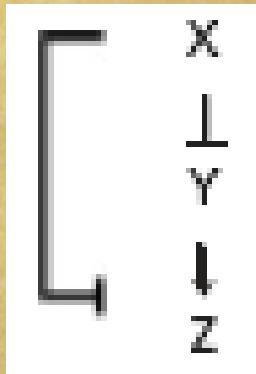


Structure and function of the feed-
forward loop network motif

The Coherent Feedforward Loop
Serves as a Sign-sensitive Delay
Element in Transcription Networks

Mangan, Zaslaver Alon

Oct, Nov 2003 PNAS&JMB



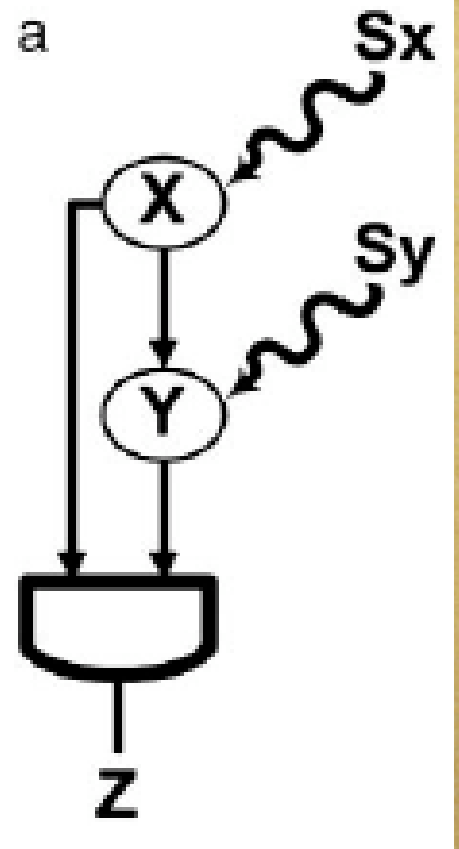
FeedForward

*Etude théorique du filtrage
dynamique*

*Vérification par mesures invivo
du module L-arabinose E.Coli*

Mangan, Zaslaver Alon

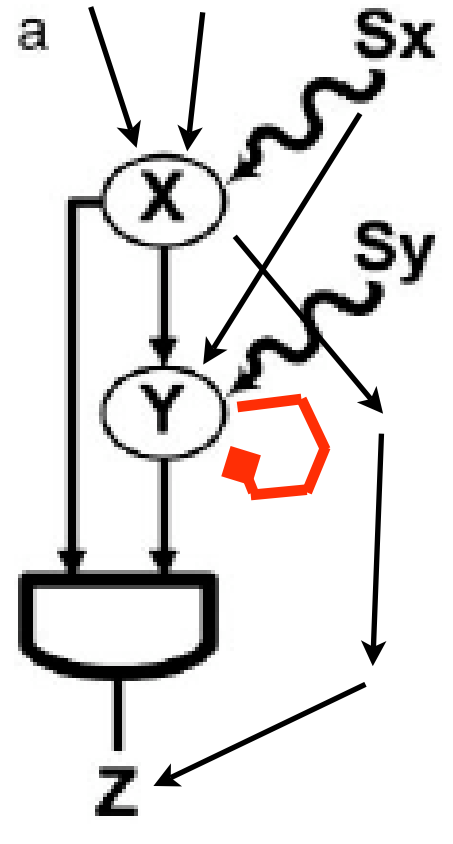
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Définitions

- X, Y régulent Z
- $+Inducteurs$
- (Activation \vee Inhibition) $\rightarrow 2^8$

Définitions-Hypothèses



- X, Y régulent Z
- $+Inducteurs$
- (Activation+ ou Inhibition-)= $2^3=8$
- *Module isolable*
- (In)Coherent: $+=+++$, $-=-+$...
- *Porte logique 'AND'*

(ou un 'OR')

	0	1
0	0	1
1	1	2

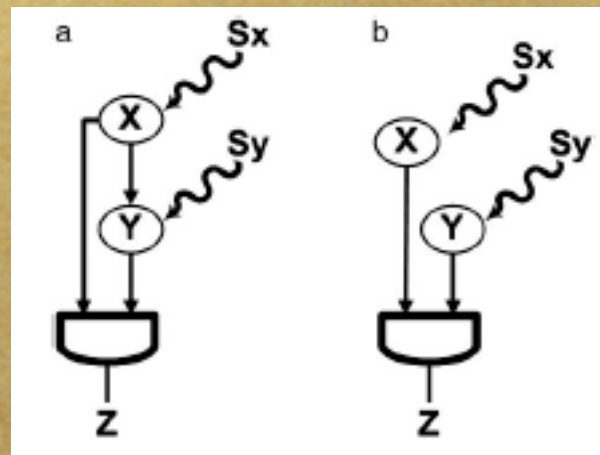
Propriété: Accélérer le switch (ON xou OFF)

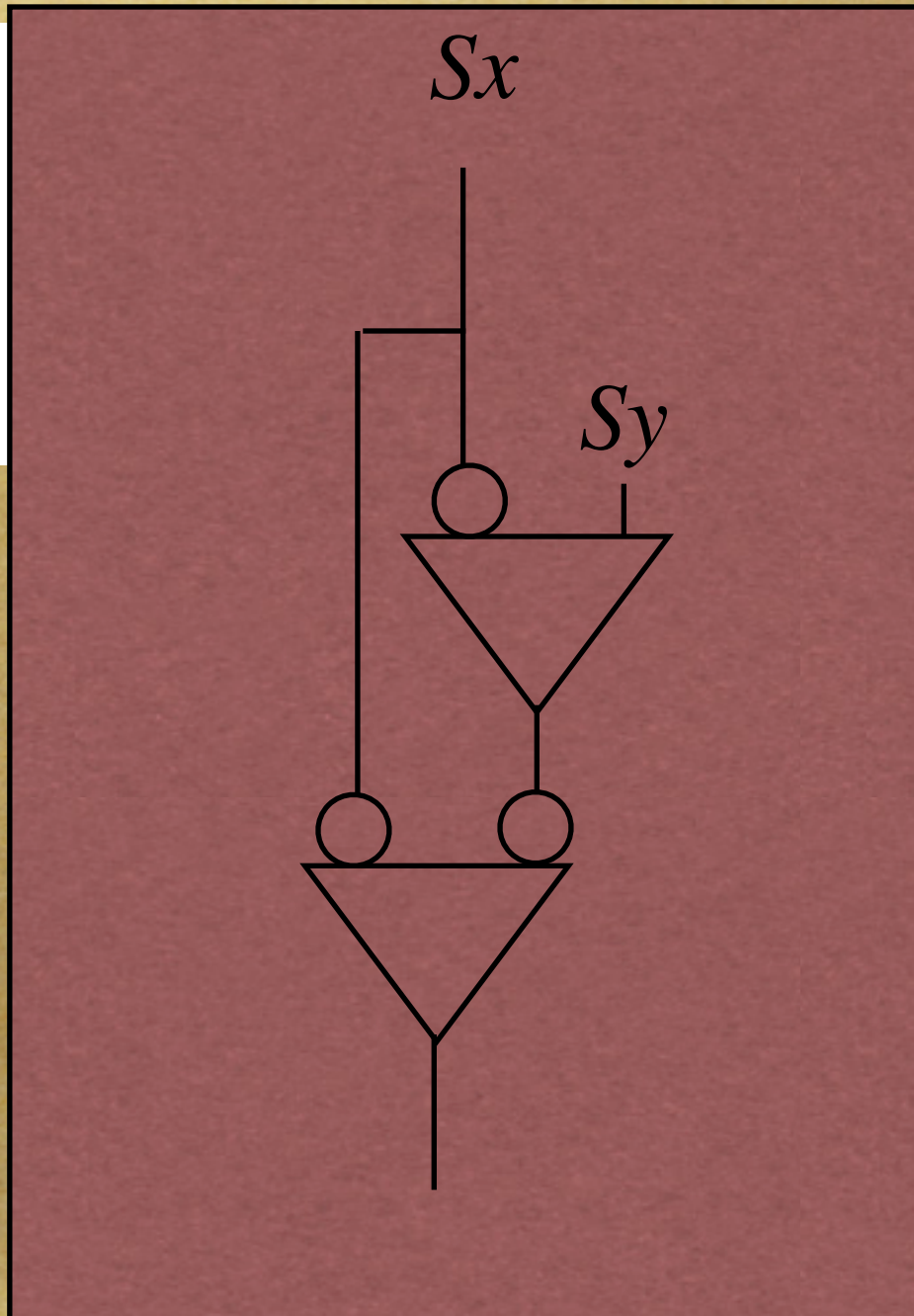
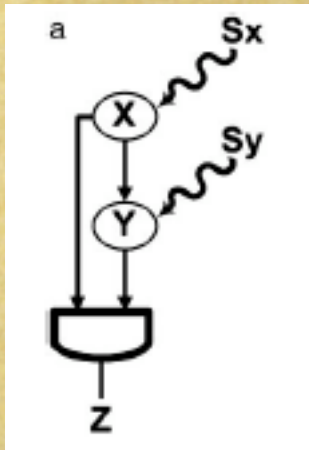
Occurrences

