Notes on modeling of growth-regulation interactions in the cAMP system

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Chapter 1

Modeling the noise transmission

1.1 Ornstein-Uhlenbeck fluctuations model

1.1.1 The model

In these notes I discuss a model which models how noise originates and transmits from and between observables respectively. I will discuss the model proposed by Philippe Nghe in the work by Kiviet et al. [3], and also possible extensions and modifications to this model.

Parameters that are considered in general are the number of enzymes (E), the rate at which these enzymes are made (p), and the growth rate of the cell (λ) . Generally, enzymes only disappear by dilution due to growth. Furthermore, there are noise sources, which add noise to these parameters. Depending on how they are implemented these source appear as N_M , N_λ and N_p or Γ_M , Γ_λ and Γ_p , the subscripts indicate to which parameter the noise source adds noise.

The goal of this model is to interpret the cross-correlations between λ , E and p (usually $R_{E,\lambda}(\tau)$, $R_{p,\lambda}(\tau)$), that are obtained from experimental data.

1.1.2 Implicit noise equations

A straight forward way to model a system with these parameters, is writing down an ordinary differential equation (ODE) for each parameter involved. The Nghe model takes a somewhat different approach (it is more concise), which will be discussed later. The ODEs below relate to the cartoon in Fig. 1.1.C and describe the dynamics per parameter:

$$\dot{M} = -\frac{(M - M_0)}{\tau} + c_M \cdot \Gamma_M + T_{M \leftarrow E} \cdot c_M \cdot (\frac{E}{E_0} - 1)$$

$$(1.1)$$

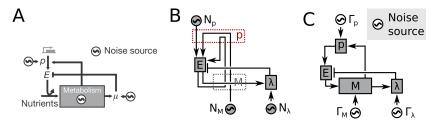


Figure 1.1: (A) Original diagram as printed in Kiviet et al. [3]. (B) More technical diagram stating the relations between parameters of interest (gray boxes), and noise sources N_P , N_M , and N_λ . Arrows indicate how ODEs or functions that describe the parameters of interest are coupled. Arrows that go trough the red dashed box correspond to terms that not only couple the two parameters connected by the arrow, but also make up the parameter p (see also main text). Arrows that go through the grey box are arrows which are thought to be biologically connected to the metabolism. (C) An alternative version of the model with more parameters modelled explicitly. Noise sources are printed white in this diagram since they are not described by separate ODEs, which was the case in Kiviet et al. [3].

$$\dot{\lambda} = -\frac{(\lambda - \lambda_0)}{\tau_{\lambda}} + c_{\lambda} \cdot \Gamma_{\lambda} + T_{\lambda \leftarrow M} \cdot c_{\lambda} \cdot (\frac{M}{M_0} - 1)$$
(1.2)

$$\dot{P} = -\frac{(P - P_0)}{\tau_P} + c_\lambda \cdot \Gamma_\lambda + T_{P \leftarrow M} \cdot c_P \cdot (\frac{M}{M_0} - 1) + R_{P \leftarrow M} \cdot c_P \cdot (\frac{M}{M_0} - 1)$$

$$(1.3)$$

$$\dot{E} = P - \lambda E \tag{1.4}$$

Where M describes the state of the metabolism, λ is the growth rate, P is the production rate, E is the amount of enzyme, τ is a dampening term (X_0 is the equilibrium value), $T_{X \leftarrow Y}$ is the noise transmission constant from X towards Y, c_X is a constant that sets the size of the fluctuations, Γ_X is a white noise source. $R_{X \leftarrow Y}$ indicates a regulatory interaction, an addition to the model by me, but this notation is currently just cosmetical, as $T_{\text{effective}} = T + R$. This model assumes all parameters have an average value from which fluctuations deviate, but always return. Hence the dampening terms. With respect to transmission, I furthermore rescale the absolute value of the noise to be comparable to the target noise (hence the M_0^{-1} and c_X terms in combination with the $T_{X \leftarrow Y}$ term).

This is similar to the model that Philipe Nghe had in Kiviet et al. [3], which was inspired by Dunlop et al. [1] (see supplement of that manuscript for a description of the Dunlop model). A difference between my equations, the Nghe equations and the Dunlop equations lies in the dampening terms (those containing τ , β or μ_E). In my model noise is effected through the ODE, and dampening occurs on the parameter of interest. In Dunlop et al., there are two dampening terms, one specifically dampening the noise and a second term dampening the parameters of interest. It seems also Nghe takes the latter approach, dampening being effected through the $(-\beta_X \cdot N_X)$ term and the μ_E parameter (see later for a more involved discussion on these parameters).

