

系统生物学

(Systems Biology)

马彬广





细胞行为建模

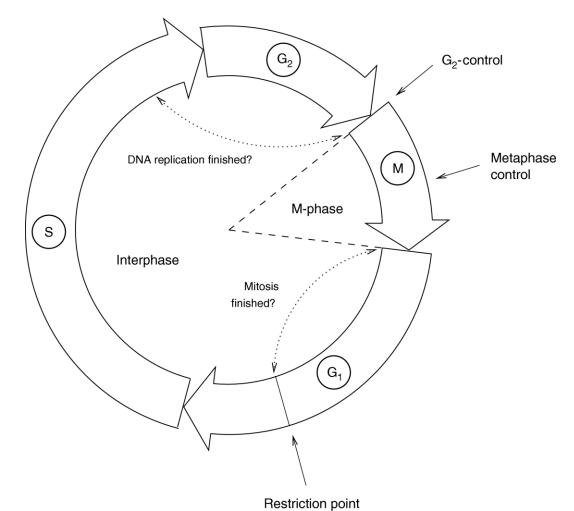
(第十六讲)











© 2010 Wiley-VCH, Weinheim Klipp - Systems Biology ISBN: 978-3-527-31874-2 fig-03-23

· 辛中震素大学 HUAZHONG AGRICULTURAL UNIVERSITY





□ G1 cyclin rises, bind to their CDK, signal the cell to prepare the chromosome replication

□ S phase promoting factor (SPF) rises, cyclin A with CDK2, enters nucleus and prepares the cell to duplicate its DNA.

 As DNA replication continues, cyclin E is destroyed and mitotic cyclin increase (in G2)

□ The M phase-promoting factor (the complex of mitotic cyclins with the Mphase CDK) initiates (i) assembly of the mitotic spindle; (2) breakdown of the nuclear envelope; (iii) condensation of the chromosomes.

□ Into metaphase, the M phase-promoting factor activate anaphase-promoting complex (APC) which allows the sister chromatids at the metaphase to separate and move to the poles (anaphase), thereby completing mitosis.

□ APC destroys mitotic cyclin and turns on the synthesis of G1 cyclin for the next cycle.







In yeast, the CDKs have been identified, as cdc2 kinase in fission yeast, and cdc28 kinase in budding yeast.

□ The minimal model was proposed by Goldbeter (Goldbeter A. PNAS 1991, 88: 9107-9111).

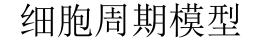
□ It is a bicyclic cascade model involving phosphorylation and dephosphorylation.

The model was used to test the hypothesis that cell cycle oscillation may arise from a negative feedback loop, i.e., the cyclin activates the Cdc2 kinase, while the Cdc 2 kinase triggers the degradation of the cyclin.

□ See the next page for the model.

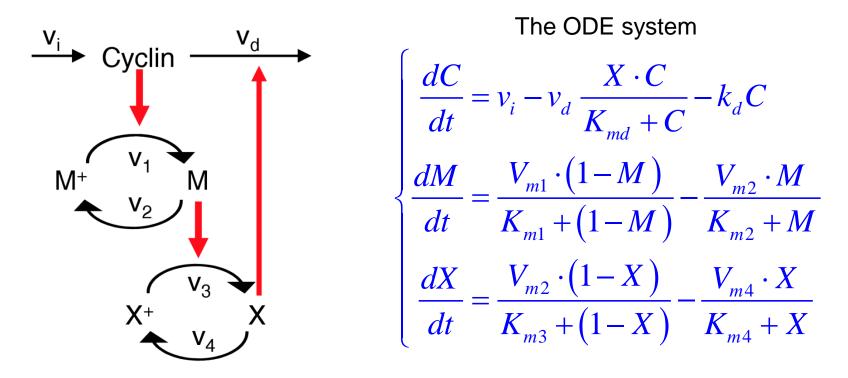








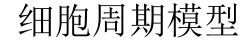
Bicyclic cascade model of yeast cell cycle