Species	Coherent type 1		Coherent type 2		Coherent type 3		Coherent type 4	
	Structure	Abundance	Structure	Abundance	Structure	Abundance	Structure	Abundance
<i>E. coli</i>		28		2		4		1
<i>S. cerevisiae</i>		26		5		0		0

	S_x	
	S_y	
$\&$	S_x	$\!S_x$
S_y	1	0
$\!S_y$	1	0

Species	Incoherent type 1		Incoherent type 2		Incoherent type 3		Incoherent type 4	
	Structure	Abundance	Structure	Abundance	Structure	Abundance	Structure	Abundance
<i>E. coli</i>		5		0		1		1
<i>S. cerevisiae</i>		21		3		1		0





	y	$!y$
x		
$!x$		

2 entrées

8 cablages

*4 transitions pour
chaque cablage
= 32 'résultats'*

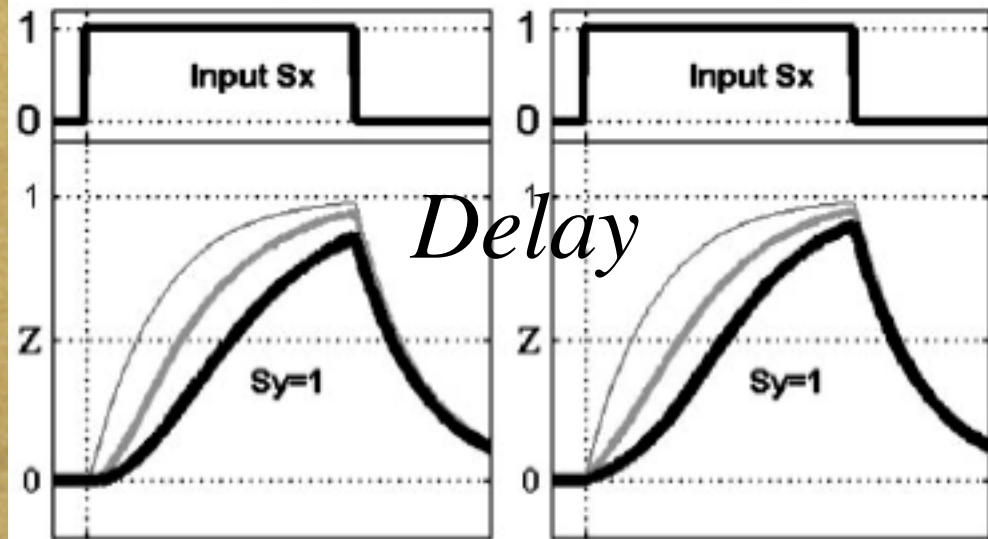
1 Sortie

Coherent Type 1 AND

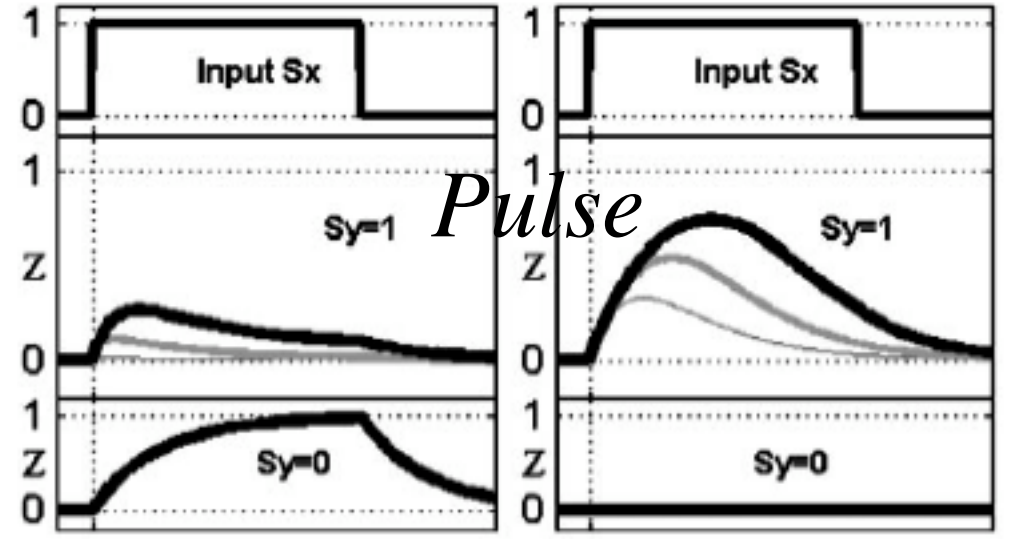
