

Supply and demand in Metabolic Control Theory

Daniel Kahn

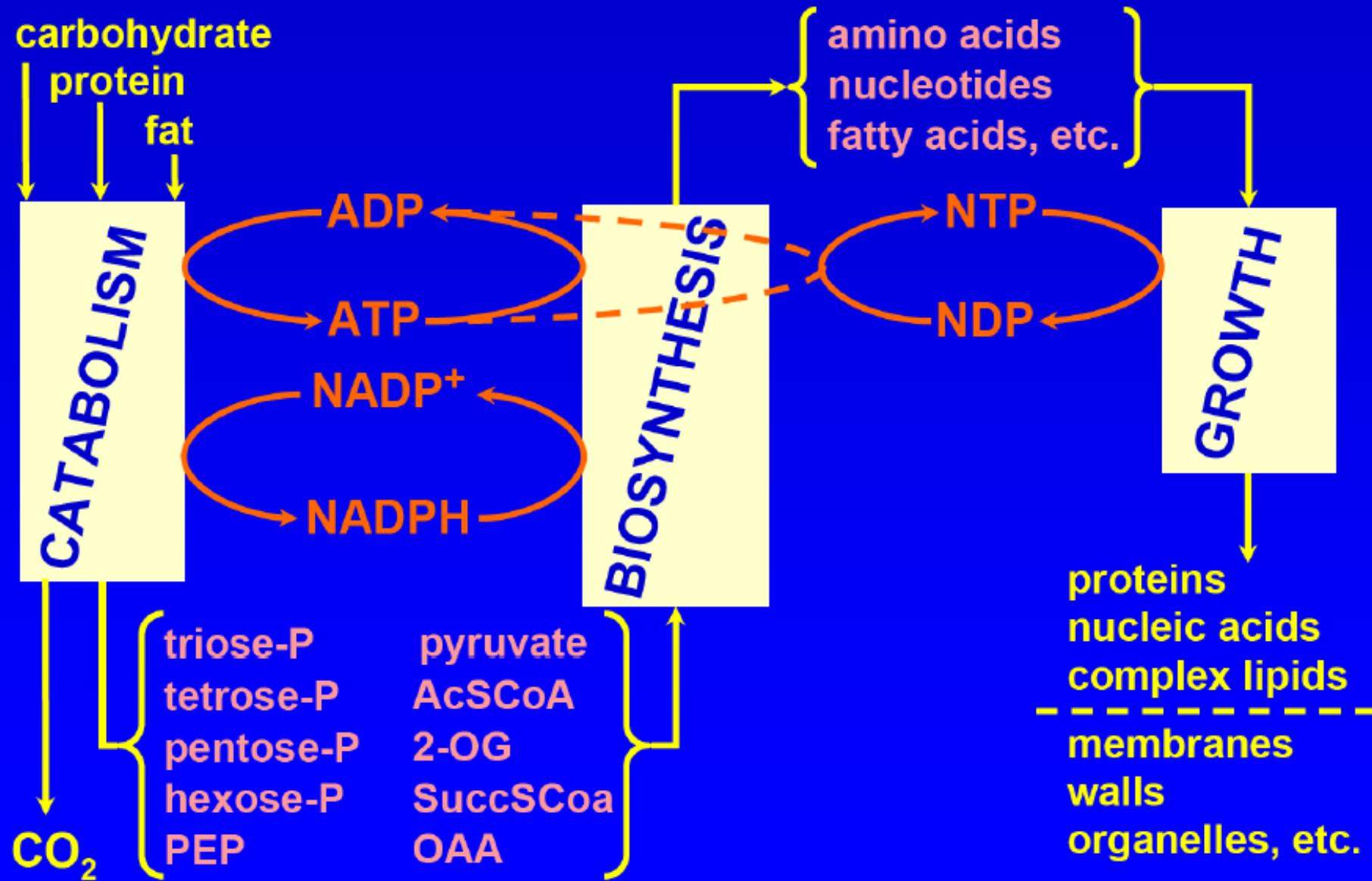
Lyon 1 University & INRA MIA Department

Slides borrowed from Jannie Hofmeyr

Stellenbosch University

Hofmeyr & Cornish-Bowden (2000) *FEBS Lett.* 476:47-51

Functional organisation of metabolism



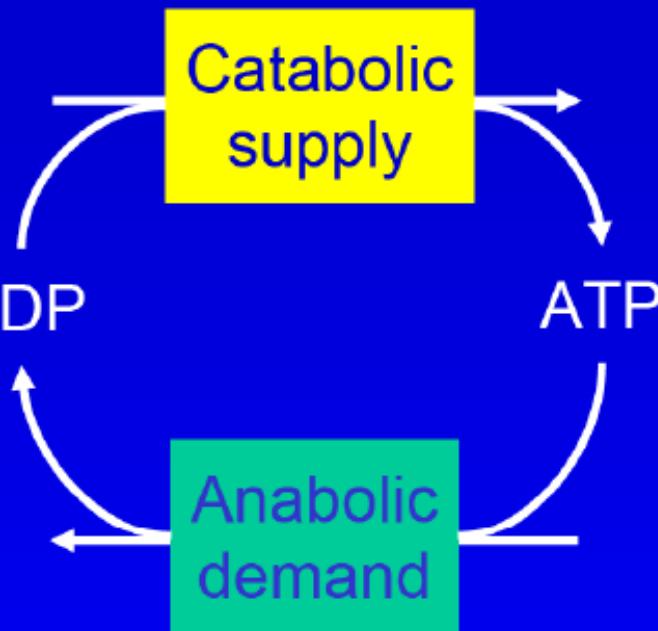
Linear couple



Metabolic end-product



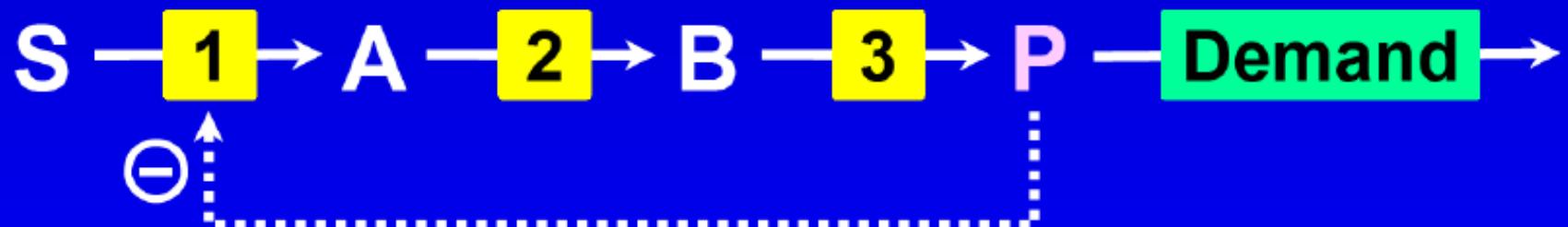
Cyclic couple

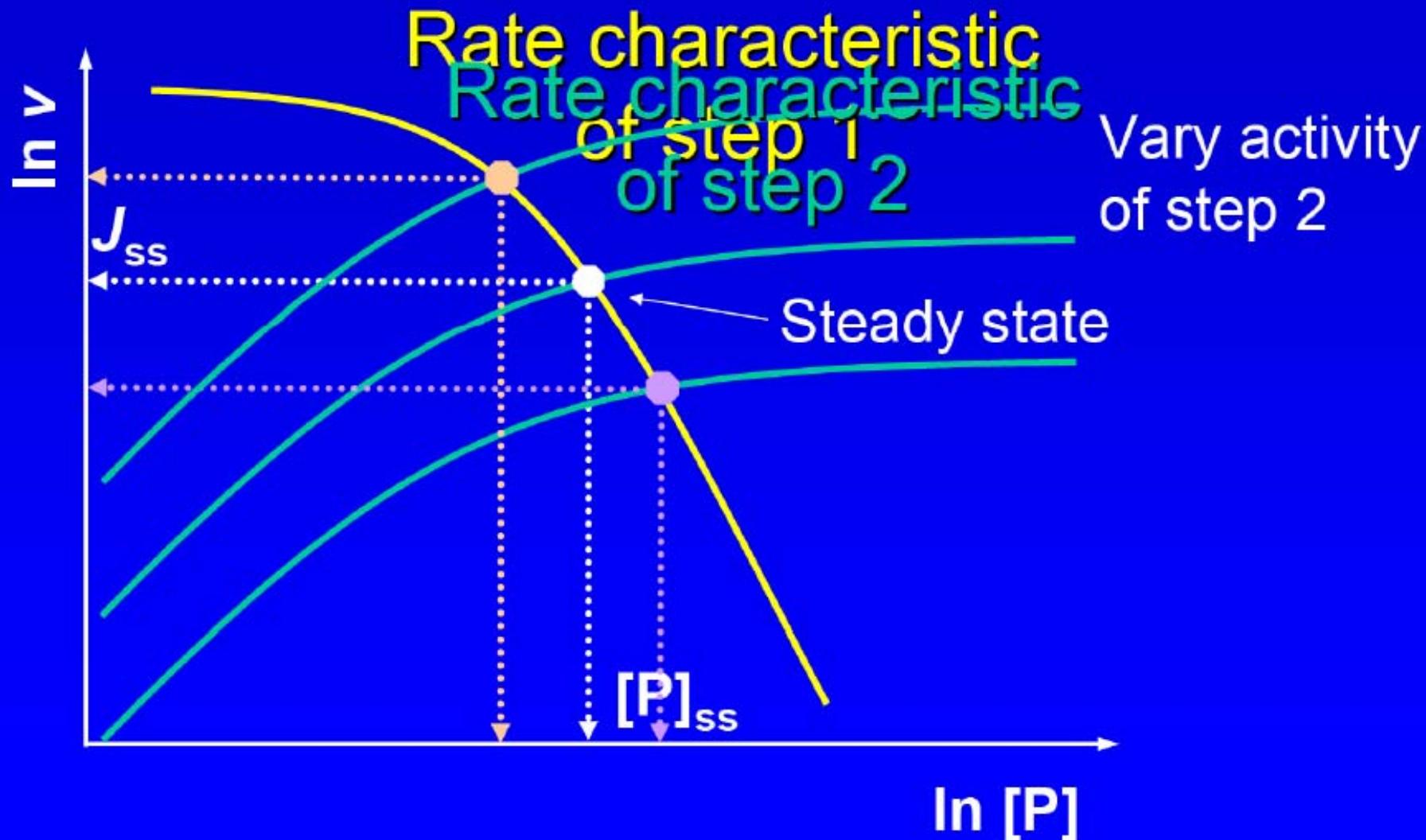
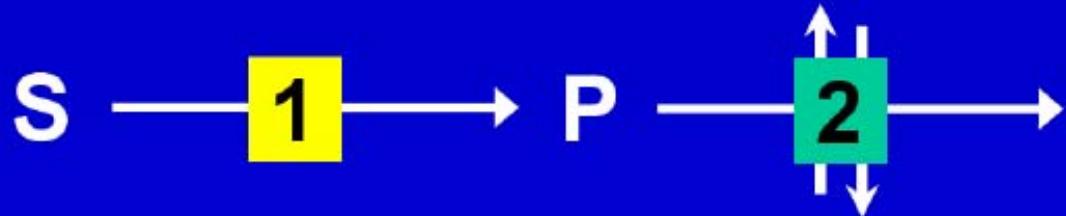