The model comprising cyclin production and degradation, phosphorylation and dephosphorylation of Cdc2 kinase, and phosphorylation and dephosphorylation of the cyclin protease.Parameter values: $K_{mi} = 0.05$ (i = 1, ..., 4), $K_{mc} = 0.5$, $k_d = 0.01$, $v_i = 0.025$, $v_d = 0.25$, $V_{m1} = 3$, $V_{m2} = 1.5$, $V_{m3} = 1$, $V_{m4} = 0.5$.

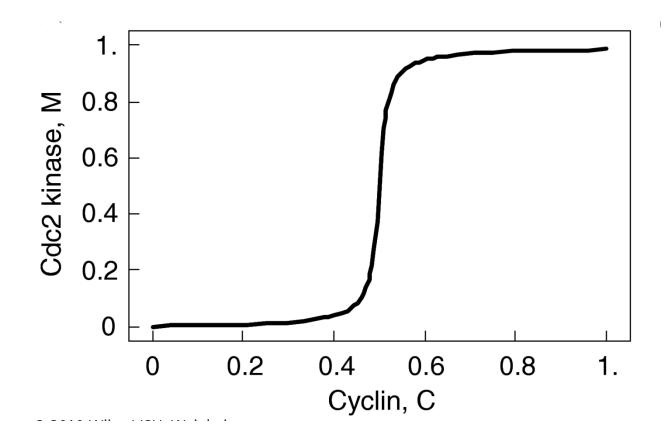








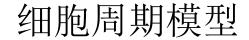
Two thresholds:



Threshold-type dependence of the fractional concentration of active Cdc2 kinase on the cyclin concentration. Initial conditions in are C(0) = M(0) = X(0) = 0.01. Units: μ M and min-1.

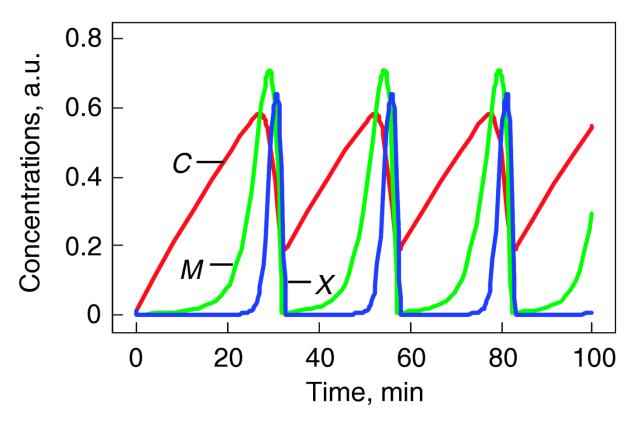








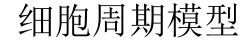
Periodicity



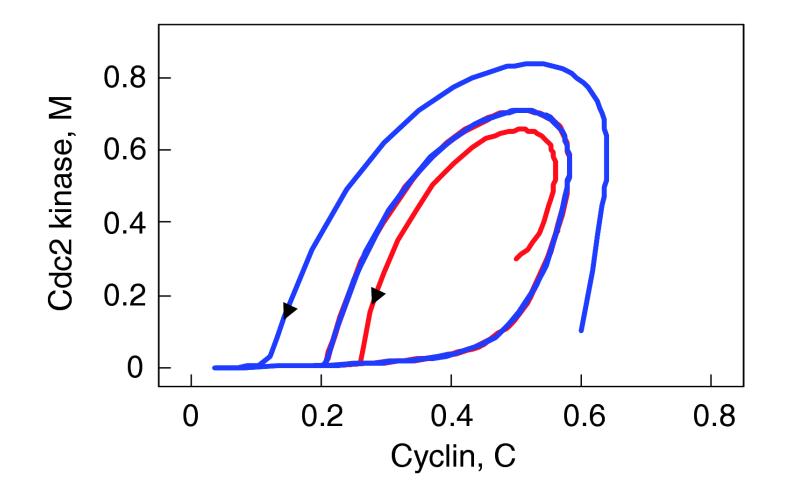
Time courses of cyclin (*C*), active Cdc2 kinase (*M*), and active cyclin protease (*X*) exhibiting oscillations. Initial conditions are X(0) = 0.01. Units: μ M and min-1.







