

## Haemolytic peptides directed against the malarial blood stages of *Plasmodium falciparum*

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Malaria affects more than 300 million people in third world countries per year, causing the annual death of at least one million children in Africa. Of particular concern is malaria caused by *Plasmodium falciparum*, of which a large number of strains are drug resistant [1]. The search for new drugs led us to investigate a number of peptides with known antimicrobial and haemolytic activity. Bilayer membranes are the primary targets of these peptides and their interaction with the cell membrane induces a permeability change followed by cell lysis [2]. Such a mechanism of action makes it difficult for organisms to develop resistance, but this non-selectivity is also highly problematic for systemic routes of drug administration. However, we found that the haemolytic activity is beneficial in killing the blood stages of the malaria parasite *P. falciparum*. Erythrocytes infected with the trophozoite stage were particularly vulnerable to lysis.

It is known the malaria parasite alters the protein [3] and lipid composition [4] of the host erythrocyte membrane, which makes the infected cell a possible target. We found that that three of the haemolytic peptides were at least ten times more active against the trophozoite blood stage of the parasite (IC<sub>50</sub> of 0.5-5 mM), than against uninfected erythrocytes. However, the highly haemolytic peptide from bee-venom, melittin, showed only a three times higher specificity. Also, two highly antimicrobial peptides, but non-haemolytic peptides from *Xenopus laevis*, were virtually inactive. We therefore conclude some haemolytic peptides may be very promising lead compounds for designing a magic bullet against *P. falciparum* infected erythrocytes.

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