Text S2

Available models for signaling cascades using one variable per unit

In the main text we have briefly reviewed some approaches followed in the literature to study signaling cascades by means of a single equation per cycle. We add here two more descriptions in the same line and summarize these models with one variable per unit in Table S2.1.

Model	Equations
Weakly	$\dot{y}_i^* = \alpha_i y_{i-1}^* - \beta_i y_i^*$
activated	
pathway	[Heinrich et al. (2002)]
Linear rates	$\dot{y}_i^* = \alpha_i y_{i-1}^* y_i - \beta_i y_i^*$
	with $y_i + y_i^* = 1$
	[Heinrich et al. (2002)]
GK-like (I)	$\dot{y}_{i}^{*} = V_{i}y_{i-1}^{*}rac{y_{i}}{K_{i}+y_{i}} - V_{i}^{\prime}rac{y_{i}^{*}}{K_{i}^{\prime}+y_{i}^{*}}$
	with $y_i + y_i^* = 1$
	[Goldbeter and Koshland (1981)]
GK-like (II)	$\dot{y}_i^* = V_i \frac{y_{i-1}^*}{K_{ai} + y_{i-1}^*} \frac{y_i}{K_i + y_i} - V_i' \frac{y_i^*}{K_i' + y_i^*}$
	with $y_i + y_i^* = 1$
	[Goldbeter (1991)]
	[Gonze and Goldbeter (2000)]
Reduced	$\dot{x}_i = V_i x_{i-1} \frac{y_i}{K_i + y_i} - V_i' \frac{x_i}{K_i'(1 + \frac{y_{i+1}}{K_{i+1}}) + x_i}$
mechanistic	with $x_i + y_i + \eta_i x_{i-1} \frac{y_i^{++}}{K_i + y_i} = 1$
	and $y_i^* = x_i \frac{K_{i+1}}{K_{i+1} + y_{i+1}}$
	(this paper)

Table S2. 1: Models for signaling cascades.

In the work of Heinrich et al [Heinrich et al. (2002)], where the linear rates model was studied, the authors also discuss the limiting case called "the weakly activated pathway", where only a small fraction of the targeted protein is activated: $y_i = 1 - y_i^* \simeq 1$. Then, the system given by Eq. (2) in the main text is turned into a system of linear equations, which have been studied extensively in different contexts [Heinrich et al. (2002), Chaves et al. (2004)]:

$$\dot{y}_i^* = \alpha_i y_{i-1}^* - \beta_i y_i^*, \qquad i = 1, \cdots, n,$$
(14)

It corresponds to the simplest possible choice, where the kinase and phosphatase reactions are assumed to follow linear rate laws with first-order rate constant, α_i and β_i , respectively. Besides that one described in the main text, there is another phenomenological extension of the GK model for one cycle that has been proposed in the following form [Goldbeter (1991), Gonze and Goldbeter (2000)]:

$$\dot{y}_{i}^{*} = V_{i} \frac{y_{i-1}^{*}}{K_{ai} + y_{i-1}^{*}} \frac{y_{i}}{K_{i} + y_{i}} - V_{i}^{\prime} \frac{y_{i}^{*}}{K_{i}^{\prime} + y_{i}^{*}}, \qquad i = 1, \cdots, n.$$
(15)

In these equations, the effective maximum rate of the kinase produced at step i - 1 is proportional to the fraction of active enzyme.

References

- [Heinrich et al. (2002)] Heinrich R, Neel BG, Rapoport TA (2002) Mathematical models of protein kinase signal transduction. Mol Cell 9:957-970.
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