Coherent Type 4 AND

Incoherent Type 1 AND

Incoherent Type 4 AND



Delay



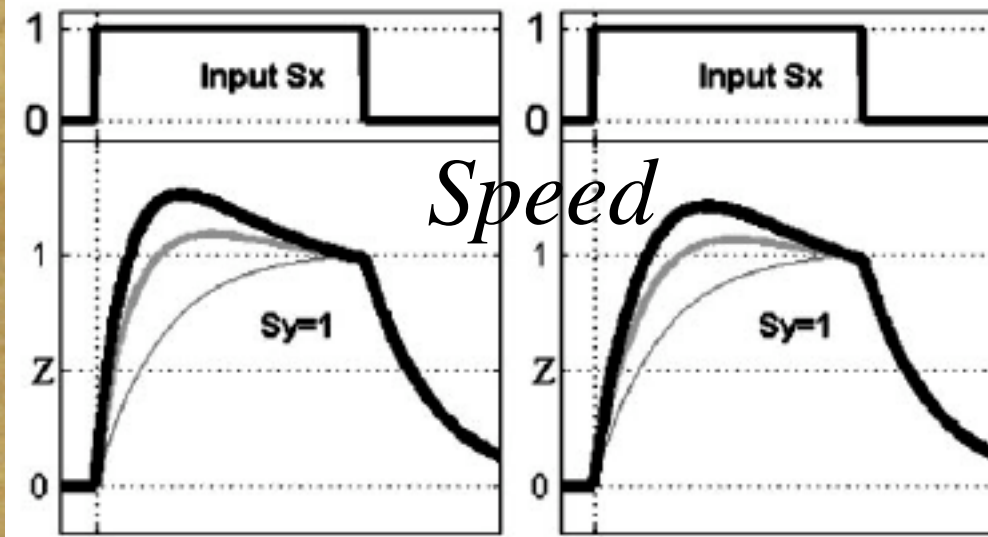
Pulse

Time [Life Time]
Incoherent Type 1 AND

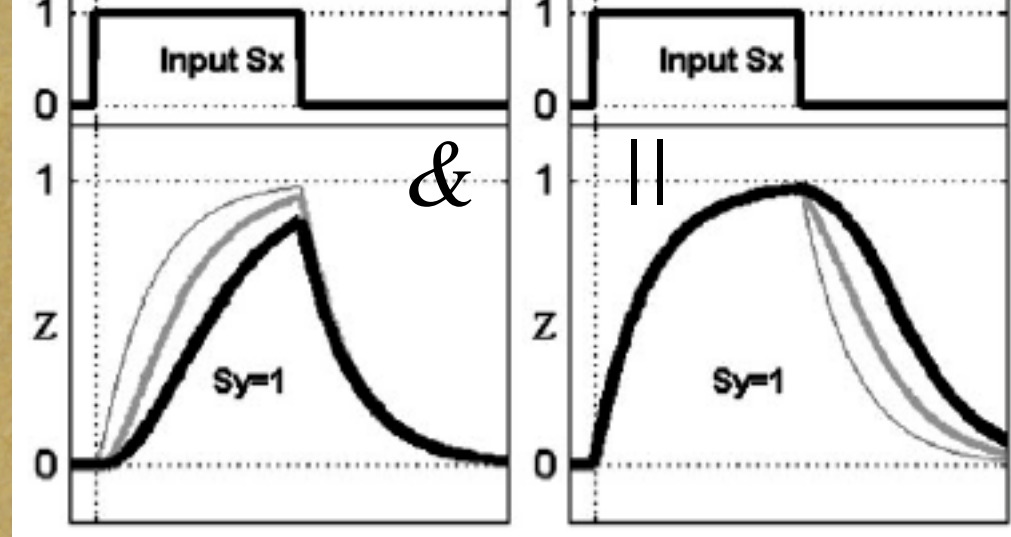
Time [Life Time]
Incoherent Type 4 AND

Time [Life Time]
Coherent Type 1 AND

Time [Life Time]
Coherent Type 1 OR



Speed



&

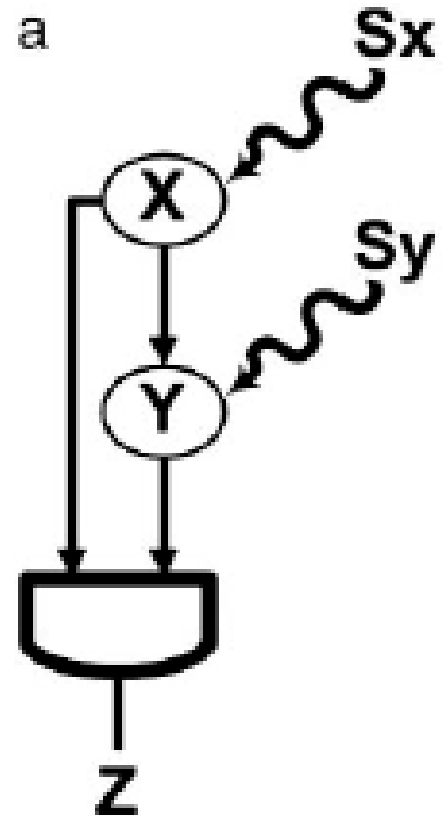
||

Time [Life Time]

Time [Life Time]

Time [Life Time]

Time [Life Time]



$$dY/dt = B_y + \beta_1 f(X, K_{xy}) - \alpha_1 Y,$$

$$dZ/dt = \beta_2 f(X, K_{xz}) f(Y, K_{yz}) - \alpha_2 Z$$

Activateur $f(u, K) = u^H / (K^H + u^H)$

Inhibiteur $f(u, K) = 1 / (1 + (u/K)^H)$

AND

$$G_z = f(X^*, K_{xz}) f(Y^*, K_{yz})$$

OR (compétitif)