Generic supply-demand couple

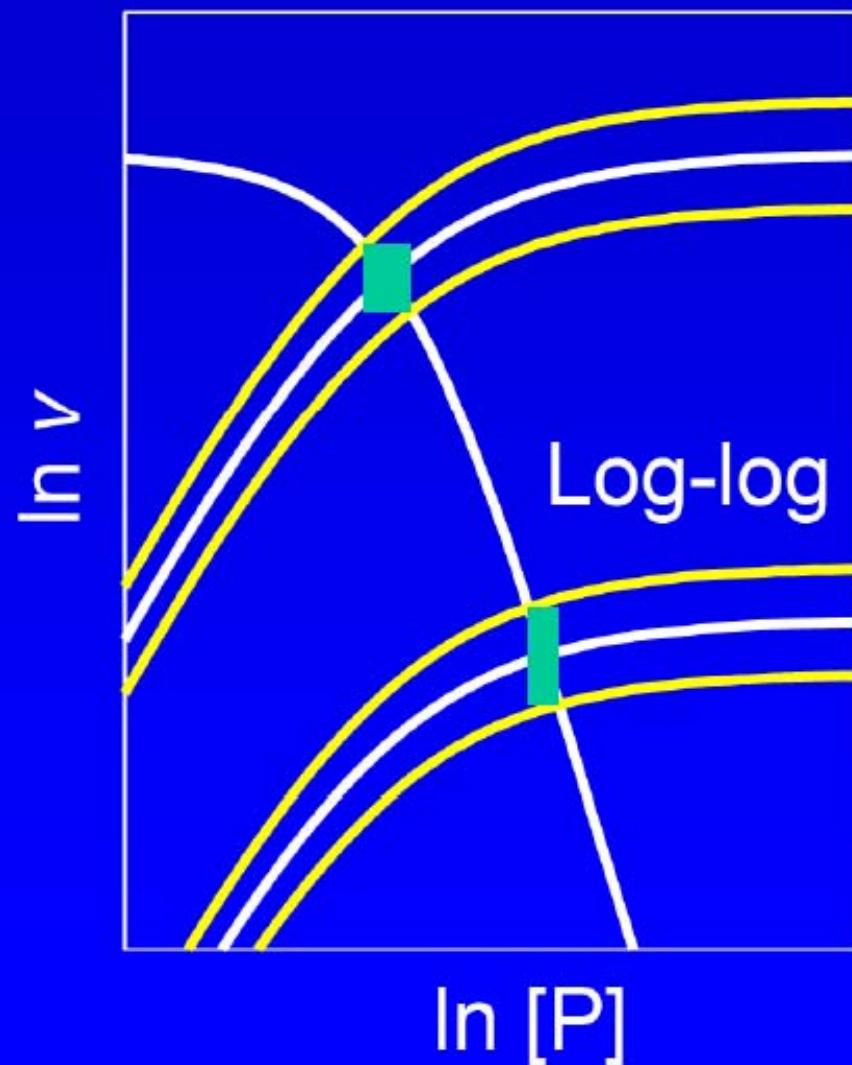
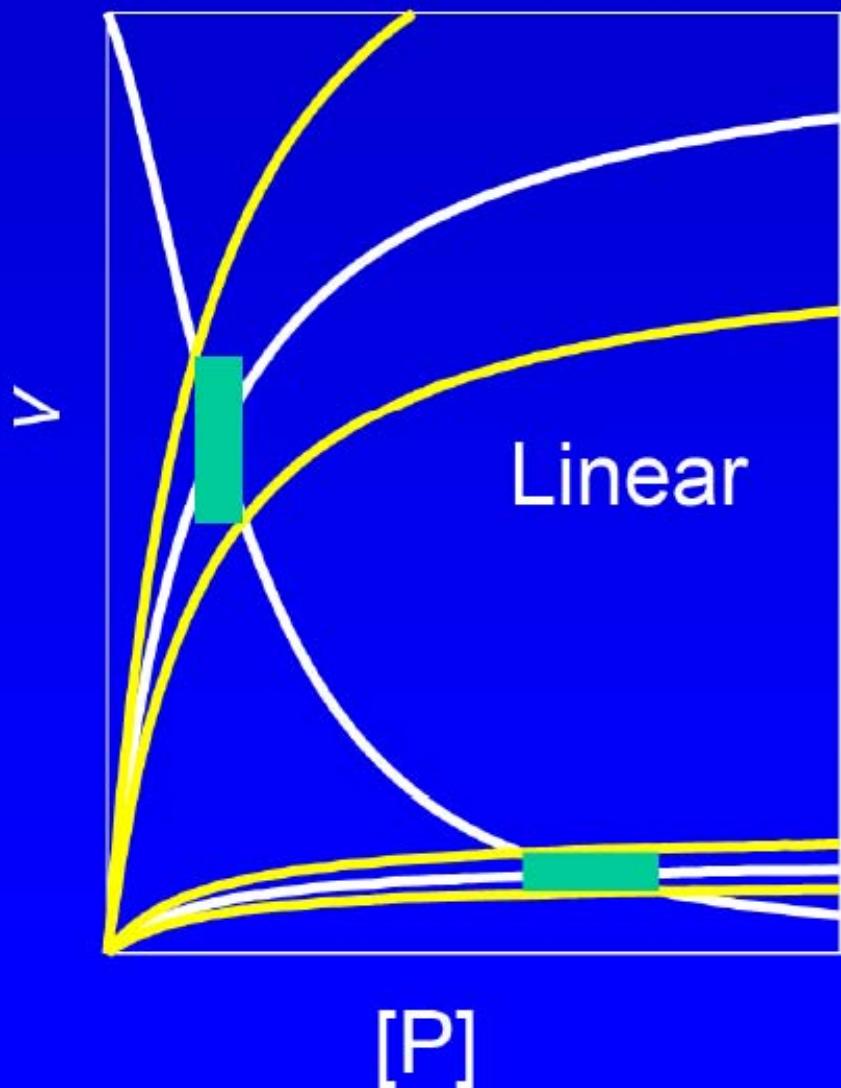
$$S - \boxed{\text{Supply}} \rightarrow \left(\frac{[P]}{[ATP]} \right) - \boxed{\text{Demand}} \rightarrow$$

Starting point: a metabolic factory

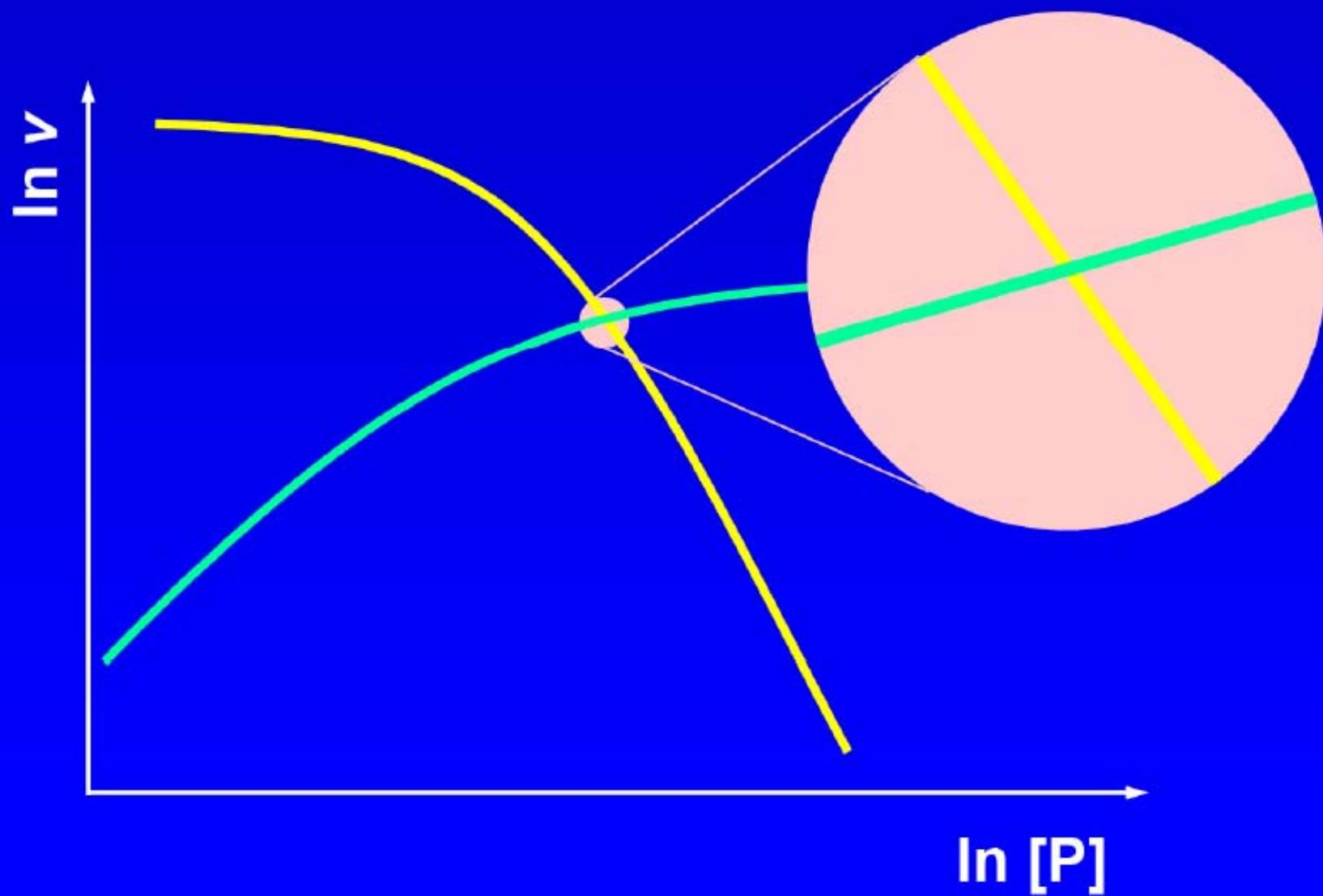




Linear vs. Logarithmic rate characteristics

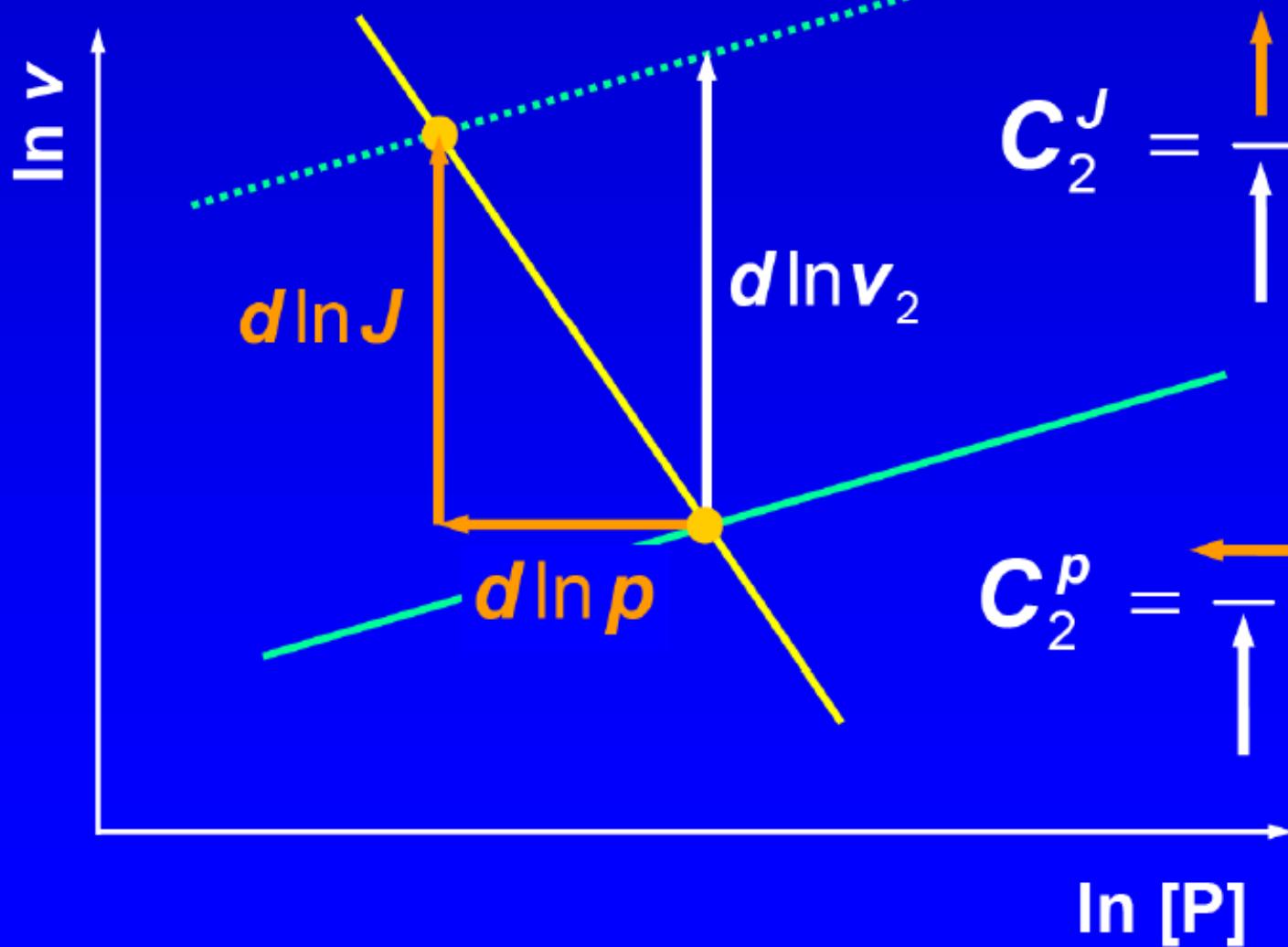


$S \xrightarrow{1} P \xrightarrow{2}$



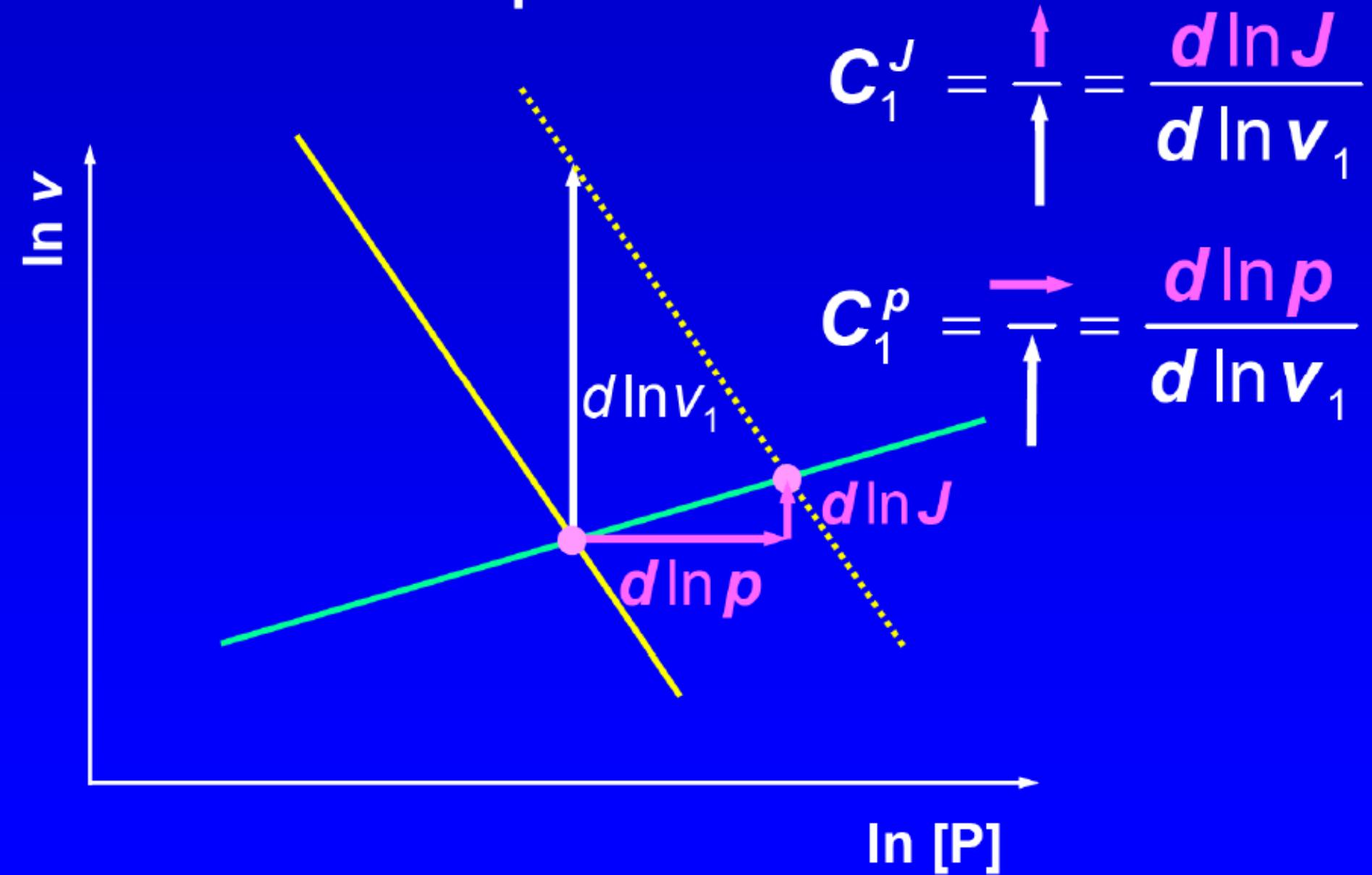
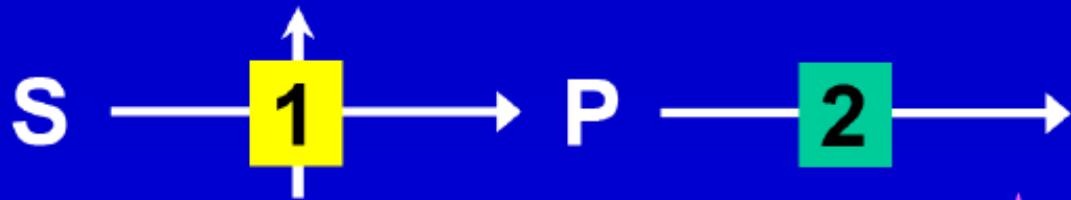


