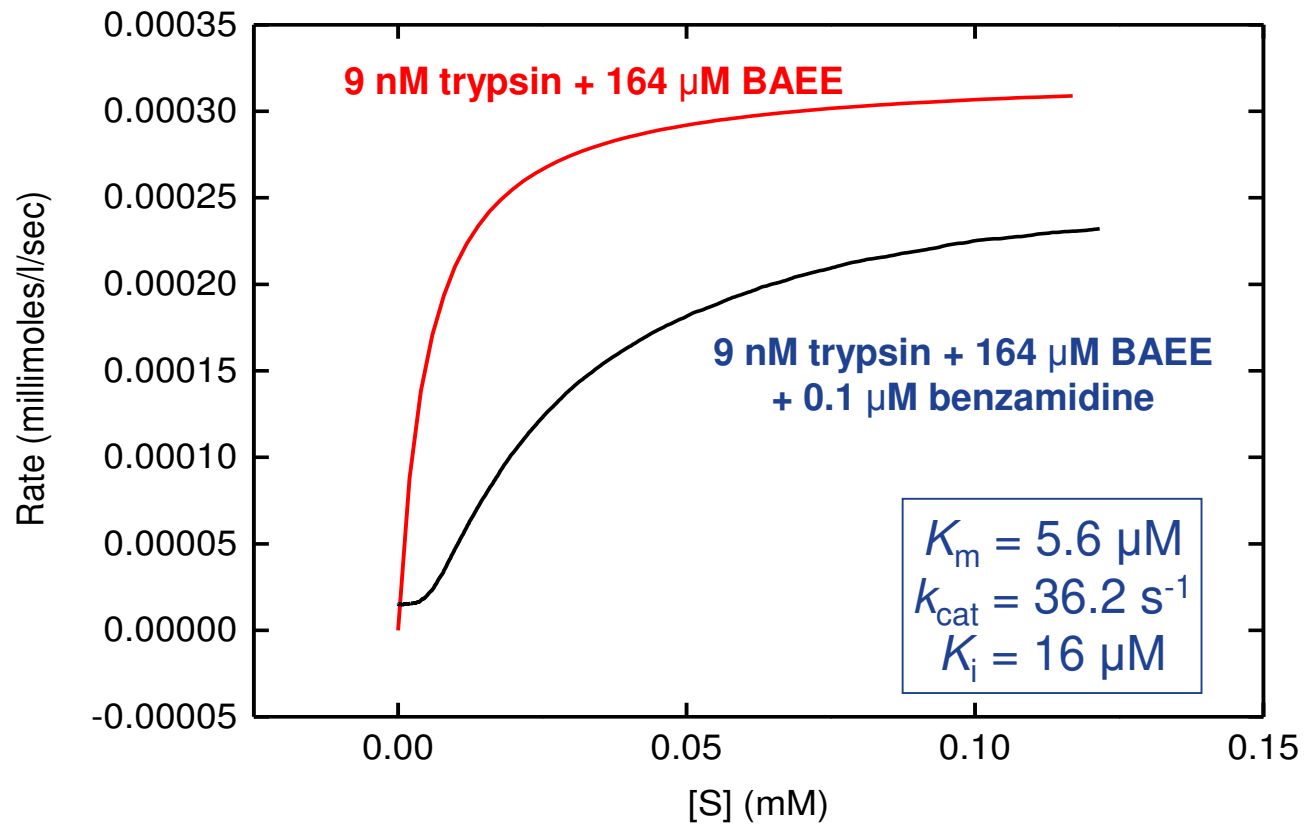
The amount of substrate left at any given time after the initial injection can be determined from the integral of the heat evolved:

$$S_t = S_{Total} - P_{(t)} = S_{Total} - \frac{\int_{t=0}^t Q_{(t)}}{\Delta H \cdot V}$$



imagination at work

Continuous Injection Method



Enzyme kinetics

Universal assay

No need to couple assay to colour change reaction

No labelling required

Modifications to substrate and/or enzyme do not effect assay

Turbid solutions can be used

Pico-femtomoles of protein required

Quick



imagination at work

Comparison of calorimetric and other assay data

Enzyme	Calorimetric		Literature Values	
	K_m	V_{max}	K_m	V_{max}
E.C. 1.5.1.3 (DHFR) ^a Substrate = DHF	1.2 μM	6 s^{-1}	6 μM	3 s^{-1} [i]
E.C. 2.7.1.1 (yeast hexokinase) ^b Substrate = glucose	72 μM	270 s^{-1}	100 μM	450 s^{-1} [ii]
E.C.3.3.2.6 <i>B. cereus</i> Penicillinase I ^c	120 μM	3600 s^{-1}	50 μM	2800 s^{-1} [iii]
E.C. 3.4.21.4 (Trypsin) ^d	4 μM	15 s^{-1}	5 μM	22 s^{-1} [iv]
E.C. 3.4.21.16 (HIV protease) ^e Substrate = KARVnLF(NO ₂)EAnL Substrate = VSQNYPIVQ	5 - 300 μM [NaCl] dependent	10 s^{-1}	15 μM	45 s^{-1} ³³
E.C. 3.5.1.5 (<i>H. pylori</i> urease) ^f	0.79 mM	1400 s^{-1}	0.17 mM	2700 s^{-1} [v]
E.C. 4.1.1.7 (<i>F. heparinum</i> heparinase) ^g	1.8 μM	0.059 s^{-1}	10.2 μM	92 s^{-1} [vi]
E.C. 4.1.1.39 (Rubisco) ^h Substrate = ribulose bis phosphate	0.15 mM	1.95 s^{-1}	0.053 mM	1.76 s^{-1} [vii]
E.C. 4.1.3.18 (Acetolactate synthase) ⁱ	4.8 mM	11 s^{-1}	5.5 mM	5.3 s^{-1} [viii]
E.C. 5.99 (GroEL) ^j	3 μM n = 2.9	0.052 s^{-1}	5 μM n = 2.5	0.08 s^{-1} ²²
E.C. 6.4.1.1 (Pyruvate carboxylase) ^k Substrate = ATP Substrate = pyruvate	85 μM 105 μM		58 μM 440 μM	[ix]



imagination at work

Todd and Gomez, Anal. Biochem. 276, 179-187 (2001)