$$G_z = f_c(X^*; K_{xz}, K_{yz}, Y^*) + f_c(Y^*; K_{yz}, K_{xz}, X^*)$$

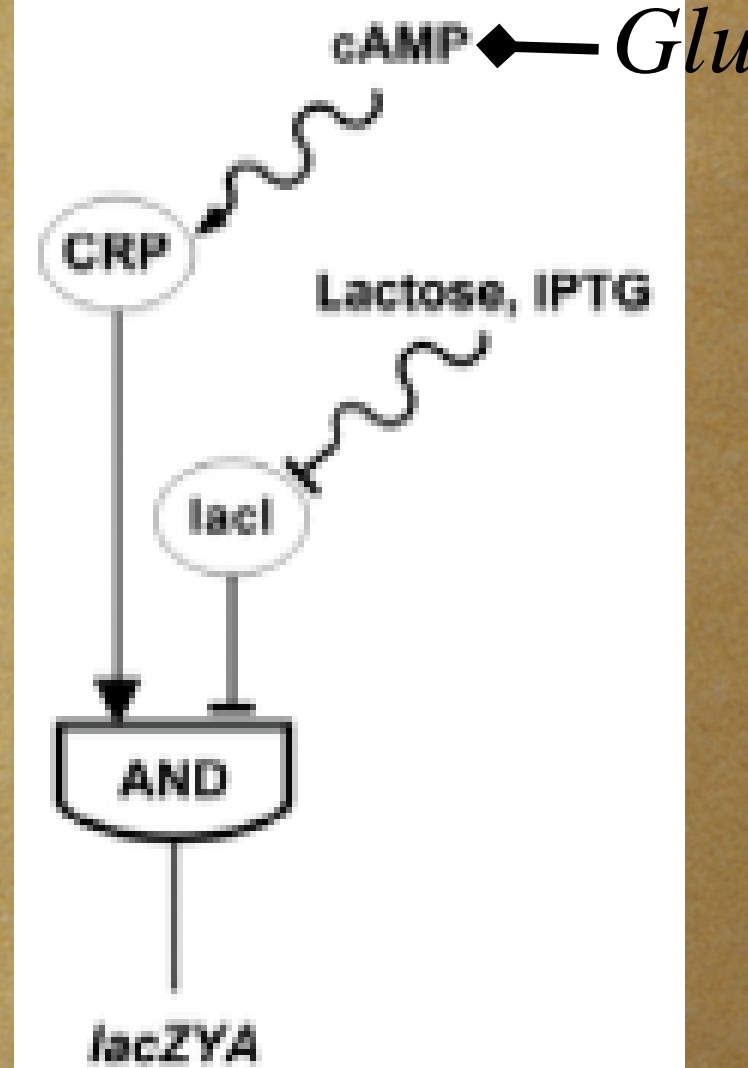
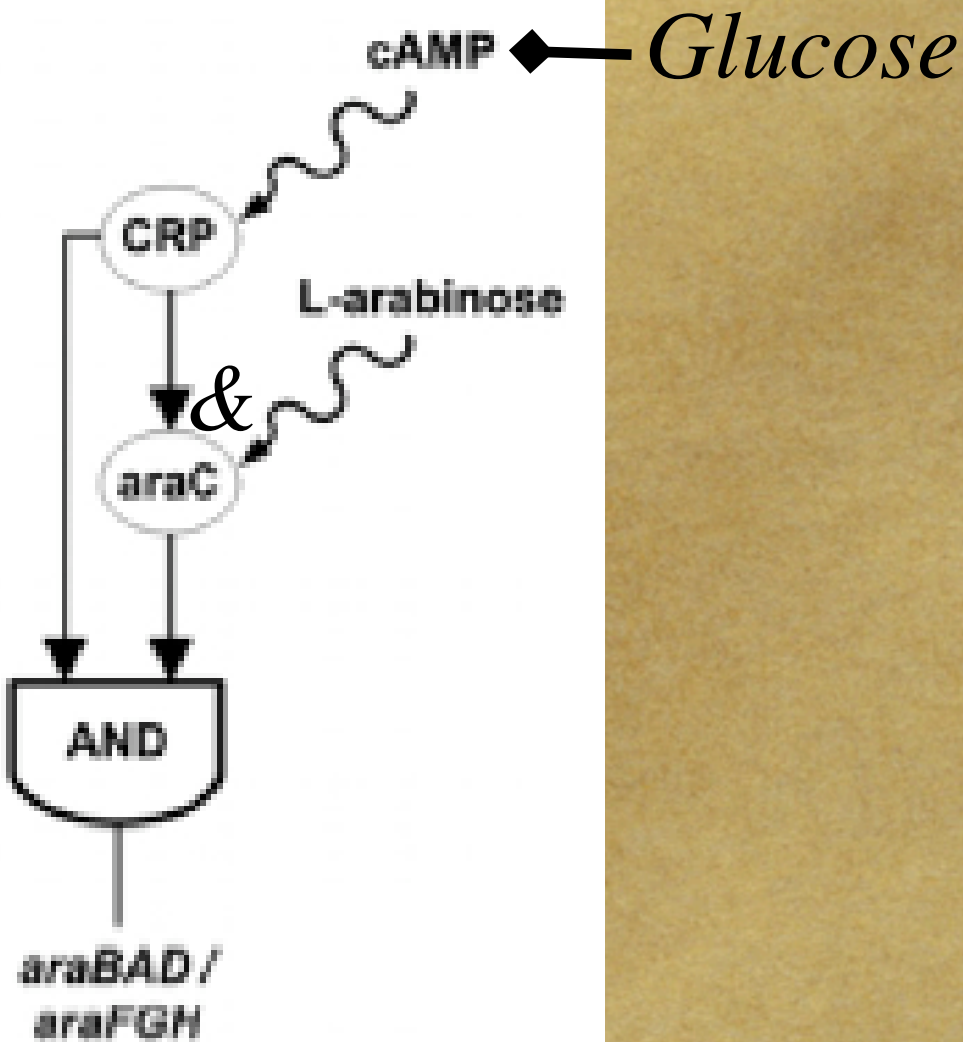
Activateur = $(u/K_u)^H / (1 + (u/K_u)^H + (v/K_v)^H)$