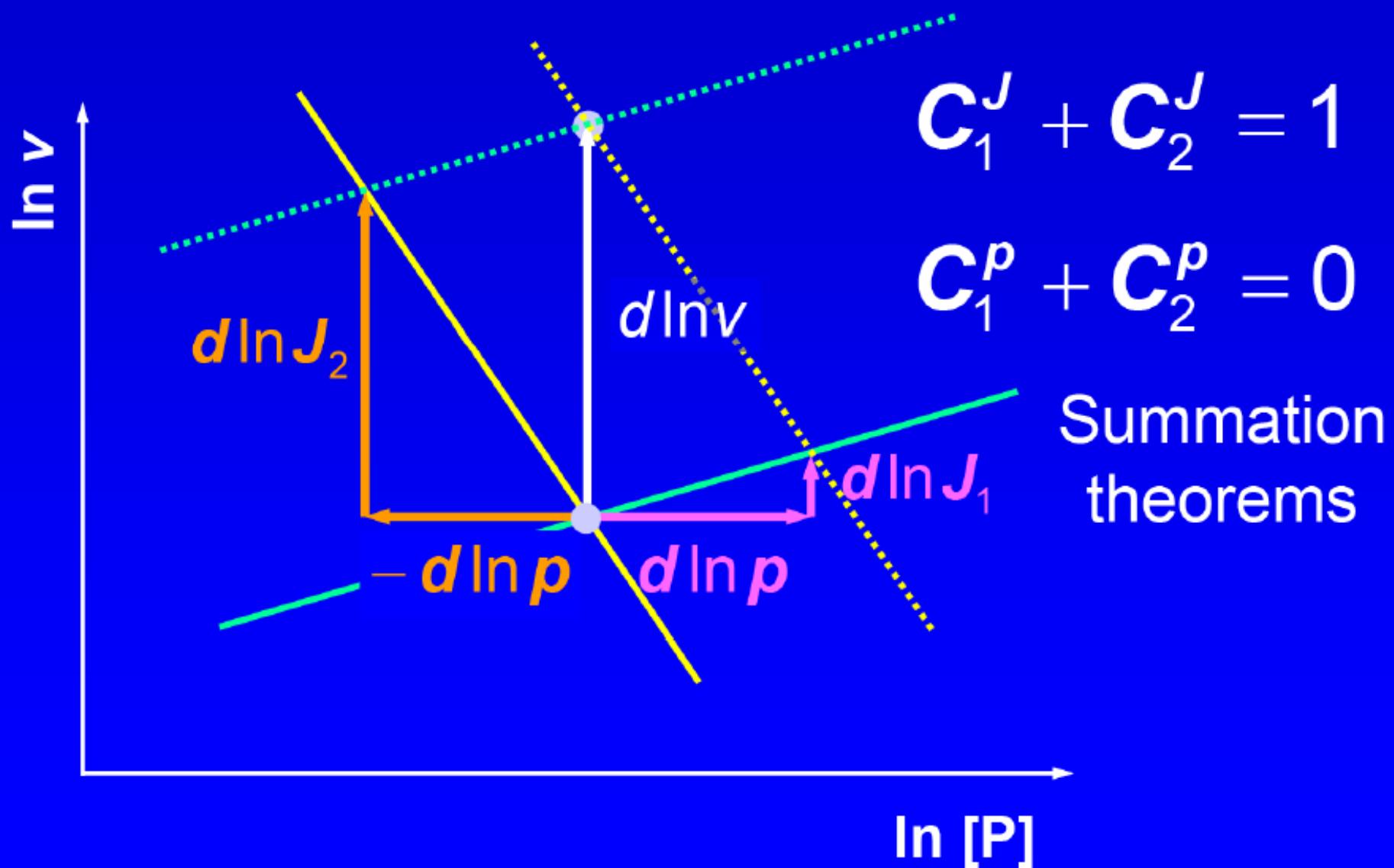
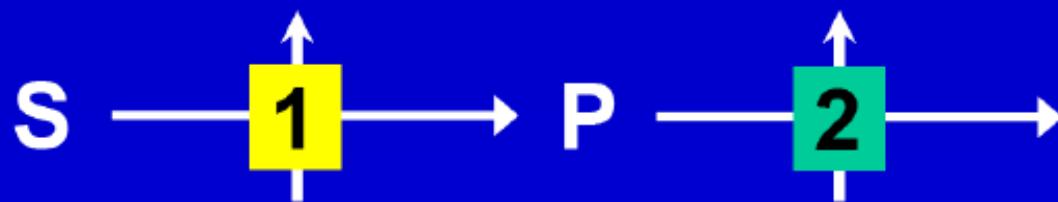
Control
coefficients

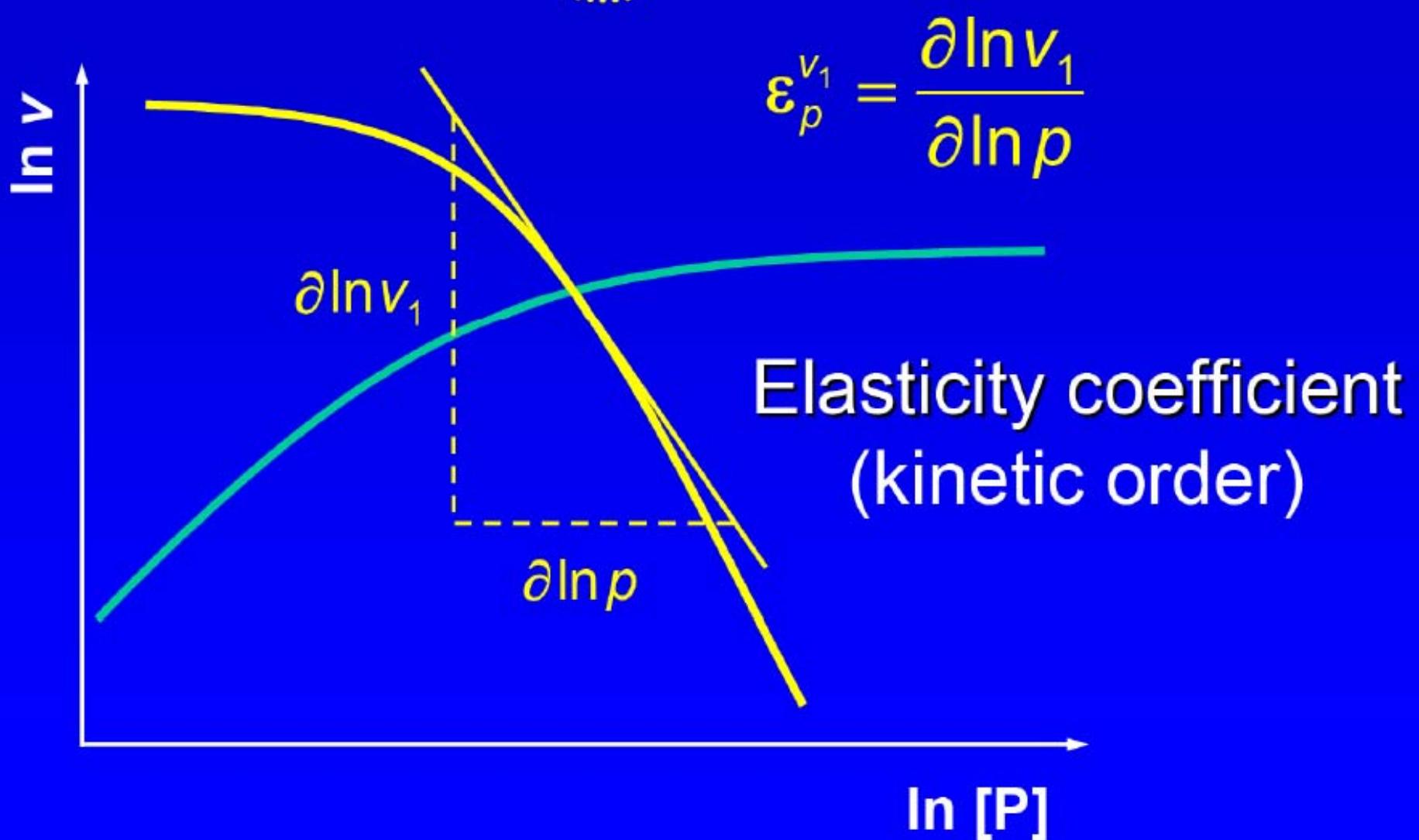


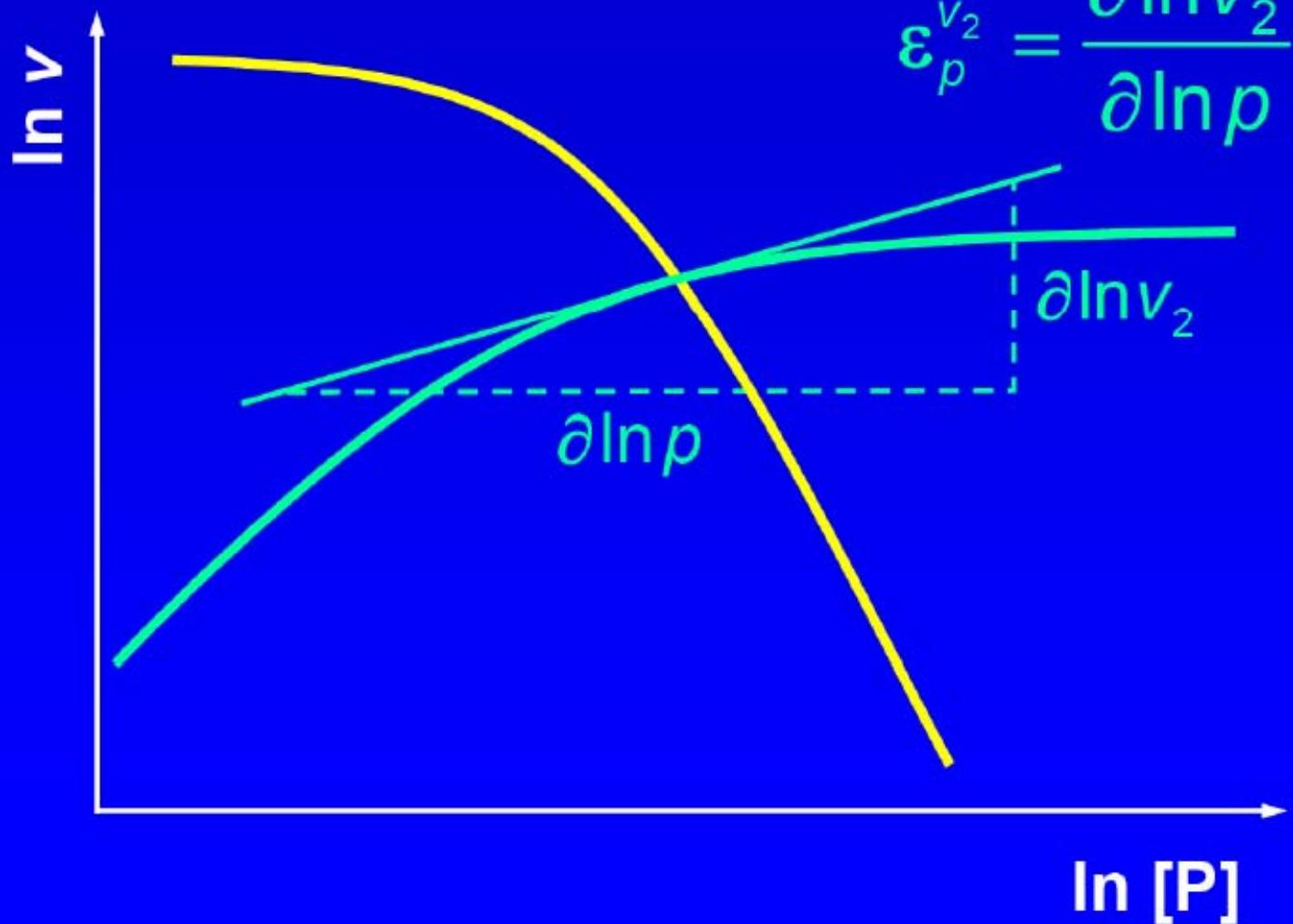
$$C_2^J = \frac{\uparrow}{\uparrow} = \frac{d \ln J}{d \ln v_2}$$

$$C_2^p = \frac{\leftarrow}{\uparrow} = \frac{d \ln p}{d \ln v_2}$$

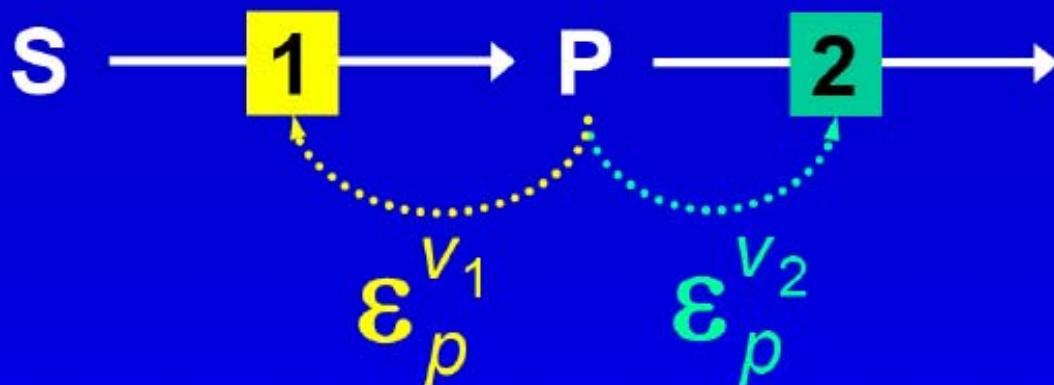








Summary of Control Properties



Summation

Connectivity

Flux

$$C_1^J + C_2^J = 1$$

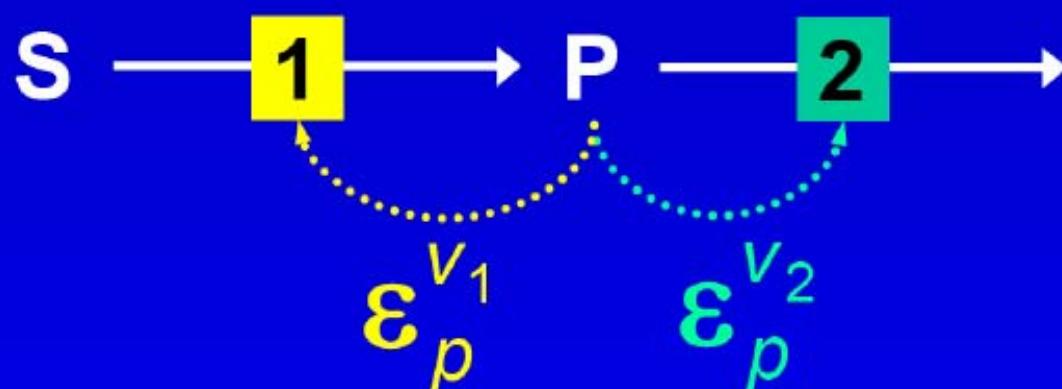
$$C_1^J \boldsymbol{\epsilon}_p^{v_1} + C_2^J \boldsymbol{\epsilon}_p^{v_2} = 0$$

