

Inhibitors of *Plasmodium Falciparum* Calcium-Dependent Protein Kinase 1 (*Pf*CDPK1), a Novel Target for the Potential Treatment of Malaria

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# Contents



- Why Malaria?
- PfCDPK1: A Novel Target
- Hit Identification and Early Lead
- Optimisation and Synthesis of Imidazopyridazines
- In Vivo Results
- Parasite Rate of Reduction Assay
- Biotinylation Experiments
- Conclusion

## Why Malaria?



- A major public health problem in more than 90 countries with combined population 3.3 billion people (50% of the world's population)
- ~220 million new infections and an estimated 660,000 deaths per year
- Mainly in Africa but incidence is growing in Asia and Latin America



### Why Malaria?

Emergence of drug resistance has compromised the therapeutic efficacy of most anti-malarial drugs

Artemisinin and derivatives, used in combination therapy, are currently the main line of defence against drug-resistant malaria



Humans can be infected by 11 species, but two account for vast majority of cases: Falciparum (~70%) & Vivax (~25%)



# PyfeDTPyCle A Novel Target



- Calcium-dependent protein kinases (CDPKs) are directly regulated by Ca<sup>2+</sup> and are found in plants and organisms in the alveolate lineage, but they are absent in humans
- PfCDPK1 is encoded by an essential gene expressed primarily in the merozoite



Blocking parasite invasion at the merozoite stage could interfere with the life cycle and potentially allow the infection to be cleared

# MRC

#### Hit Identification



- Hit series of Imidazopyridazines showed sub 100 nM IC<sub>50</sub> values against CDPK1 and strong inhibition of parasite growth
- Concerns over high log D values, poor microsome stability and selectivity over human kinases

Kinase Selectivity Panel:





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#### Optimisation of Imidazopyridazines



- A structure-based design approach using the homology model of *Pf*CDPK1 was used to attempt to gain increased binding affinity against the target and correspondingly increase the cellular potency
- The model suggested that the binding pocket occupied by the alkyl chain was not optimally filled

Virtual libraries were enumerated with a range of groups at this position and examined through docking





- Additional VDW interactions and fewer clashes
- Improved electrostatic complementarity and additional H-bond
- Reduced entropic penalty due to conformational focussing

# Synthesis of Imidazopyridazines





<5 steps – A prerequisite for MMV collaboration</p>

#### Testing the Model





 $EC_{50} < 9$  mm  $EC_{50} = 465$  nM

 $EC_{50} = 83 \text{ nM}$ 

 $IC_{50} < 8 \text{ nM}$  $EC_{50} = 12 \text{ nM}$ 

- Introduction of a hydrogen bond acceptor *ortho* to amine showed a significant cellular efficacy boost
- A further boost obtained by the introduction of a fluorine at the 3 position
- Poor permeability, could be due to diaminocyclohexane?

#### Optimisation of Left Hand Side

# MRC





 $IC_{50} = < 8 \text{ nM}$   $EC_{50} = 13 \text{ nM}$ Cytotox  $IC_{50} = >20 \mu M$ Log D = 0.2 MLM 85 % rem HLM 93 % rem Pampa = 3.9 nm/s  $IC_{50} = 13 \text{ nM}$   $EC_{50} = 40 \text{ nM}$ Cytotox  $IC_{50} = 5.9 \mu M$ Log D = 1.4 MLM 85 % rem HLM 98 % rem Pampa = 1.0 nm/s



 $IC_{50} = 44 \text{ nM}$   $EC_{50} = Inactive$ Cytotox  $IC_{50} = 6.7 \mu M$ Log D = 2.5 MLM 92 % rem HLM 82 % rem Pampa = 137 nm/s

- Variation of left hand side to improve ADME properties
- Basic side chain required on the left hand side
- Attempts to remove resulted in loss of cellular efficacy
- A range of basic groups with varying pKa explored (lower pKa improved permeability)
- Trade off between good ADME and good efficacy!

#### **Optimisation of Linker**









 $IC_{50} = 13 \text{ nM}$   $EC_{50} = 40 \text{ nM}$ Cytotox  $IC_{50} = 5.9 \mu \text{M}$ Log D = 1.4 MLM 85 % rem HLM 98 % rem Pampa = 1.0 nm/s  $IC_{50} = 9 \text{ nM}$   $EC_{50} = 388 \text{ nM}$ Cytotox  $IC_{50} = 0.5 \mu\text{M}$ Log D = 3.1 MLM 57 % rem HLM 77 % rem Pampa = 63 nm/s  $IC_{50} = 16 \text{ nM}$   $EC_{50} = 406 \text{ nM}$ Cytotox  $IC_{50} = 3.2 \mu\text{M}$ Log D = 3.2 MLM 47 % rem HLM 80 % rem Pampa = 170 nm/s

#### **Increasing Permeability**

**Decreasing Efficacy** 

# In Vivo Results



	IC50 (μM)	Cytotox v-50 (µM)	EC50 (µM)	HLM (% rem)	MLM (% rem)	LogD	PAMPA Papp (nm/s)	In vivo reduction in parasitaemia (%)
Z Z Z Z Z Z Z Z Z Z Z Z Z Z Z Z Z Z Z	0.013	9.0	0.400	63	85	3.4	81	46
	0.016	3.2	0.390	80	47	3.2	171	44
	0.012	2.0	0.078	86	61	3.7	48	51
H <sub>2</sub> N H H H H H H H H H H H H H	<0.008	>20	0.013	93	85	0.2	4	4

#### Parasite Rate of Reduction Assay



- Quantifies number of parasites that remain viable after drug treatment
- Parasites are treated with the selected drug at a concentration corresponding to 10 x IC<sub>50</sub> over 48 hours
- Dilutions are made and cultured for 21 days
- Samples taken at 21 and 28 days to confirm number of viable parasites
- Can group according to control compounds
  - Fast killers (chloroquine)
  - Moderate killers (pyrimethamine)
  - Slow killers (atovaquone)

#### Parasite Rate of Reduction Assay MRC 6 Compound A Controls 5 ▲ Compound B +1) +4 $\times$ Compound C log (viable parasites Slow/Moderate Chloroquine 3 Killers + Atovaquone 2 Pyrimethamine 1 - Untreated ŇΗ Fast Killer 0

#### **Biotinylation Experiments**



- Used to determine other targets that a compound may interact with
- Biotin is tethered to compound by a short linker and exposed to potential target proteins
- (Strept)Avidin has a high affinity for the biotin portion and can be used to "pull-out" the proteins for which the compound has an affinity.



**Biotinylation Experiments** 





- Assuming a similar binding mode biotin was tethered to the basic side chain
- Be However, a new target could mean a new binding mode

#### Conclusion



- A series of *Pf*CDPK1 inhibitors was identified from a high throughput screen
- Subsequent optimisation yielded potent, selective inhibitors with suitable ADME and pharmacokinetic profiles
- Testing in an *in vivo* efficacy model showed insufficient reduction in parasitaemia for the project to progress further
- On going investigation into off-target effects through biotinylation experiments

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Figure 5. Classical antimalarial killing rate profiles.



Sanz LM, Crespo B, De-Cózar C, Ding XC, et al. (2012) P. falciparum In Vitro Killing Rates Allow to Discriminate between Different Antimalarial Mode-of-Action. PLoS ONE 7(2): e30949. doi:10.1371/journal.pone.0030949 http://www.plosone.org/article/info:doi/10.1371/journal.pone.0030949