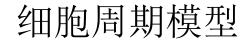




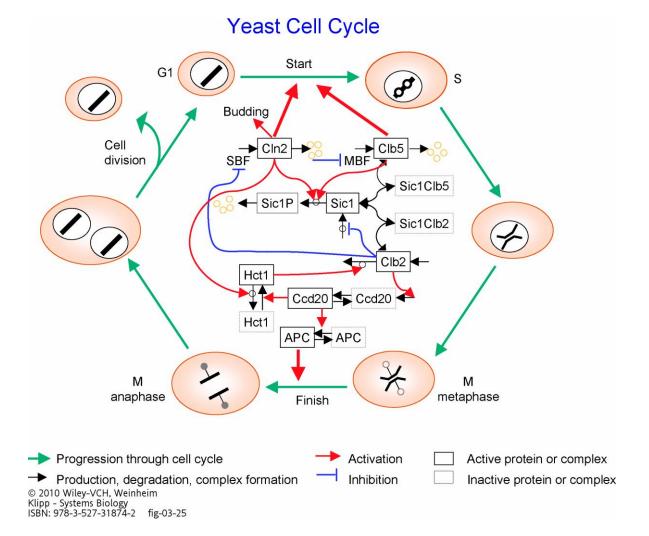
Limit cycle behavior, represented for the variables C and M.











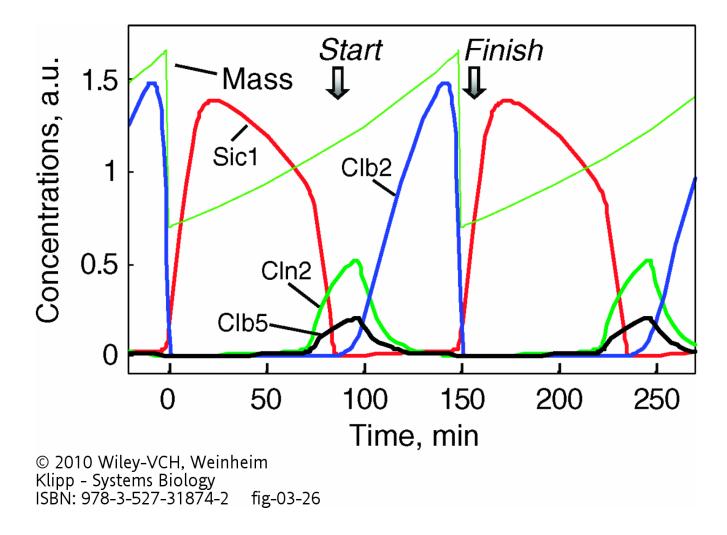
More complicated model











Chen K.C. et al. (2000) Molecular Biology of the Cell, 11. 369-391.