Concen-
tration

$$C_1^p + C_2^p = 0$$

$$C_1^p \boldsymbol{\epsilon}_p^{v_1} + C_2^p \boldsymbol{\epsilon}_p^{v_2} = -1$$

Control analytic expressions



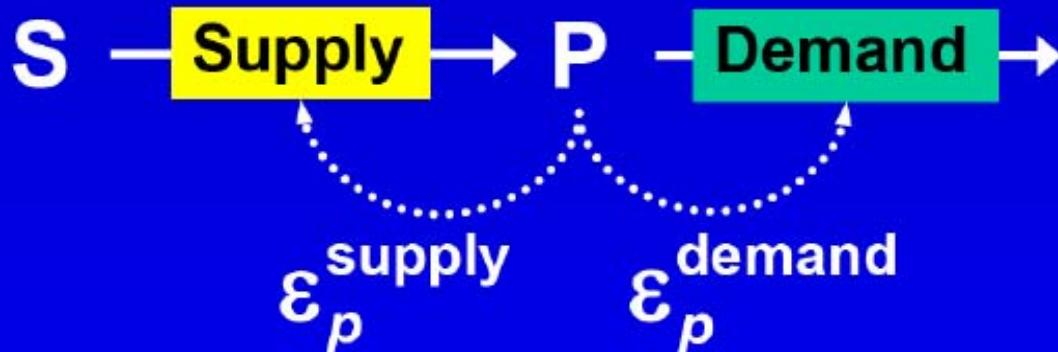
$$C_1^J = \frac{\epsilon_p^{v_2}}{\epsilon_p^{v_2} - \epsilon_p^{v_1}}$$

$$C_2^J = \frac{-\epsilon_p^{v_1}}{\epsilon_p^{v_2} - \epsilon_p^{v_1}}$$

$$C_1^P = \frac{1}{\epsilon_p^{v_2} - \epsilon_p^{v_1}}$$

$$C_2^P = \frac{-1}{\epsilon_p^{v_2} - \epsilon_p^{v_1}}$$

Control analysis of supply and demand



$$C_J^J_{\text{supply}} = \frac{\varepsilon_p^{\text{demand}}}{\varepsilon_p^{\text{demand}} - \varepsilon_p^{\text{supply}}}$$

$$C_P^P_{\text{supply}} = \frac{1}{\varepsilon_p^{\text{demand}} - \varepsilon_p^{\text{supply}}}$$

$$C_J^J_{\text{demand}} = \frac{-\varepsilon_p^{\text{supply}}}{\varepsilon_p^{\text{demand}} - \varepsilon_p^{\text{supply}}}$$

$$C_P^P_{\text{demand}} = \frac{-1}{\varepsilon_p^{\text{demand}} - \varepsilon_p^{\text{supply}}}$$

Distribution of flux control is determined by the RATIO of supply and demand elasticities

$$\frac{C_J^{\text{demand}}}{C_J^{\text{supply}}} = \frac{-\varepsilon_p^{\text{supply}}}{\varepsilon_p^{\text{demand}}}$$

Case

$$\varepsilon_p^{\text{demand}} \ll |\varepsilon_p^{\text{supply}}|$$

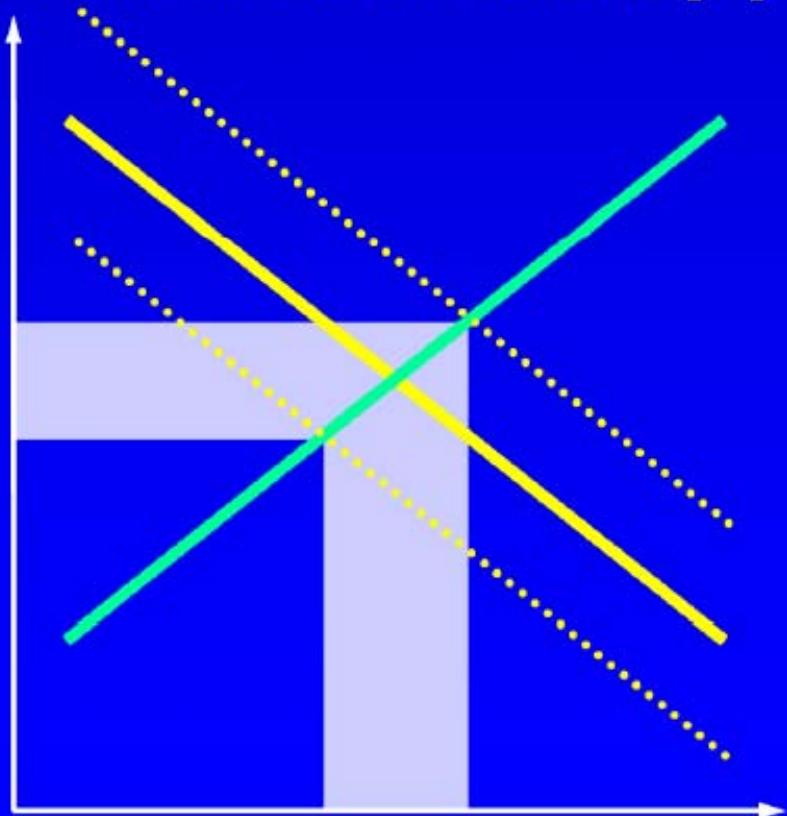
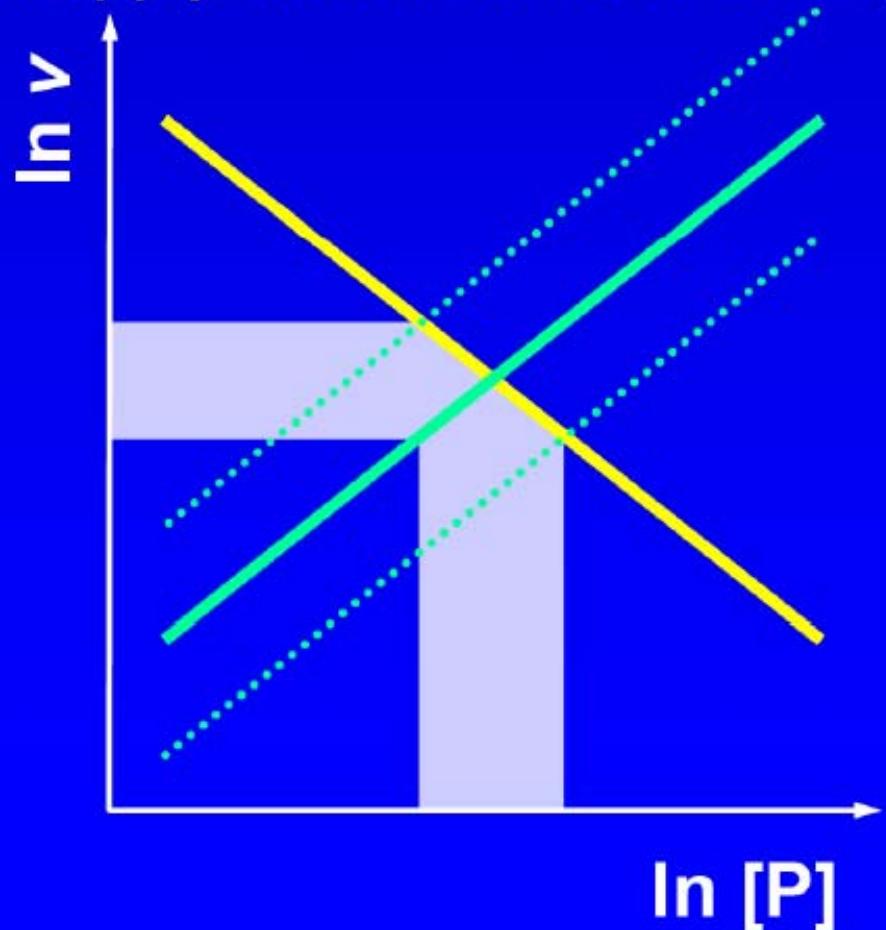
$$\Rightarrow C_J^{\text{demand}} \gg C_J^{\text{supply}}$$

