

Advanced Special Topics I

Genetic Circuit Design

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Overview

- 1 Literature Review
- 2 Project

Literature Review: Overview

- 1 Hardware vs Wetware Analogy
- 2 Logic Gates
- 3 Gate Implementation
- 4 Composability
- 5 Continuous Response Variables
- 6 Applications
- 7 Challenges

Logic Gates

YES



INPUT		OUTPUT
A		
0		0
1		1

NOT



INPUT		OUTPUT
A		
0		1
1		0

AND



INPUT		OUTPUT
A	B	
0	0	0
1	0	0
0	1	0
1	1	1

OR



INPUT		OUTPUT
A	B	
0	0	0
1	0	1
0	1	1
1	1	1

XOR



INPUT		OUTPUT
A	B	
0	0	0
1	0	1
0	1	1
1	1	0

NAND



INPUT		OUTPUT
A	B	
0	0	1
1	0	1
0	1	1
1	1	0

NOR



INPUT		OUTPUT
A	B	
0	0	1
1	0	0
0	1	0
1	1	0

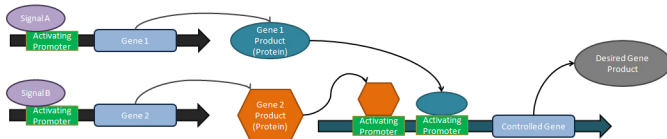
XNOR



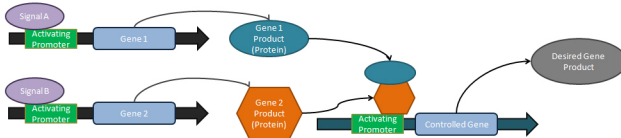
INPUT		OUTPUT
A	B	
0	0	1
1	0	0
0	1	0
1	1	1

Gate Implementation

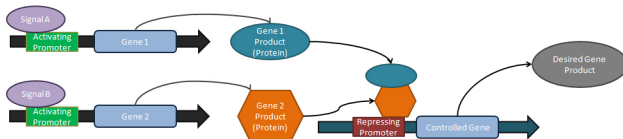
OR Gate (© Public Domain)



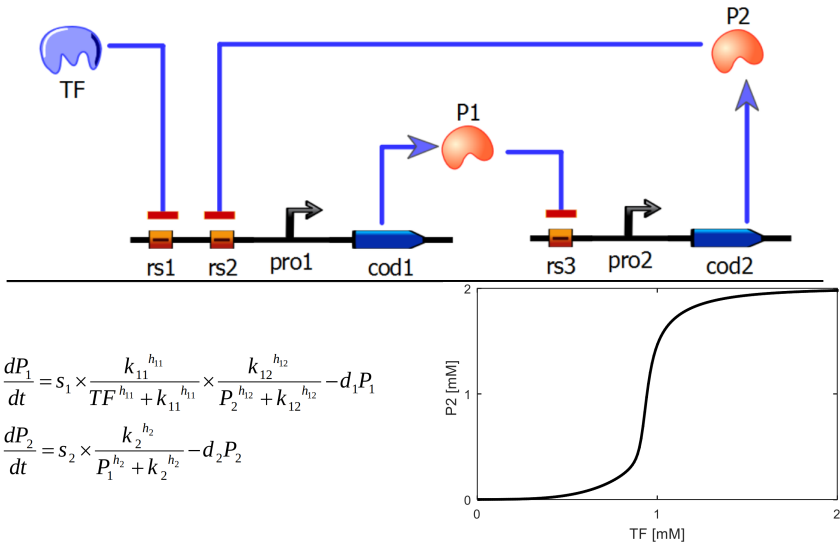
AND Gate (© Public Domain)



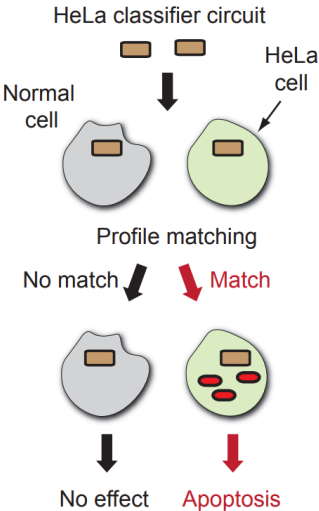
NAND Gate (© Public Domain)



Continuous Response Variables



- Disease diagnosis
- Environmental sensing
- Research and development
- Material and chemical production



(Xie et al 2011)

Challenges

- ① Non-orthogonality of components.
 - Undesired behaviour.
- ② Metabolic load.
 - Mortality.
 - Evolutionary pressure.

