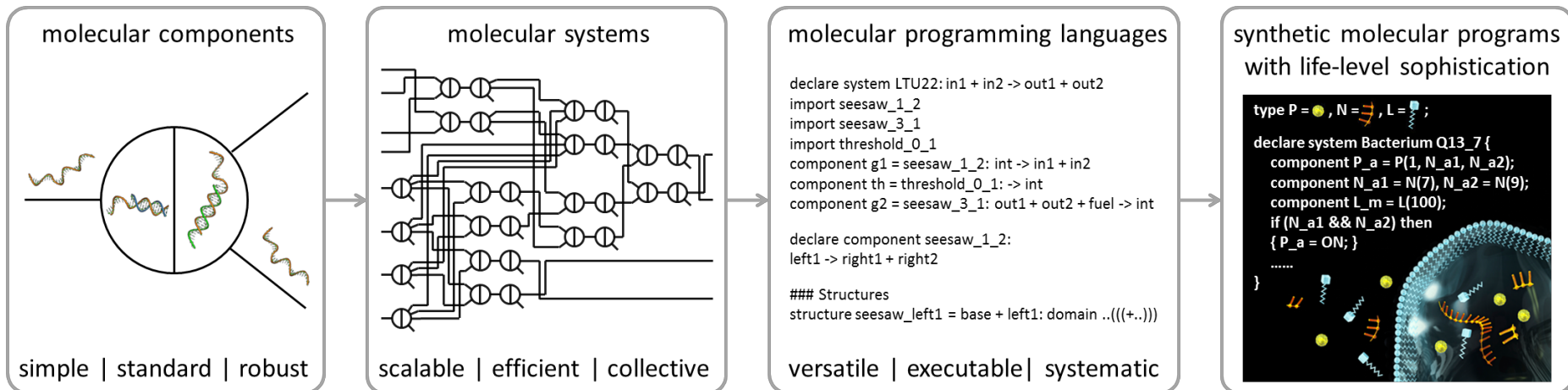
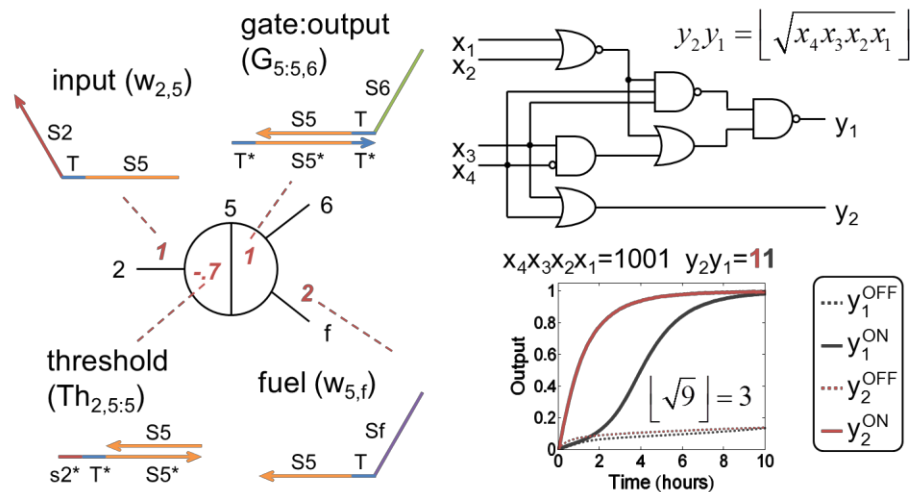
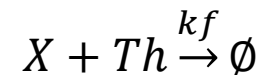
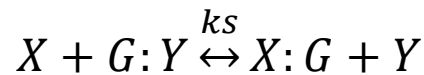
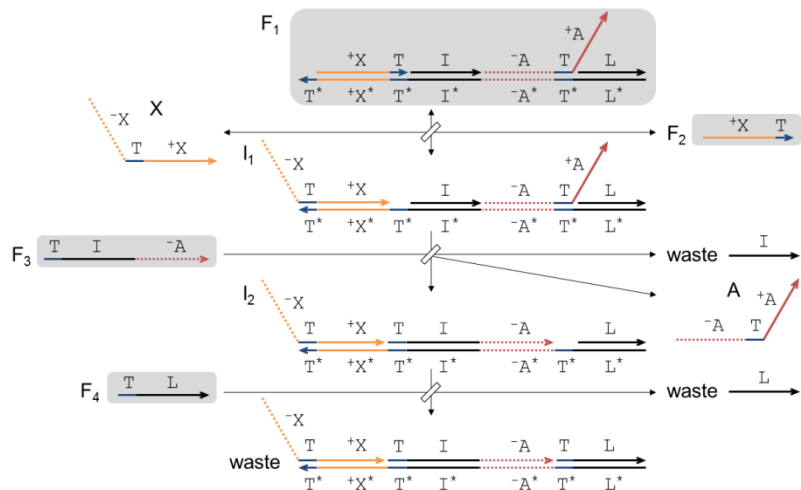
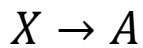


# Implementing complex CRNs with modular DNA components

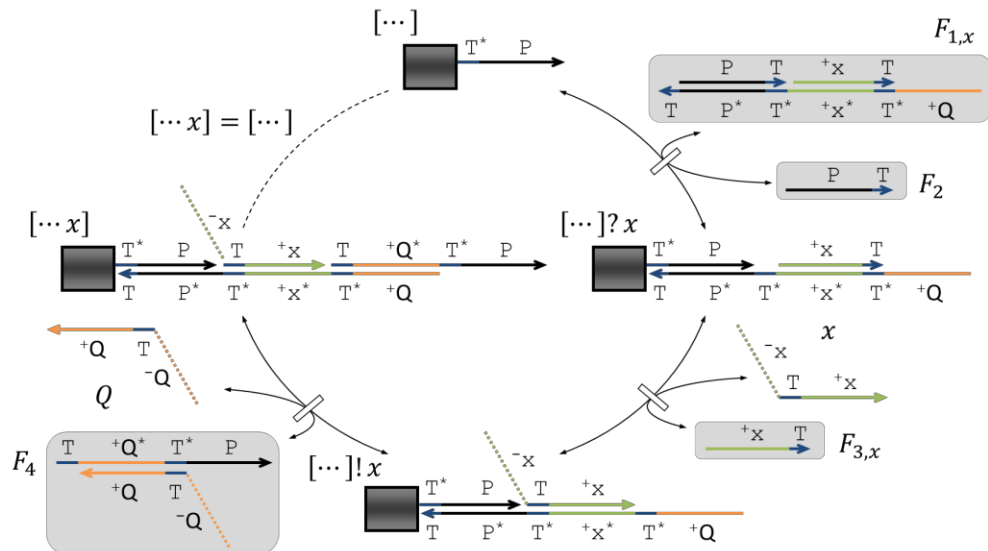
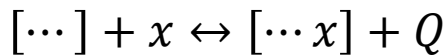


Lulu Qian  
Bioengineering  
Caltech

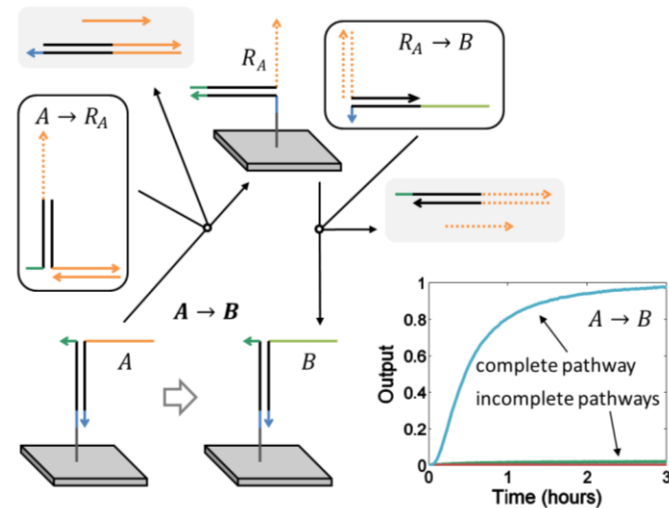
## Well-mixed CRNs



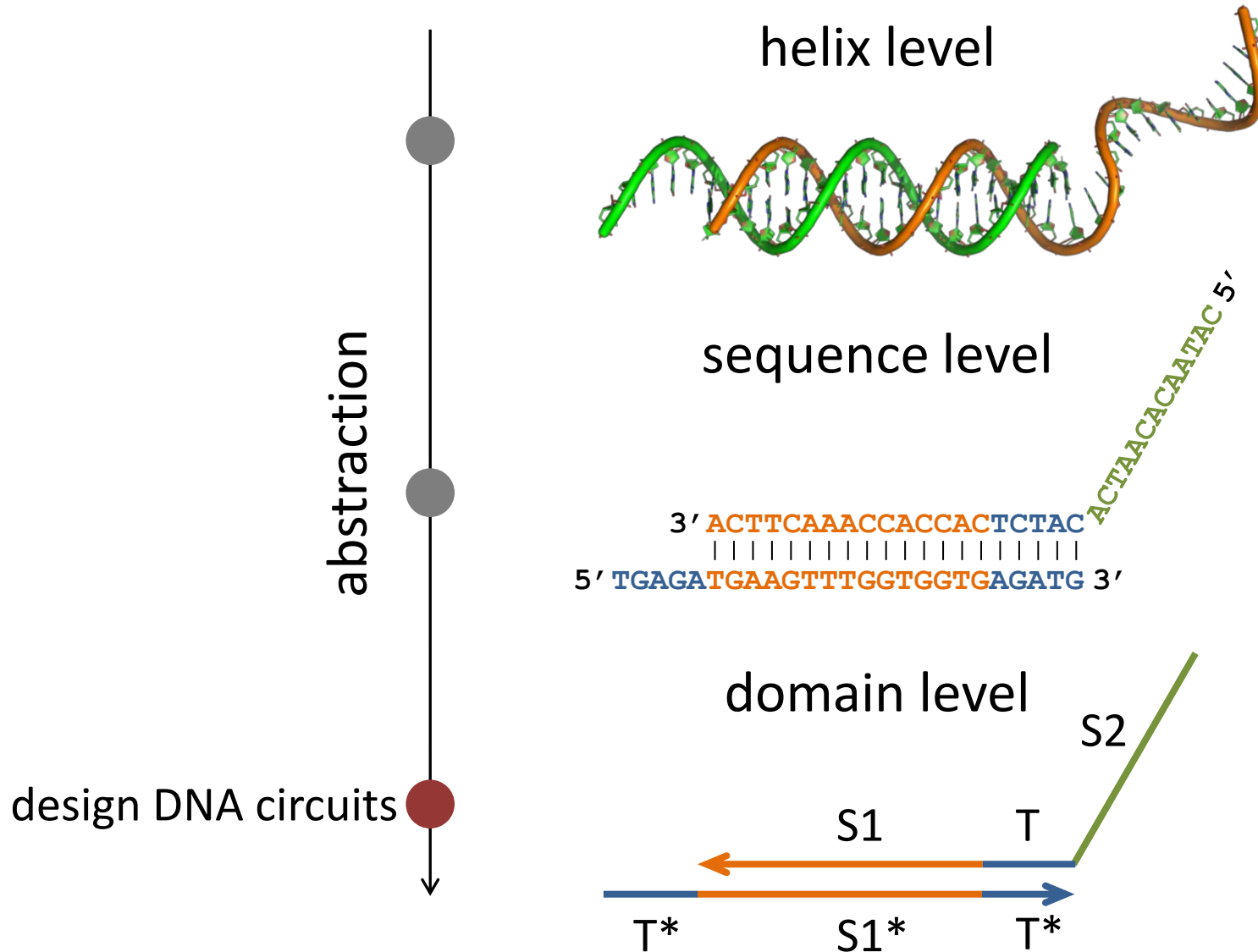
# Polymer CRNs



## Surface CRNs



# Representations of DNA molecules

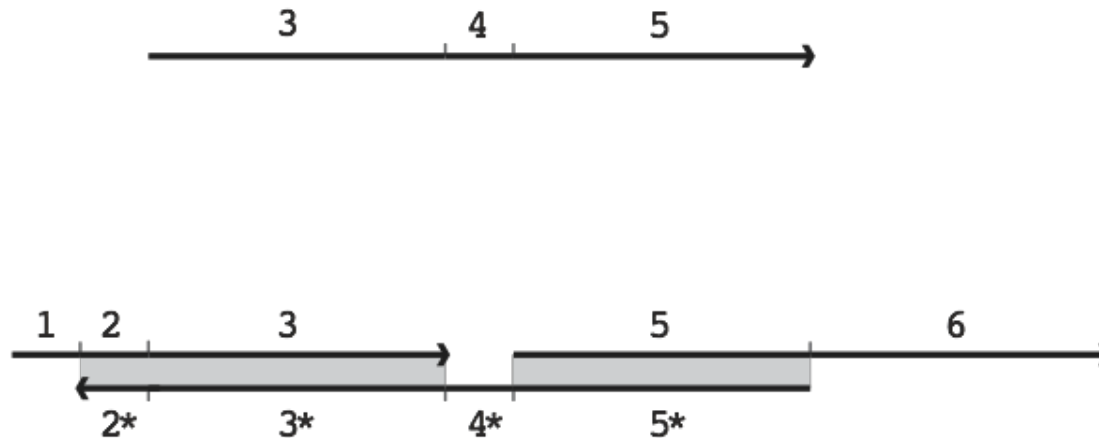


# Principles of DNA strand displacement circuits

**bind:** two complementary domains can bind.

**unbind:** any strands held by only a short domain can unbind.

**displace:** a domain can displace an identical domain if this extends existing binding.

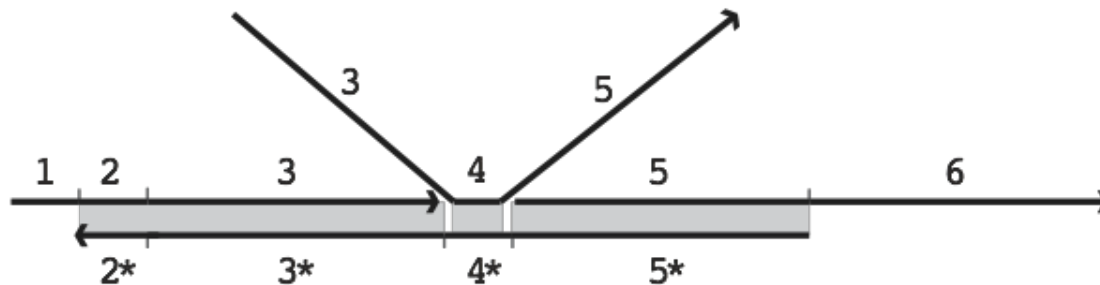


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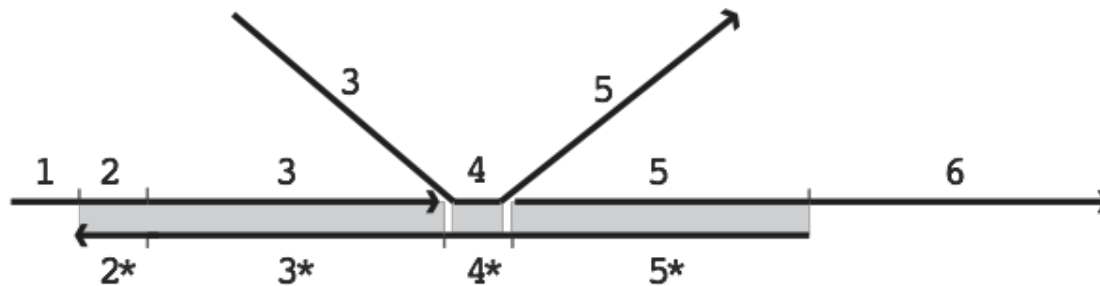


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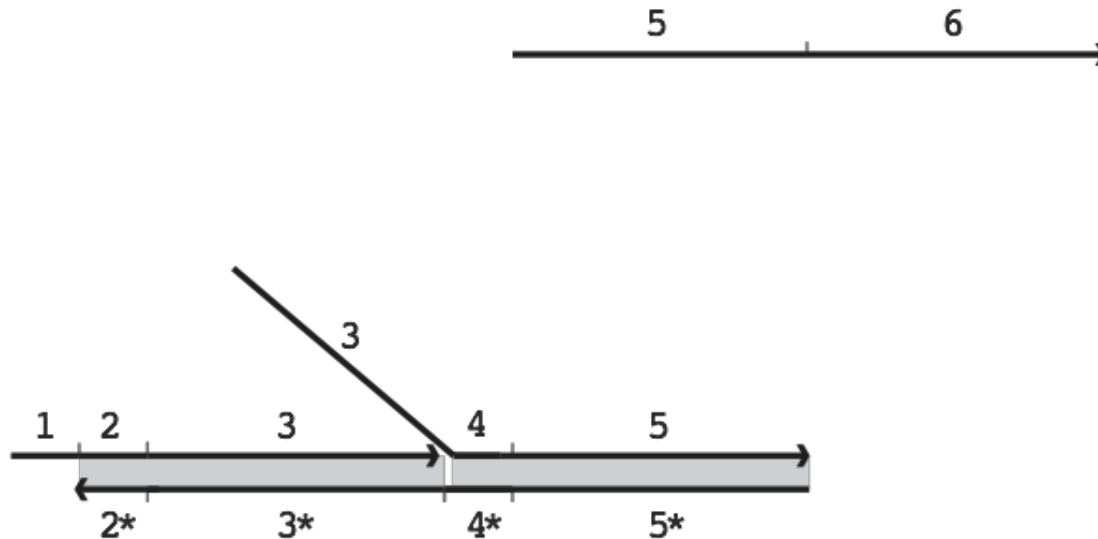


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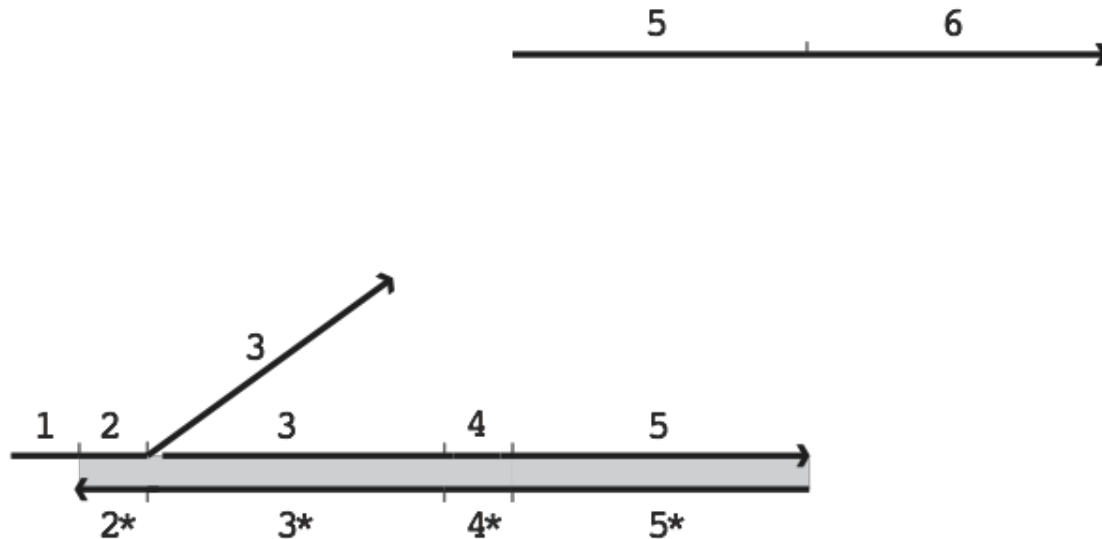


