

Steady-State Analysis of a Reversible Covalent Modification Cycle Jeremy A. Owen and Jeremy Gunawardena

ABSTRACT

The properties of biological switches are shaped by competing selection pressures. Switches must be sensitive to their inputs, and their "on" and "off" states should be well-distinguished. At the same time, it is important for switches to exhibit a certain robustness to noise. A simple biological network motif that gives rise to switch-like behaviour is the Goldbeter-Koshland (GK) loop, in which the interconversion of two substrate forms is catalysed by two biased enzymes. In the context of the GK loop, it is possible to conduct a simple algebraic analysis to reveal that the degree of bias of each enzyme is the primary determinant of how switch-like the system is. The same analysis suggests a way that a switch might make itself robust to changes in enzyme concentrations—a trick that may underlie the importance of enzyme bifunctionality.

REACTION NETWORK

In the GK loop, two enzymes, *E* and *F*, catalyse the interconversion of substrates S_0 and S_1 . We can summarize the salient features of each branch of the loop using a few numbers that describe how reversible the enzyme in question is and to what extent product rebinding occurs.

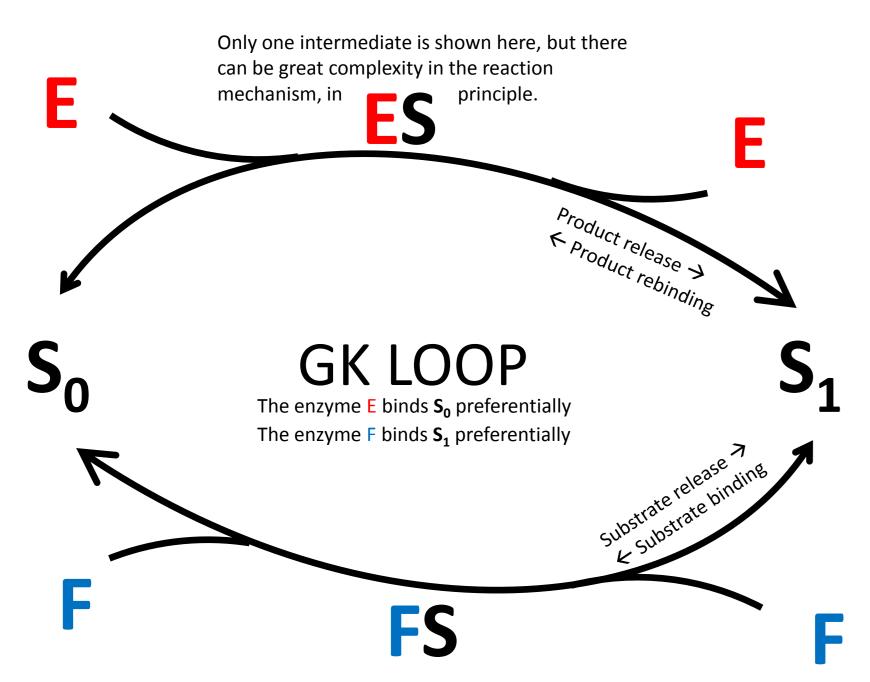
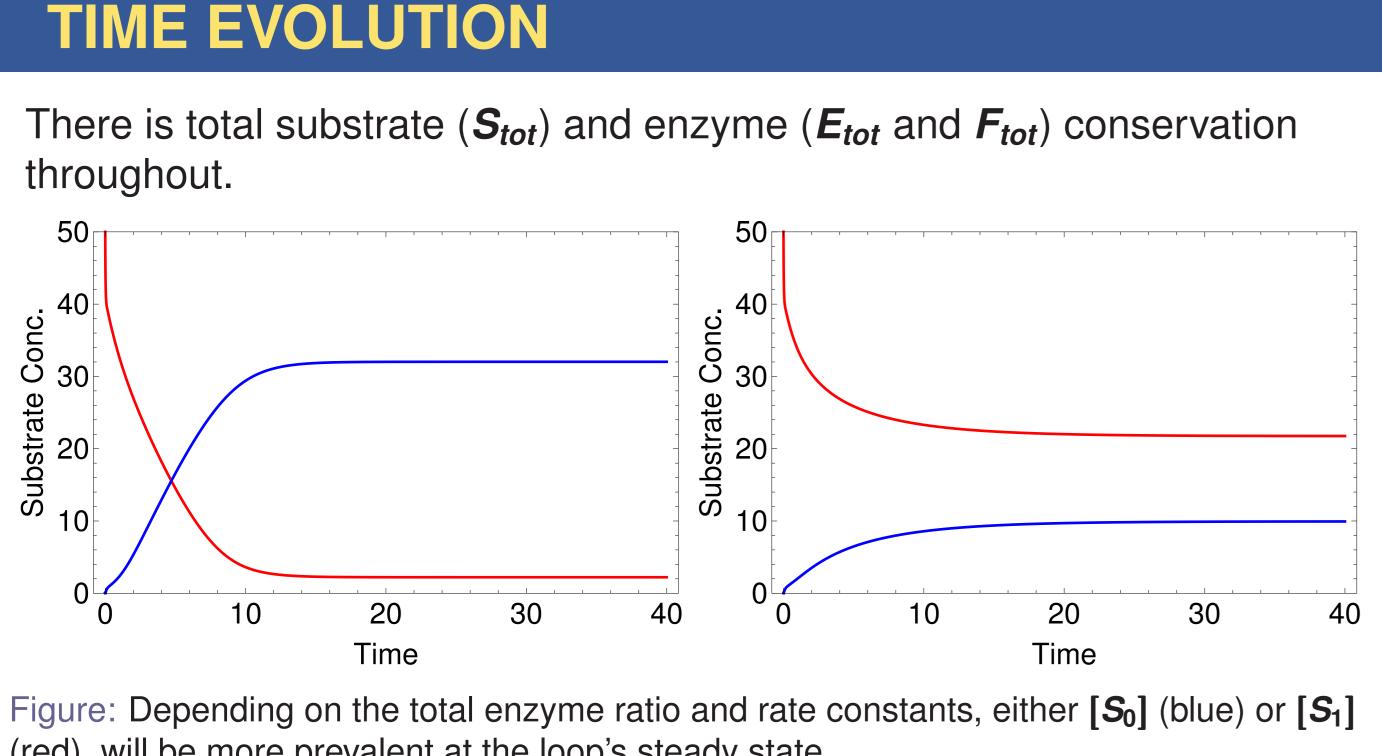