Inhibiteur = $1 / (1 + (u/K_u)^H + (v/K_v)^H)$

FFL, L-arabinose

vs

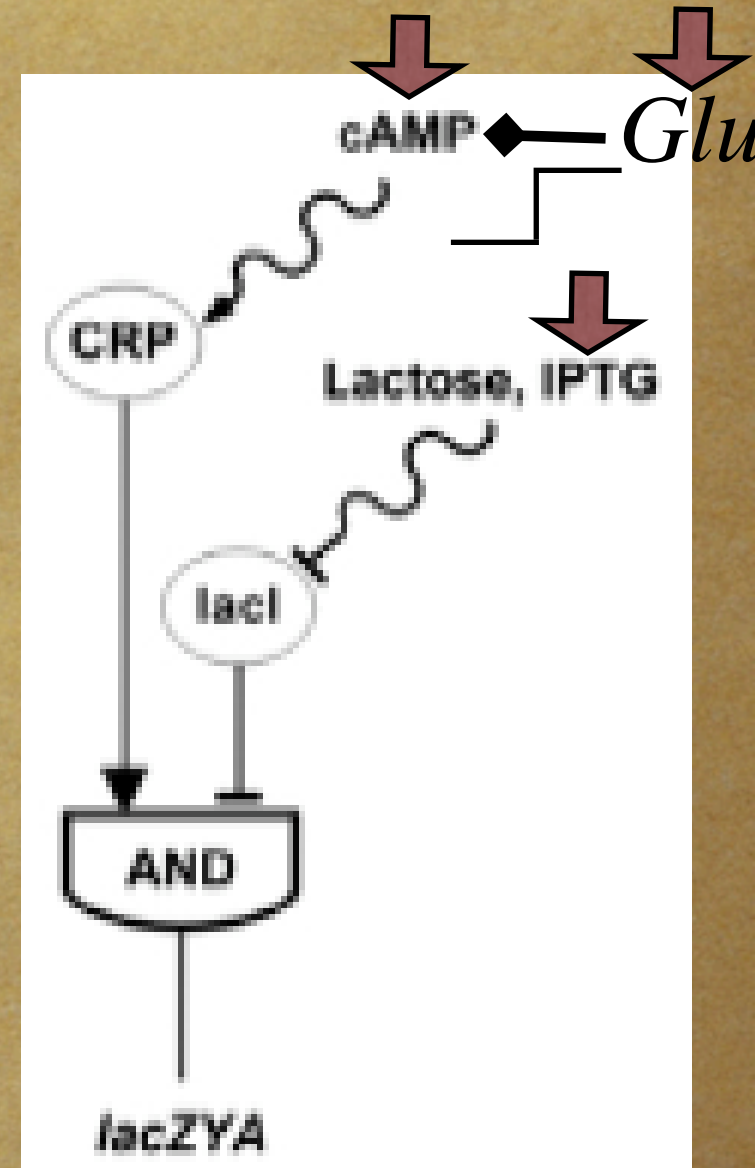
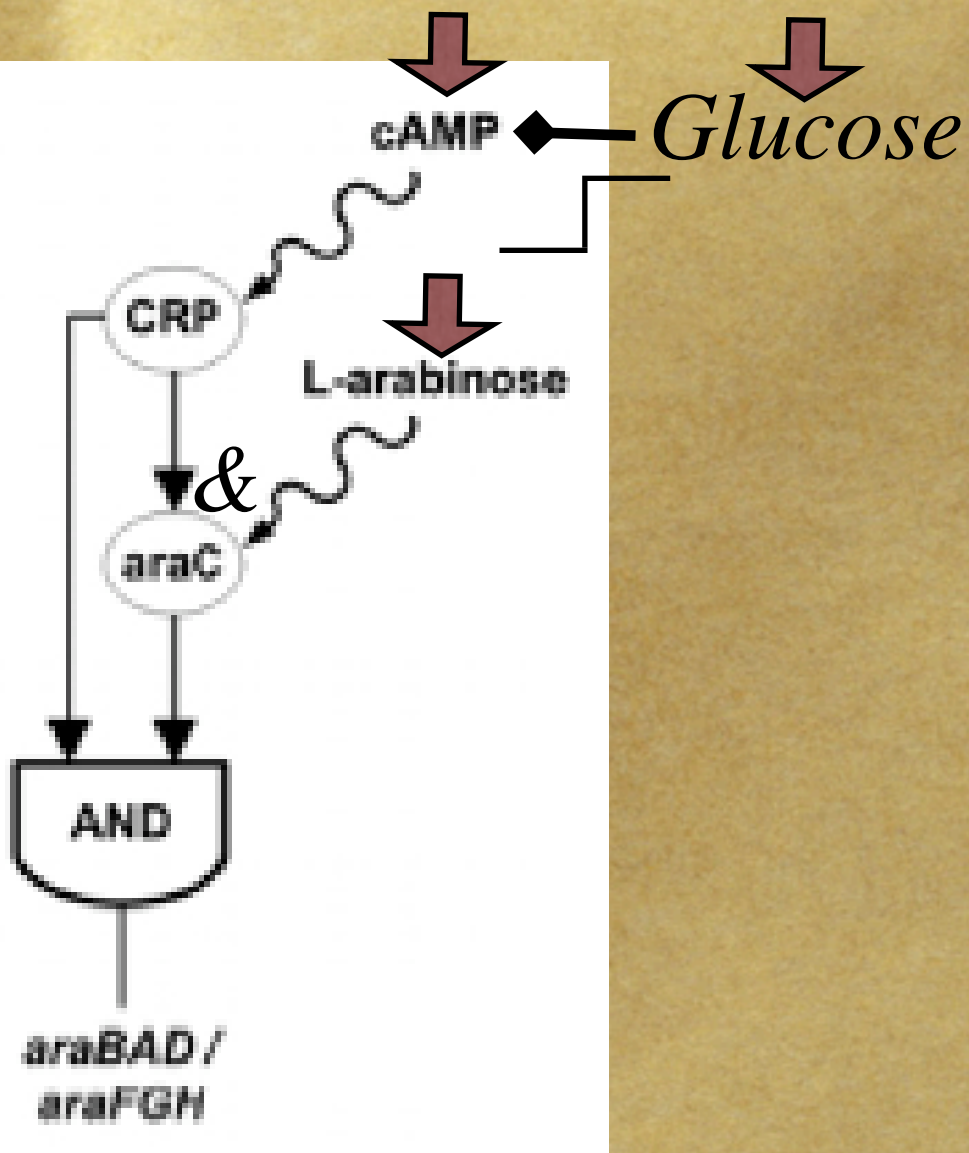
Test