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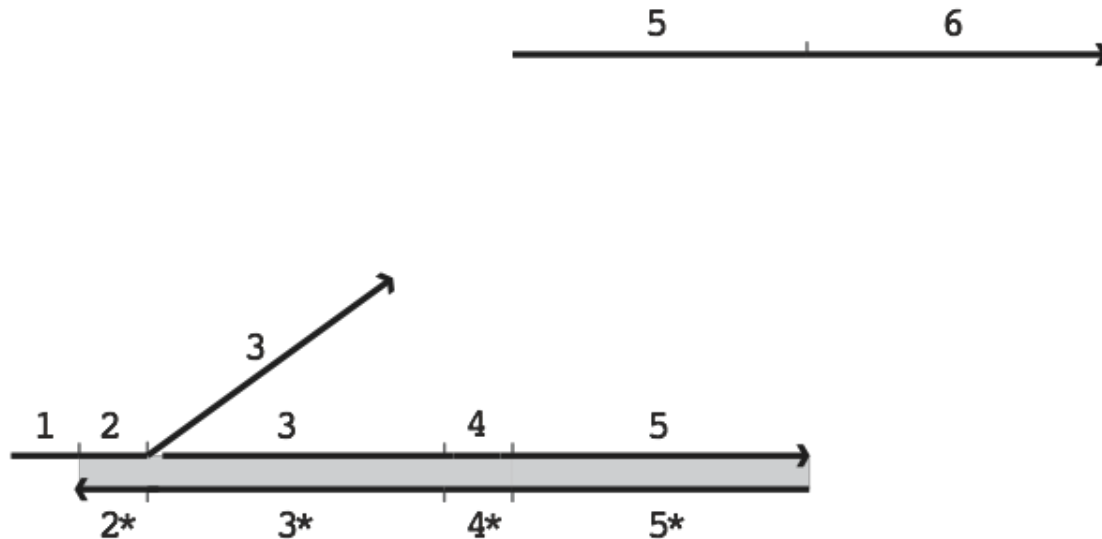


# Principles of DNA strand displacement circuits

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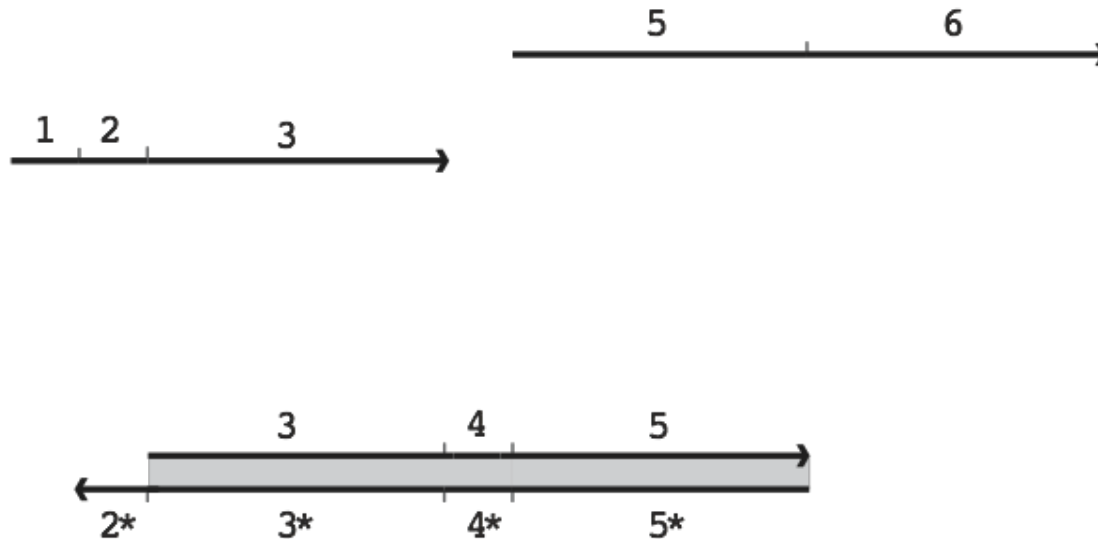


# Principles of DNA strand displacement circuits

**bind:** two complementary domains can bind.

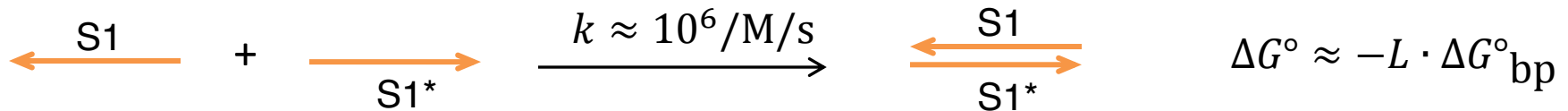
**unbind:** any strands held by only a short domain can unbind.

**displace:** a domain can displace an identical domain if this extends existing binding.

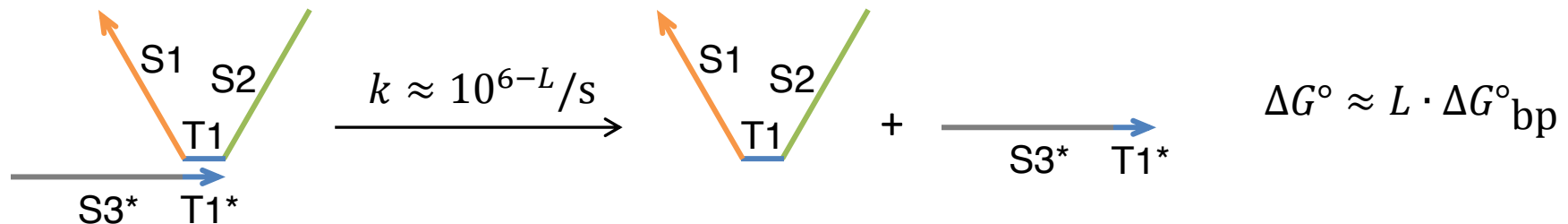


# Principles of DNA strand displacement circuits

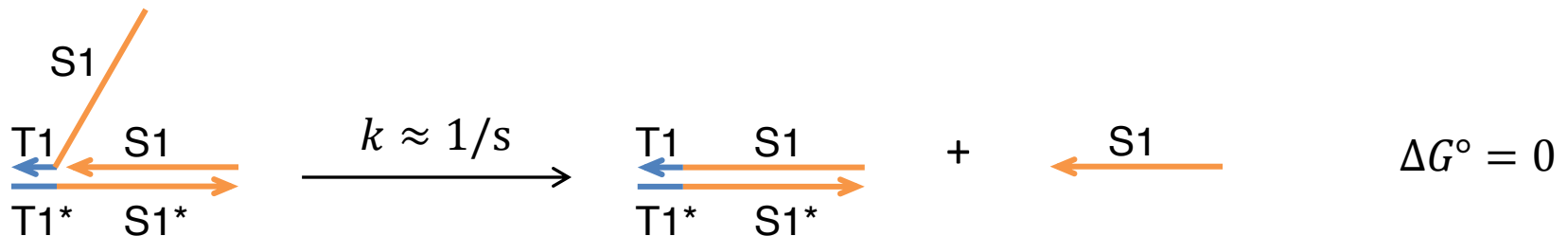
**bind:** two complementary domains can bind.



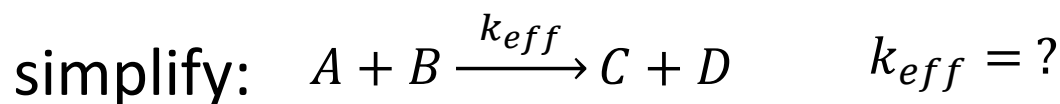
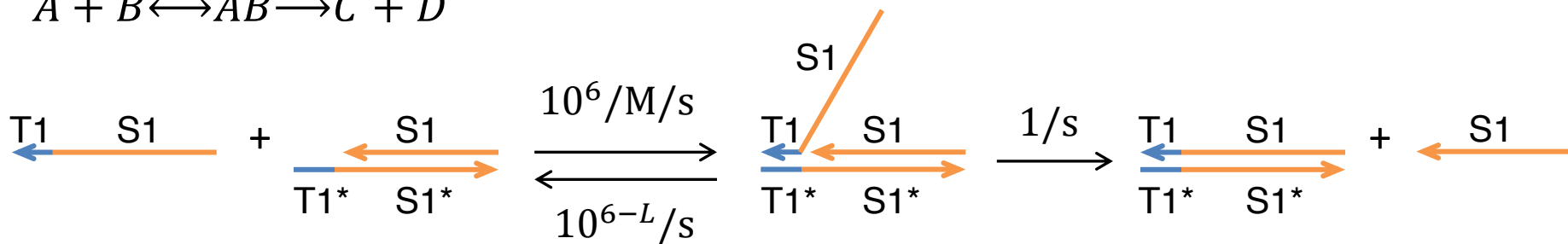
**unbind:** any strands held by only a short domain can unbind.



**displace:** a domain can displace an identical domain if this extends existing binding.



# Kinetics of toehold-mediated strand displacement



collision rate:  $10^6 [A][B]$

collision success probability:  $\frac{1/s}{1/s + 10^{6-L}/s}$

net rate of success:  $10^6 \cdot \underbrace{\frac{1}{1 + 10^{6-L}}}_{k_{eff}} [A][B]$

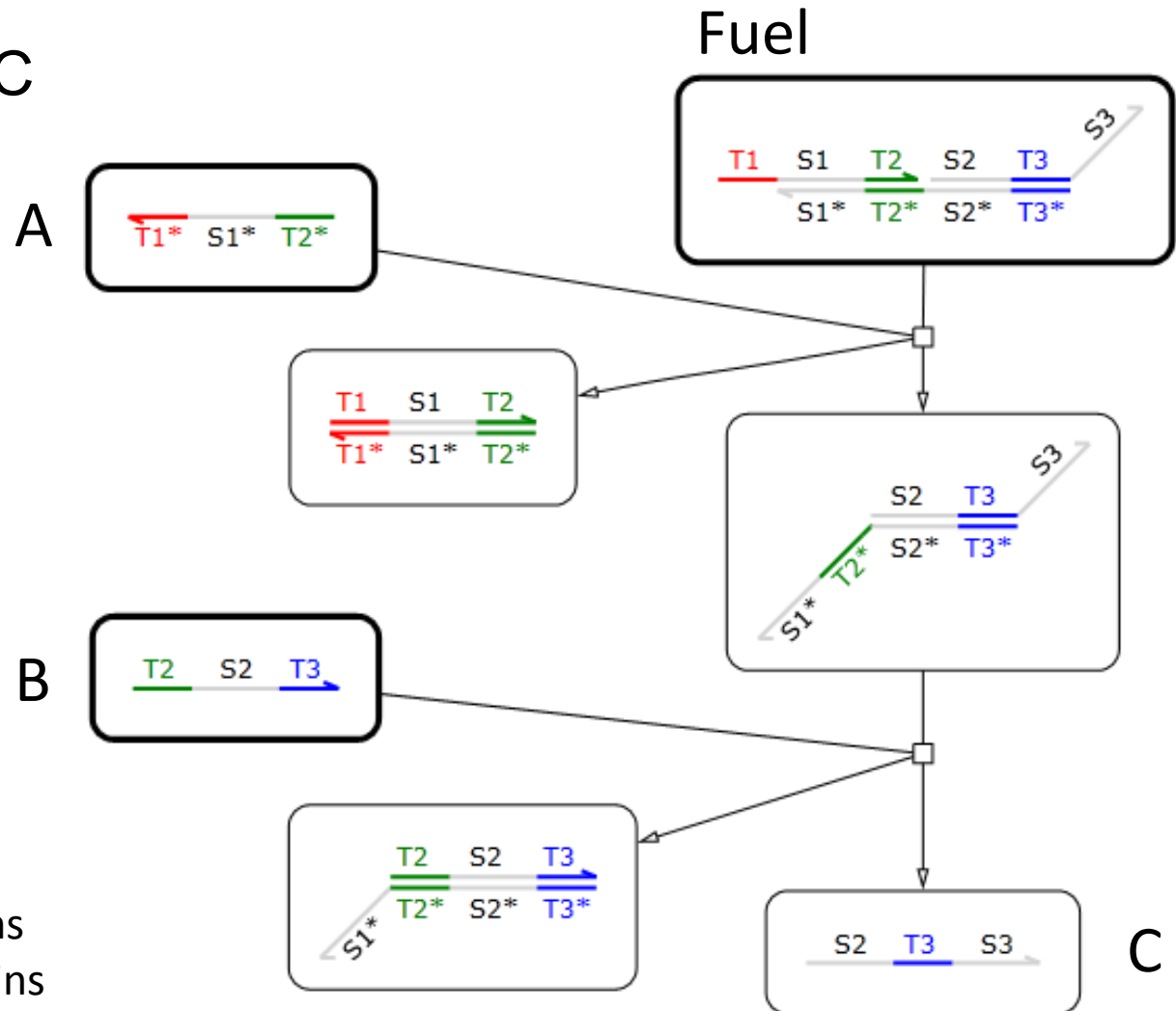
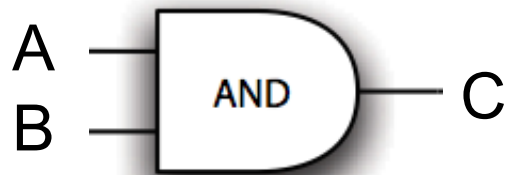
$k_{eff} \approx 10^L \text{ M/s}$  when  $L \leq 6$   
 otherwise  $k_{eff} \approx 10^6 \text{ M/s}$   
 $L$ : toehold length  $|T1|$

Zhang et al, *JACS* 2009

Srinivas et al, *NAR* 2013

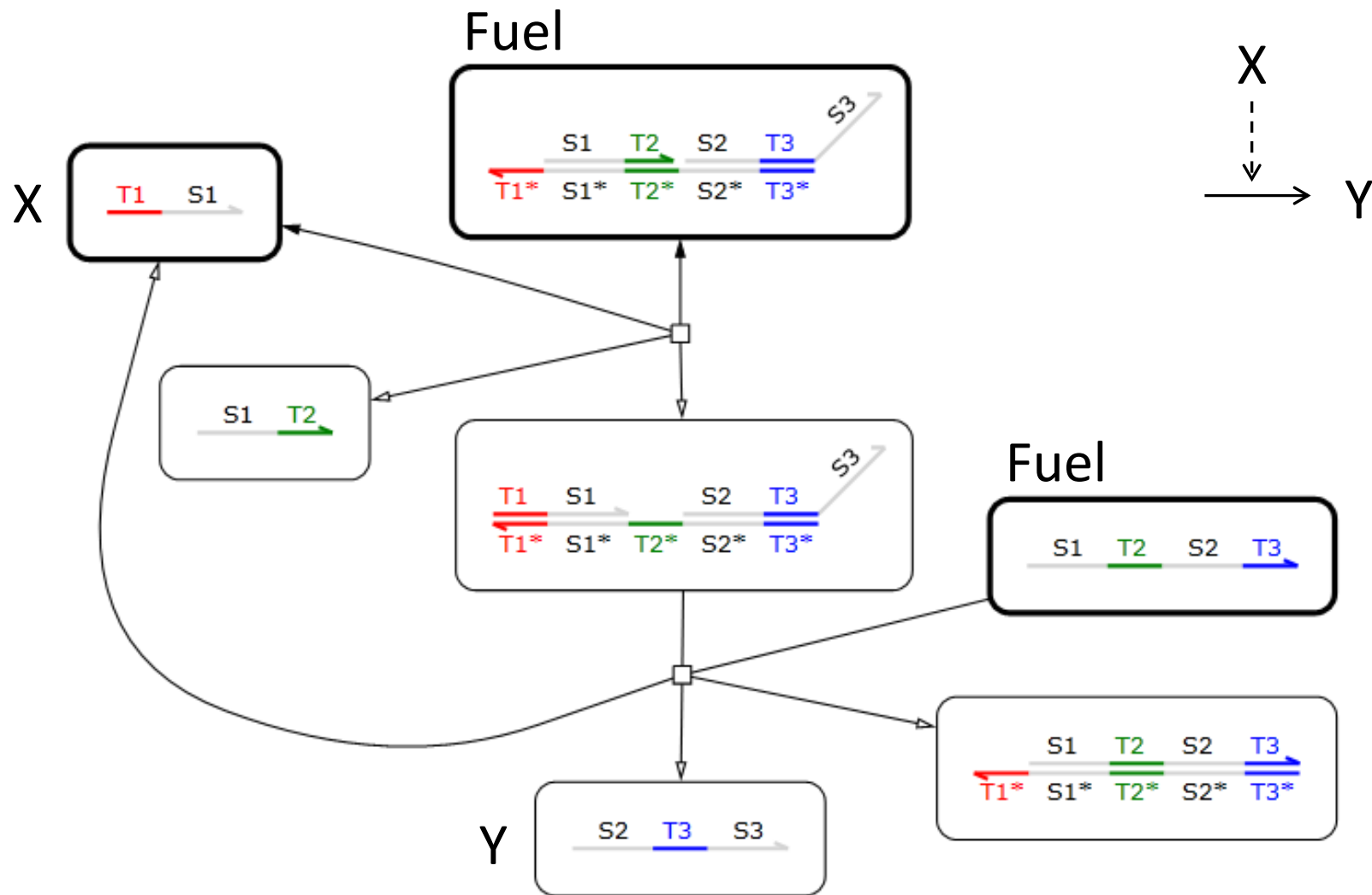
This approximation is valid for low concentrations of A and B (e.g.  $[A]=[B]=100 \text{ nM}$ ) such that the unimolecular reaction is sufficiently faster than the bimolecular reaction.