Figure: Schematic of a GK loop with a single intermediate form for each enzyme.

This work uses a mathematical framework laid out in: Y. Xu, J. Gunawardena, J. Theor. Biol. (2012).



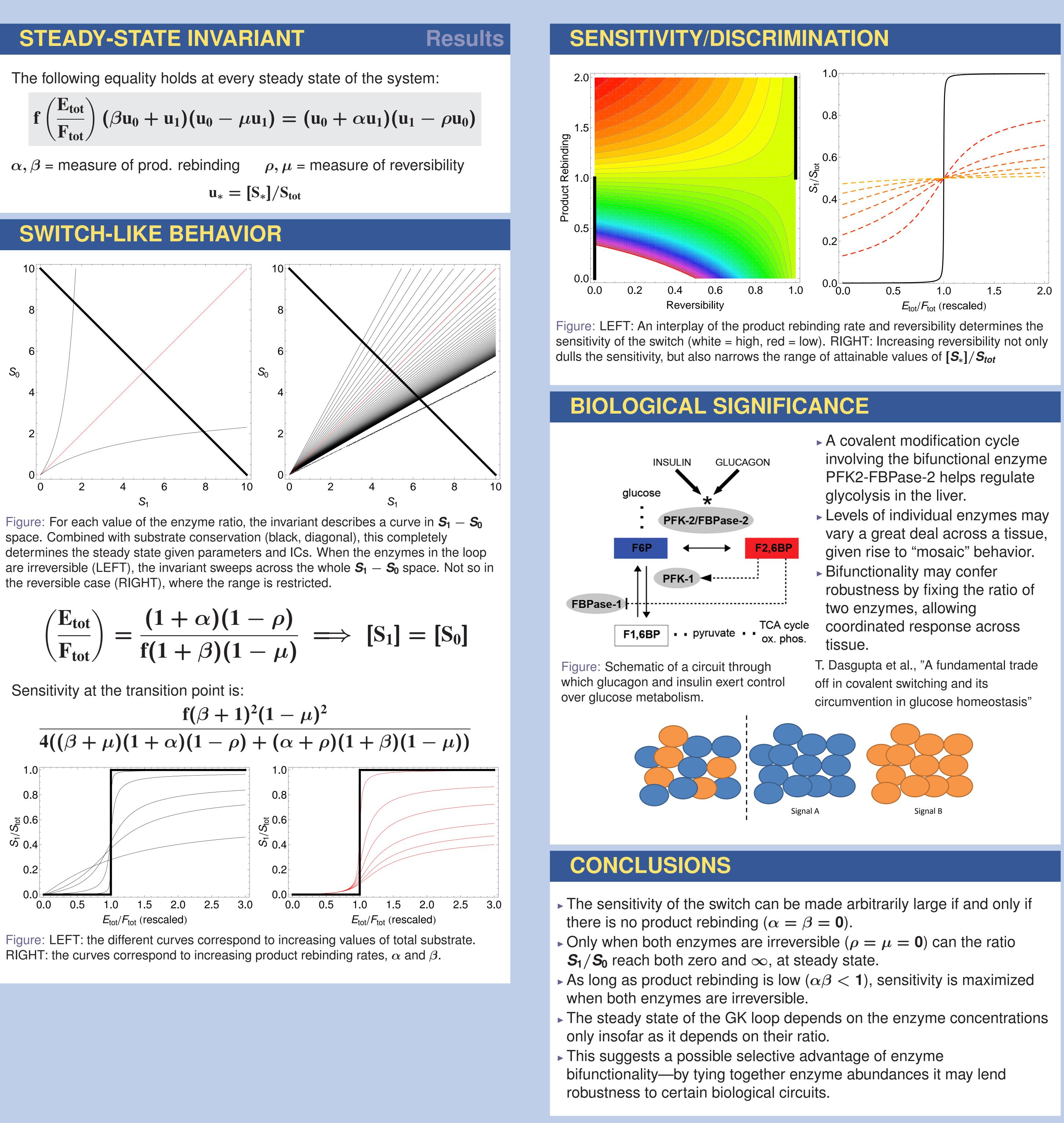
(red), will be more prevalent at the loop's steady state.

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Background

$$f\left(\frac{E_{tot}}{F_{tot}}\right)\left(\beta u_0 + u_1\right)\left(u_0 - \mu u_1\right) =$$

 α, β = measure of prod. rebinding



$$\begin{pmatrix} E_{tot} \\ F_{tot} \end{pmatrix} = \frac{(1+\alpha)(1-\rho)}{f(1+\beta)(1-\mu)}$$