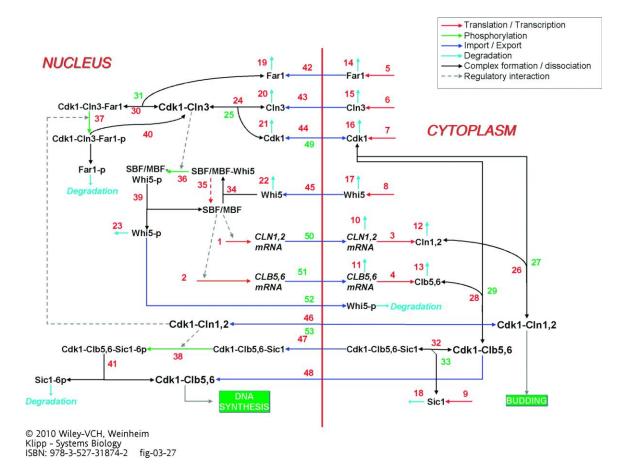




细胞周期模型

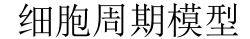


Modeling of Nucleo/Cytoplasmatic Compartmentalization

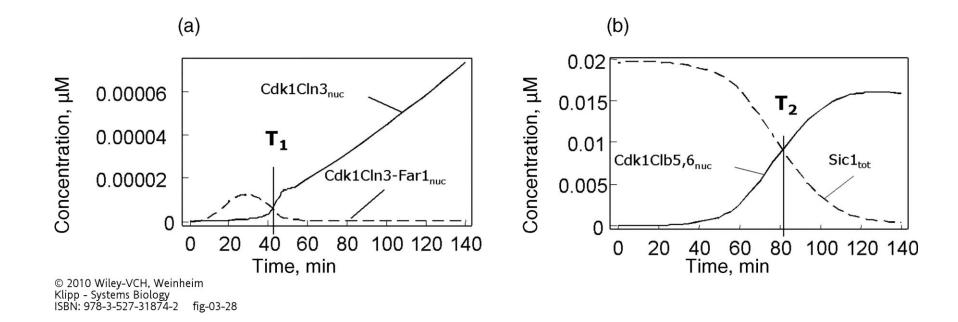


華中震業大学 HUAZHONG AGRICULTURAL UNIVERSITY Barberis M, et al. (2007) PLoS Comput. Biol. 3, e64









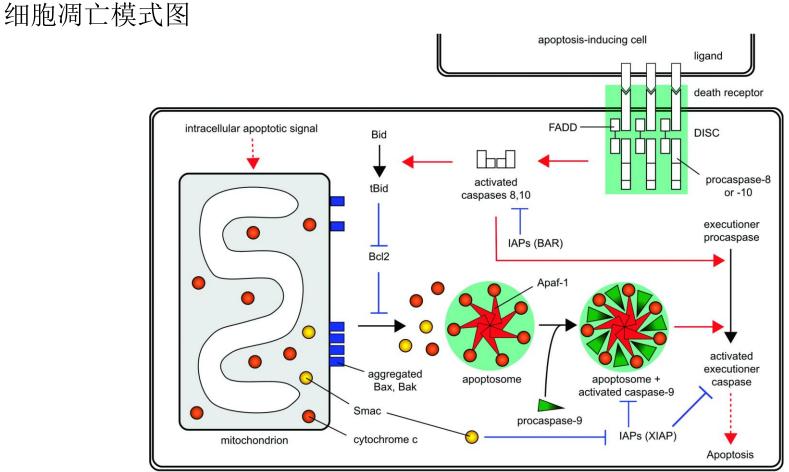
Barberis M, et al. (2007) PLoS Comput. Biol. 3, e64











© 2010 Wiley-VCH, Weinheim Klipp - Systems Biology ISBN: 978-3-527-31874-2 fig-03-34









Table 3.5 Predicted effects of combined therapies based onsimultaneous extrinsic- and intrinsic-induced apoptosis^a.

	Overexpression				Disruption or mutation	
	Bcl-2/Bcl-XL	Bax/Bad/Bik	FLIPs	IAPs	FADD	P53
Bcl-2/Bcl-X _L	_	-	+	+	+	-
Bax/Bad/Bik	-	-	-	+	-	-
FLIPs	+	-	-	+	-	+
IAPs	+	+	+	+	+	+
FADD	+	and _ storad in the		+	-	+
P53		-	-	+	+	-

^aEntries in the diagonal denote therapies with a single target; others are combinations of potential therapies. A plus sign (+) denotes therapies with decreased activation of executioner caspase, and the minus sign (-) denotes the opposite [65].