# Examples of simple strand displacement circuits

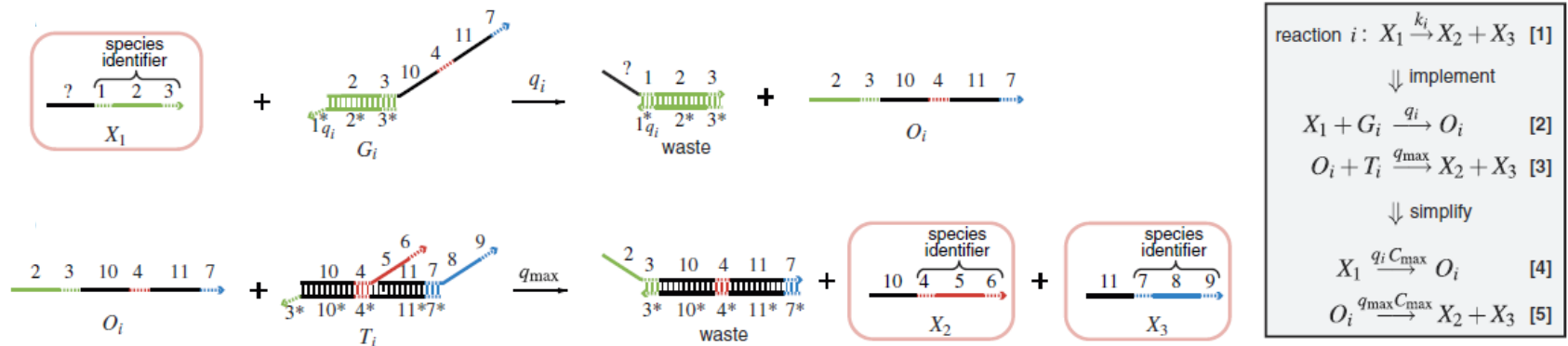


S1, S2, S3 are long domains  
T1, T2, T3 are short domains

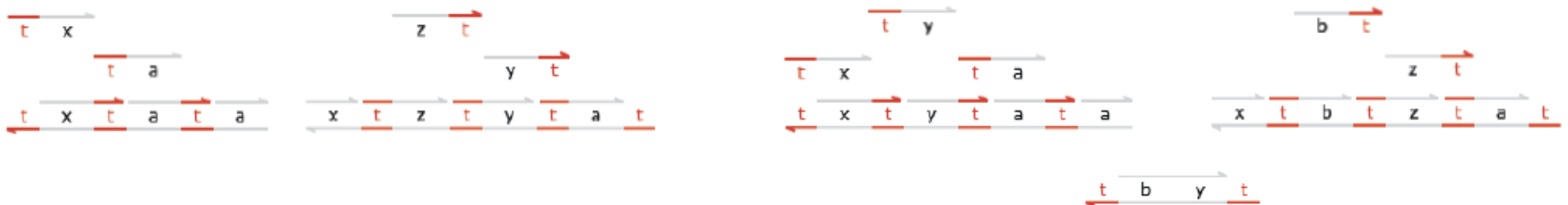
# Examples of simple strand displacement circuits



# Can one implement arbitrary CRNs with DNA strand displacement circuits?



Soloveichik et al, *PNAS* 2010



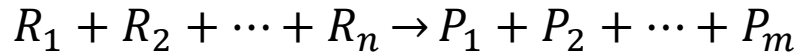
Fork  $F_{xyz} \mid tx \rightarrow ty \mid tz$

Join  $J_{xyz} \mid tx \mid ty \rightarrow tz$

Cardelli, *Math. Struct. Comput. Sci.* 2013

# Conditions of a successful CRNs implementation

## 1. logical conditions



- a. The reaction pathway must first consume a molecule of each reactant, and then produce a molecule of each product.
- b. The reaction pathway must first become irreversible after all reactants have been consumed and before any product has been produced.

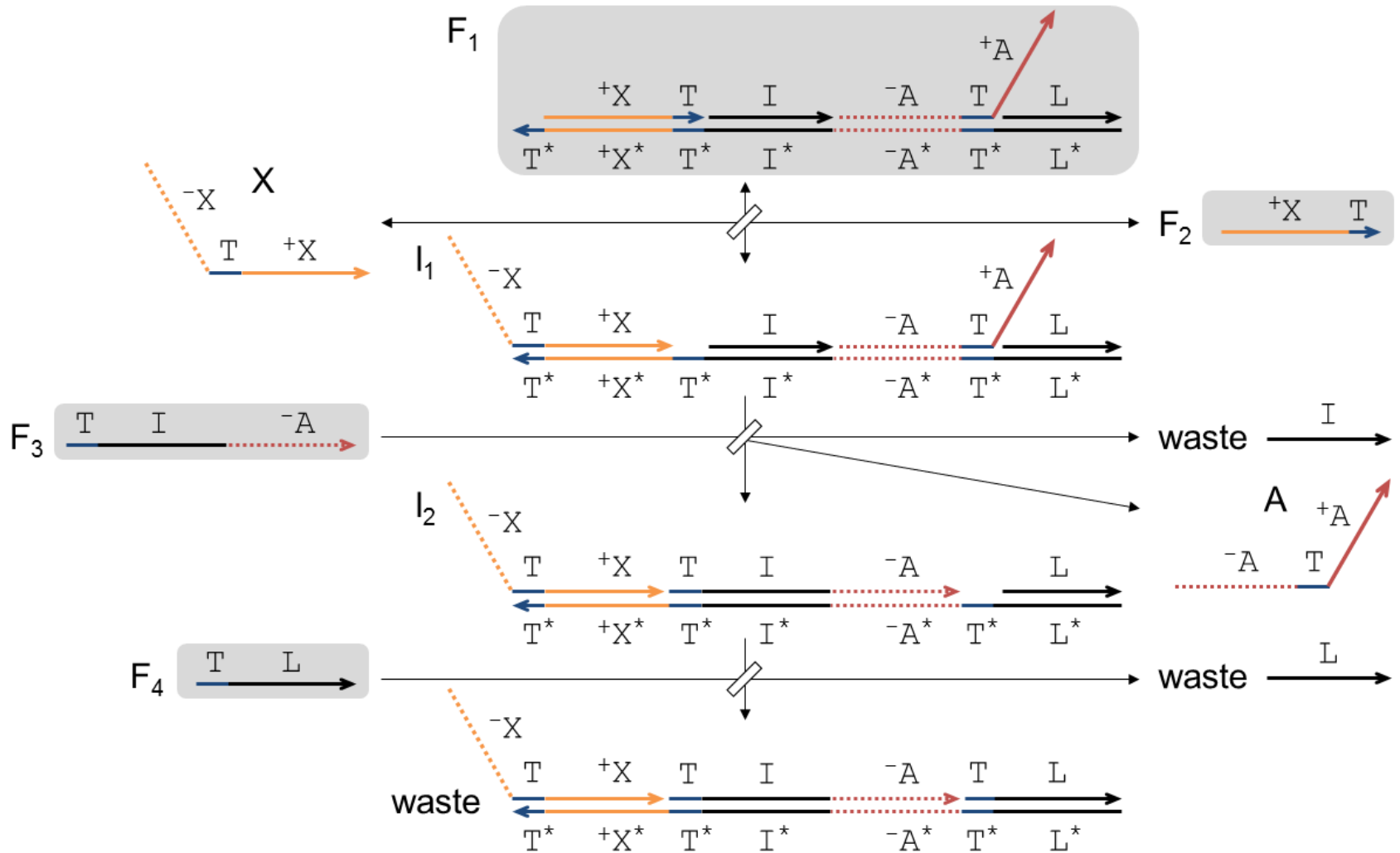
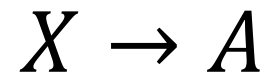
## 2. kinetics conditions

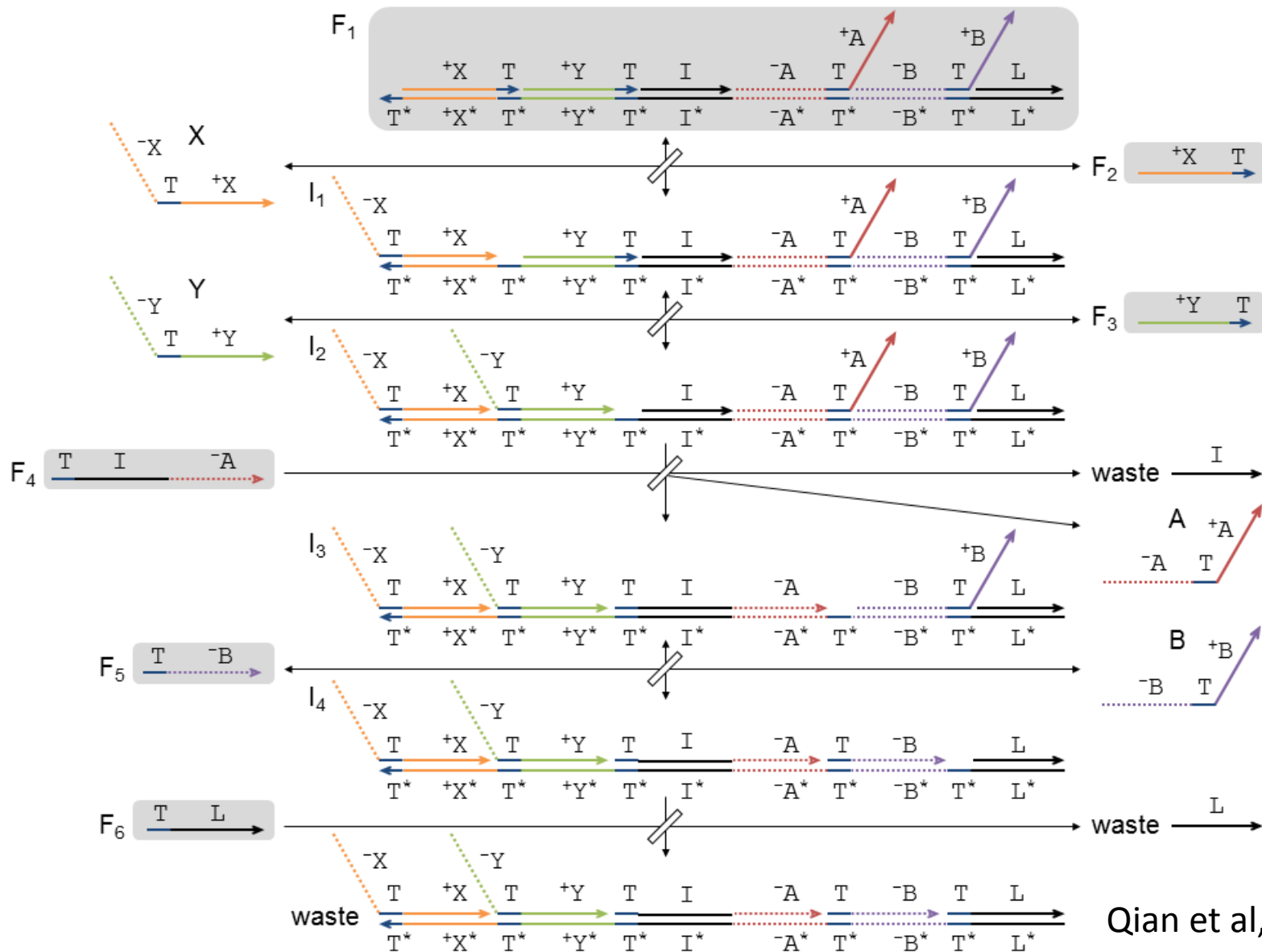
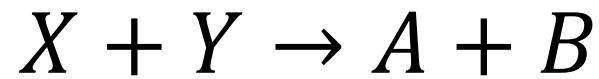
The rate of a reaction scales with the concentration of all reactants.

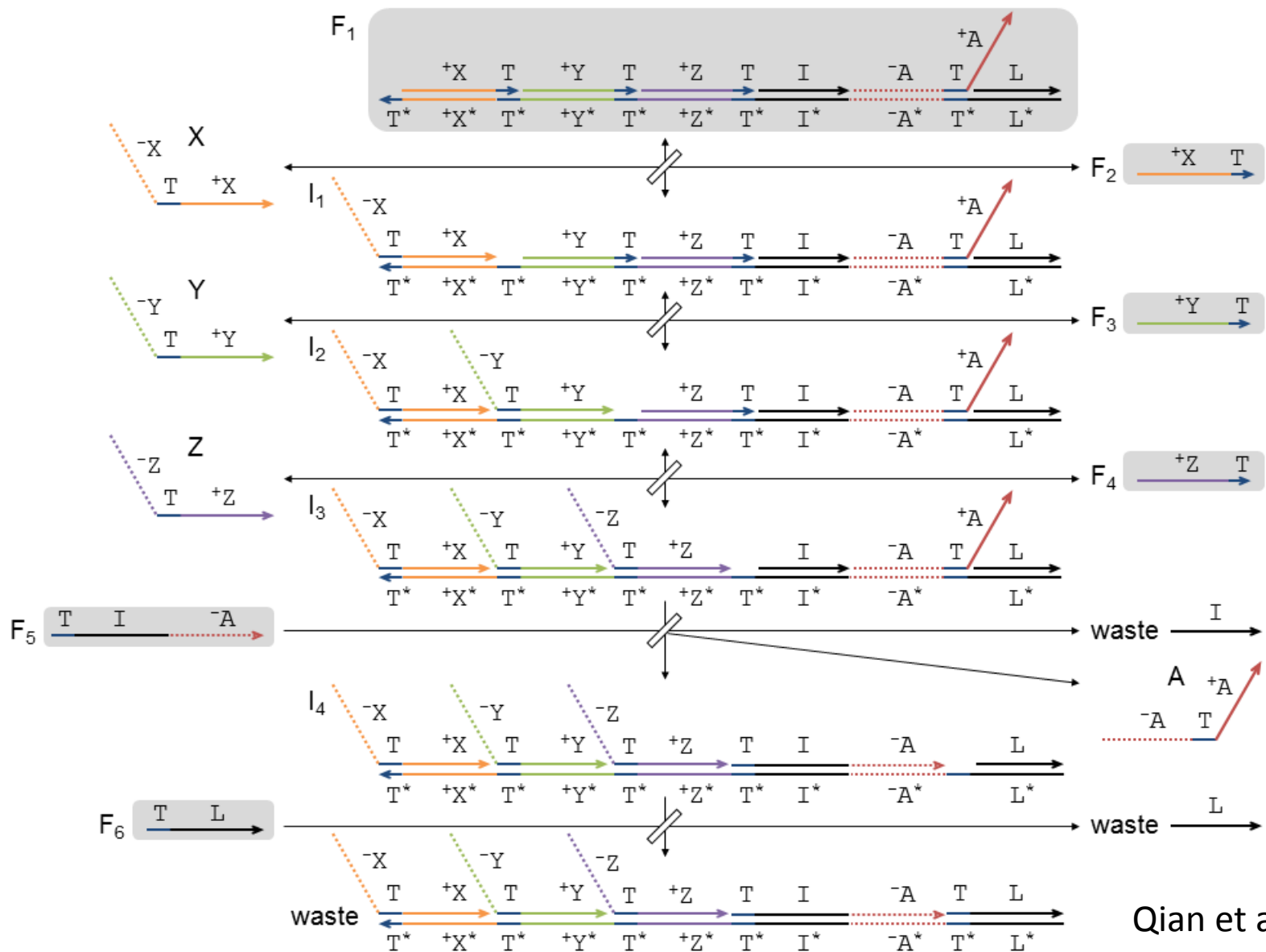
## 3. composable conditions

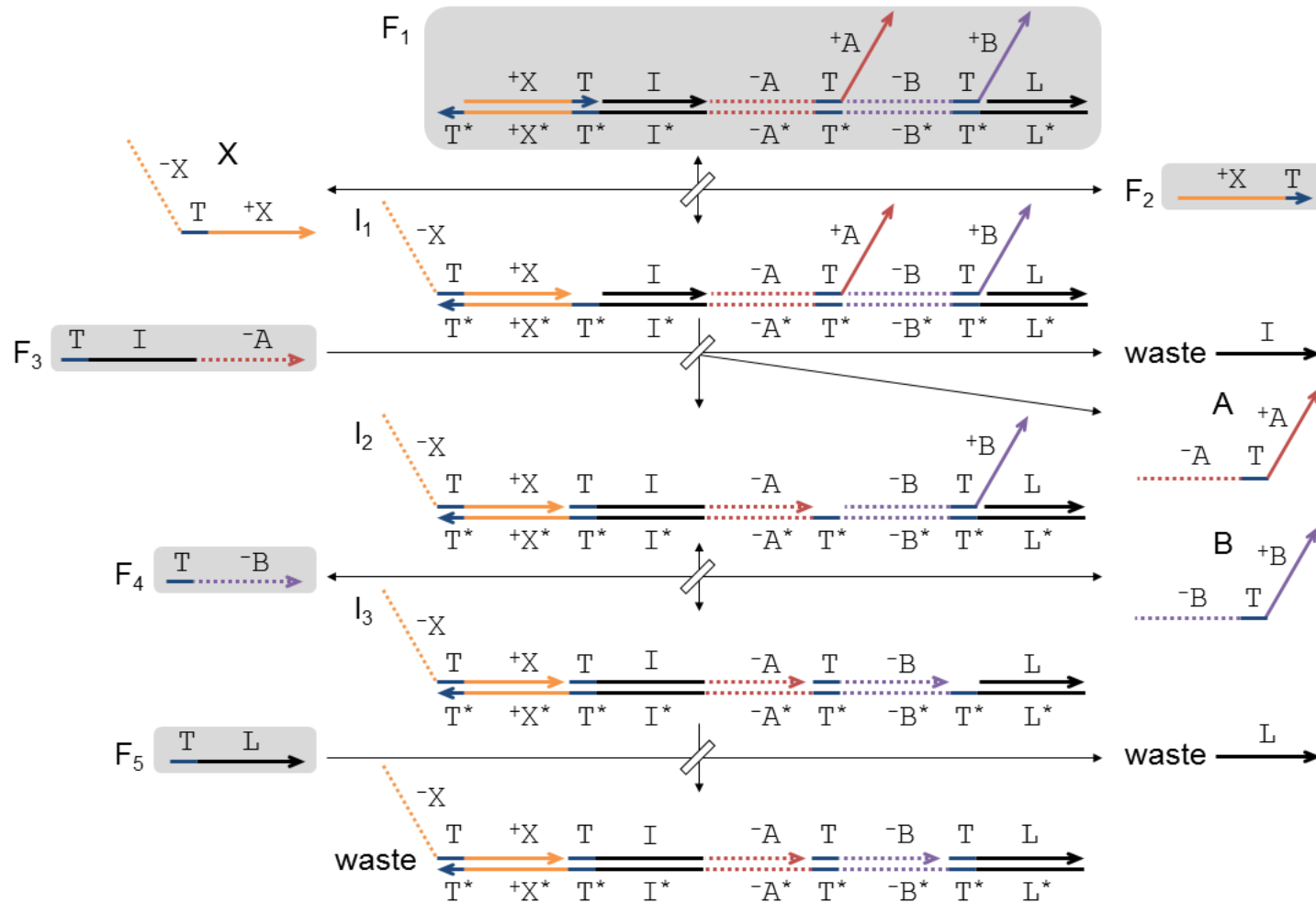
- a. All chemical species are implemented with the same form.
- b. No fuel or intermediate species crosstalk.

