

# A Systems Biology approach to model the Glutathione pathway in *Plasmodium falciparum*.

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## Background

- Malaria morbidity and mortality is a severe burden, no where is this more apparent than in sub-Saharan Africa (SSA).
- There is great efforts to control Malaria in SSA but this faces great challenges notably to drug resistance, especially to classical and affordable antimalarials, such as Chloroquine (CQ).
- Glutathione (GSH) has its metabolism greatly associated with the mode of action of CQ.
- This mathematical model is a series of biochemical reactions of this pathway in terms of its biosynthesis, reduction, consumption, and efflux over a specified period of time.

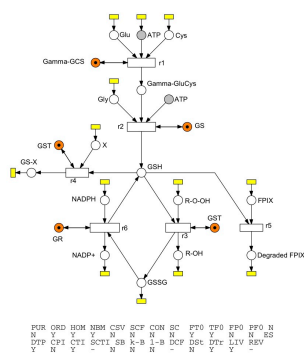


## Objectives

- Broad:
  - To construct a mathematical model that elucidates the metabolism of the GSH pathway.
- Specific:
  - To collect kinetic data of metabolites involved in GSH metabolism from published literature.
  - To construct and validate of a Petri net (PN) model of the GSH pathway using the software Snoopy and Charlie respectively.
  - To construct an ordinary differential equation (ODE) based model using MATLAB and its annotation using the software CellDesigner.
  - To subject the ODE based model to analysis by performing inhibition studies.

## Methods

### PN model construction & Validation



### ODE based model with MATLAB

• Biosynthesis:  

$$V_{\gamma\text{-GCS}} = \frac{V_{\max}[\text{Glu}][\text{Cys}][\text{ATP}]}{(K_m + [\text{Glu}])(K_m + [\text{Cys}](K_m + [\text{ATP}]})}$$

$$V_{\text{CS}} = \frac{V_{\max}[\gamma\text{-GluCys}][\text{Clv}][\text{ATP}]}{(K_m + [\text{GluCys}](K_m + [\text{Gly}](K_m + [\text{ATP}]}))}$$
 • Reduction:  

$$V_{\text{GR}} = \frac{V_{\max}[\text{GSSG}][\text{NADPH}]}{(K_m + [\text{NADPH}] + (K_m + [\text{GSSG}]})}$$

$$V_{\text{NADP}^+} = K[\text{NADP}^+]$$
 • Consumption:  

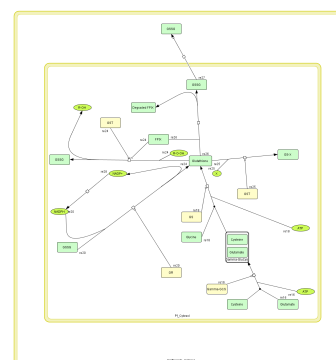
$$V_{\text{GST}} = \frac{V_{\max}[\text{X}][\text{GSH}]}{(K_m + [\text{X}])(K_m + [\text{GSH}])}$$

$$V_{\text{GST}} = \frac{V_{\max}[\text{GSH}][\text{R-O-OH}]}{(K_m + [\text{GSH}](K_m + [\text{R-O-OH}]))}$$

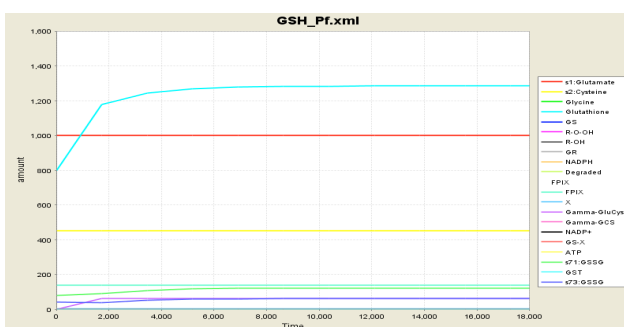
$$V_{\text{FPX}} = k[\text{FPX}][\text{GSH}]$$
 • Excretion/Efflux:  

$$V_{\text{GSSG Efflux}} = \frac{V_{\text{GSSG}}/K_{\text{mGSSGPF}}}{1 + \frac{V_{\text{GSSG}}/K_{\text{mGSSGPF}}}{K_{\text{mGSSGRBC}}} + \frac{V_{\text{GSSG}}/K_{\text{mGSSGRBC}}}{K_{\text{mGSSGPF}}}}$$

### Annotation of the ODE based model using CellDesigner



## Results



- Inhibition studies (50  $\mu\text{M}$  of Methylene Blue (MB)):
  - Concentration of GSH increased by 4-folds (1290 to 350 $\mu\text{M}$ ).
  - Concentration of oxidized Glutathione (GSSG) in Pf increased by 200 folds (122 to 28,234 $\mu\text{M}$ ).
  - Concentration of GSSG (in RBC) increased by over 100 folds (61 to 7671 $\mu\text{M}$ ).

## Discussion & Conclusion

- 50  $\mu\text{M}$  of MB affects the GSH/GSSG ratio to such levels that are deleterious to *Plasmodium falciparum*.
- The reduction reaction of the GSH pathway/metabolism has the potential for the development of a drug target.
- MB has effective antimalarial activity and hence could be developed into an antimalarial drug.
- More research has to be conducted on this, and it should focus on how MB can be carefully administered so that it only affects *Plasmodium falciparum* and not its human host.