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1. Model

The model for our AND-Gate is based on the model of the iGEM team PKU Beijing 2009 for their AND-Gate1. We modified the equations such that the change in tRNA and Aa-tRNA does not include the degradation of the mRNA which caused negativity of some concentrations in our model.[1]

For more details see the paper by Anderson [2].

the dye output was an adaption of the model propose in the paper by Yildirim[3].



2. Equations

tRNA	\dot{x}_1	=	$k_t \frac{\left(\frac{OmpR-P}{K1}\right)^2}{\left(1+\frac{OmpR-P}{K1}\right)^2} - (\gamma_1 + k_a)x_1 + \gamma_{2p}x_2 + 2k_{7p}x_3\left(\frac{\gamma_3}{k_{7m}}\right)\left(\frac{x_1}{\gamma_0 + x_1}\right)^2$
Aa - tRNA	\dot{x}_2	=	$k_a x_1 - 2k_{7p} x_3 \left(rac{\gamma_3}{k_{7m}}\right) \left(rac{x_1}{\gamma_0 + x_1}\right)^2 - \gamma_2 x_2$
$T7RNAP_{mRNA}$	\dot{x}_3	=	$k_{7m} \left(1 - rac{\left(rac{Y cg E}{K3} ight)^2}{\left(1 + rac{Y cg E}{K3} ight)^2} ight) - \gamma_3 x_3$
T7RNAP	\dot{x}_4	=	$k_{7p}x_3\left(rac{\gamma_3}{k_{7m}} ight)\left(rac{x_1}{\gamma_0+x_1} ight)^2-\gamma_4x_4$
$lacZ_{mRNA}$	\dot{x}_5	=	$\alpha_M \left(1 - \frac{\left(\frac{x_4}{K_5}\right)^2}{\left(1 + \frac{x_4}{K_5}\right)^2} \right) - \gamma_M x_5$
$\beta-Galactosidase$	\dot{x}_6	=	$\alpha_B x_5 - \gamma_B x_6$
dye	\dot{x}_7	=	$lpha_A x_6$

Parameter	Value	Unit	it Name	
k_t	$\frac{46.67}{60}$	$\frac{nM}{s}$	$\frac{nM}{s}$ max transcription rate tRNA	
ka	$\frac{0.08}{60}$	$\frac{1}{s}$	synthesis rate Aa-tRNA	[1]
k_{7p}	$\frac{1.5625}{60}$	$\frac{nM}{s}$	max transcription rate T7RNAP	[1]
k_{7m}	$\frac{268*0.05}{60}$	$\frac{1}{s}$	max translation rate T7RNAP	[1]
k_S	0.3	$\frac{1}{nM}$	AND Gate rate	[1]
γ_0	1	-	threshold Aa-tRNA	guessed
γ_1	$\frac{1}{60*60}$	$\frac{1}{s}$	degradation of tRNA	[1]
γ_2	$\frac{1}{40*60}$	$\frac{1}{s}$	degradation of Aa-tRNA	[1]
γ_3	$\frac{1}{4.4*60}$	$\frac{1}{s}$	degradation of T7RNAP mRNA	[1]
γ_4	$\frac{46.67}{40*60}$	$\frac{1}{s}$	degradation of T7RNAP	[1]
<i>K</i> 1	5	nM	nM response param. OmpR-P,tRNA	
K3	600	nM	response param. YcgE,T7RNAP	guessed
K5	$\frac{k7p}{4*\gamma_4}$	nM	response param T7RNAP,lacZ	guessed
α_M	$\frac{0.997}{60}$	$\frac{nM}{s}$	max transcription rate lacZ	[3]
α_B	$\alpha_B \qquad \frac{1.661e-5}{60} \qquad \frac{1}{s} \qquad \text{max translation rate lacZ}$		[3]	

3. PARAMETERS

Parameter	Value	e Unit Name		Source
α_A	$\frac{20}{60}$	$\frac{1}{s}$	enzymatic reaction rate	[3]
γ_M	$\frac{0.411}{60}$	$\frac{1}{s}$	degradation lacZ mRNA	[3]
γ_B	$\tfrac{8.331e-4}{60}$	$\frac{1}{s}$	degradation β -Galactosidase	[3]

4. INITIAL DATA

Name	Variable	Initial Value	Comment	Source
tRNA	x_1	0		
Aa - tRNA	x_2	0		
$T7RNAP_{mRNA}$	x_3	0		
T7RNAP	x_4	0		
$lacZ_{mRNA}$	x_5	0		
$\beta - Galactosidase$	$\overline{x_6}$	0		
dye	x_7	0		

5. SIMULATION

Since the AND-Gate is only an intermediate part, no additional simulation was done here.

References

- 1. PKU Beijing 2009, And gate 1, 2009.
- 2. J Christopher Anderson, Christopher A Voigt, and Adam P Arkin, *Environmental signal integration by a modular and gate*, Mol Syst Biol **3** (2007).
- 3. N Yildirim, M Santillan, D Horike, and MC Mackey, *Dynamics and bistability in a reduced model of the lac operon*, CHAOS **14** (2004), no. 2, 279–292 (English).