$$\Rightarrow C_J^{\text{demand}} \approx 1$$

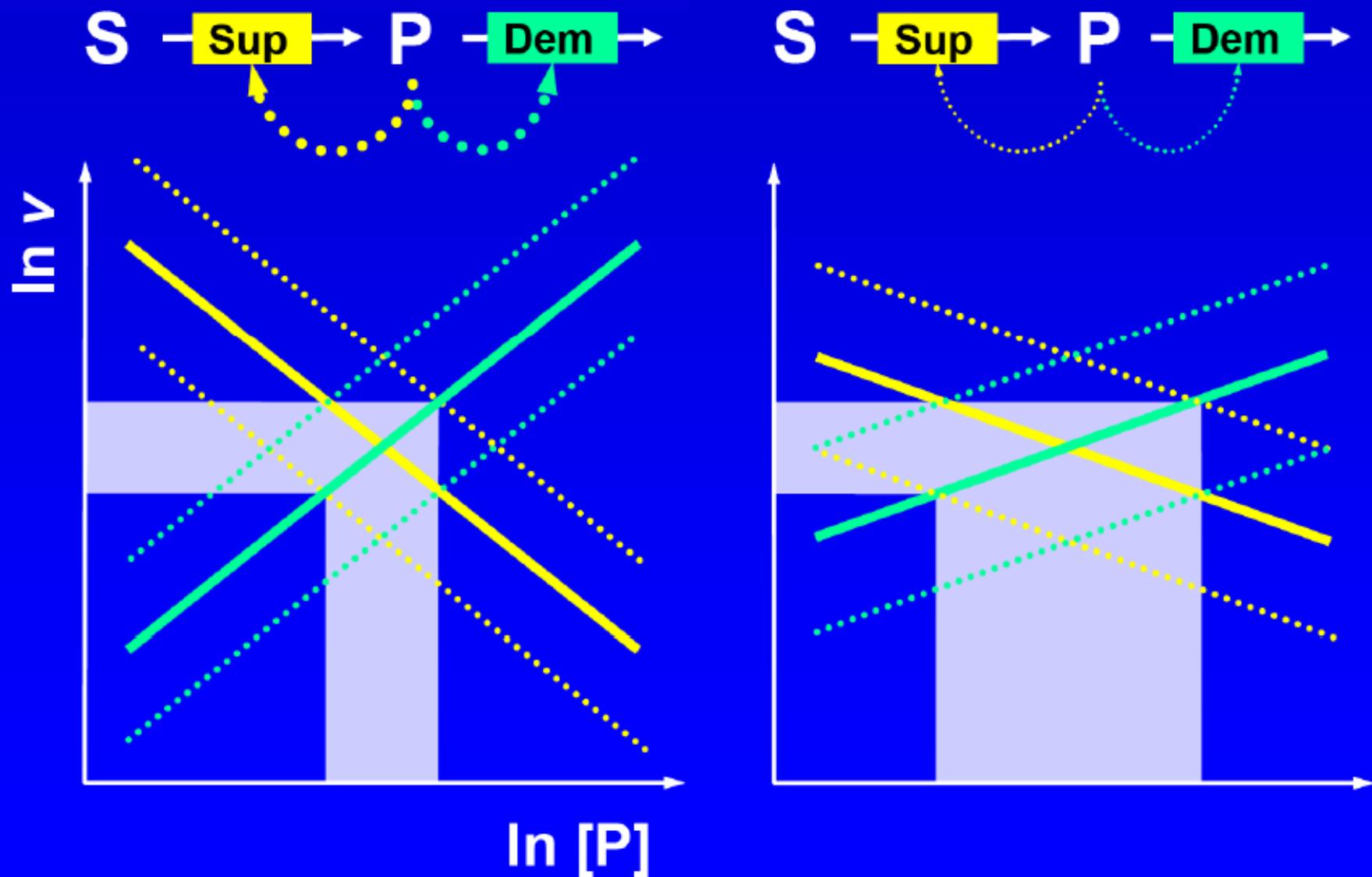
A functionally undifferentiated system



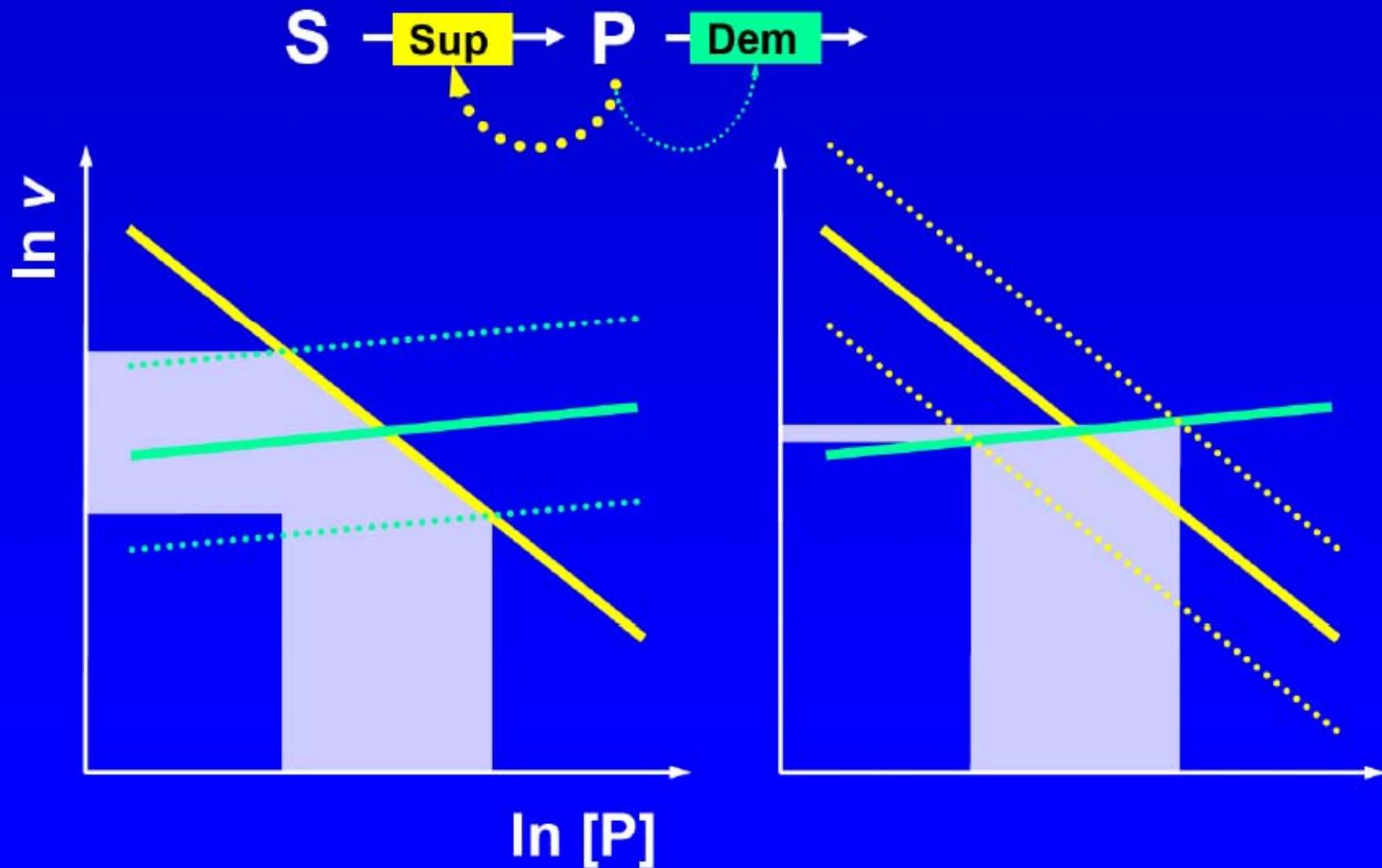
Supply and demand have equal control over flux and [P]



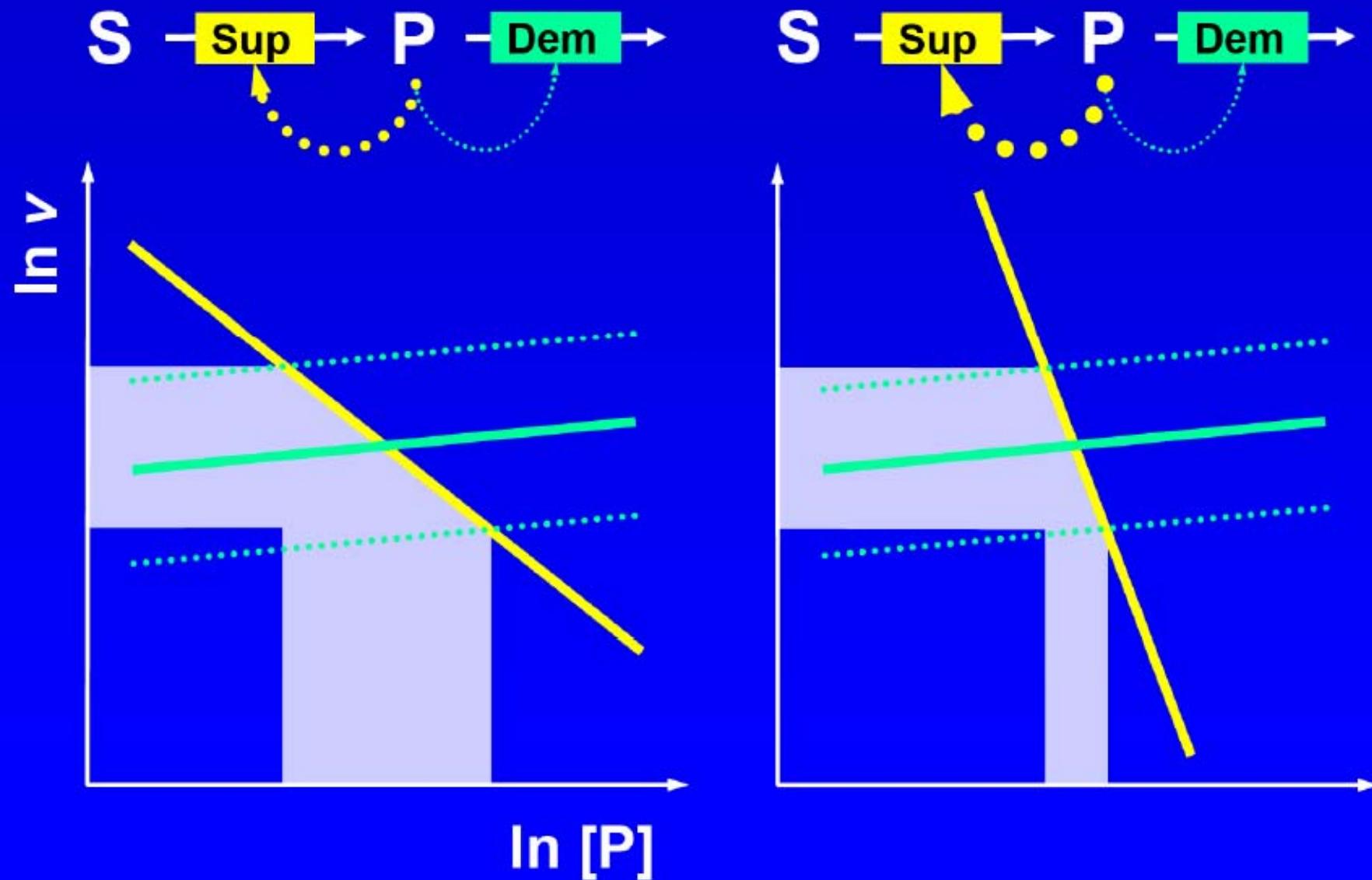
The sum of slopes determines the magnitude of concentration control



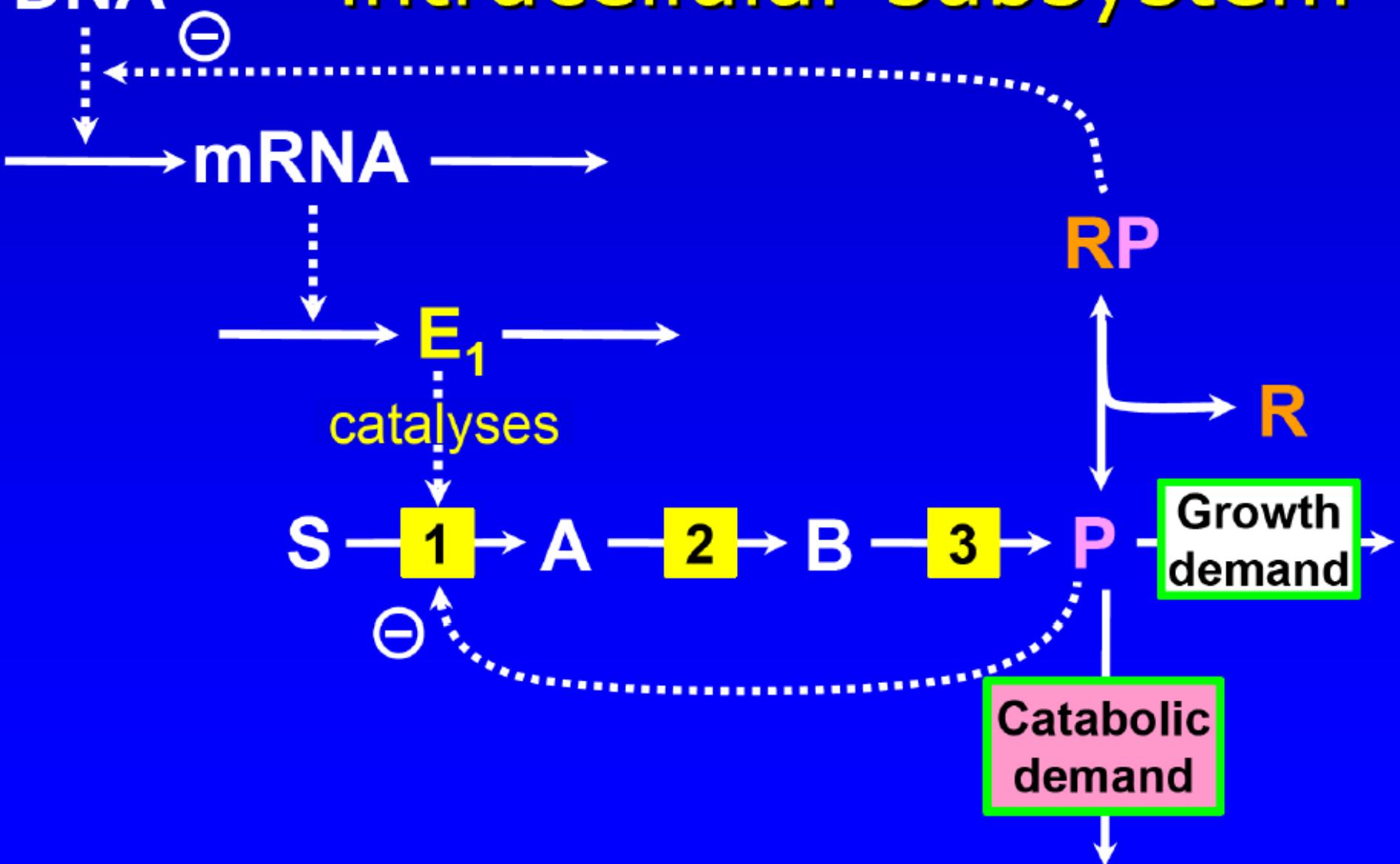
If the supply/demand slope ratio > 1,
then flux-control shifts to the demand



The more one block controls the flux, the more the other determines the magnitude of P-control