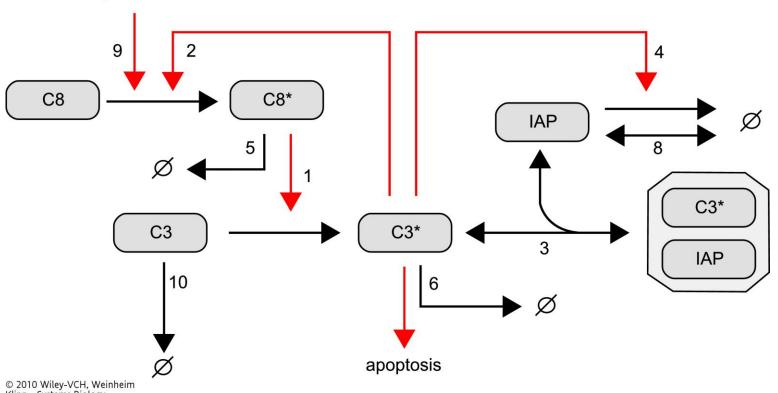






Modeling of Apoptosis

receptor stimulus



© 2010 Wiley-VCH, Weinheim Klipp - Systems Biology ISBN: 978-3-527-31874-2 fig-03-35

Outline of the apoptotic model developed by Eissing *et al.* (JBC 279: 36892-36897). It comprises the components of the extrinsic pathway of apoptosis. The asterisk denotes the activated form of a caspase.









动力学方程 $\frac{d[C8]}{dt} = -v_2 - v_9$ $\frac{d[C8^*]}{dt} = v_2 - v_5$ $\frac{d[C3]}{dt} = -v_1 - v_{10}$ $\frac{d[C3^*]}{dt} = v_1 - v_3 - v_6$ $\frac{d[IAP]}{dt} = -v_3 - v_4 - v_8$ $\frac{d[C3^* \sim IAP]}{dt} = -v_2 - v_9$ $\frac{d[BAR]}{d} = -v_{11} - v_{13}$ $\frac{d[C8^* \sim BAR]}{dt} = -v_{11} - v_{13}$



$$\begin{cases} v_{1} = k_{1} \cdot [C8^{*}] \cdot [C3] \\ v_{2} = k_{2} \cdot [C3^{*}] \cdot [C8] \\ v_{3} = k_{3} \cdot [C3^{*}] \cdot [IAP] - k_{-3} \cdot [C3^{*} \sim IAP] \\ v_{4} = k_{4} \cdot [C3^{*}] \cdot [IAP] \\ v_{5} = k_{5} \cdot [C8^{*}] \\ v_{5} = k_{5} \cdot [C8^{*}] \\ v_{6} = k_{6} \cdot [C3^{*}] \\ v_{7} = k_{7} \cdot [C3^{*} \sim IAP] \\ v_{8} = k_{8} \cdot [IAP] - k_{-8} \\ v_{9} = k_{9} \cdot [C8] - k_{-9} \\ v_{10} = k_{10} \cdot [C3] - k_{-10} \\ v_{11} = k_{11} \cdot [C8^{*}] \cdot [BAR] - k_{-11} \cdot [C8^{*} \sim BAR] \\ v_{12} = k_{12} \cdot [BAR] - k_{-12} \\ v_{13} = k_{13} \cdot [C8^{*} \sim BAR] \end{cases}$$