Possible Solution: Distributed circuits.

Project: Overview

- ① Hill Equation
- ② YES Gate
- ③ NOT Gate
- ④ Time-Dependent NAND Gate
- ⑤ Limitations
- ⑥ Learning Outcomes

Hill Equation

$$\theta = \frac{1}{1 + \left(\frac{K_A}{[L]}\right)^n}$$

- θ is the fraction of protein bound by ligand.
- $[L]$ is the total ligand concentration.
- K_A is the half-occupation concentration of ligand.
- n is the Hill coefficient.
 - $n < 1 \implies$ negatively-cooperative binding.
 - $n > 1 \implies$ positively-cooperative binding.
 - $n = 1 \implies$ non-cooperative binding.

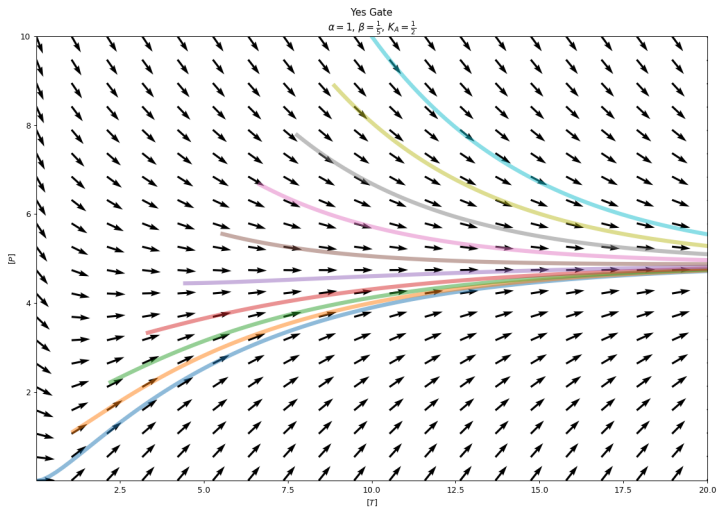
Example: YES Gate

$$\frac{d[P]}{d[T]} = \frac{\alpha}{1 + \left(\frac{K_A}{[T]}\right)} - \beta[P]$$

- $[P]$: Concentration of protein product.
- $[T]$: Concentration of transcription factor.
- α : Maximum rate of synthesis of P .
- β : Protein degradation rate.
- K_A : Half-occupation concentration transcription factor.

$$[P] = \frac{\alpha}{\beta} \frac{[T]}{K_A + [T]}$$

Example: YES Gate



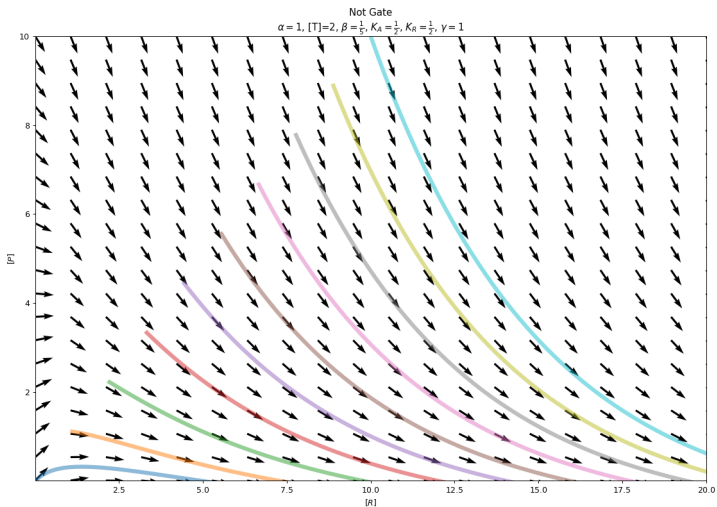
Example: NOT Gate

$$\frac{d[P]}{d[R]} = \frac{\alpha}{1 + \left(\frac{K_A}{[T]}\right)} - \frac{\gamma}{1 + \left(\frac{K_R}{[R]}\right)} - \beta[P]$$

