Modelling RNA Oscillatory Circuits

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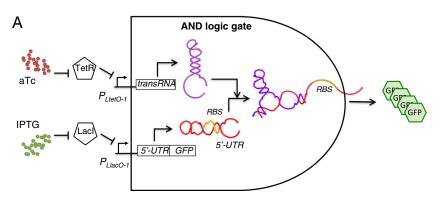
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Outline

- Description of RNA Oscillatory system
 - Recent Experimental data
 - ODE Model for system
- Parameter Estimation in ODE Model
 - Methodology least squares error minimisation
 - Results Inestimable parameters, model simplifications
 - Future Work

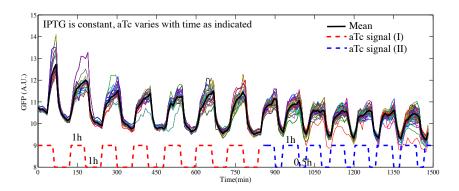
The RNA Oscillatory System



- mRNA produced 'self repressed' tail folded over the RBS
- sRNA binds to it new complex has RBS uncovered.



Recent Experimental data



• Time series of individual cell fluorescence, forced with aTc.



ODE Model for system

$$\frac{ds}{dt} = \frac{N\alpha_T}{f_T}y(t) - (\mu + \delta_s)s - k_{\text{on}}sm + k_{\text{off}}s : m$$

$$\frac{dm}{dt} = \frac{N\alpha_L}{f_L}x(t) - (\mu + \delta_m)m - k_{\text{on}}sm + k_{\text{off}}s : m$$

$$\frac{ds : m}{dt} = k_{\text{on}}sm - (k_{\text{off}} + k_{\text{hyb}})s : m - (\mu + \delta_{sm})s : m$$

$$\frac{dc}{dt} = k_{\text{hyb}}s : m - (\mu + \delta_c)c$$

 Hybridization of sRNA(s) and mRNA (m), into unstable complex (s: m) then stable one (c).

ODE Model for system

$$\frac{dp}{dt} = \beta m + f_s \beta c - (\gamma + \mu + \delta_g) p - \frac{v_z p}{K_z + p + g}$$

$$\frac{dg}{dt} = \gamma p - (\mu + \delta_g) g - \frac{v_z g}{K_z + p + g}$$

$$z = z_0 + \frac{g}{\Theta}$$

- Translation of stable complex (c) into immature GFP (p).
- maturation of GFP (g), machine calibration giving measured fluorescence (z).

Parameter Estimation in ODE Model - Methodology

Goal

Estimate unknown parameters in model using time series data.

Least squares error minimization approach

$$\arg\min_{\theta} \sum_{i=1}^{n} (z_{\exp,\text{mean}}(t_i) - z(t_i, \theta))^2.$$

Used evolutionary algorithm, CMA-ES, to perform minimisation.

Parameters to be estimated in our model

$$\frac{ds}{dt} = \frac{N\alpha_T}{f_T}y(t) - (\mu + \delta_s)s - k_{on}sm + k_{off}s: m$$

$$\frac{dm}{dt} = \frac{N\alpha_L}{f_L}x(t) - (\mu + \delta_m)m - k_{on}sm + k_{off}s: m$$

$$\frac{ds:m}{dt} = k_{on}sm - (k_{off} + k_{hyb})s: m - (\mu + \delta_m)s: m$$

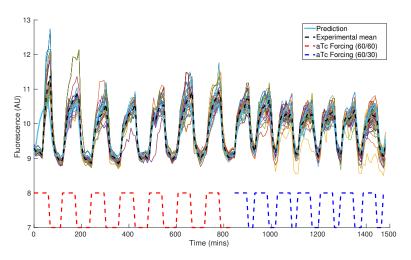
$$\frac{dc}{dt} = k_{hyb}s: m - (\mu + \delta_m)c$$

$$\frac{dp}{dt} = \beta m + f_s\beta c - (\gamma + \mu + \delta_g)p - \frac{v_zp}{K_z + p + g}$$

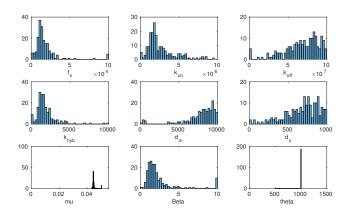
$$\frac{dg}{dt} = \gamma p - (\mu + \delta_g)g - \frac{v_zg}{K_z + p + g}$$

$$z = z_0 + \frac{g}{\Theta}$$

Initial Minimization Results



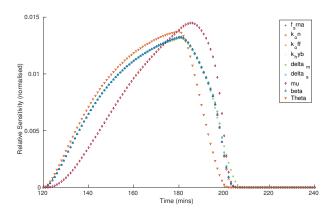
Initial Minimization Results



Many parameters poorly estimated.

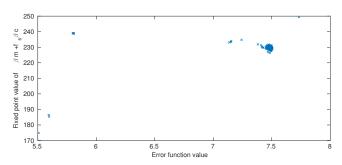


Sensitivity Analysis



 Sensitivity analysis suggests perturbations of several parameters all give similar effects on model output - hard to resolve

Sensitivity Analysis



- Similar fixed point values of translation forcing term, $\beta m + f_s \beta c$, across estimated parameter sets.
- Suggests translation may be rate limiting step.



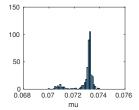
Simplified Model

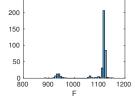
$$egin{aligned} rac{dp}{dt} &= \emph{F}y(t) - (\gamma + \mu + \delta_g)p - rac{\emph{v}_{\emph{Z}}\emph{p}}{\emph{K}_{\emph{Z}} + \emph{p} + \emph{g}} \ rac{dg}{dt} &= \gamma \emph{p} - (\mu + \delta_g)g - rac{\emph{v}_{\emph{Z}}\emph{g}}{\emph{K}_{\emph{Z}} + \emph{p} + \emph{g}} \ z &= \emph{z}_0 + rac{\emph{g}}{\emph{G}} \end{aligned}$$

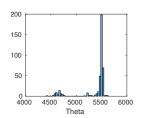
- Model only rate limiting steps force translation term directly.
- 3 unknown parameters



Simplified Model Results



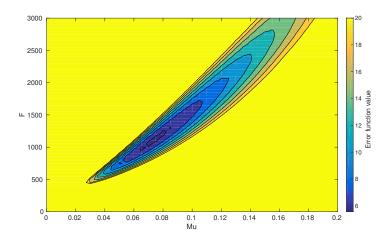




- Error values as low as full model
- Clearer parameter estimation results
- Two peak structure due to local minima

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Simplified Model Results



• Error landscape, $\Theta = 1000$.



Future Work

- Fluorescence data not enough to estimate all unknown parameters
 - Further experiments?
 - Bounds on unknown parameter values.
- Improve modelling of translation step
- Change methodology Bayesian Methods?