The correlations between these equations can be found by linearizing them, writing the correlations in Fourier space, and back-transforming them using residue integration techniques. This document currently does not fully explores this, but this will partially be discussed later. First, a comparison with the Nghe model is made.

1.2 Separate noise equations

Both Nghe and Dunlop define separate ODEs for the noise terms:

$$\dot{N}_X = \sqrt{C_X} \cdot \Gamma_X - N_X / \tau, \tag{1.5}$$

though their notation might be slightly different (I used Daniel Gillespie's notation [2]; a capital C is used here to follow Gillespie's square root notation, $\sqrt{C_X} = c_x$). With for our case X equaling λ , M or P. Note that $\tau^{-1} = \beta$ (β is used in Nghe and Dunlop).

Not so relevant for our case, but noteworthy, is that in the Dunlop model, which models a completely different process than the one described here [1], the *solutions* of the ODEs describing the noise are plugged into the ODEs describing the protein dynamics. This leads to an additional memory effect. That is:

$$\dot{X} = N_X + F(X) + X/\tau, \tag{1.6}$$

with F(X) some arbitrary function of X. Note that the N_X function also contains a τ term (see Eq. 1.5), which is effectively integrated, thus leading to effects of the fluctuations much longer timescales than τ . This effect is (partially) countered by the third term in Eq. 1.6, which also contains the τ term.

1.3 Nghe model

As mentioned, the Nghe model takes a different approach. The formulae that follow correspond to Fig. 1.1.B (Fig. 1.1.A contains a more polished version of that cartoon that was published in Kiviet et al.). The starting point,

$$\dot{E} = P - \lambda E,\tag{1.7}$$

is the same in my model, but after linearization (defined as $X = X_0 + \delta X$) this leads to only one ODE (see written notes from 28.9.2016 for explicit linearization):

$$\frac{\delta \dot{E}}{E_0 \mu_0} + \frac{\delta E}{E_0} = \frac{\delta p}{E_0 \mu_0} + T_{E \leftarrow \mu} \frac{\delta \mu}{\mu_0}. \tag{1.8}$$

Noise terms are introduced with ODEs that are also shown above in Eq. 1.5. Additionally, two functions are defined for p and λ . These are not ODEs, as the effects on these parameters are thought to happen on fast timescales. The parameters are however linearized (and thus written as δX). The equation

$$\frac{\delta\mu}{\mu_0} = T_{\mu\leftarrow E} \frac{\delta E}{E_0} + T_{\mu\leftarrow G} N_G + N_\mu \tag{1.9}$$

simply defines the evolution of $\delta\mu$. There is also an equation for δp

$$\frac{\delta p}{E_0 \mu_0} = T_{E \leftarrow E} \frac{\delta E}{E_0} + T_{E \leftarrow G} N_G + N_E, \tag{1.10}$$

which plays a bit more complicated role. It is defined using terms that pertain to E, like $T_{E\leftarrow G}$, such that it can be directly plugged into Eq. 1.8. Indeed, plugging Eq. 1.10 into Eq. 1.8 leads to equation:

$$\frac{\delta \dot{E}}{E_0 \mu_0} + \frac{\delta E}{E_0} = \left[T_{E \leftarrow E} \frac{\delta E}{E_0} + T_{E \leftarrow G} N_G + N_E \right] + T_{E \leftarrow \mu} \frac{\delta \mu}{\mu_0} \tag{1.11}$$

which corresponds to equation 5 in the Kiviet et al. [3] manuscript. I say it is a bit complicated, since Eq. 1.10 has no role in the model (we could also just have defined Eq. 1.11 immediately), except that it shows us which part of the model can be interpreted as being the production rate. This is also the reason why p is depicted as a red dashed box in Fig. 1.1.

Note however, that this results in the absence of dampening terms on the parameters X themselves, which leads to non-steady state behavior of the parameters. The term X/τ could be added to each of the equations to resolve this issue; the term $-N_X/\tau$ from the noise ODEs could then be dropped. Note that a μ_E term appears eventually in Nghe's equations, which might play the role of a second dampening term.

1.4 Obtaining the cross correlations

TODO: Process below such that it correctly handles Philippe's and my model (text above has been updated after phone call w. PN). Discuss w. ST/others whether and how to expand model.