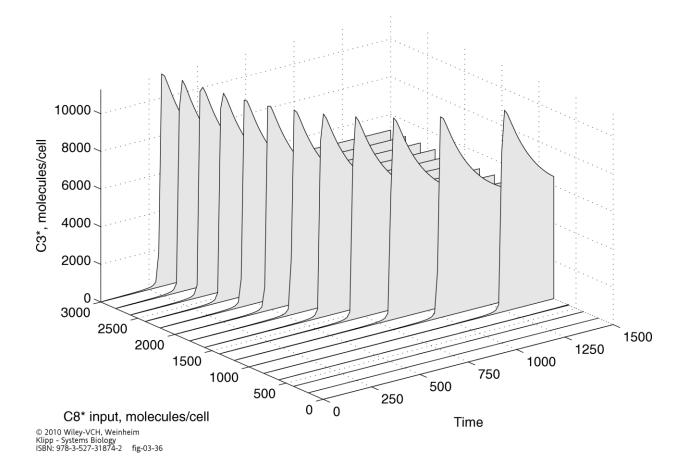
Parameter	Value	Reverse parameter	Value
k ₁	5.8×10^{-5} cell min ⁻¹ mo ⁻¹	K_1	0
k ₂	10^{-5} cell min ⁻¹ mo ⁻¹	k_2	0
k3	5×10^{-4} cell min ⁻¹ mo ⁻¹	k_3	$0.21 \mathrm{min}^{-1}$
k4	3×10^{-4} cell min ⁻¹ mo ⁻¹	k_4	0
ks	$5.8 \times 10^{-3} \text{ min}^{-1}$	k_s	0
k ₆	$5.8 \times 10^{-3} \min^{-1}$	k_6	0
k7	$1.73 \times 10^{-2} \min^{-1}$	K_7	0
k ₈	$1.16 \times 10^{-2} \min^{-1}$	k_8	$464 \mathrm{mo}\mathrm{cell}^{-1}\mathrm{min}^{-1}$
kg	$3.9 \times 10^{-3} \min^{-1}$	k_9	$507 \mathrm{mo}\mathrm{cell}^{-1}\mathrm{min}^{-1}$
k10	$3.9 \times 10^{-3} \min^{-1}$	k_10	$81.9 \mathrm{mo} \mathrm{cell}^{-1} \mathrm{min}^{-1}$
k ₁₁	$5 \times 10^{-4} \text{min}^{-1}$	k_11	$0.21 {\rm min}^{-1}$
k12	$10^{-3} \min^{-1}$	k_12	$40 \mathrm{mo} \mathrm{cell}^{-1} \mathrm{min}^{-1}$
k ₁₃	$1.16 \times 10^{-2} \min^{-1}$	k_13	0











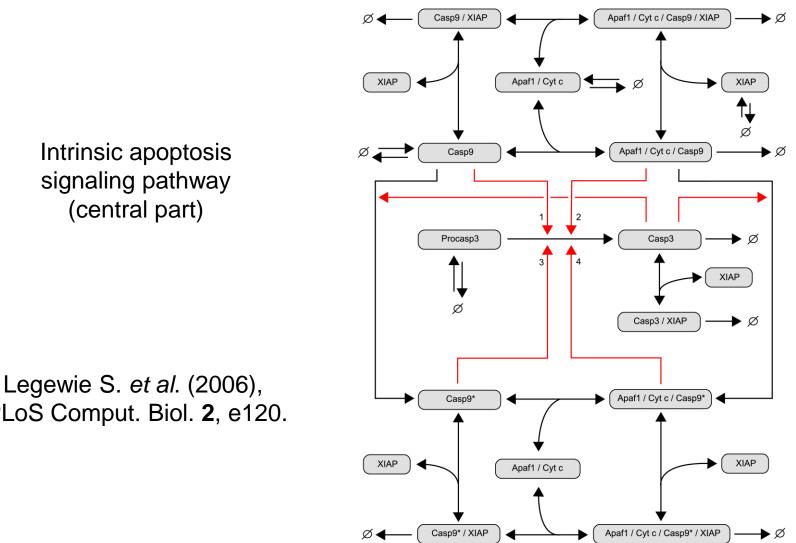
Bistable behavior of the extrinsic apoptosis model versus varying input signals. The input signal is modeled by the initial concentration of the activated caspase-8.













PLoS Comput. Biol. 2, e120.