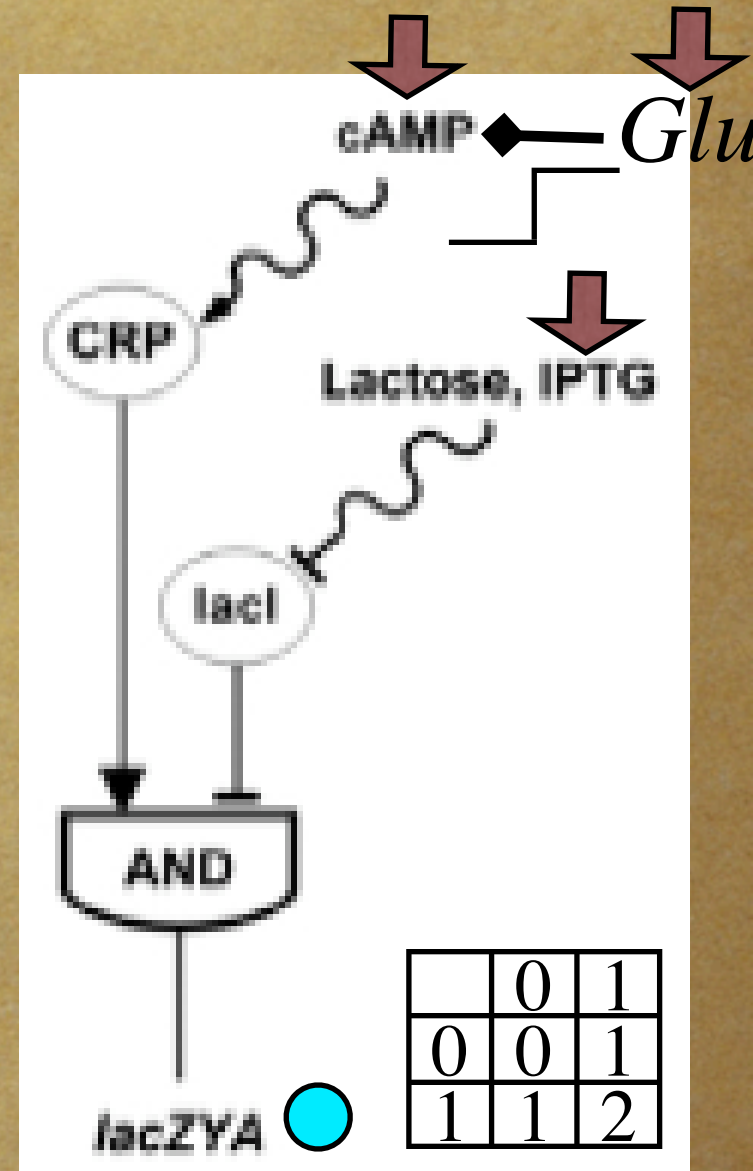
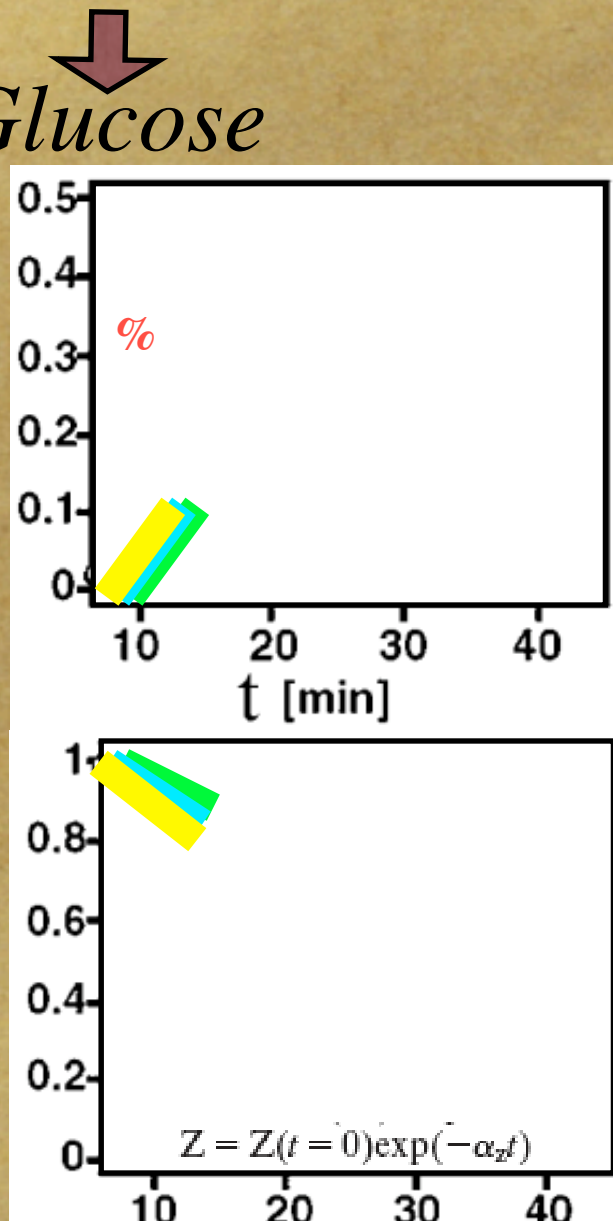
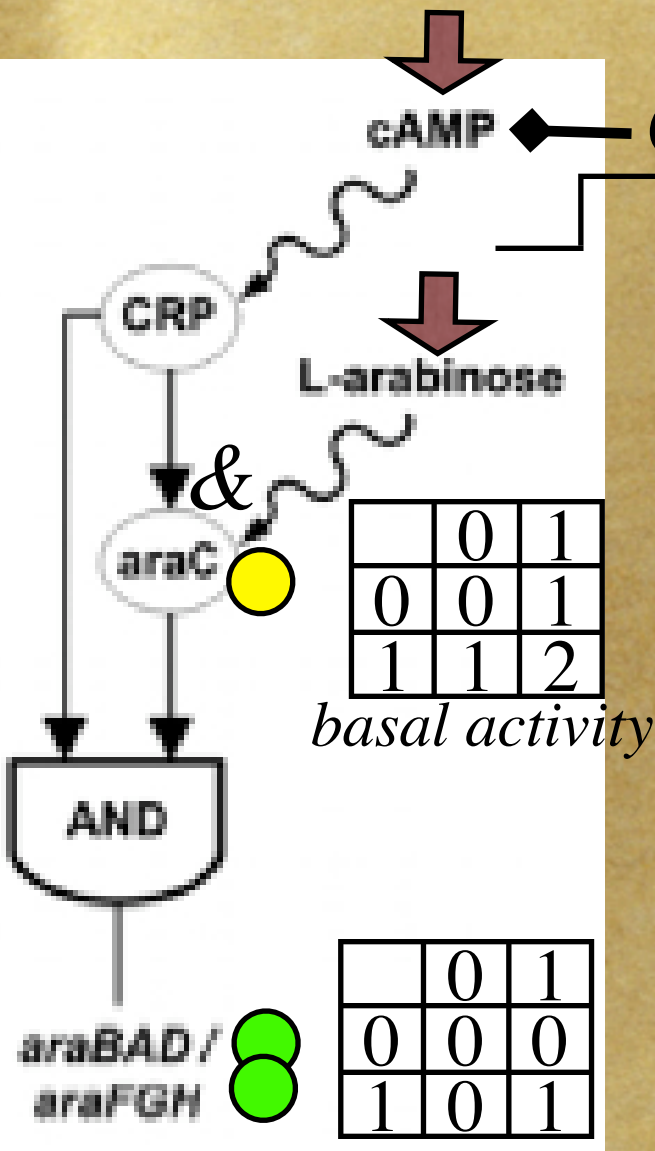
FFL, L-arabinose