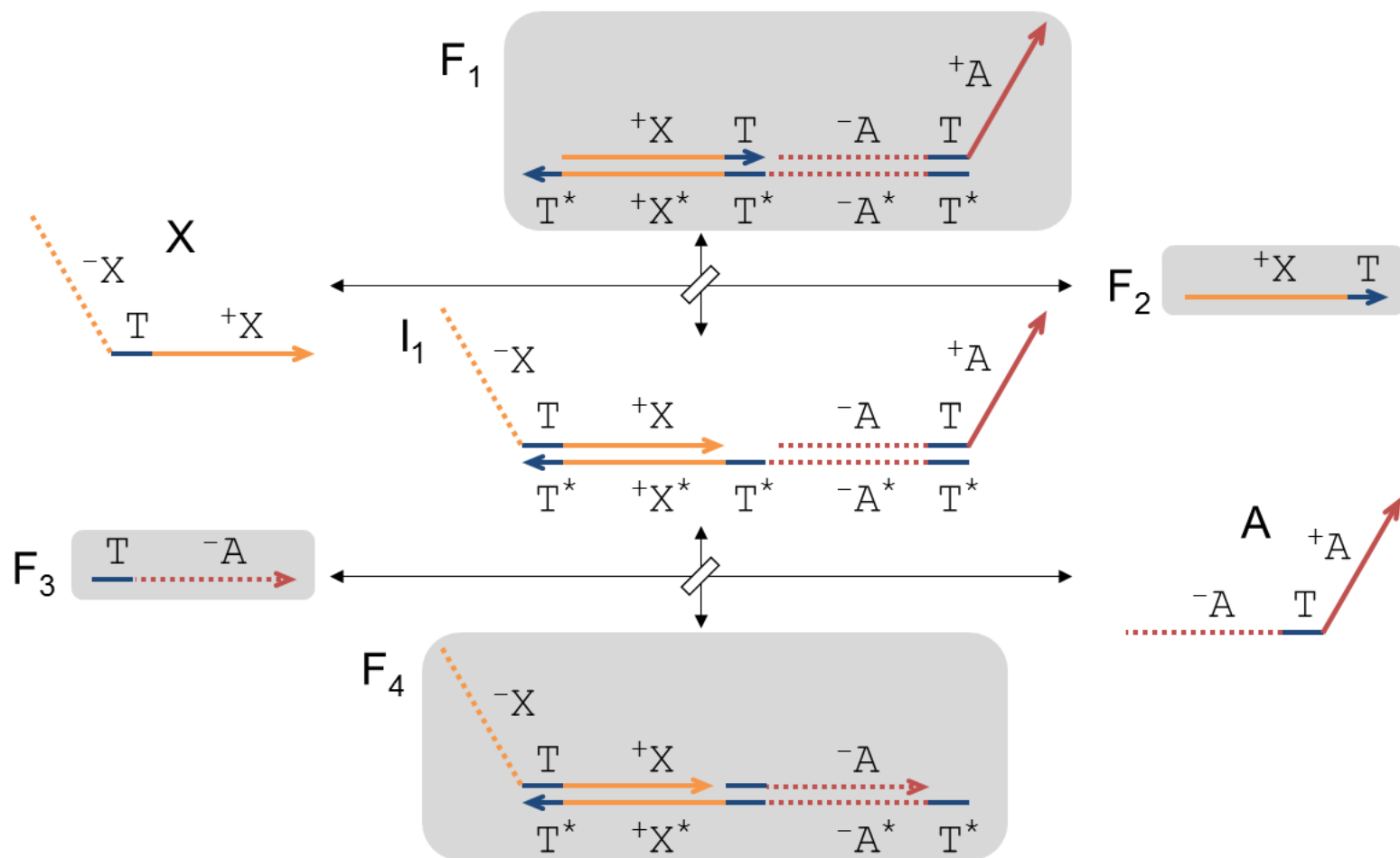
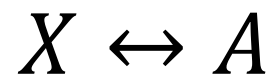


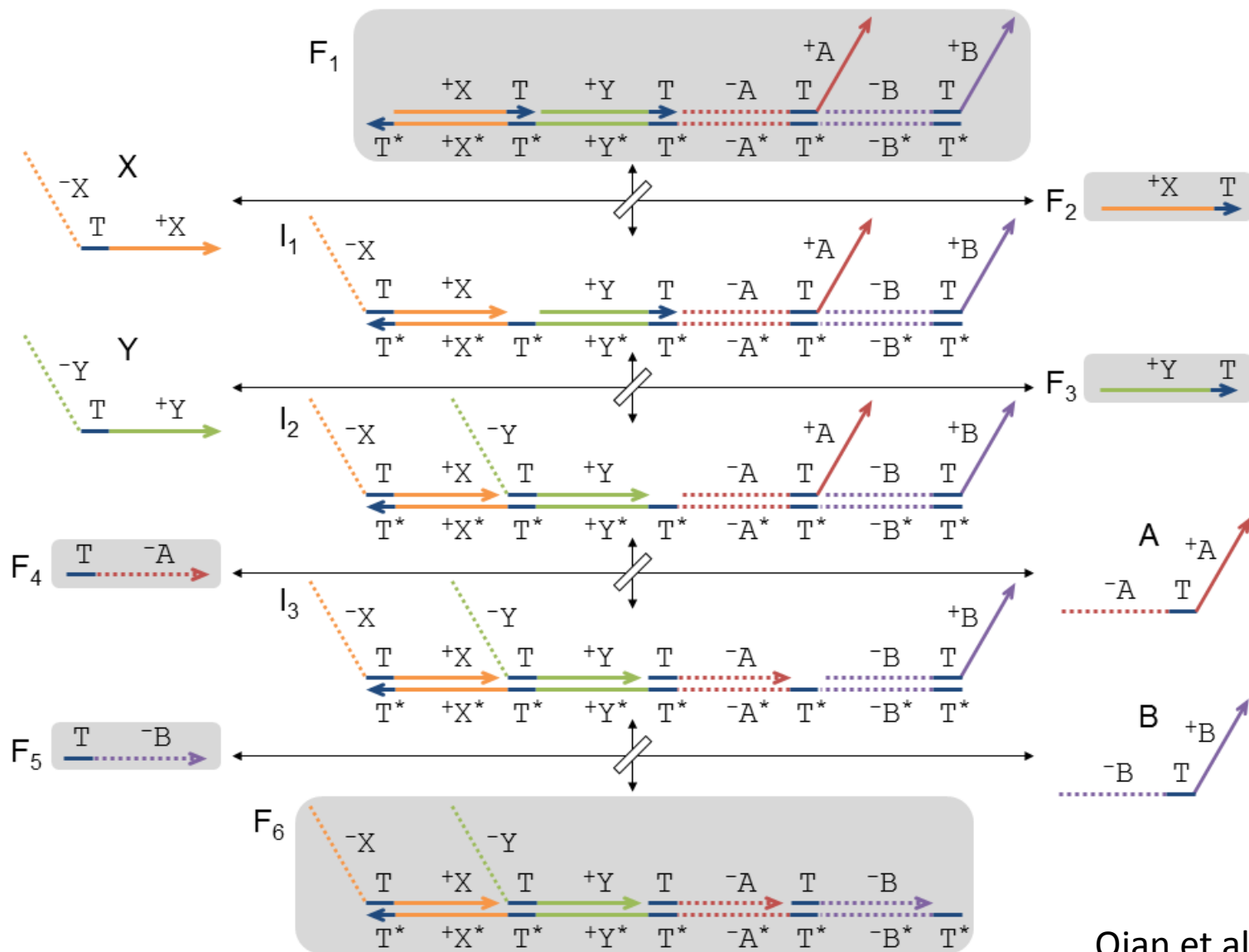
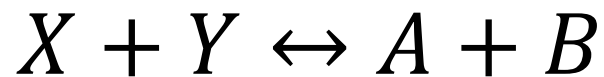




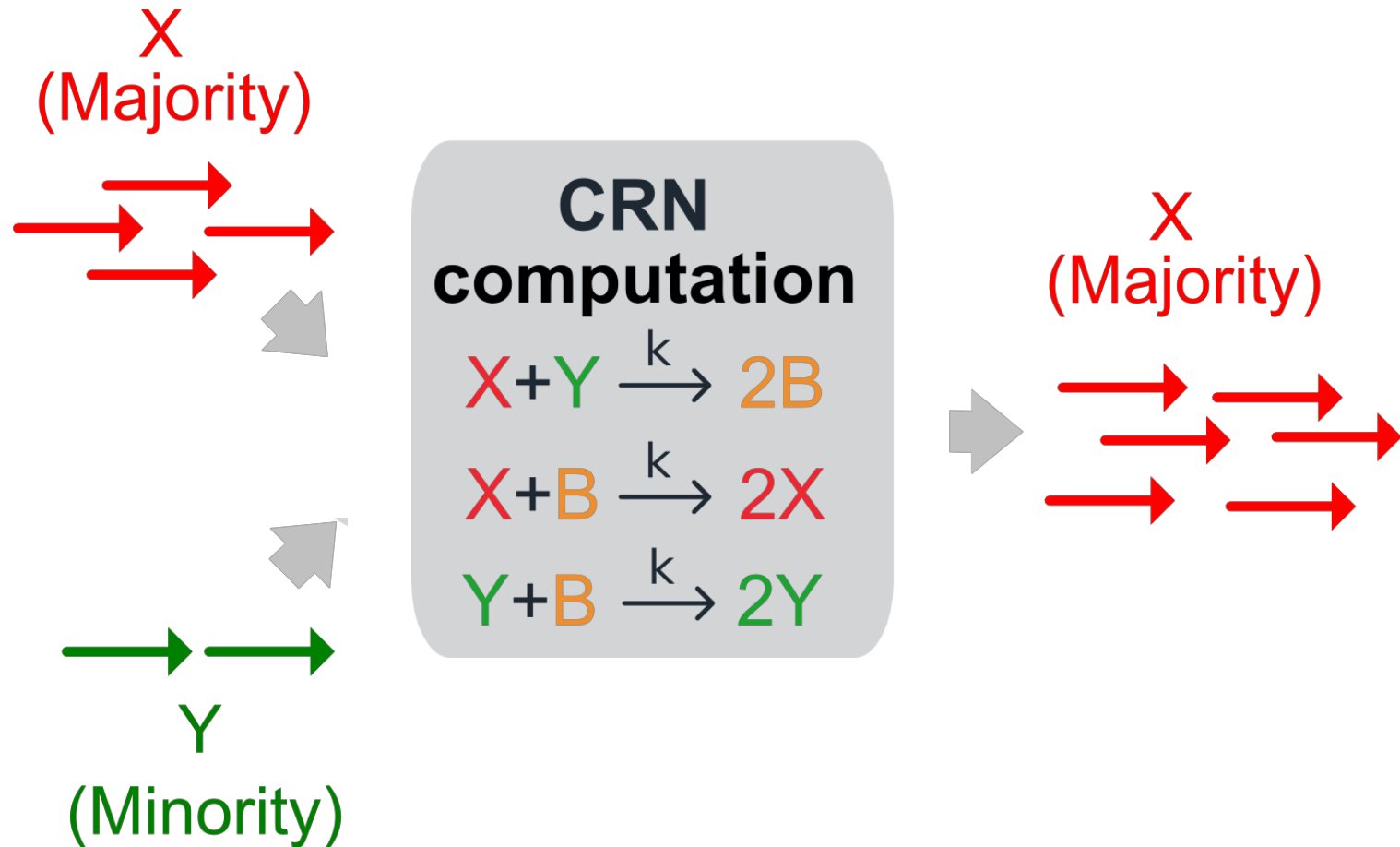




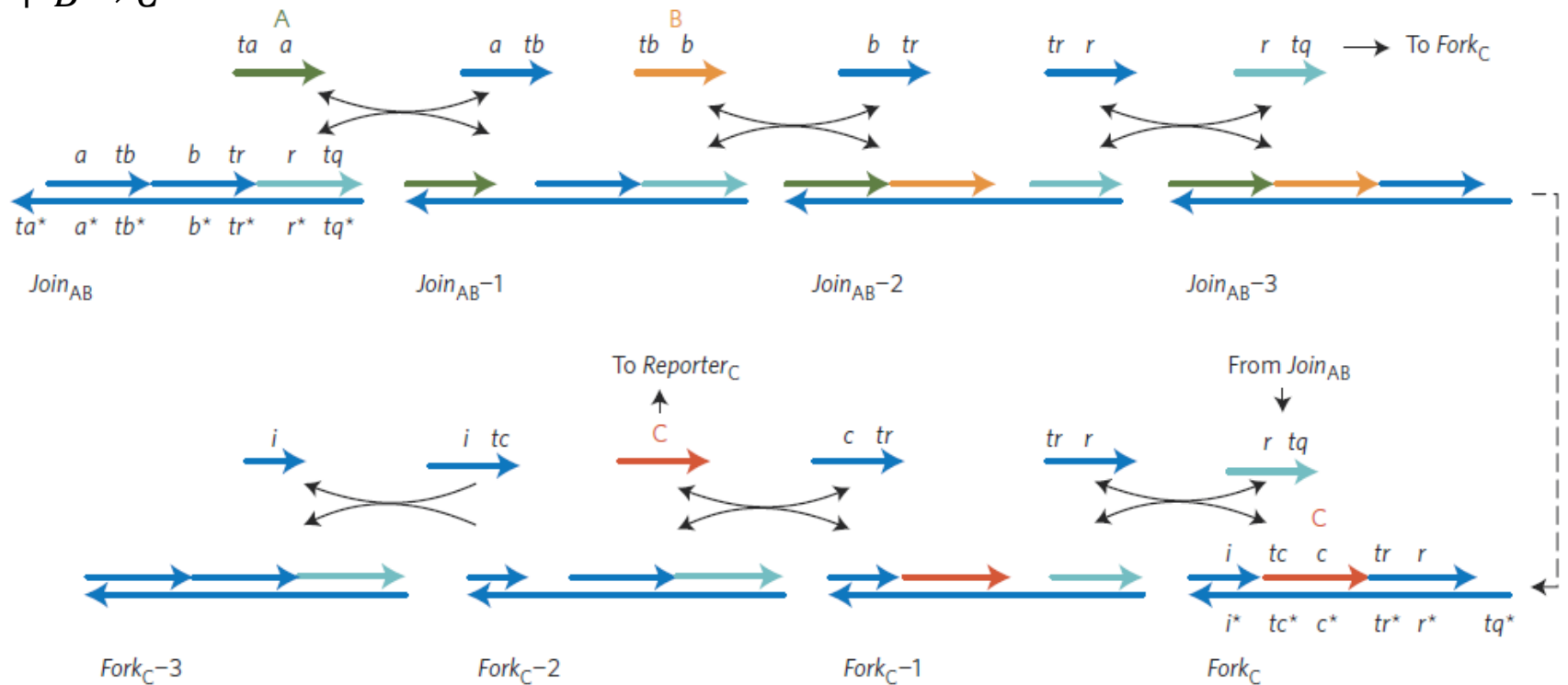
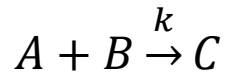




# CRNs to DNA implementation: a consensus network



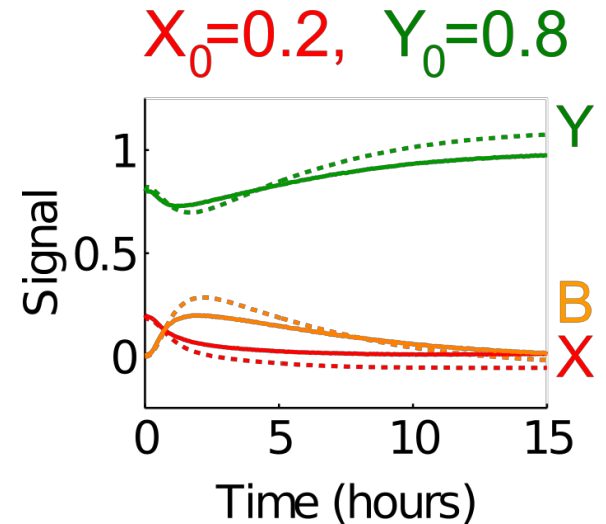
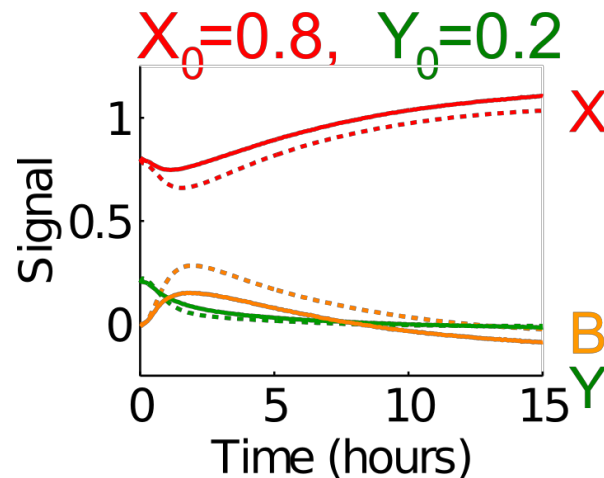
# CRNs to DNA implementation: a consensus network





# CRNs to DNA implementation: a consensus network

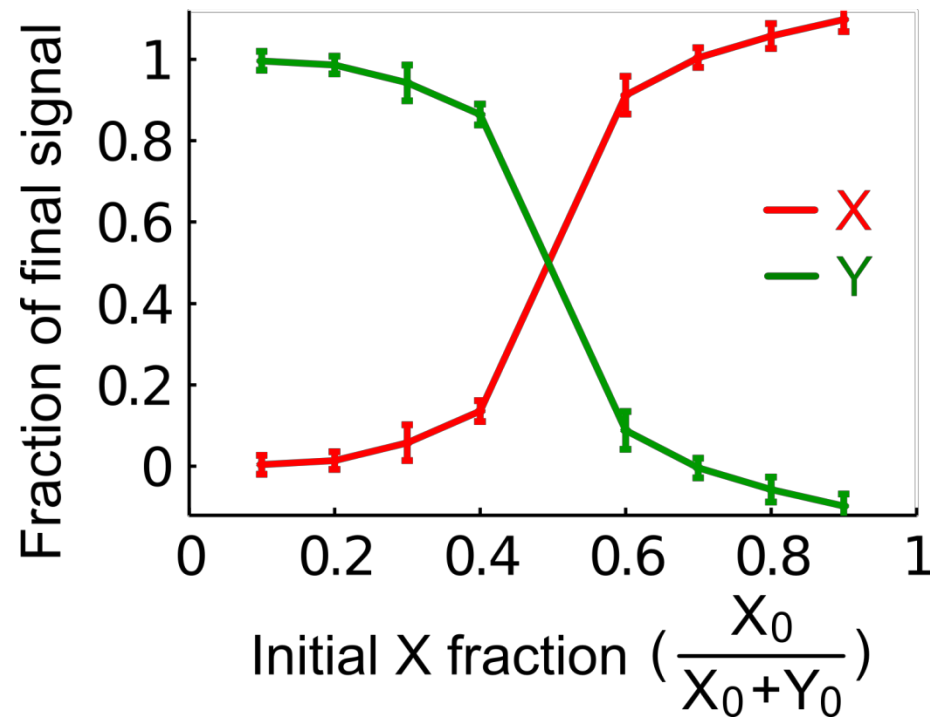
## CRN computation



Dashed lines: simulations  
Solid lines: experiments

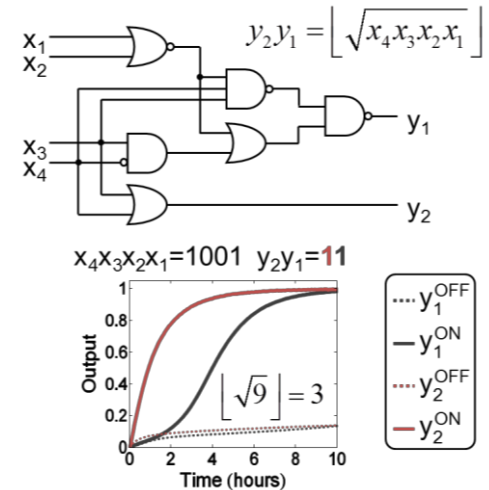
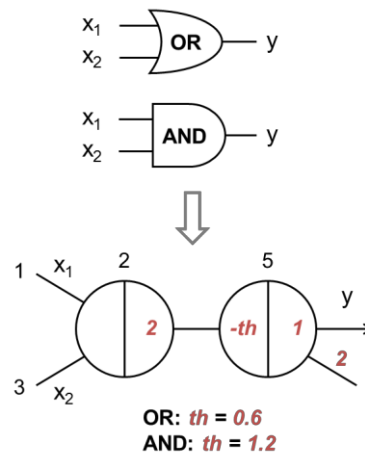
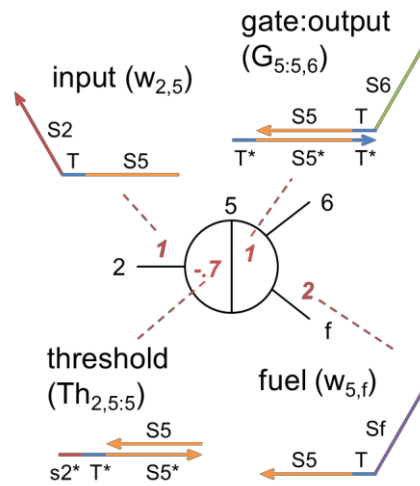
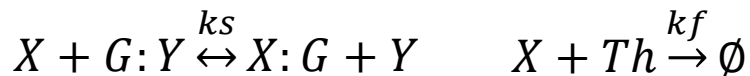
# CRNs to DNA implementation: a consensus network