1.4.1 Obtaining solutions in Fourier space

How do we get the cross correlations? A way to define correlations is 1:

$$R_{f,g}(\tau) = f \star g = \int_{\tau = \infty}^{\infty} \bar{f}(\tau)g(t+\tau)\delta\tau \tag{1.12}$$

With the \bar{f} denoting the complex conjugate. This is equal to the convolution of $f^*(-t)$ and g(t),

 $^{^1{\}rm From}$ Wikipedia, https://en.wikipedia.org/wiki/Cross-correlation and Weisstein, Eric W. "Cross-Correlation Theorem." From MathWorld–A Wolfram Web Resource. http://mathworld.wolfram.com/Cross-CorrelationTheorem.html

$$f \star g = f * (-t) * g.$$
 (1.13)

It can be derived that:

$$\mathcal{F}(f \star g) = \overline{\mathcal{F}(f)}\mathcal{F}(g). \tag{1.14}$$

This property can be exploited to find the cross correlation that we are interested in:

$$R_{\mu,E}(\tau) = \mathcal{F}^{-1}\left(\overline{\mathcal{F}(\mu)}\mathcal{F}(E)\right). \tag{1.15}$$

First we find the solutions to the ODEs in Fourier space:

$$\mathcal{F}(\dot{E}) = \mathcal{F}(P - \lambda E)$$
$$i\omega \tilde{E} = \tilde{P} - \tilde{\lambda}\tilde{E}$$
(1.16)

$$\tilde{E} = \frac{1}{i\omega + \tilde{\lambda}} \tilde{P} \tag{1.17}$$

For Eq. 1.5 the solution in Fourier space is:

$$\tilde{N}_X = \frac{\sqrt{c_x}}{i\omega - 1/\tau} \tilde{\Gamma}_x \tag{1.18}$$

which however will not be directly used to solve Eq. 1.1-1.4. With regard to Eq. 1.1-1.3, the solutions in general terms can be found from taking the Fourier transform:

$$i\omega\tilde{X} = -\frac{\tilde{X} - X_0}{\tau_X} + c_X\tilde{\Gamma}_X + T_{X \leftarrow Y}c_X(\frac{\tilde{Y}}{X_0} - 1)$$
(1.19)

which, solving for X gives:

$$\tilde{X} = \left(i\omega + \frac{1}{\tau_X}\right)^{-1} \left(\frac{X_0}{\tau_X} + c_X \tilde{\Gamma}_X + T_{X \leftarrow Y} c_X (\frac{\tilde{Y}}{Y_0} - 1)\right),\tag{1.20}$$

X being either λ , M or P. The solution for E in Fourier space (i.e. to Eq. 1.4) without P or λ terms is very involved. This can be seen by plugging Eq. 1.20 into Eq. 1.16, which results in:

$$i\omega \tilde{E} = \left(i\omega + \frac{1}{\tau_P}\right)^{-1} \left(\frac{P_0}{\tau_P} + c_P \tilde{\Gamma}_P + T_{P \leftarrow M} c_P \left(\frac{\tilde{M}}{M_0} - 1\right)\right) - \left(i\omega + \frac{1}{\tau_\lambda}\right)^{-1} \left(\frac{\lambda_0}{\tau_\lambda} + c_\lambda \tilde{\Gamma}_\lambda + T_{\lambda \leftarrow M} c_\lambda \left(\frac{\tilde{M}}{M_0} - 1\right)\right) \tilde{E}$$
(1.21)

Where M is defined as:

$$\tilde{M} = \left(i\omega + \frac{1}{\tau_M}\right)^{-1} \left(\frac{M_0}{\tau_M} + c_M \tilde{\Gamma}_M + T_{M \leftarrow E} c_M \left(\frac{\tilde{E}}{E_0} - 1\right)\right)$$
(1.22)

Plugging Eq. 1.22 into Eq. 1.21 results in an equation where the only time-dependent parameter is E. Solving that formula for E gives a very involved formula, which is not displayed here.

1.4.2 Actually getting the cross-correlations

As Dunlop et al. point out, we are now looking at the expectation values of the correlation functions (since we're talking about noise), so in general terms:

$$\langle R_{X,Y}(\tau) \rangle = \left\langle \mathcal{F}^{-1} \left[\overline{\mathcal{F}(X)} \mathcal{F}(Y) \right] \right\rangle$$
 (1.23)

1.5 Notes

Note that Philippe Nghe's scripts can be found at:

Bibliography

- [1] Mary J Dunlop, Robert Sidney Cox, Joseph H Levine, Richard M Murray, and Michael B Elowitz. Regulatory activity revealed by dynamic correlations in gene expression noise. *Nature genetics*, 40(12):1493–8, dec 2008.
- [2] Dt Gillespie. Exact numerical simulation of the Ornstein-Uhlenbeck process and its integral. *Physical review. E, Statistical physics, plasmas, fluids, and related interdisciplinary topics*, 54(2):2084–2091, aug 1996.
- [3] Daniel J. Kiviet, Philippe Nghe, Noreen Walker, Sarah Boulineau, Vanda Sunderlikova, and Sander J. Tans. Stochasticity of metabolism and growth at the single-cell level. *Nature*, sep 2014.