vs

Test

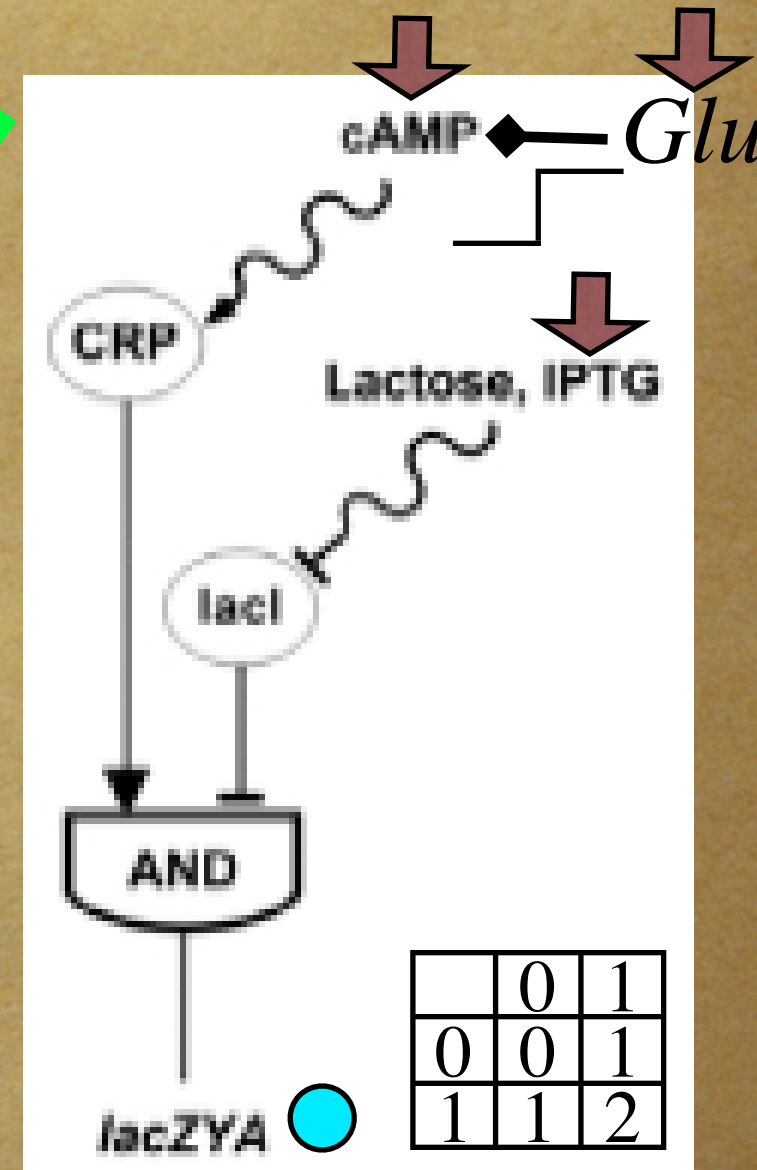
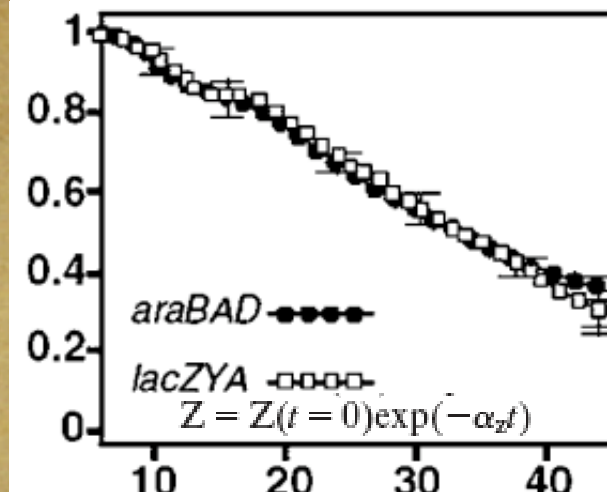
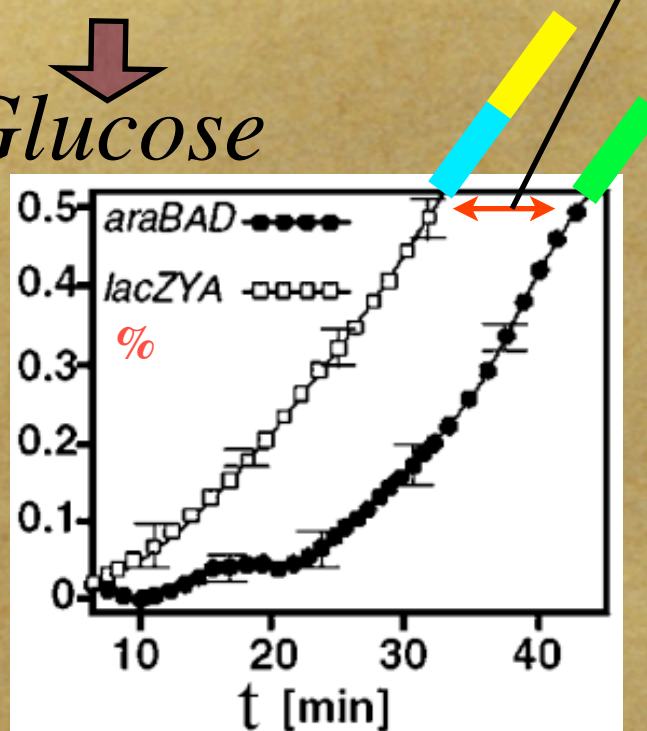
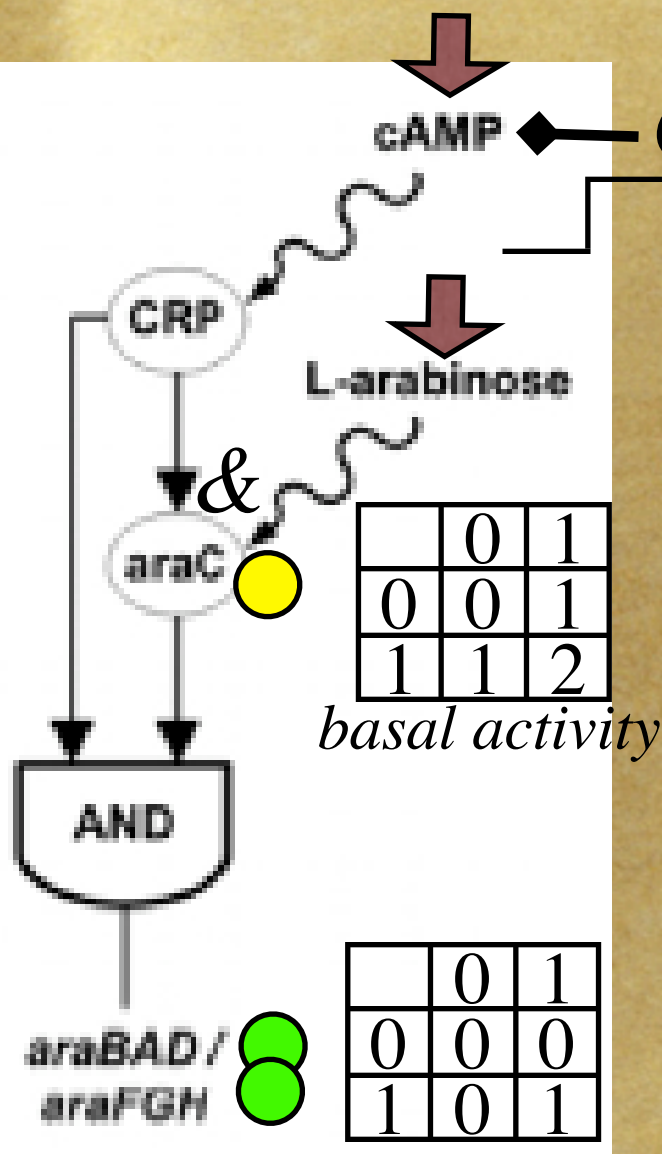


