

PART 1 Original Full Model

At our first step, we wanted to describe the system thoroughly without leaving out any seemingly unimportant actions and factors. As a result, the description of the system contains every possible mass actions as well as some hill kinetics, Henri-Michaelis-Menten. We came up a set of ODEs with 19 equations.

Construction of ODE equation

CELL I

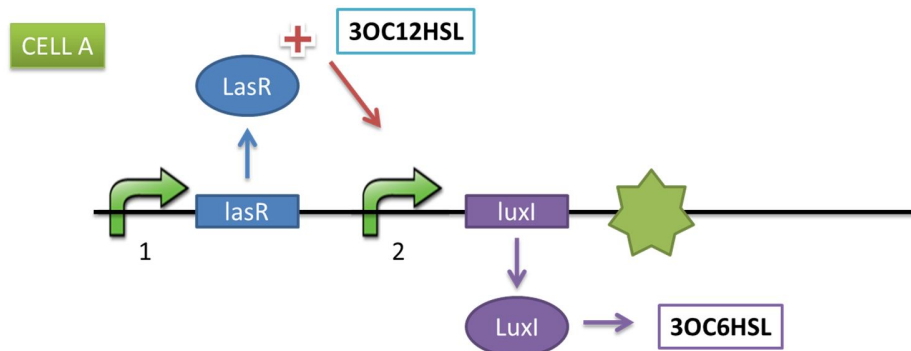


Figure 1 designed circuit of cell I

CELL II

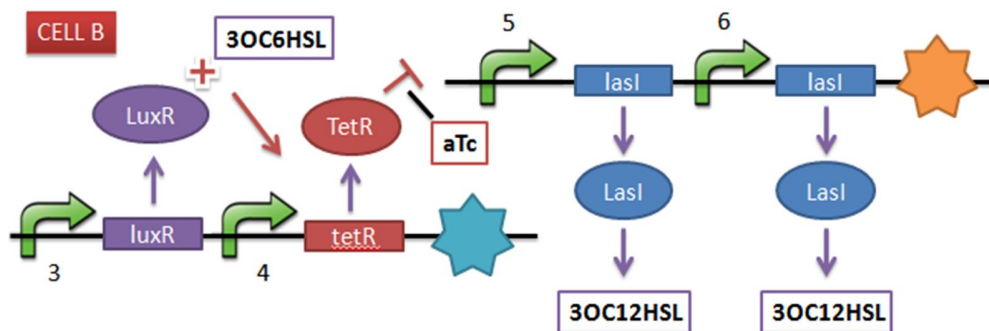


Figure 2 designed circuit of cell II

Promoter 1 and promoter 2 preceding *lasR* and *luxR* genes respectively are constant promoters, which will transcribe and translate into protein *PlasR* and *PluxR*. $LA1$ is the binding association of *lasR* and $3OC12HSL(A2C1)$ and it can affect the subsequent promoter 2 which can be described by Hill Equation. The same goes to $LA2$. Gene *luxI* will be translated into protein *PluxI* which would generate $3OC6HSL(A1C1)$ through enzymatic reaction. The AHL will diffuse through the membrane to the environment ($A1e$) and finally enter into Cell 2 ($A1C2$). Protein *PtetR* which is translated from gene *tetR* represses promoter 5 which is responsible for transcription of gene *lasI*. Promoter 6 is constant for translation of protein *PlasI*. $3OC12HSL(A2C2)$ is generated from Protein *PlasI* through enzymatic reaction. $3OC12HSL$ in the environment is called $A2e$ which will diffuse to Cell 1. *aTc* is

added manipulatively to change the phase of oscillation by binding the protein PTetR. Therefore, we have these following ODEs:

$$\frac{dM_{lasR}}{dt} = v_{M_{lasR}} - d_{M_{lasR}} \times M_{lasR} \quad (1)$$

$$\frac{dM_{luxI}}{dt} = k_{M_{luxI}} \times [\beta_{M_{luxI}} + (1 - \beta_{M_{luxI}}) \times \frac{LA1^{n1}}{K_{M1}^{n1} + LA1^{n1}}] - d_{M_{luxI}} \times M_{luxI} \quad (2)$$

