

Supplementary Materials

Parameterising Translational Feedback Models of Autoregulatory RNA-Binding Proteins in *Saccharomyces cerevisiae*

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Text S1: Calculation of best-case predicted feedback dynamics.

Using the equations when $n = 1$ (negative feedback):

$$\left\{ \begin{array}{l} \frac{dm}{dt} = k_1 - k_2 m \\ \frac{dP}{dt} = \frac{k_3 m}{1 + K_a P} - k_4 P \end{array} \right. \quad (1)$$

At the steady state:

$$\left\{ \begin{array}{l} \frac{dm_{ss}}{dt} = 0 \\ \frac{dP_{ss}}{dt} = 0 \end{array} \right. \quad (2)$$

$$\left\{ \begin{array}{l} m_{ss} = \frac{k_1}{k_2} \\ k_3 m_{ss} - k_4 P_{ss} - K_a k_4 P_{ss}^2 = 0 \end{array} \right. \quad (3)$$

$$K_a = \frac{k_3 m_{ss} - k_4 P_{ss}}{k_4 P_{ss}^2} \quad (4)$$

And likewise, for positive feedback when $n = 1$:

$$\left\{ \begin{array}{l} \frac{dm}{dt} = k_1 - k_2 m \\ \frac{dP}{dt} = \frac{k_3 m P}{K_d + P} - k_4 P \end{array} \right. \quad (5)$$

So that again,

$$\left\{ \begin{array}{l} m_{ss} = \frac{k_1}{k_2} \\ k_3 m P - K_d k_4 P_{ss} + k_4 P_{ss}^2 = 0 \end{array} \right. \quad (6)$$

$$\left\{ \begin{array}{l} m_{ss} = \frac{k_1}{k_2} \\ K_d = \frac{k_3 m P - k_4 P_{ss}^2}{k_4 P_{ss}} \end{array} \right. \quad (7)$$