## CRN computation



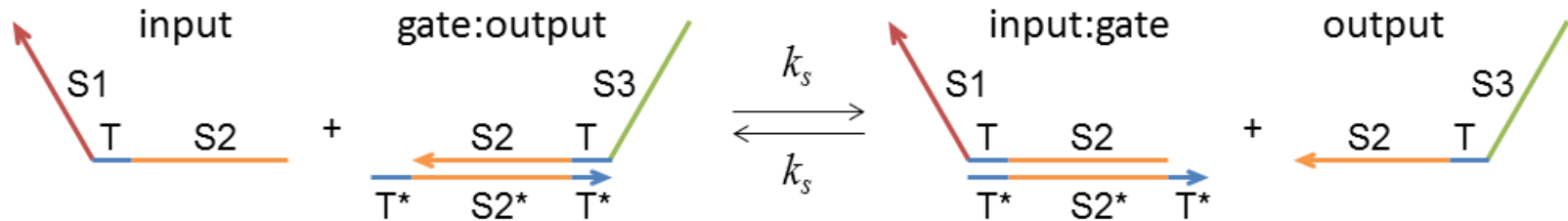
# What is the simplest DNA building blocks for creating CRNs with complex behaviors?

## How robustly can DNA-based CRNs scale up?

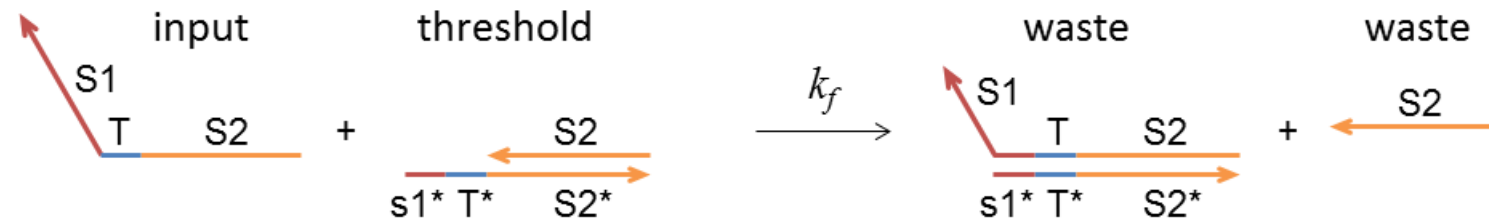


# Basic reactions in seesaw networks

## 1. seesawing



## 2. thresholding



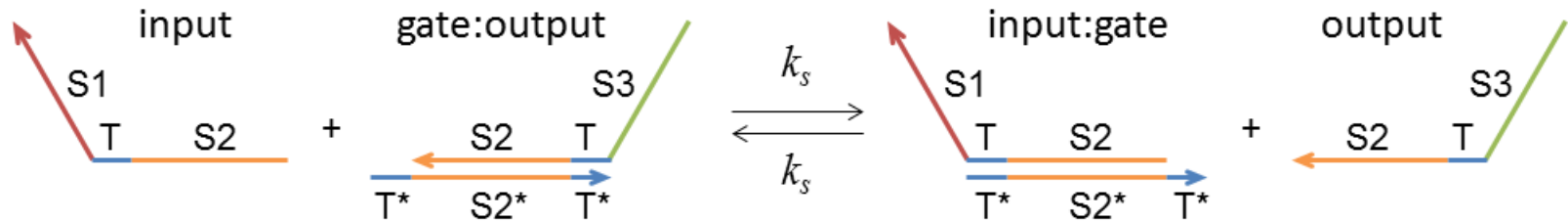
$$\begin{aligned}
 |T^*| &= 5nt & k_s &\approx 10^5/\text{M/s} \\
 |s1^* T^*| &= 8nt & k_f &\approx 10^6/\text{M/s} \\
 & & k_f &\gg k_s
 \end{aligned}$$

$$\begin{cases}
 \text{if } [\text{input}]|_{t=0} < [\text{threshold}]|_{t=0} & [\text{output}]|_{t \rightarrow \infty} \approx 0 \\
 \text{if } [\text{input}]|_{t=0} > [\text{threshold}]|_{t=0} & [\text{output}]|_{t \rightarrow \infty} \gg 0
 \end{cases}$$

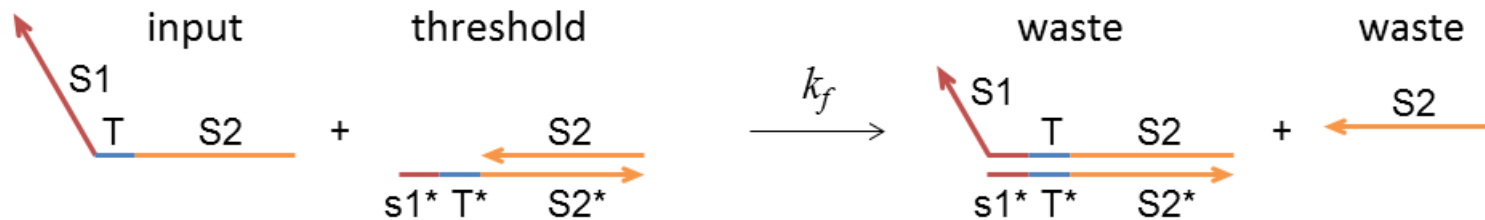
$$\begin{aligned}
 k_{eff} &\approx 10^L/\text{M/s} \text{ when } L \leq 6 \\
 &\text{otherwise } k_{eff} \approx 10^6/\text{M/s} \\
 &L: \text{toehold length}
 \end{aligned}$$

# Basic reactions in seesaw networks

## 1. seesawing

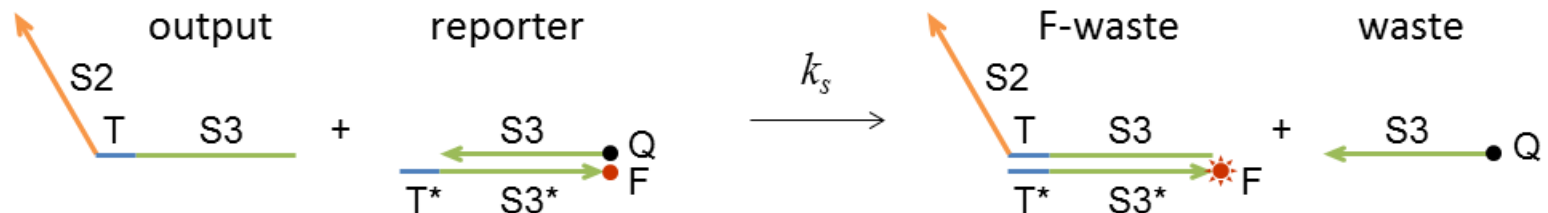


## 2. thresholding



## 3. reporting

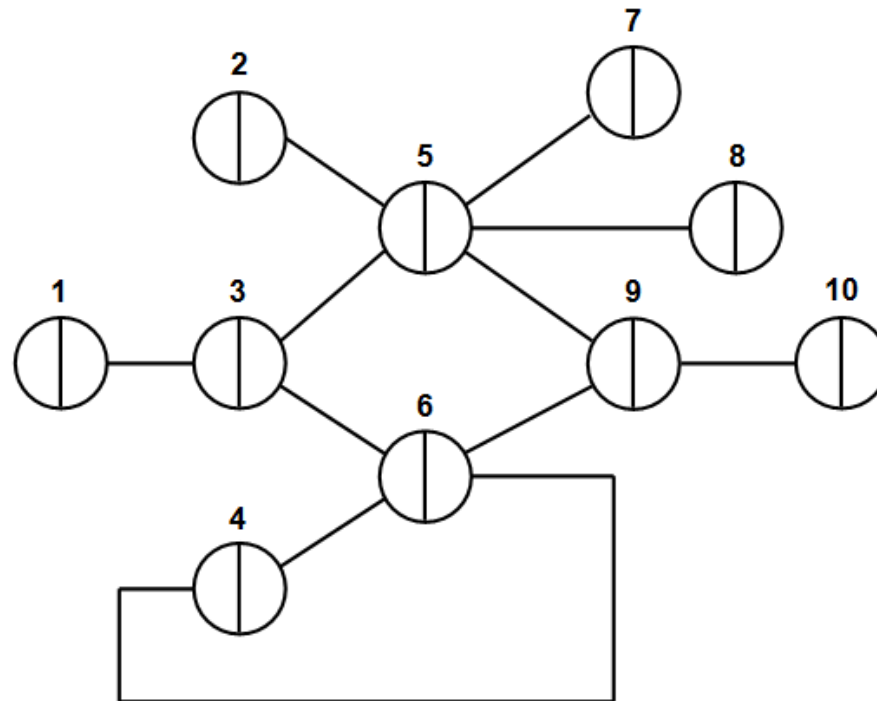
read the output of the computation with fluorescence signal



# Seesaw abstraction

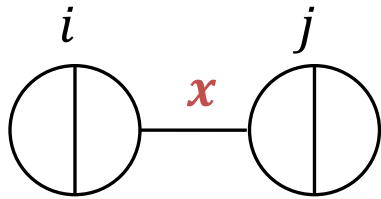
A seesaw network has a number of two-sided nodes and a number of wires. Each node can be connected to any number of wires on each side. Each wire connects exactly two nodes. Each node has an identity:  $i, i \in \{1, 2, 3, \dots\}$

Example:



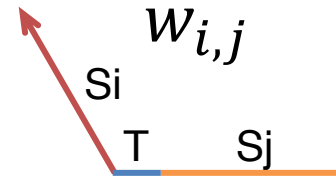
# Seesaw abstraction

## 1. free signal

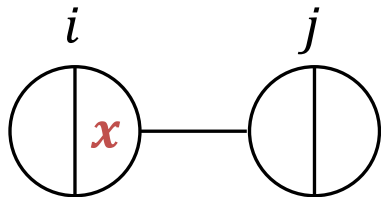


$x$  is relative to a standard concentration (e.g. 100nM)

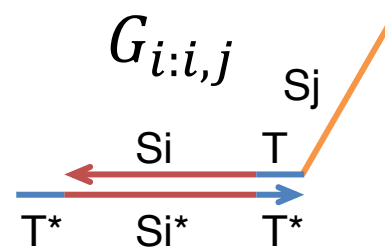
$$[w_{i,j}]|_{t=0} = x$$



## 2. bound signal

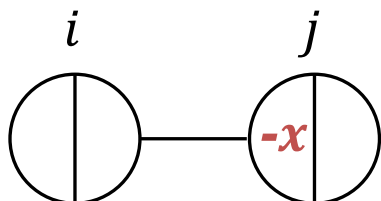


$$[G_{i:i,j}]|_{t=0} = x$$

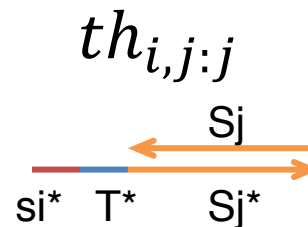


signal  $w_{i,j}$  bound to the right side of gate  $i$

## 3. threshold



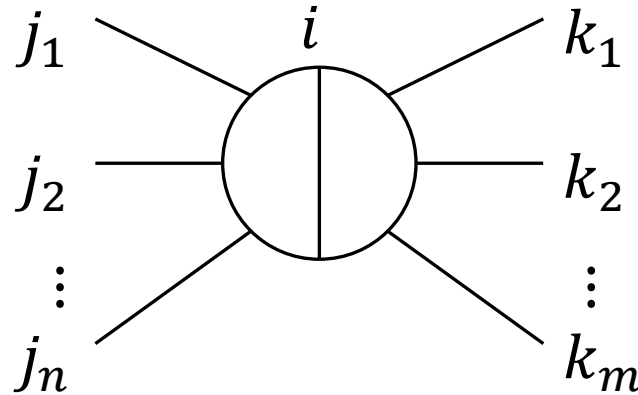
$$[th_{i,j:j}]|_{t=0} = x$$



threshold on gate  $j$  to absorb signal  $w_{i,j}$

# Seesaw abstraction

For any node  $i$  in a seesaw network:



for all  $j \in \{j_1, j_2, \dots, j_n\}$  and  $k \in \{k_1, k_2, \dots, k_m\}$

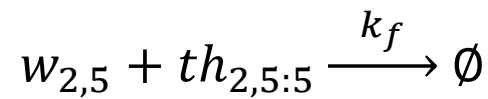
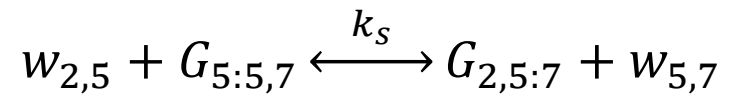
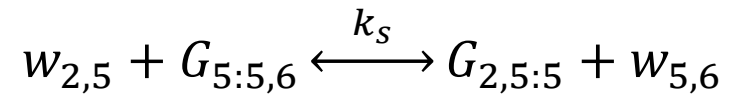
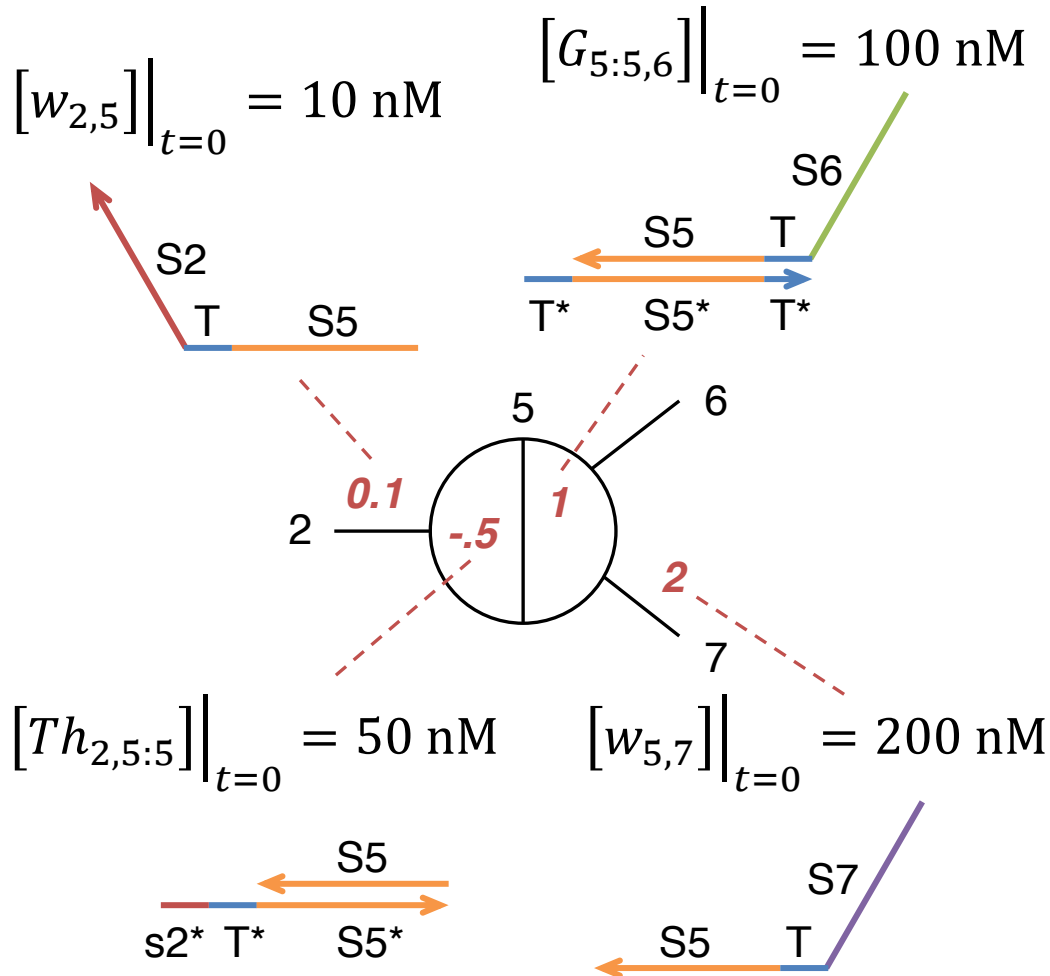
$$w_{j,i} + G_{i:i,k} \xleftrightarrow{k_s} G_{j,i:i} + w_{i,k}$$

$$w_{j,i} + th_{j,i:i} \xrightarrow{k_f} \emptyset$$



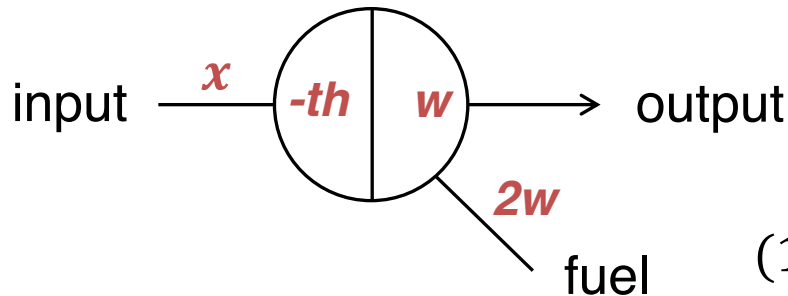
# Seesaw abstraction

Example: standard concentration is 100 nM



# Two types of seesaw gates

## 1. amplifying gate



$$[\text{input}]|_{t=0} = x \quad [\text{threshold}]|_{t=0} = th$$

$$[\text{gate:output}]|_{t=0} = w \quad [\text{fuel}]|_{t=0} = 2w$$

$$(1) \text{ input} + \text{threshold} \xrightarrow{k_f} \emptyset$$

$$(2) \text{ input} + \text{gate:output} \xrightarrow{k_s} \text{input:gate} + \text{output}$$

$$(3) \text{ input} + \text{gate:fuel} \xleftrightarrow{k_s} \text{input:gate} + \text{fuel}$$

pathway (net effect) of (2) and (3) with the above initial conditions:

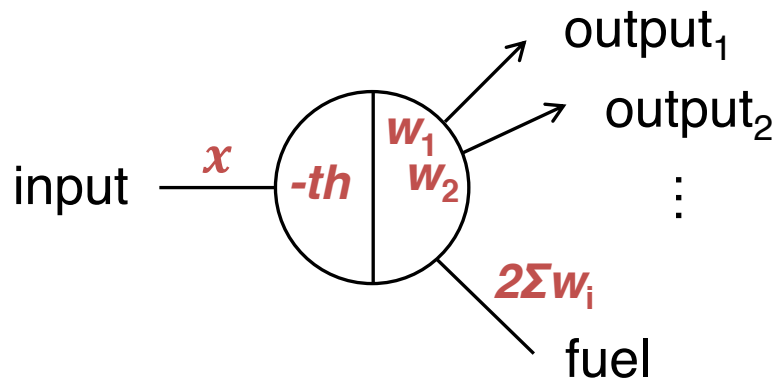
$$\text{input} + \text{gate:output} + \text{fuel} \longrightarrow \text{input} + \text{gate:fuel} + \text{output}$$

simplify:  $\text{input} \longrightarrow \text{input} + \text{output}$  when supply of gate:output and fuel last

$$\begin{cases} \text{if } x \leq th, [\text{output}]|_{t \rightarrow \infty} = 0 \\ \text{if } x > th, [\text{output}]|_{t \rightarrow \infty} = w \end{cases}$$

# Two types of seesaw gates

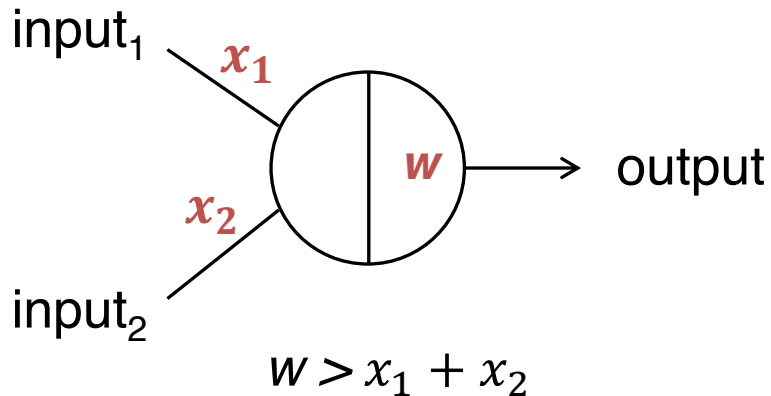
## 1. amplifying gate



$$\begin{cases} \text{if } x \leq th, [\text{output}_1]|_{t \rightarrow \infty} = 0, [\text{output}_2]|_{t \rightarrow \infty} = 0, \dots \\ \text{if } x > th, [\text{output}_1]|_{t \rightarrow \infty} = w_1, [\text{output}_2]|_{t \rightarrow \infty} = w_2, \dots \end{cases}$$

# Two types of seesaw gates

## 2. integrating gate



$$[\text{input}_1] \Big|_{t=0} = x_1$$

$$[\text{input}_2] \Big|_{t=0} = x_2$$

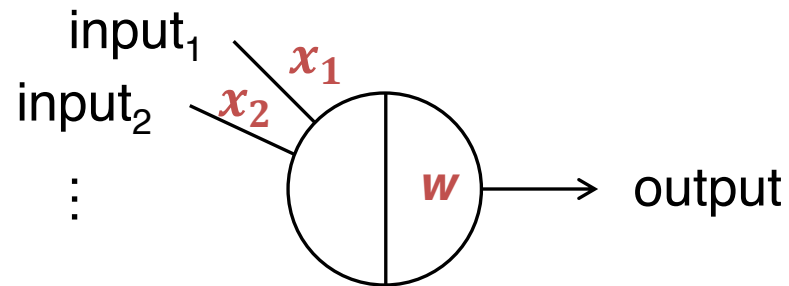
$$[\text{gate:output}] \Big|_{t=0} = w$$

$$\left\{ \begin{array}{l} \text{input}_1 + \text{gate:output} \xrightarrow{k_s} \text{input}_1:\text{gate} + \text{output} \\ \text{input}_2 + \text{gate:output} \xrightarrow{k_s} \text{input}_2:\text{gate} + \text{output} \end{array} \right.$$

$$\Rightarrow [\text{output}] \Big|_{t \rightarrow \infty} = x_1 + x_2$$

# Two types of seesaw gates

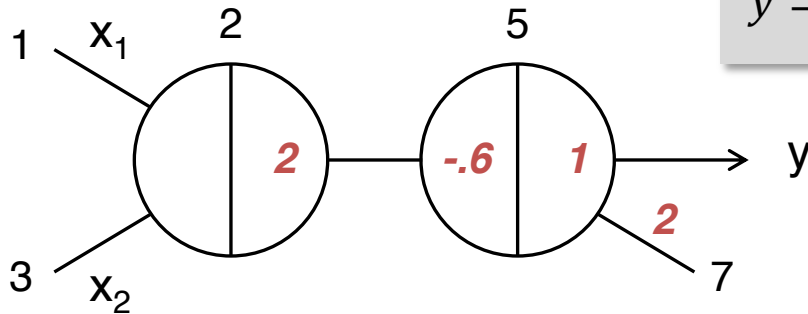
## 2. integrating gate



$$w > \sum x_i$$

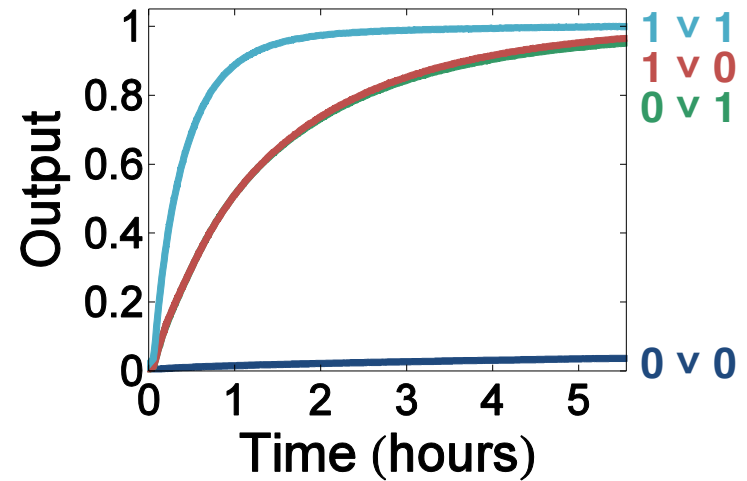
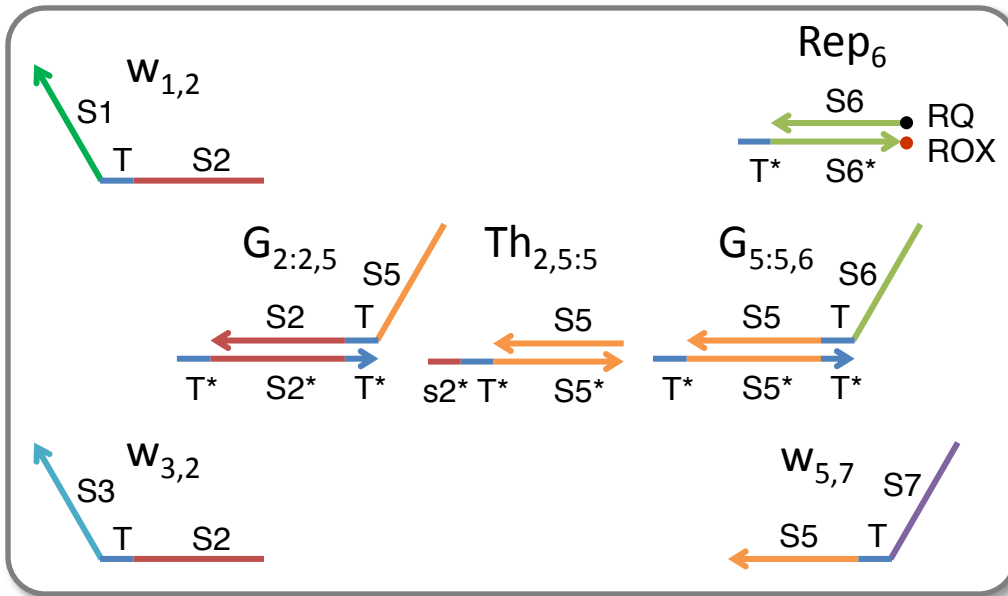
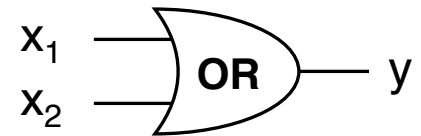
$$[\text{output}] \Big|_{t \rightarrow \infty} = \sum x_i$$

# Logic gates



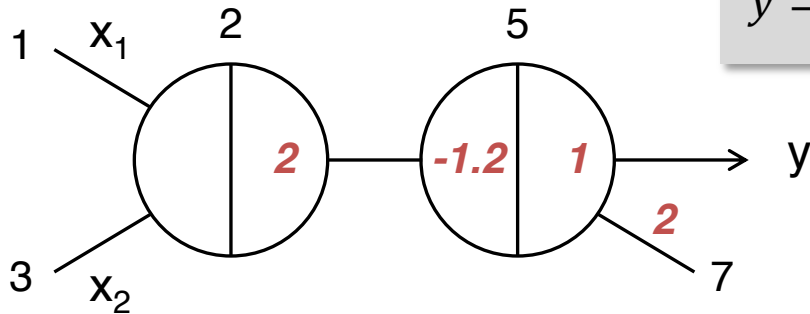
$$y = \begin{cases} 1 & x_1 + x_2 > 0.6 \\ 0 & x_1 + x_2 \leq 0.6 \end{cases}$$

OFF: 0 ~ 0.2  
ON: 0.8 ~ 1



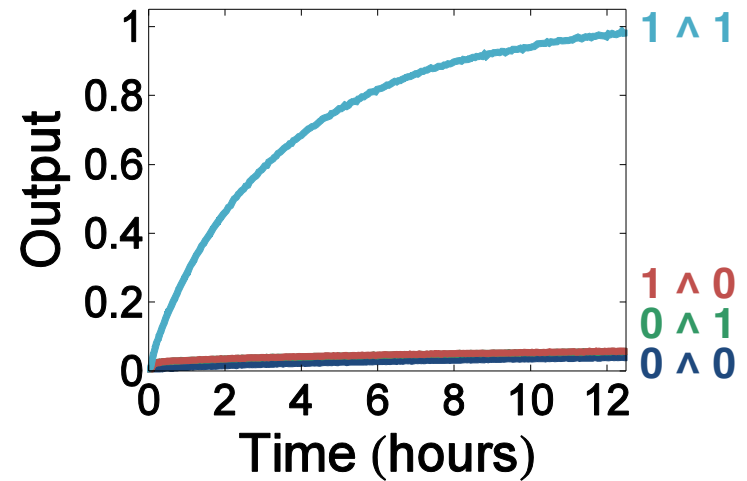
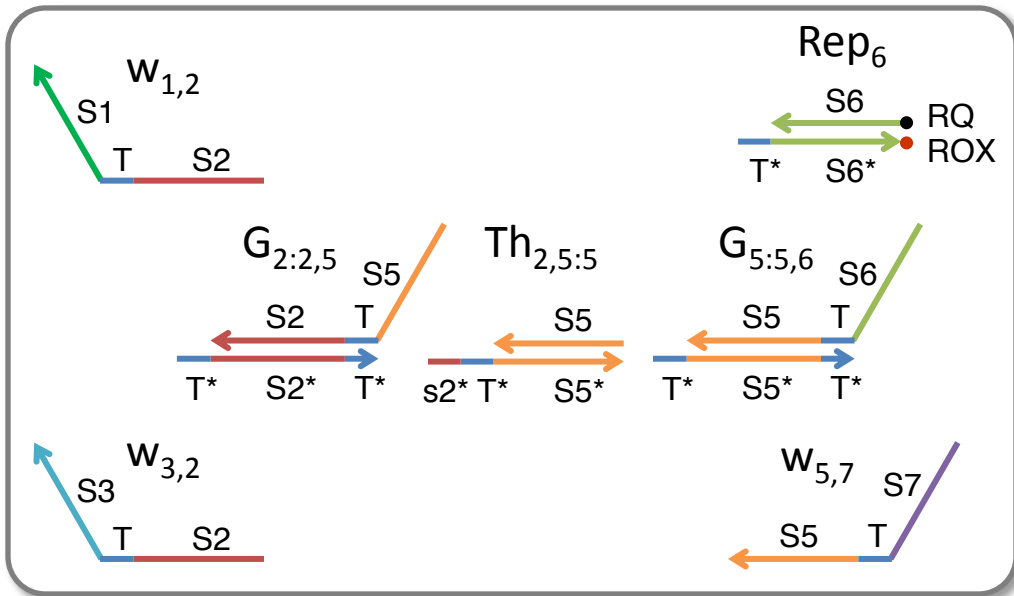
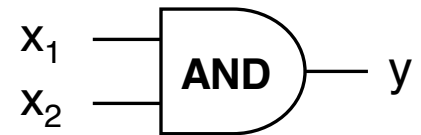
0=0.1x 1=0.9x 1x = 100 nM

# Logic gates



$$y = \begin{cases} 1 & x_1 + x_2 > 1.2 \\ 0 & x_1 + x_2 \leq 1.2 \end{cases}$$

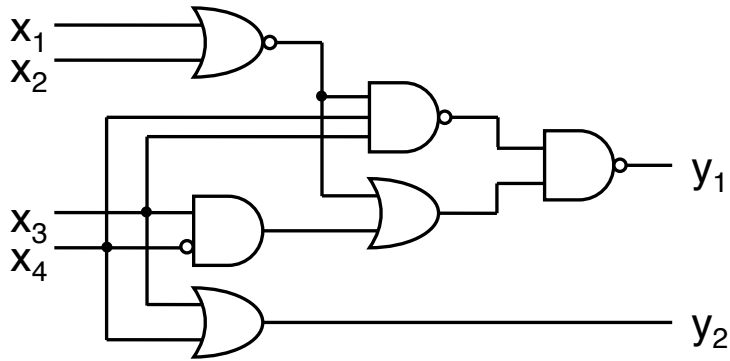
OFF: 0 ~ 0.2  
ON: 0.8 ~ 1



0=0.1x 1=0.9x 1x = 100 nM

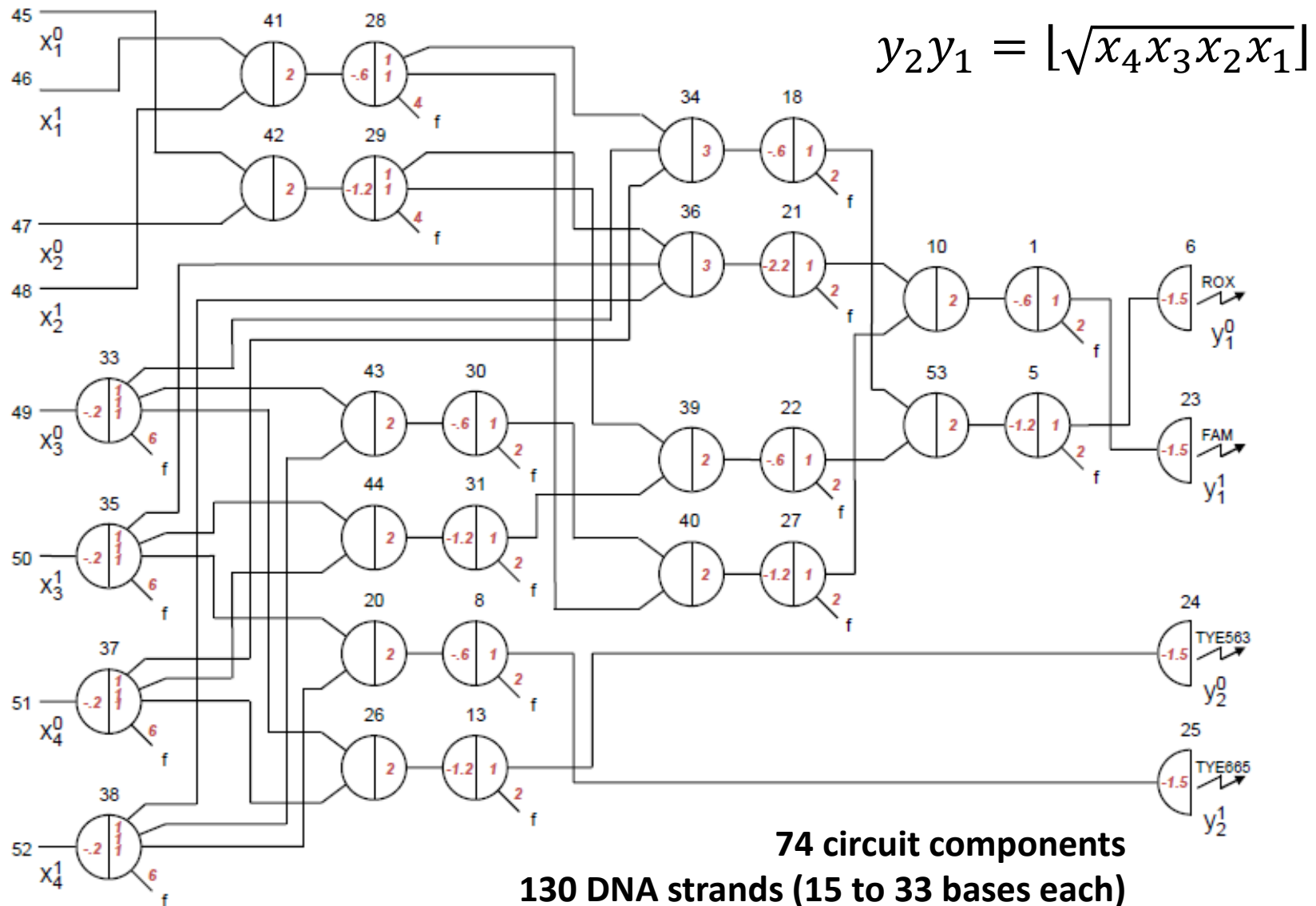
# A four-bit square root circuit

$$y_2 y_1 = \lfloor \sqrt{x_4 x_3 x_2 x_1} \rfloor$$



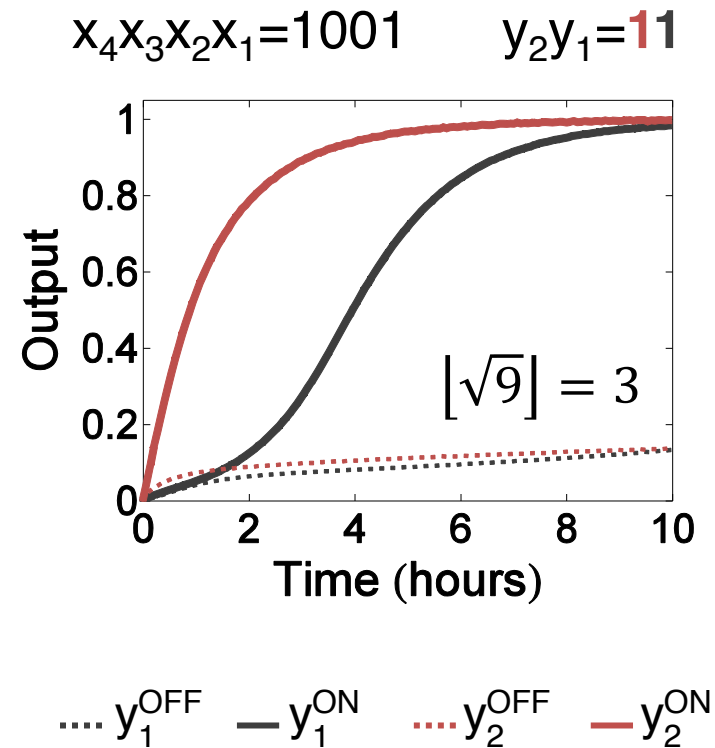
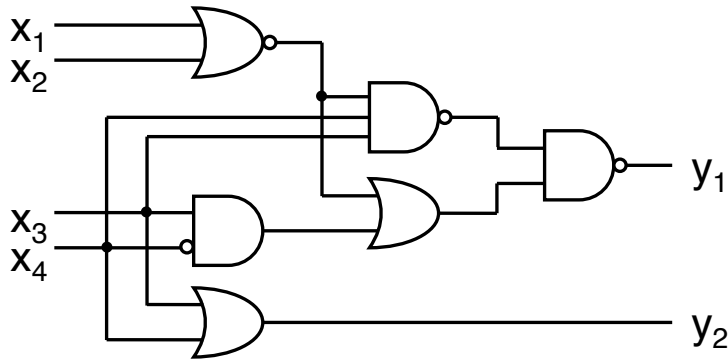