$$\frac{dP_{lasR}}{dt} = k_{TL1} \times M_{lasR} - d_{P_{lasR}} \times P_{lasR} - k1 \times A1_{c1} \times P_{lasR} + k2 \times LA1 \quad (3)$$

$$\frac{dP_{luxI}}{dt} = k_{TL2} \times M_{luxI} - d_{P_{luxI}} \times P_{luxI} \quad (4)$$

$$\frac{dLA1}{dt} = k1 \times A1_{c1} \times P_{lasR} - k2 \times LA1 \quad (5)$$

$$\frac{dA1_{c1}}{dt} = -k1 \times A1_{c1} \times P_{lasR} + k2 \times LA1 + \gamma \cdot (A1_e - A1_{c1}) \quad (6)$$

$$\frac{dA2_{c1}}{dt} = \lambda_1 P_{luxI} + \gamma \cdot (A2_e - A2_{c1}) \quad (7)$$

$$\frac{dM_{luxR}}{dt} = v_{M_{luxR}} - d_{M_{luxR}} \times M_{luxR} \quad (8)$$

$$\frac{dM_{TetR}}{dt} = k_{MTetR} \times [\beta_{MTetR} + (1 - \beta_{MTetR}) \times \frac{LA2^{n2}}{K_{M2}^{n2} + LA2^{n2}}] - d_{MTetR} \times M_{TetR} \quad (9)$$

$$\frac{dM_{lasI}}{dt} = k_{M_{lasI}} \times [\beta_{M_{lasI}} + (1 - \beta_{M_{lasI}}) \times \frac{K_{MT}^{n3}}{K_{M3}^{n3} + TetR^{n3}}] - d_{M_{lasI}} \times M_{lasI} \quad (10)$$

$$\frac{dP_{luxR}}{dt} = k_{TL3} \times M_{luxR} - d_{P_{luxR}} \times P_{luxR} - k3 \times A2_{c2} \times P_{luxR} + k4 \times LA2 \quad (11)$$

$$\frac{dP_{lasI}}{dt} = k_{TL4} \times M_{lasI} - d_{P_{lasI}} \times P_{lasI} \quad (12)$$

$$\frac{dP_{TetR}}{dt} = k_{TL5} \times M_{TetR} - d_{PTetR} \times P_{TetR} - k5 \times P_{TetR} \times aTc + k6 \times TetR^* \quad (13)$$

$$\frac{dLA2}{dt} = k3 \times A2_{c2} \times P_{luxR} - k4 \times LA2 \quad (14)$$

$$\frac{dA2_{c2}}{dt} = -k3 \times A2_{c2} \times P_{luxR} + k4 \times LA2 + \gamma \cdot (A2_e - A2_{c2}) \quad (15)$$

$$\frac{dTetR^*}{dt} = k5 \times P_{TetR} \times aTc - k6 \times TetR^* \quad (16)$$

$$\frac{dA1_{c2}}{dt} = \lambda_2 P_{lasI} + \gamma \cdot (A1_e - A1_{c2}) \quad (17)$$

$$\frac{dA1_e}{dt} = -\gamma \frac{1 - p \cdot (1 + n_{12})}{p \cdot n_{12}} \cdot (A1_e - A1_{c1}) - \gamma \cdot \frac{1 - p \cdot (1 + n_{12})}{p} \cdot (A1_e - A1_{c2}) - \mu A1_e \quad (18)$$

$$\frac{dA2_e}{dt} = -\gamma \frac{1 - p \cdot (1 + n_{12})}{p \cdot n_{12}} \cdot (A2_e - A2_{c1}) - \gamma \cdot \frac{1 - p \cdot (1 + n_{12})}{p} \cdot (A2_e - A2_{c2}) - \mu A2_e \quad (19)$$

Parameters

The parameters are inherent factors determining the behaviors, properties of a system. We selected the quantities thoughtfully from previous iGEM teams and some others were found from published papers.