A typical regulated intracellular subsystem

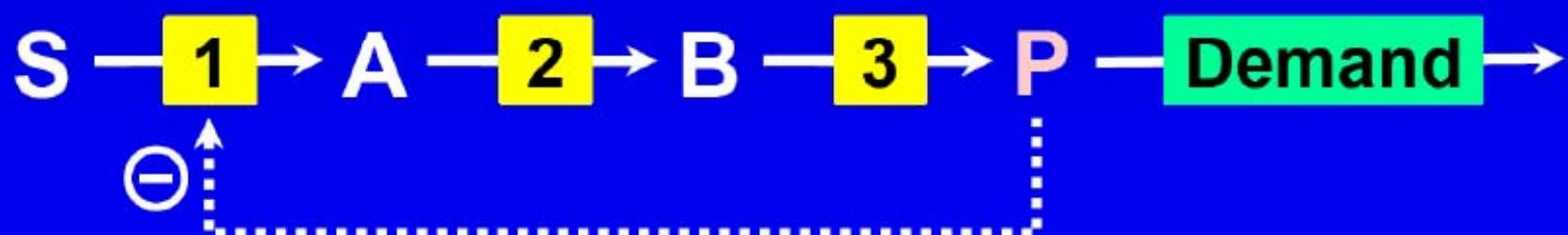


Modelling the biosynthetic supply

Reversible
Hill-equation
with modifier P

Reversible
Michaelis-Menten
equations

Irreversible
Michaelis-Menten
equation



$$K_{\text{eq}} = 400$$

$$K_{\text{eq}} = 10$$

$$K_{\text{eq}} = 10$$

Equilibrium concentrations

1

400

4000

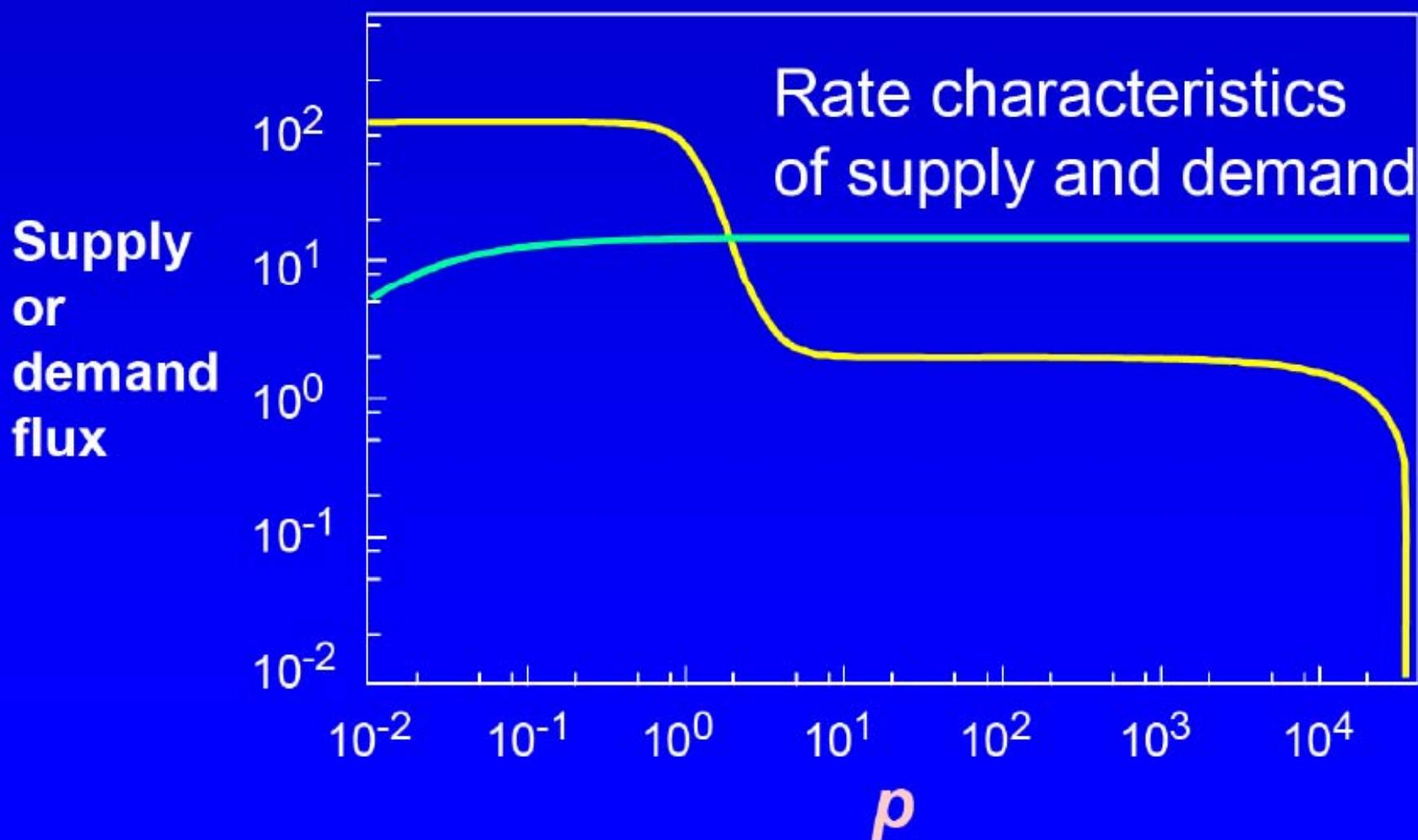
40000

Reversible Hill equation with allosteric modifier

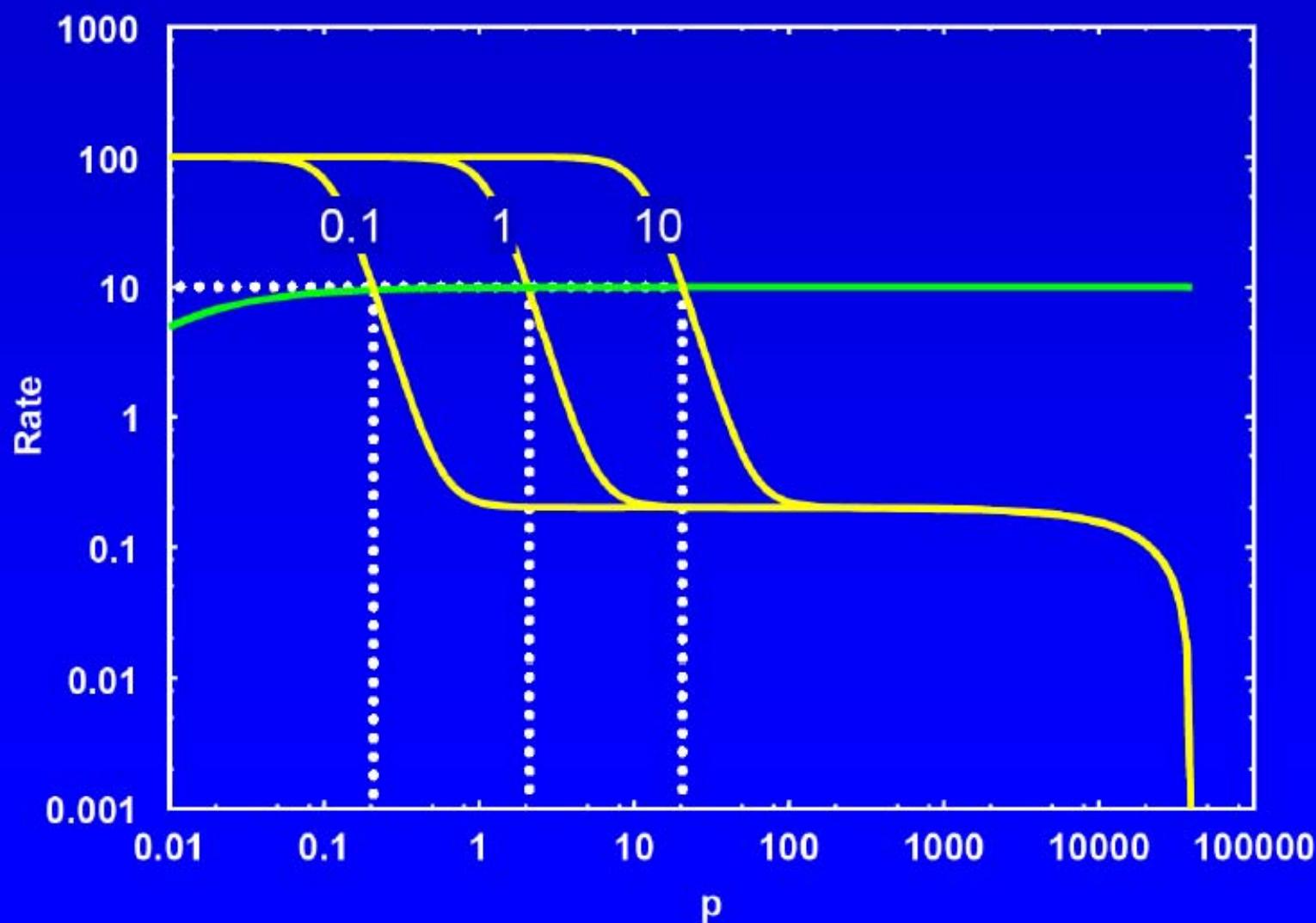
(Hofmeyr & Cornish-Bowden, 1997)



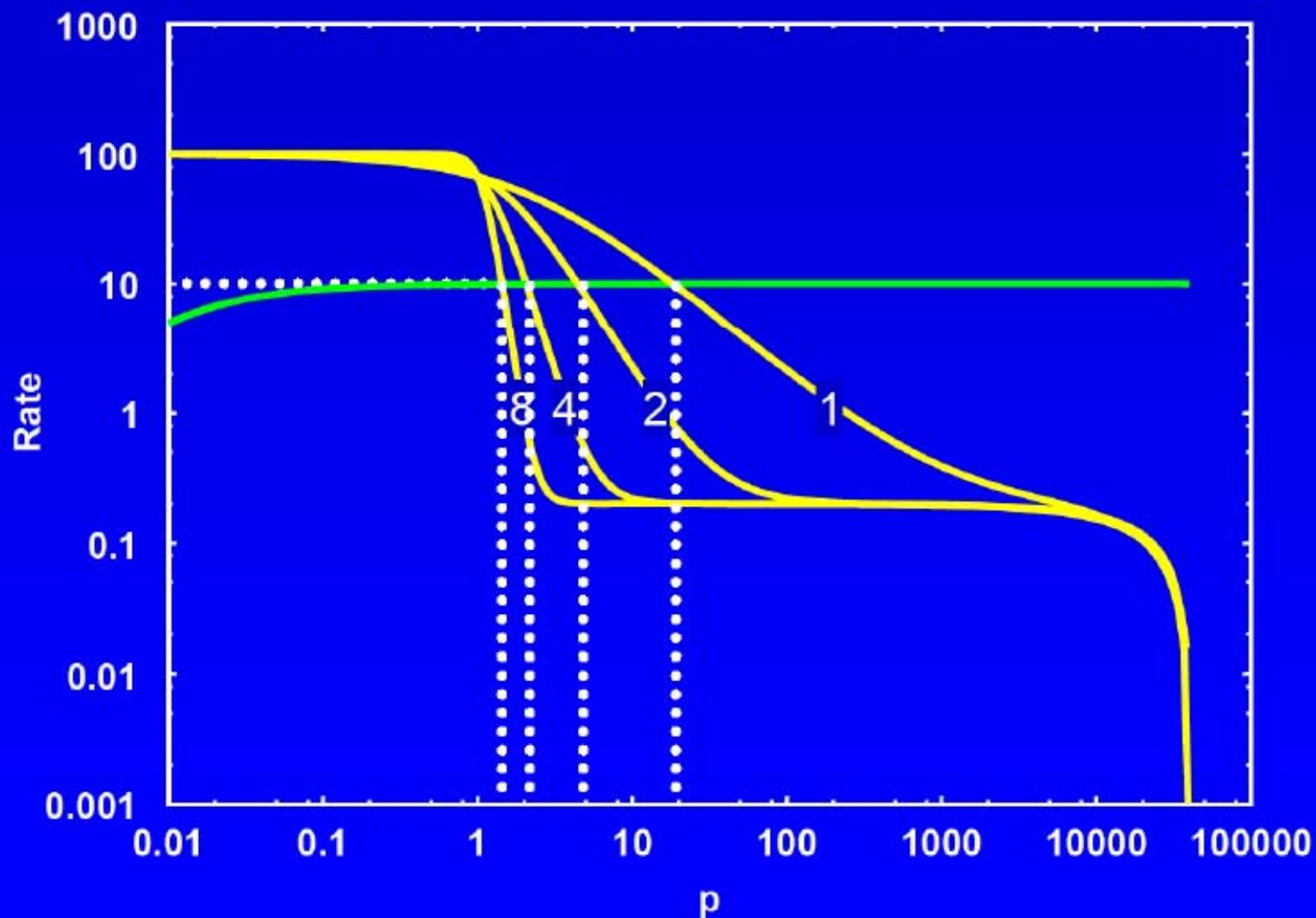
$$V = \frac{k_{\text{cat}}}{S_{0.5}} \cdot e_1 \cdot \frac{\left(\frac{s}{S_{0.5}} + \frac{a}{a_{0.5}} \right)^{h-1}}{1 + \alpha \left(\frac{p}{p_{0.5}} \right)^h} \cdot \left(s - \frac{a}{K_{\text{eq}}} \right)$$



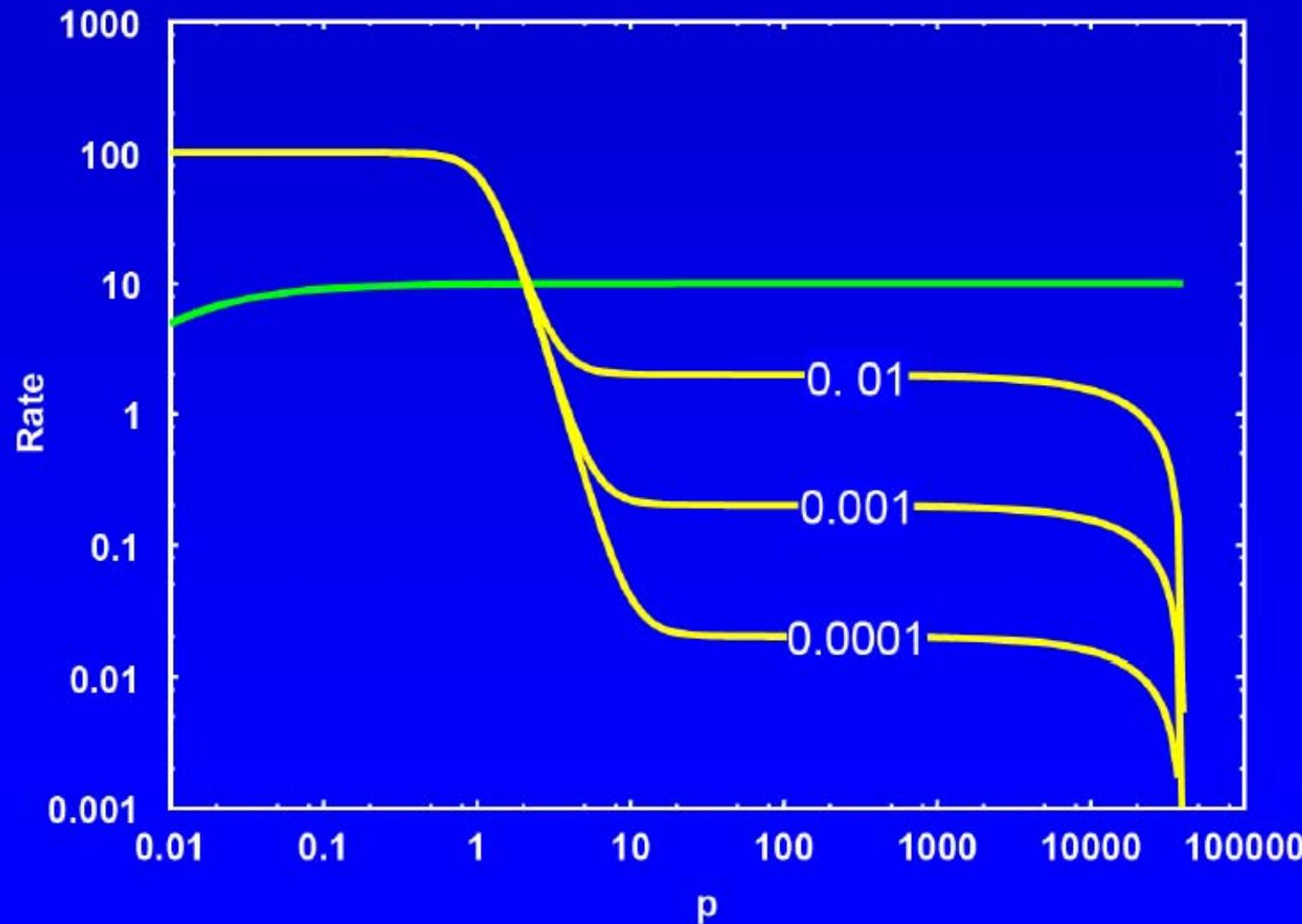
Varying the binding strength ($p_{0.5}$) of allosteric effector P