# A four-bit square root circuit



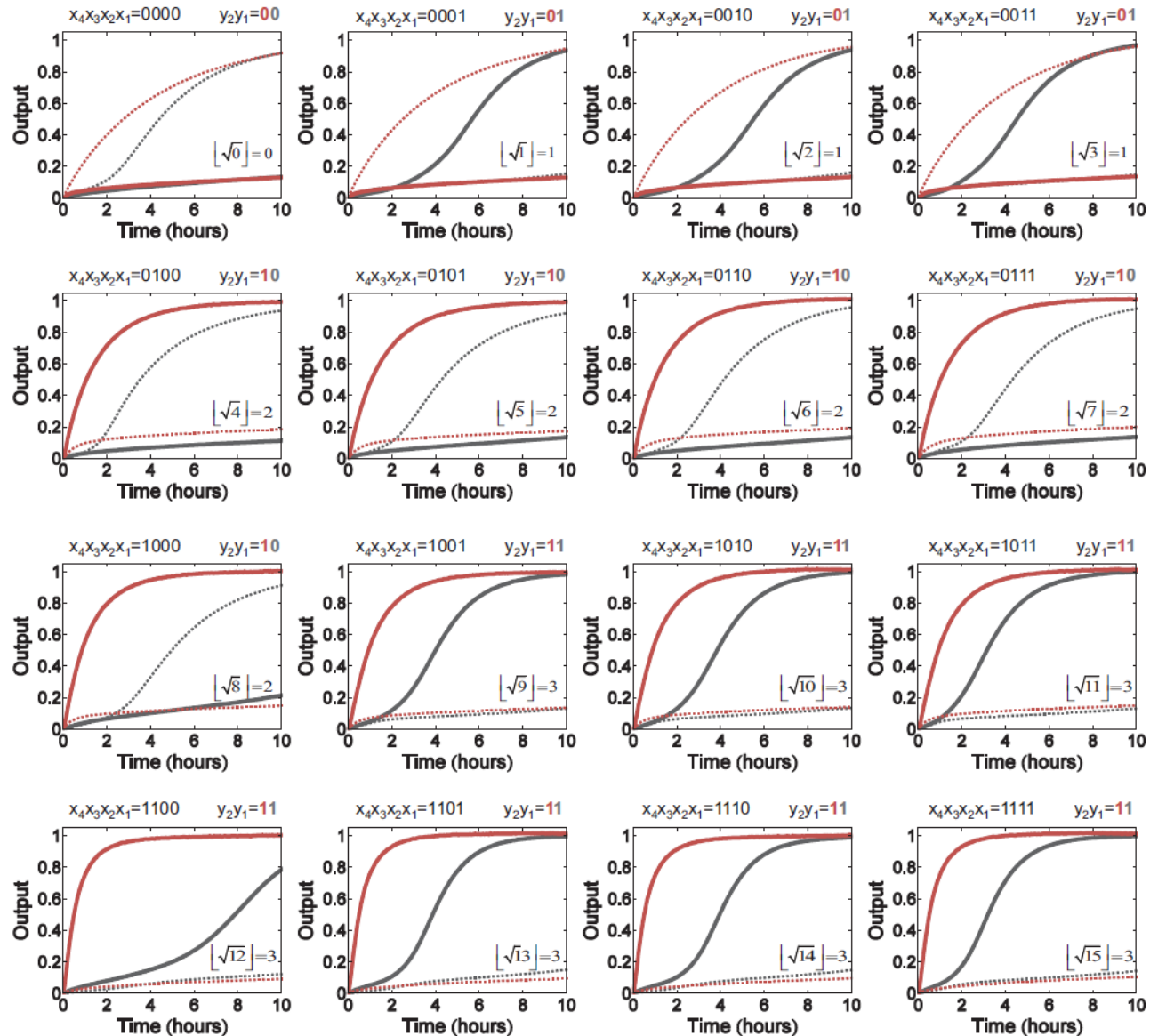
# A four-bit square root circuit

$$y_2 y_1 = \lfloor \sqrt{x_4 x_3 x_2 x_1} \rfloor$$



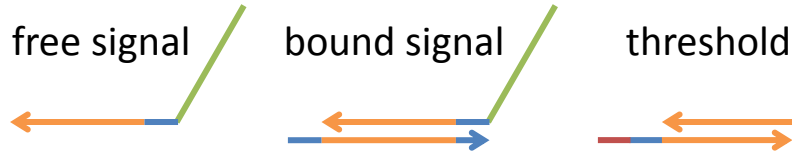
0=0.1x 1=0.9x 1x = 50 nM

# A four-bit square root circuit

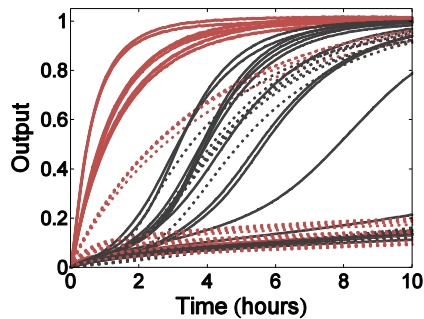


# Simplicity and robustness

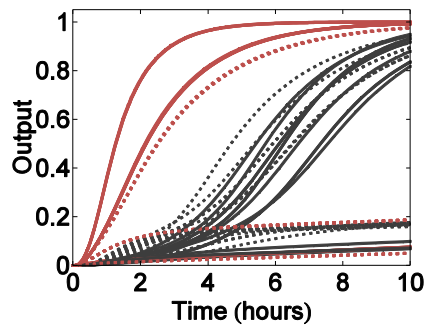
single- or double-stranded  
circuit components



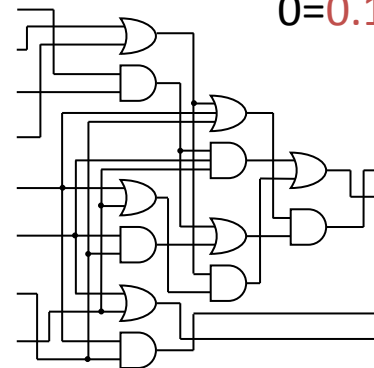
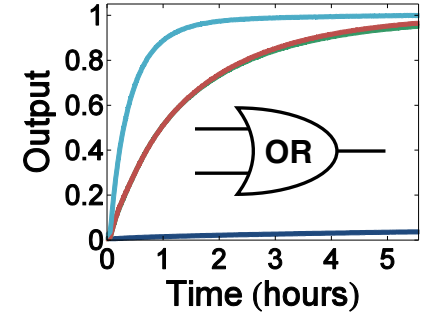
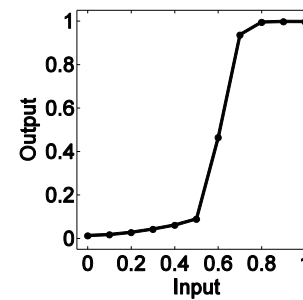
experiments



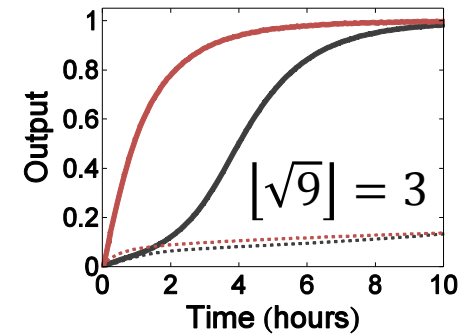
simulations



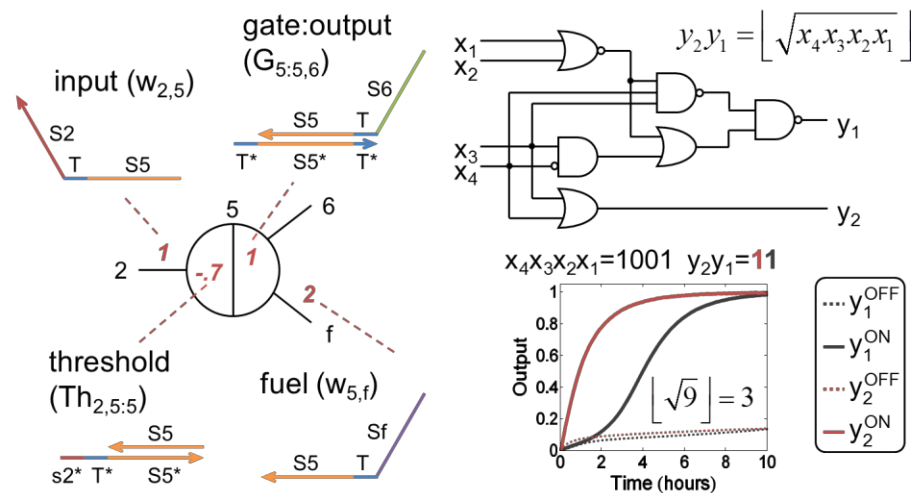
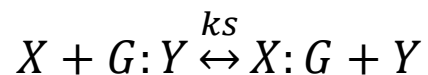
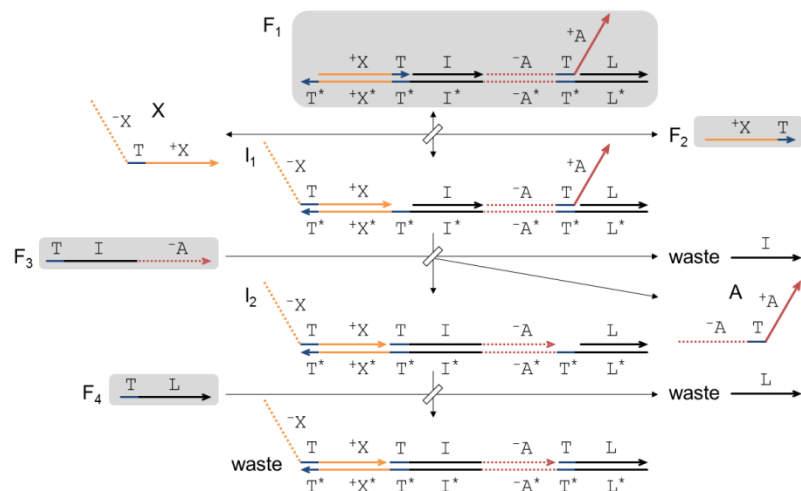
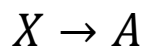
circuit performance  
stays roughly the same with size



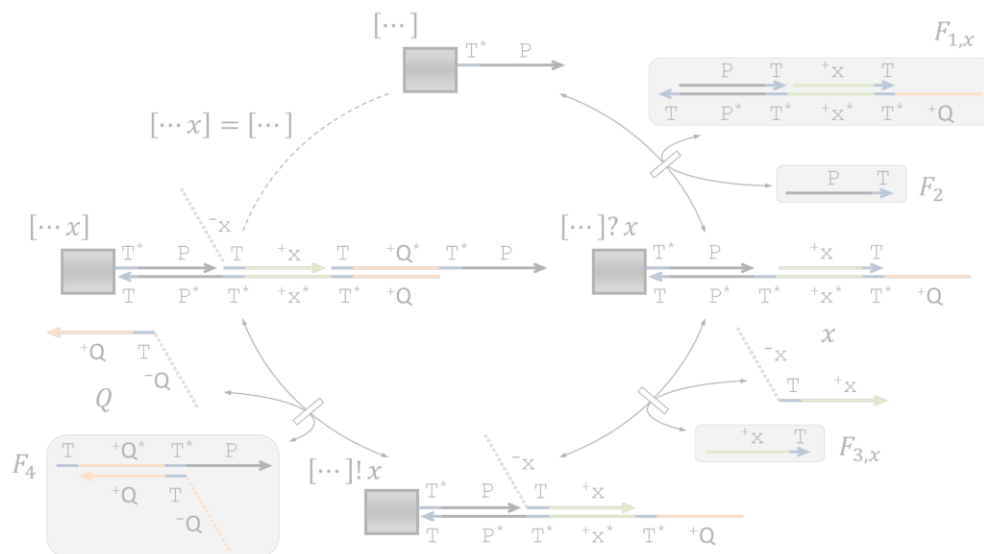
$$0 = 0.1x \quad 1 = 0.9x$$



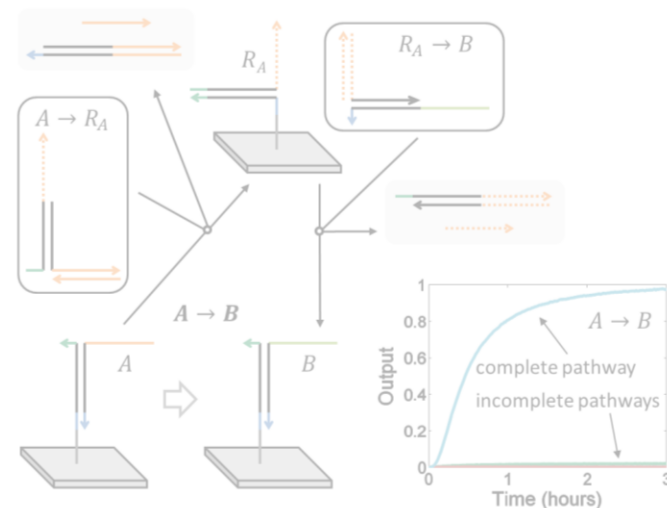
## Well-mixed CRNs



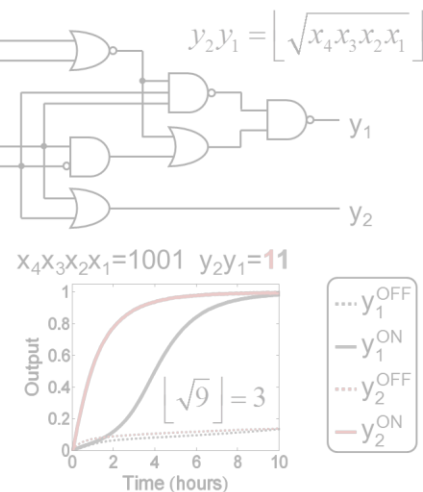
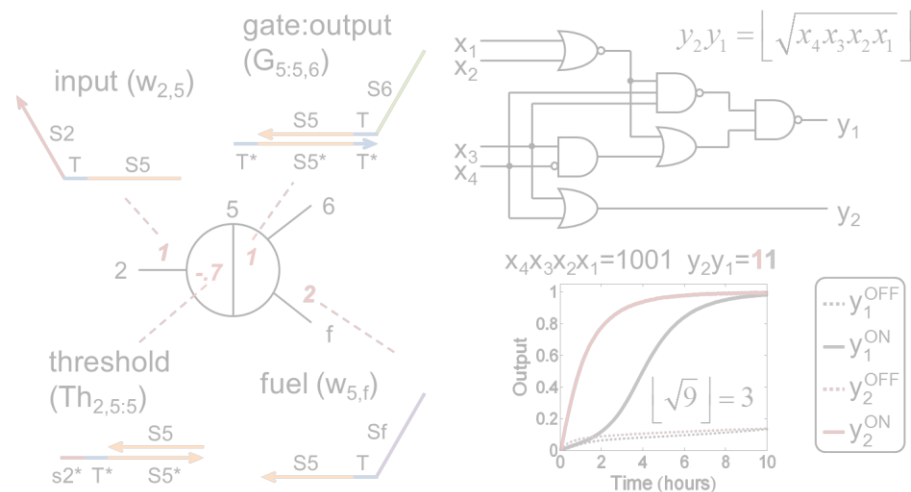
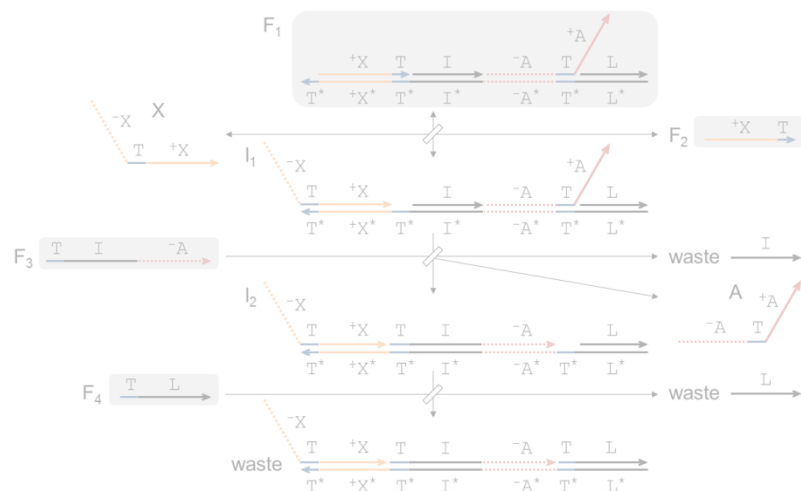
## Polymer CRNs



