

# Development of peptide-based antimalarial drugs

Nicole Lawrence

Institute for Molecular Bioscience The University of Queensland

n.lawrence@imb.uq.edu.au







## Malaria: a problem worth solving

~ three billion people from > 90 countries are at risk of contracting the disease

Each year > 200 million people are infected with deadly *Plasmodium falciparum* parasites, and > 400 thousand people die





## Malaria: the treatment arsenal needs expansion

- control measures including insecticides and use of bed nets have decreased the incidence in many regions
- resurgence of disease due to spread of drug resistant parasites is a major concern
- new classes of antimalarial drugs that act via different mechanisms compared to existing drugs are urgently needed





### Drug resistance (small molecule drugs)



Malaria parasites have developed resistance to every class of small molecule drug that has been developed

Blasco, Leroy, Fidock Nature Medicine (2017)



### Peptides as a new class of drugs – selective targeting inside cells









# Human defence molecules





Larger, multifunctional and organised into active domains



### Platelets can kill blood stage malaria parasites





### Platelet factor 4 (PF4)-derived peptides



#### Stable structure

Resistant to breakdown Kill blood stage malaria parasites Non-toxic to red blood cells





 ➡ PF4P<sub>57-70</sub>
➡ PF4PD 1.8  $\mathbf{t}_{1}(h)$ % Stability in human serum >>24 - cPF4PD 80 60 40 20 0 24 0 8 16 Time (h)



Lawrence et al. *Cell Chem Biol* (2018)



## Selective membrane-active mechanism of action



The positively charged peptide recognises the negatively charged surface of infected red blood cells It crosses the host membrane without damage, enters the parasite and disrupts the digestive vacuole (DV) The peptide does not enter or harm uninfected cells



### Improving potency and drug-like properties of the peptide





1. charged and/or hydrophobic residues

We are working to improve the antimalarial activity of the peptide while maintaining stable structure and non-toxic to healthy cells

#### ACKNOWLEDGEMENTS



#### Funding and research infrastructure



AUSTRALIAN RESEARCH COUNCIL CENTRE OF EXCELLENCE FOR INNOVATIONS IN PEPTIDE AND PROTEIN SCIENCE







#### **Scientific collaborators**

University of Queensland Institute for Molecular Bioscience

David Craik



Australian National University Research School of Chemistry

Lara Malins



John Curtin School of Medical Research

Brendan McMorran



Australian Defence Force Malaria and Infectious Disease Institute

Mike Edstein