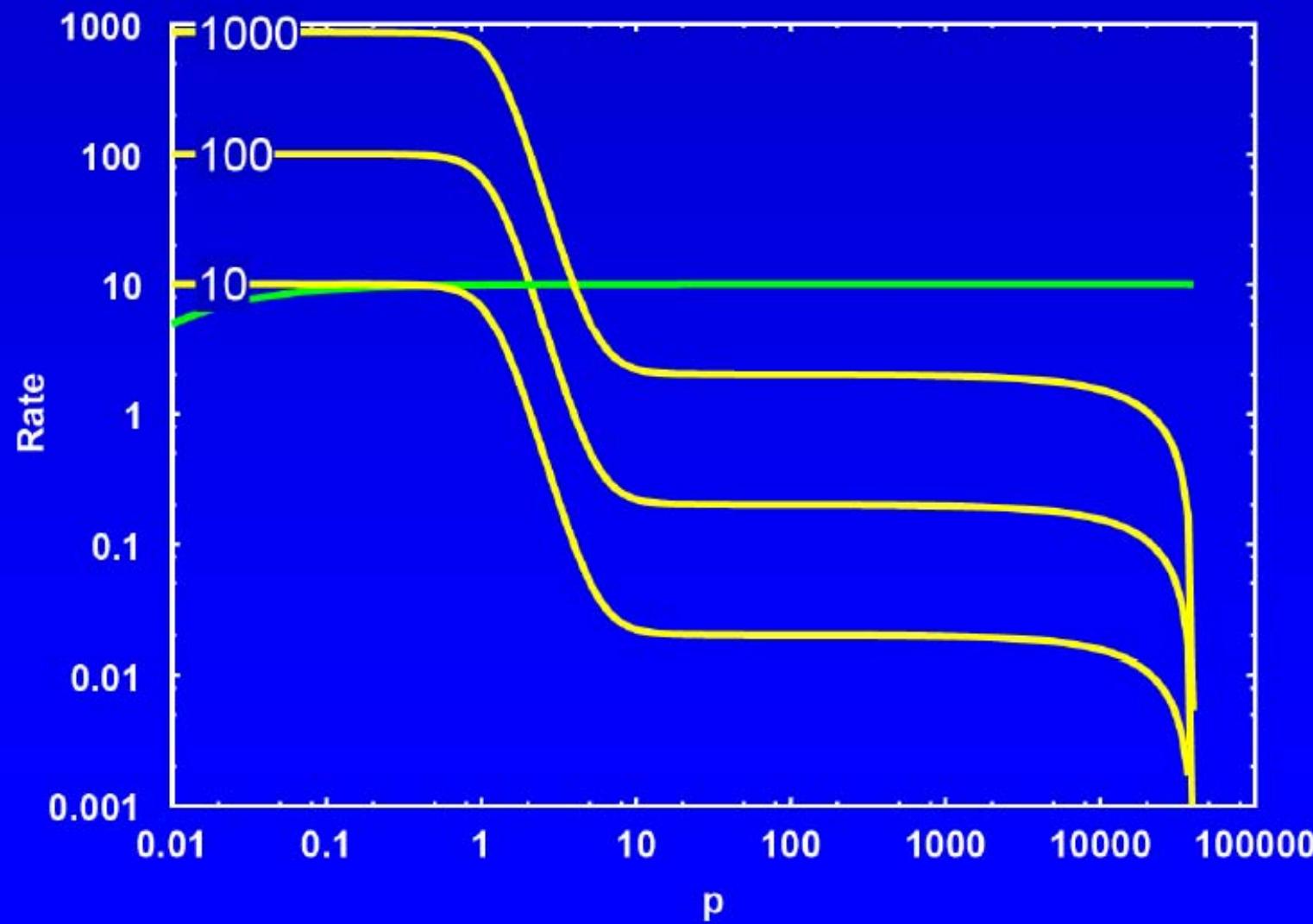
Varying the degree of cooperativity (h)

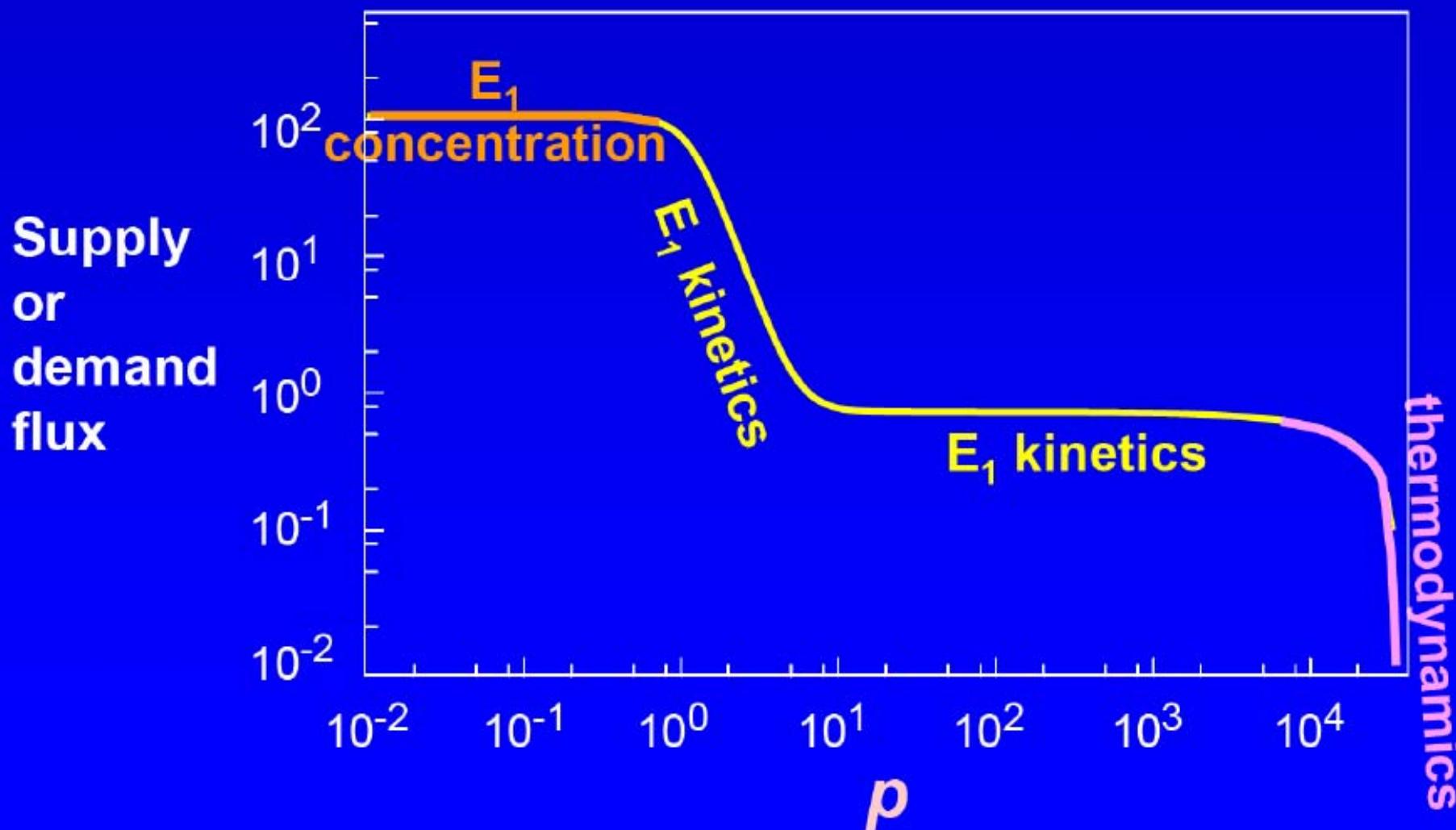


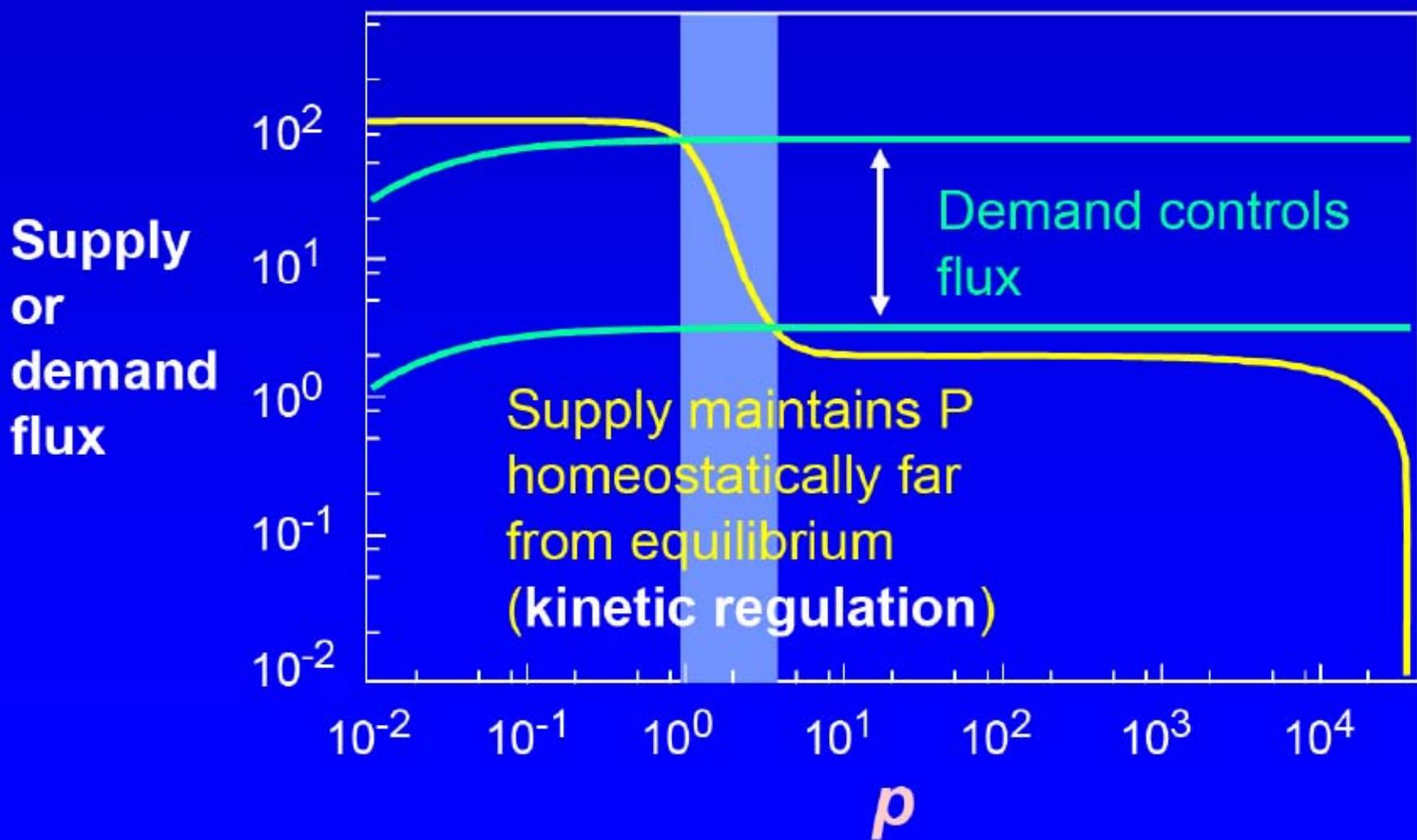
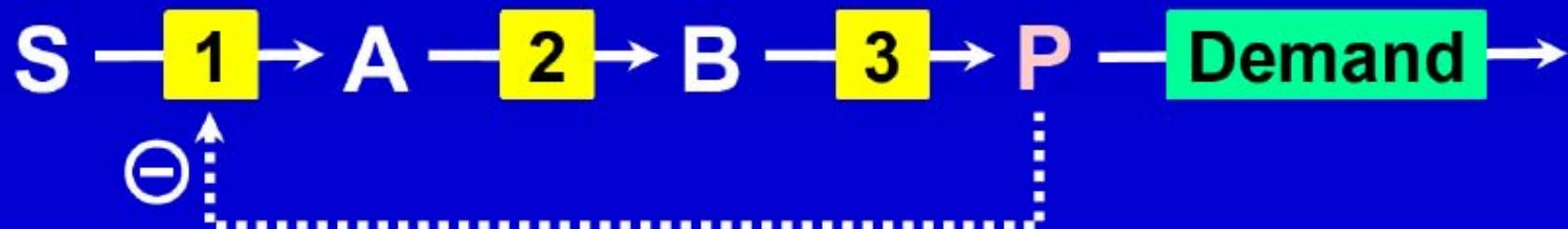
Varying the strength of allosteric inhibition of supply by P (α)