Parameter Name	Value	Description	Reference	
n_1	2	Parameters of hill equation	Assumption	
n_2	2	Parameters of hill equation	Assumption	
n_3	2	Parameters of hill equation	Assumption	
K_{M1}	40nM	Parameters of hill equation	Assumption	
K_{M2}	40nM	Parameters of hill equation	Assumption	
K_{M3}	40nM	Parameters of hill equation	Assumption	
k_{MluxI}	5.25nM/min	Strength decide by R0079	Peking 2009 All tunable for test Just estimate as standard	
k_{MTetR}	5.25nM/min	Strength decide by R0062		
k_{MlasI}	5.25nM/min	Strength decide by R0040		
k_{TL1}	42	Translation rate, connecting with strength of RBS(tunable)		
k_{TL2}	42	Translation rate, connecting with strength of RBS(tunable)		
k_{TL3}	42	Translation rate, connecting with strength of RBS(tunable)		
k_{TL4}	42	Translation rate, connecting with strength of RBS(tunable)		
k_{TL5}	42	Translation rate, connecting with strength of RBS(tunable)		
v_{MlasR}	5.25nM/min	Transcription rate(tunable)		Peking 2009
v_{MluxR}	5.25nM/min	Transcription rate(tunable)		Peking 2009
β_{MluxI}	0.01	Basal expression in hill equation	Assumption	
β_{MTetR}	0.01	Basal expression in hill equation	Assumption	
β_{MlasI}	0.01	Basal expression in hill equation	Assumption	
γ	2.5min^{-1}	Diffusion rate of AHL through membrane.	published paper	

λ_1	0.06	Generation rate of 30C6HSL	published paper
λ_2	0.06	Generation rate of 30C12HSL	published paper
d_{MlasR}	0.0173min^{-1}	Degradation constant of mRNA	published paper
d_{MluxI}	0.0173min^{-1}	Degradation constant of mRNA	published paper
d_{MluxR}	0.0173min^{-1}	Degradation constant of mRNA	published paper
d_{MTetR}	0.0173min^{-1}	Degradation constant of mRNA	published paper
d_{PluxR}	$2.31 \times 10^{-2}\text{min}^{-1}$	Degradation constant of luxR protein.	2010 MIT
d_{PluxI}	$1.67 \times 10^{-2}\text{min}^{-1}$	Degradation constant of luxI protein.	2010 MIT
d_{PlasI}	0.01min^{-1}	Degradation constant of lasI protein.	2010 MIT
d_{PlasR}	$1.88 \times 10^{-2} \text{min}^{-1}$	Degradation constant of lasR protein.	published paper
d_{PtetR}	$1.67 \times 10^{-2}\text{min}^{-1}$	Degradation constant of tetR protein.	assumption
k_1	$9.6 \times 10^{-3}\text{nM}^{-1}\text{min}^{-1}$	Rate constant of binding reaction between LasR and 30C12HSL	published paper
k_2	15min^{-1}	Rate constant of dissociation reaction between LasR and 30C12HSL	published paper
k_3	$0.14232\text{nM}^{-1}\text{min}^{-1}$	Rate constant of binding reaction between LuxR and 30C6HSL	2008 KULEuven
k_4	60min^{-1}	Rate constant of dissociation reaction between LuxR and 30C6HSL	2008 KULEuven
k_5	$0.06\text{nM}^{-1}\text{min}^{-1}$	Rate constant of binding reaction between tetR and aTc	published paper
k_6	50min^{-1}	Rate constant of dissociation reaction between tetR and aTc	published paper
p	0.2	Ratio of cell2 volume to total volume	For test
n_{12}	1	Ratio of cell1 volume to cell2 volume	For test
μ	10min^{-1}	Dilution rate of C12 and C6 in environment (tunable)	For test

Table 1 Parameters of ODEs

Results

We simulated this system by SIMBIOLOGY, a toolbox embedded in MATLAB. However, unaware of the key parameters to which the system is sensitive, we felt difficult to control or adjust properly, and the simulation result of the system came into a damped oscillation. We ascribed the inability of our model to the fact that the precise descriptions contain too many equations and parameters and we felt obliged to establish a simplified model in place of the precise one for simulation and further analysis.