FFL, L-arabinose vs Test

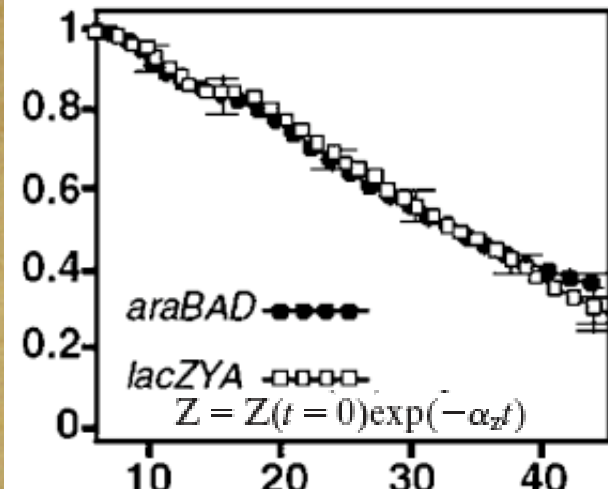
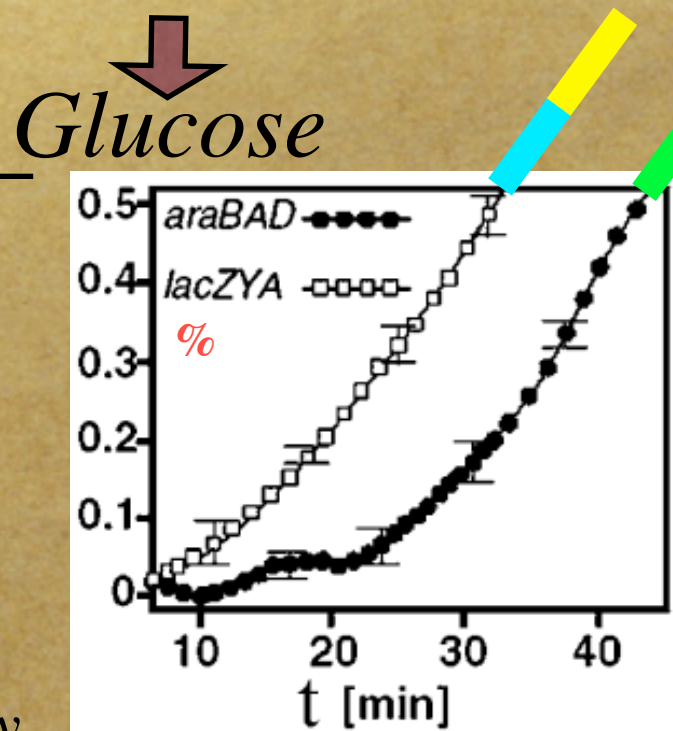
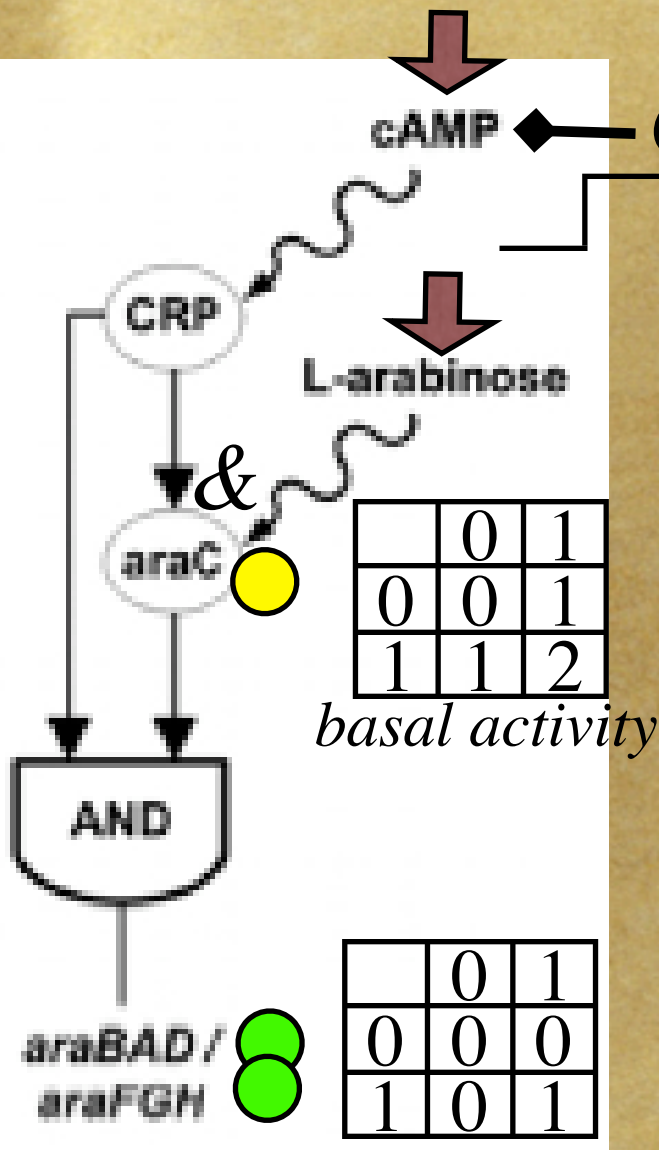


FFL, L-arabinose vs Test

$$t_D = \alpha_y^{-1} \log[(Y_{\max} - Y_{\min}) / (Y_{\max} - K_{yz})]$$



FFL, L-arabinose



1) Dans intestin
 L-arabinose(++)
 + pulse de Glucose
 -> Stop rapide ~
 -> réenclenchement lent
 mais stock de ara

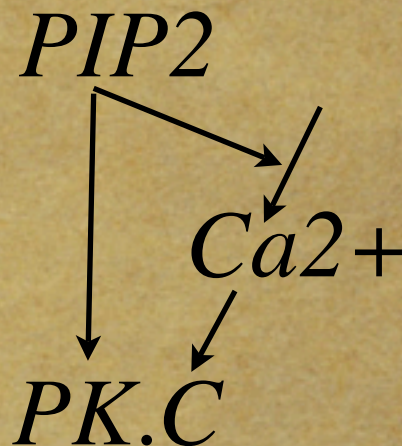
Fructose(++)(FFL)
 Maltose(++)(FFL)
 Lactose(--)

Filtre signal de glucose bruité
 2)?

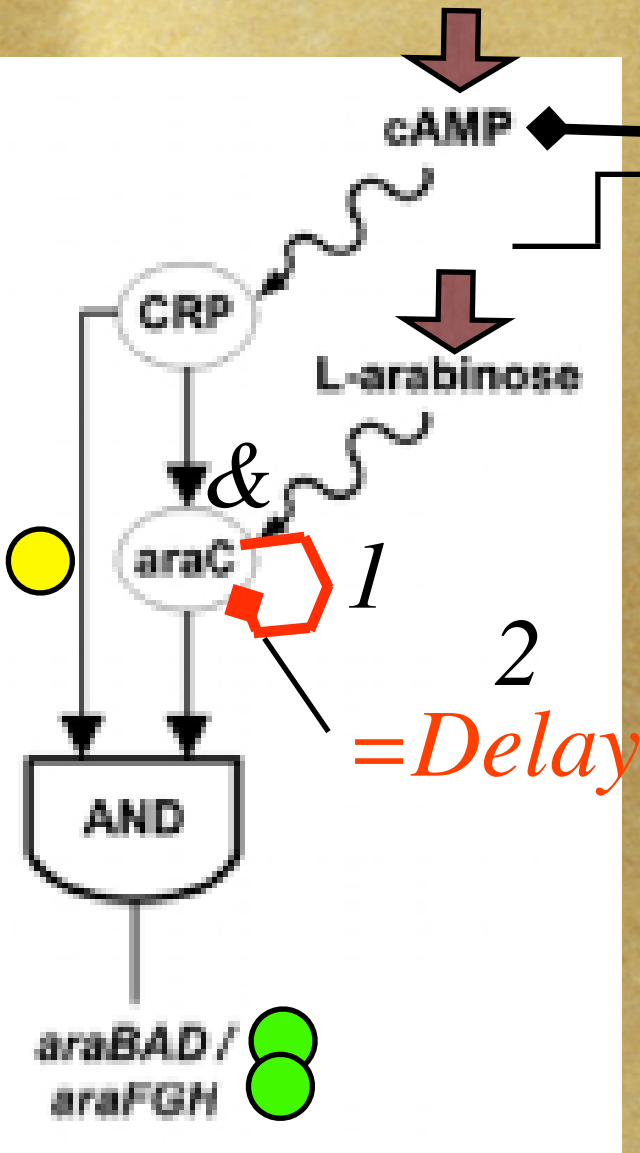
Où sont les FFL ?

- *FFL E.coli*: Arabinose, Fructose, Maltose, Azote, drugs.
- *FFL S.cerevisiae*: Même module, (même fonction), mais protéines différentes.
- *C.elegans*, Oursin, développement, neurones, Homme...

- *PPI net*:



Autres informations connues



3

T.M. Dunn, S. Hahn, S. Ogden and R.F. Schleif, An operator at -280 base pairs that is required for repression of araBAD operon promoter: addition of DNA helical turns between the operator and promoter cyclically hinders repression. *Proc. Natl Acad. Sci. USA* **81** (1984),