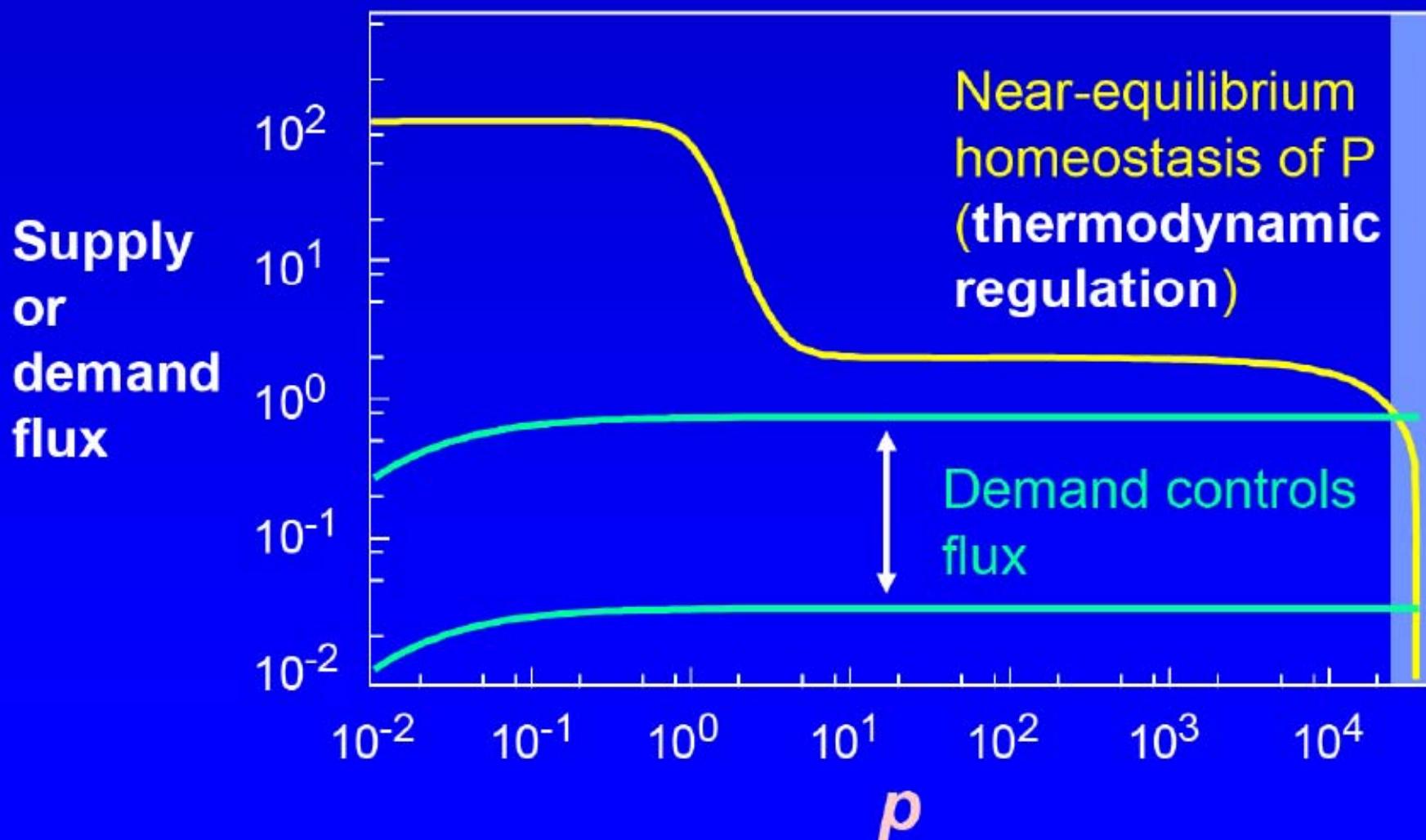


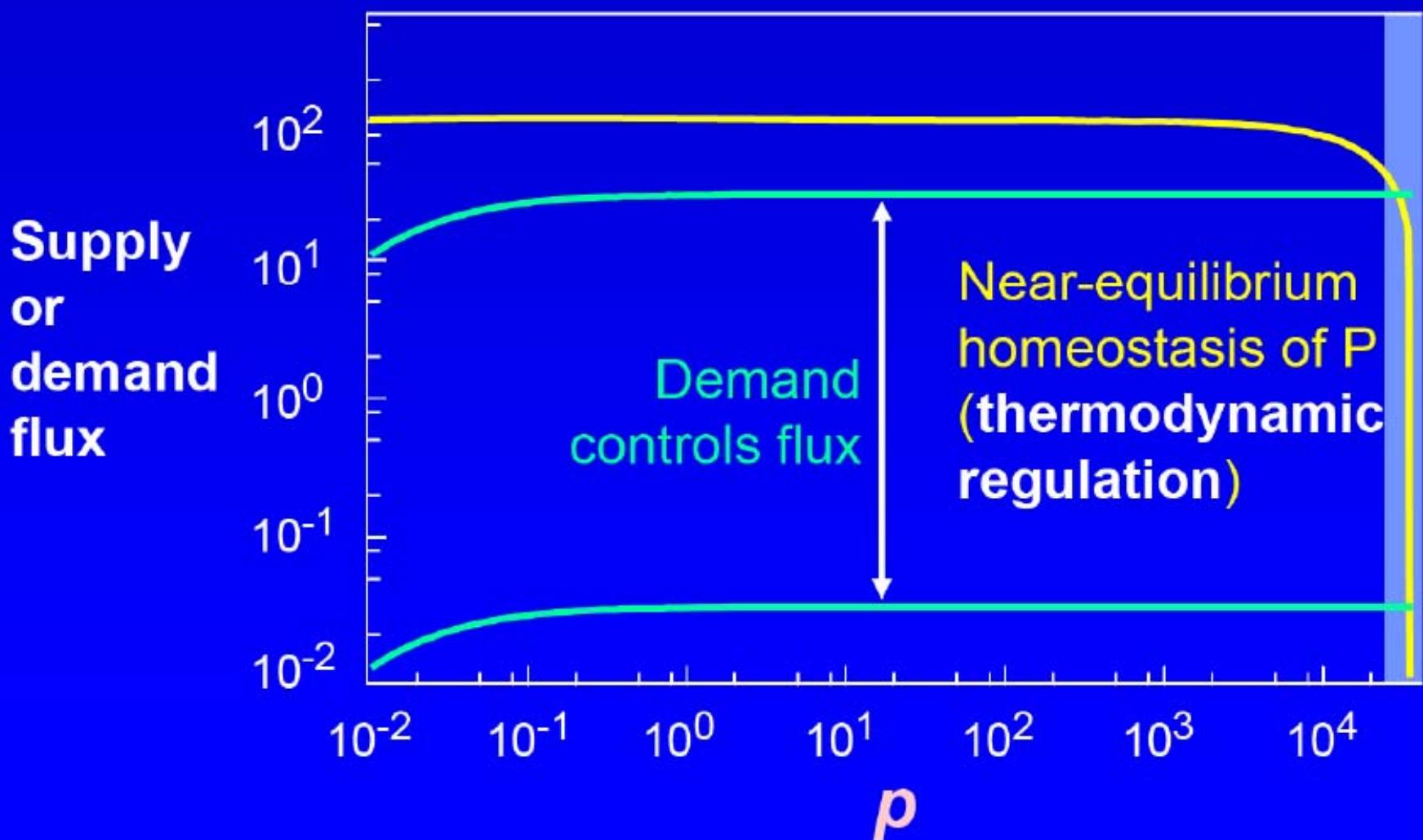
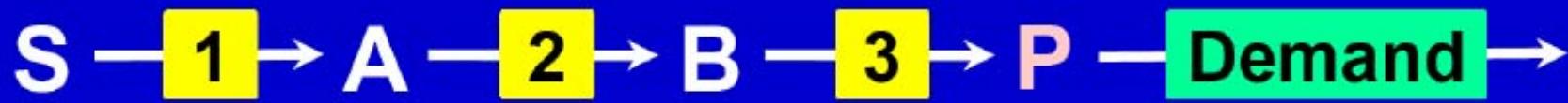
Varying the maximal capacity (V_{\max}) of the supply through $[E_1]$

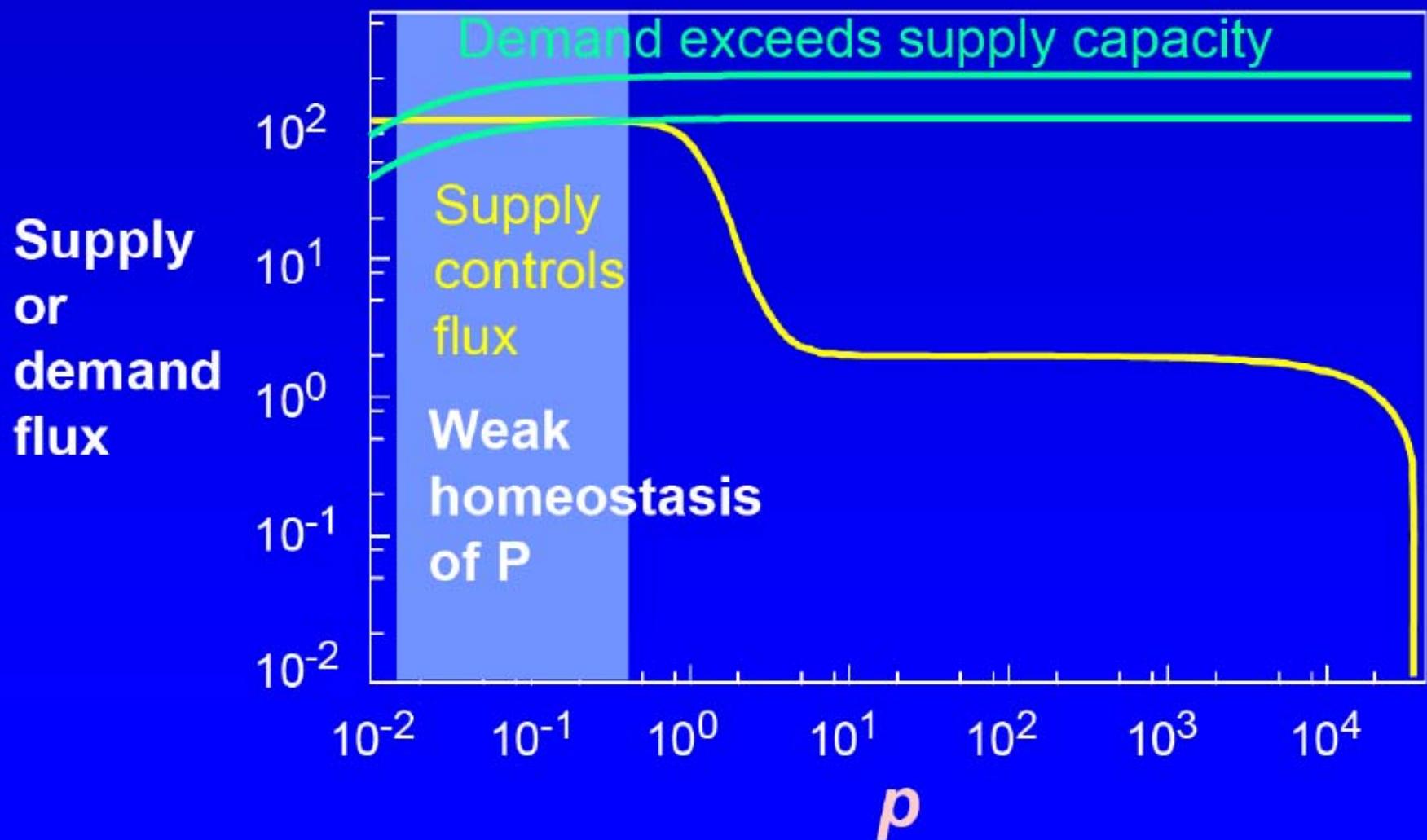




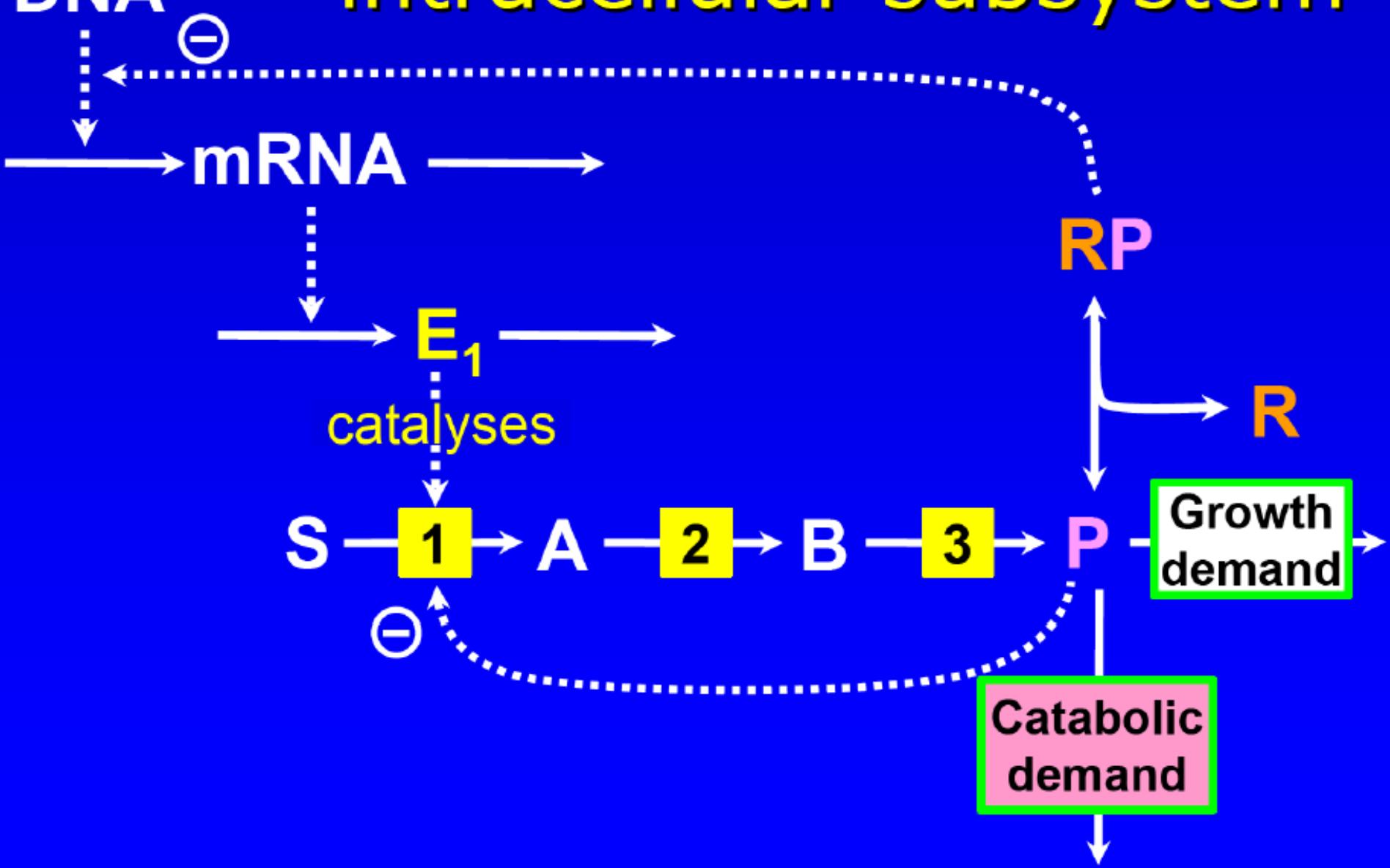








A typical regulated intracellular subsystem



Simplified core model