- $[P]$: Concentration of protein product.
- $[R]$: Concentration of repressor.
- $[T]$: Concentration of transcription factor.
- α : Maximum rate of synthesis of P .
- γ : Maximum rate of inhibition of synthesis of P .
- β : Protein degradation rate.
- K_A : Half-occupation concentration transcription factor.
- K_R : Half-occupation concentration inhibition factor.

$$[P] = \left(C_1 - \frac{\int \frac{K_A[R]\gamma e^{[R]\beta}}{K_R + [R]} d[R] + \int \left(-\frac{K_R[T]\alpha e^{[R]\beta}}{K_R + [R]} \right) d[R] + \int \left(-\frac{[R][T]\alpha e^{[R]\beta}}{K_R + [R]} \right) d[R] + \int \frac{[R][T]\gamma e^{[R]\beta}}{K_R + [R]} d[R]}{K_A + [T]} \right) e^{-[R]\beta}$$

Example: NOT Gate



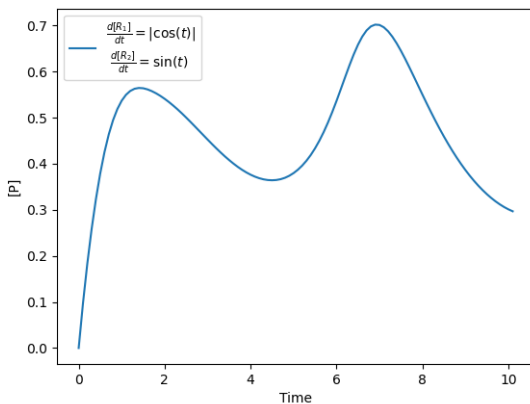
Time-Dependent NAND

$$\frac{d[P](t, [R_1], [R_2])}{dt} = \left(1 - \left[\frac{\alpha}{1 + \frac{K_{A_1}}{[R_1](t)}} \right] \left[\frac{\gamma}{1 + \frac{K_{A_2}}{[R_2](t)}} \right] \right) - \beta[P](t, [R_1], [R_2])$$

- $[P]$: Concentration of protein product.
- $[R_1]$: Concentration of repressor.
- $[R_2]$: Concentration of repressor.
- α : Maximum rate of synthesis of P .
- γ : Maximum rate of inhibition of synthesis of P .
- β : Protein degradation rate.
- K_{A_1} : Half-occupation concentration inhibition factor 1.
- K_{A_2} : Half-occupation concentration inhibition factor 2.

Time-Dependent NAND: Non-Monotonicity

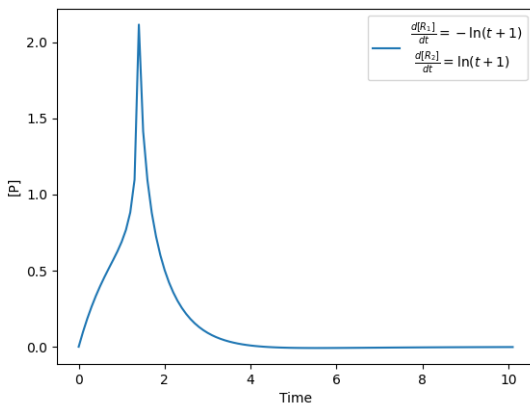
What happens over time?



Letting $\alpha = \beta = \gamma = 1$, $[P]_0 = 0$, and $K_{A_1} = K_{A_2} = \frac{1}{2}$.

Time-Dependent NAND: Discontinuity

What happens over time?



Letting $\alpha = \beta = \gamma = 1$, $[P]_0 = 0$, and $K_{A_1} = K_{A_2} = \frac{1}{2}$.

Limitations

- Our models are possibly too simple for a real cell.
- Parameters in real life cannot be arbitrary tweaked; depend on choice of molecules.
- Untested predictive power.

Learning Outcomes

- Learned about genetic circuit design.
- Learned about solving elementary ODE's.
 - Symbolically: Separation of variables
 - Numerically: Euler's method
- Learned about software packages.
 - SymPy
 - py-PDE