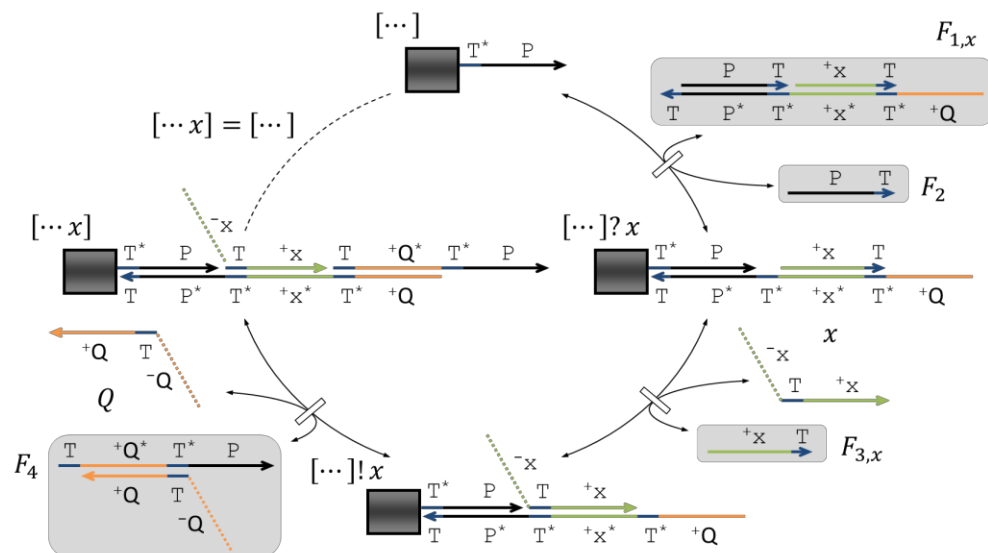
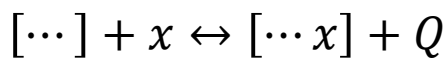
## Surface CRNs



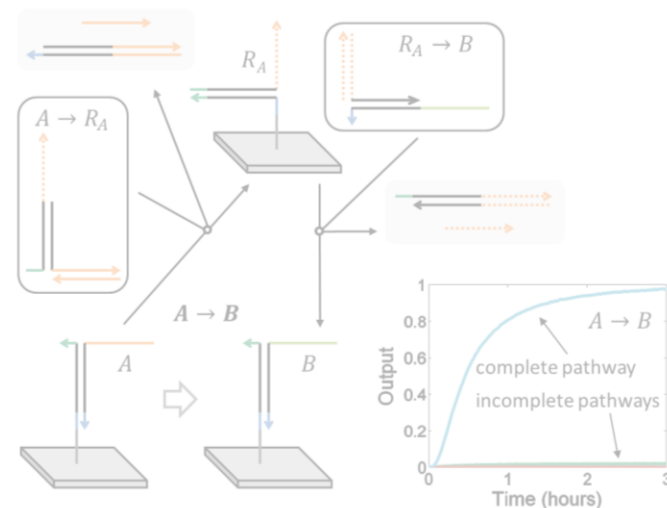
## Well-mixed CRNs



## Polymer CRNs



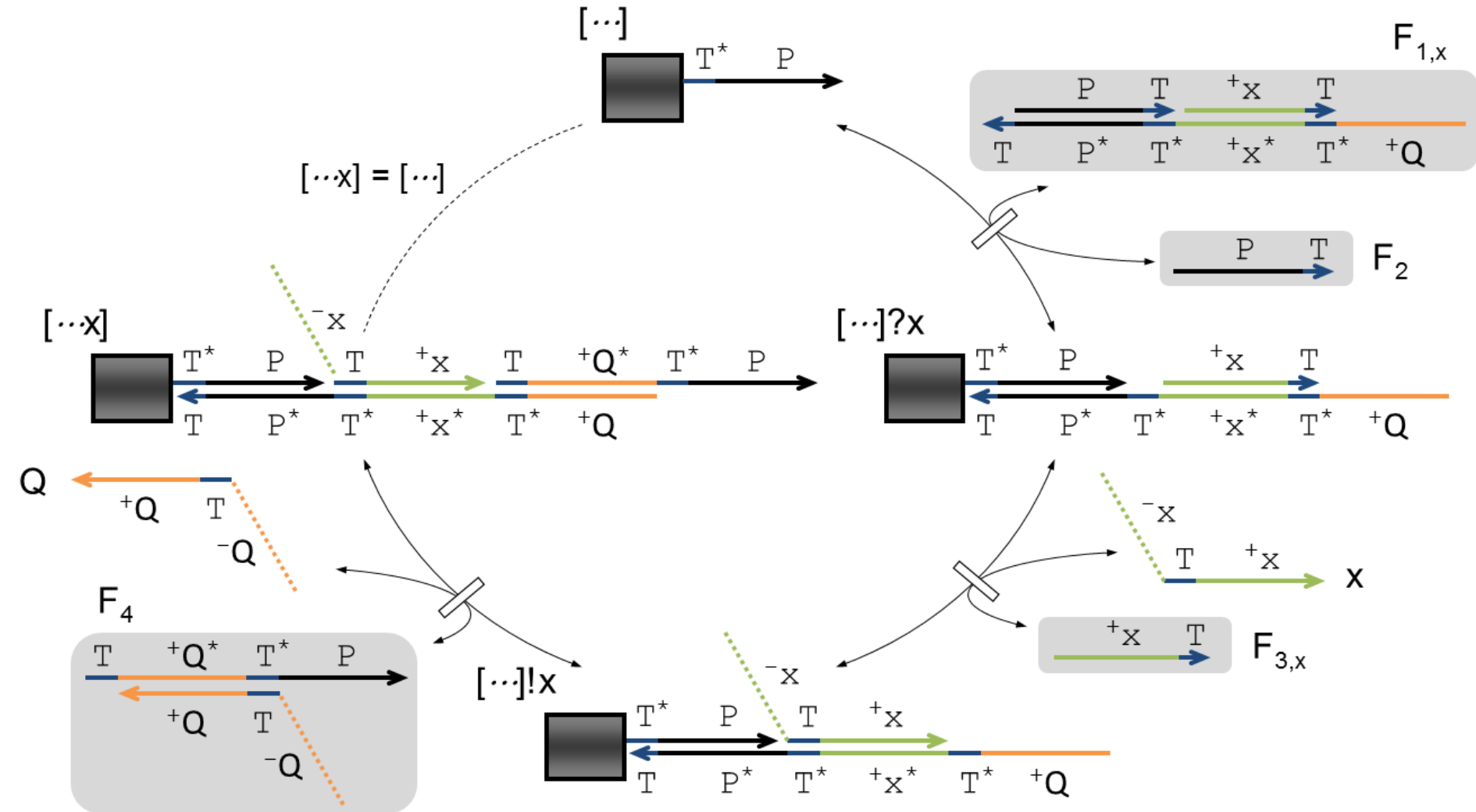
## Surface CRNs



# A DNA polymer reaction $[\dots] + x \leftrightarrow [\dots x] + Q$

push:  $[\dots] + x \rightarrow [\dots x] + Q$

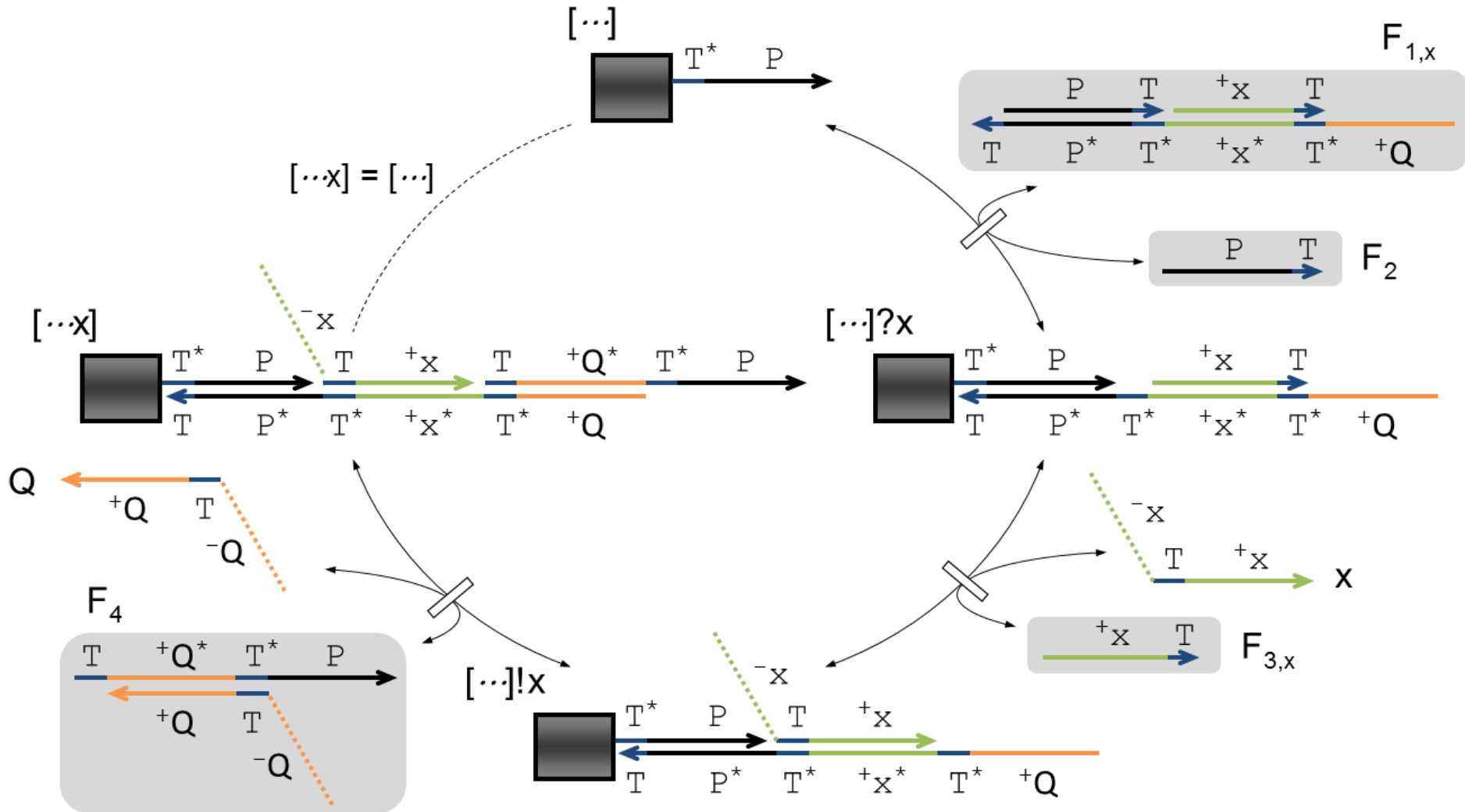
$Q$  is a confirmation signal



# A DNA polymer reaction $[\dots] + x \leftrightarrow [\dots x] + Q$

pop:  $[\dots x] + Q \rightarrow [\dots] + x$

$Q$  is a query signal





# A DNA stack machine implementation

transition rules:

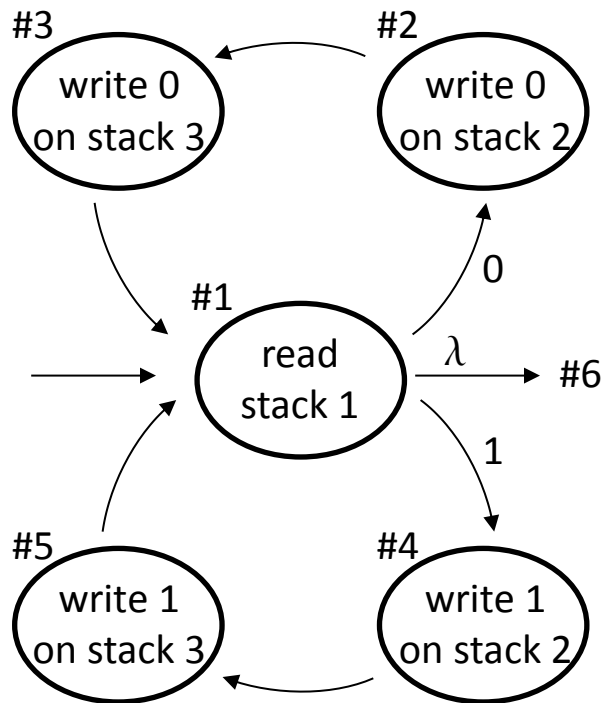
current state, pop symbol, stack number --> next state, push symbol, stack number

$$Q \rightleftharpoons Q_i$$

$$[\cdots]_i + x_i \rightleftharpoons [\cdots x]_i + Q_i$$

1.  $\alpha \ x \ i \longrightarrow \beta \ y \ j \quad \Rightarrow \quad S_\alpha + x_i \rightarrow S_\beta + y_j$
2.  $\alpha \ x \ i \longrightarrow \beta \quad \Rightarrow \quad S_\alpha + x_i \rightarrow S_\beta + Q$
3.  $\alpha \quad \longrightarrow \beta \ y \ j \quad \Rightarrow \quad S_\alpha + Q \rightarrow S_\beta + y_j$
4.  $\alpha \ \lambda \ i \longrightarrow \beta \ \lambda \ i \quad \Rightarrow \quad S_\alpha + \perp_i \rightarrow S_\beta + \perp_i$

# A DNA stack machine implementation



#1 0 1  $\rightarrow$  #2

#1 1 1  $\rightarrow$  #4

#1  $\lambda$  1  $\rightarrow$  #6  $\lambda$  1

#2  $\rightarrow$  #3 0 2

#3  $\rightarrow$  #1 0 3

#4  $\rightarrow$  #5 1 2

#5  $\rightarrow$  #1 1 3

1. (#1, 00111,  $\lambda$ ,  $\lambda$ )

2. (#4, 0011,  $\lambda$ ,  $\lambda$ )

3. (#5, 0011, 1,  $\lambda$ )

4. (#1, 0011, 1, 1)

5. (#4, 001, 1, 1)

6. (#5, 001, 11, 1)

7. (#1, 001, 11, 11)

8. (#4, 00, 11, 11)

9. (#5, 00, 111, 11)

10. (#1, 00, 111, 111)

11. (#2, 0, 111, 111)

12. (#3, 0, 1110, 111)

13. (#1, 0, 1110, 1110)

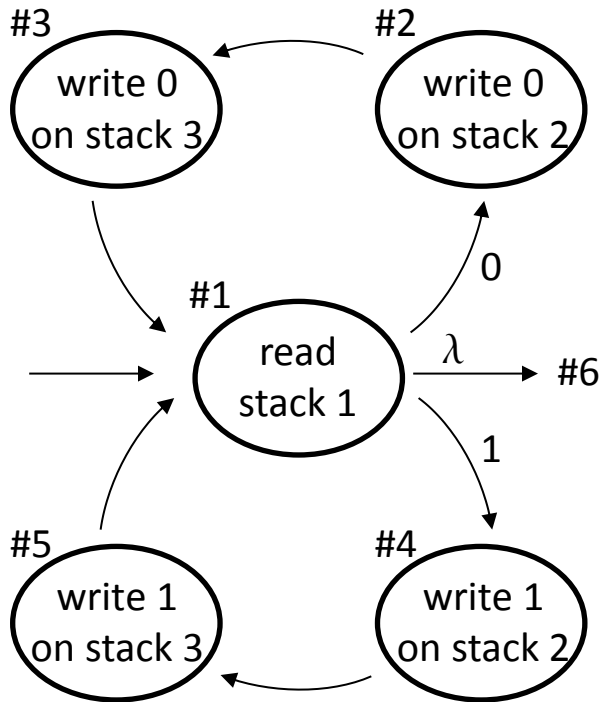
14. (#2,  $\lambda$ , 1110, 1110)

15. (#3,  $\lambda$ , 11100, 1110)

16. (#1,  $\lambda$ , 11100, 11100)

17. (#6,  $\lambda$ , 11100, 11100)

# A DNA stack machine implementation



$\#1 \ 0 \ 1 \longrightarrow \#2$   
 $\#1 \ 1 \ 1 \longrightarrow \#4$   
 $\#1 \ \lambda \ 1 \longrightarrow \#6 \ \lambda \ 1$   
 $\#2 \longrightarrow \#3 \ 0 \ 2$   
 $\#3 \longrightarrow \#1 \ 0 \ 3$   
 $\#4 \longrightarrow \#5 \ 1 \ 2$   
 $\#5 \longrightarrow \#1 \ 1 \ 3$

$S_{\#1} + 0_1 \longrightarrow S_{\#2} + Q$   
 $S_{\#1} + 1_1 \longrightarrow S_{\#4} + Q$   
 $S_{\#1} + \perp_1 \longrightarrow S_{\#6} + \perp_1$   
 $S_{\#2} + Q \longrightarrow S_{\#3} + 0_2$   
 $S_{\#3} + Q \longrightarrow S_{\#1} + 0_3$   
 $S_{\#4} + Q \longrightarrow S_{\#5} + 1_2$   
 $S_{\#5} + Q \longrightarrow S_{\#1} + 1_3$

$Q_1 \longleftrightarrow Q$

$Q_2 \longleftrightarrow Q$

$Q_3 \longleftrightarrow Q$

$[...]_1 + 0_1 \longleftrightarrow [...0]_1 + Q_1$

$[...]_1 + 1_1 \longleftrightarrow [...1]_1 + Q_1$

$[...]_2 + 0_2 \longleftrightarrow [...0]_2 + Q_2$

$[...]_2 + 1_2 \longleftrightarrow [...1]_2 + Q_2$

$[...]_3 + 0_3 \longleftrightarrow [...0]_3 + Q_3$

$[...]_3 + 1_3 \longleftrightarrow [...1]_3 + Q_3$

# A DNA stack machine implementation

$$S_{\#1} + 0_1 \longrightarrow S_{\#2} + Q$$

$$S_{\#1} + 1_1 \longrightarrow S_{\#4} + Q$$

$$S_{\#1} + \perp_1 \longrightarrow S_{\#6} + \perp_1$$

$$S_{\#2} + Q \longrightarrow S_{\#3} + 0_2$$

$$S_{\#3} + Q \longrightarrow S_{\#1} + 0_3$$

$$S_{\#4} + Q \longrightarrow S_{\#5} + 1_2$$

$$S_{\#5} + Q \longrightarrow S_{\#1} + 1_3$$

$$Q_1 \longleftrightarrow Q$$

$$Q_2 \longleftrightarrow Q$$

$$Q_3 \longleftrightarrow Q$$

$$[\dots]_1 + 0_1 \longleftrightarrow [\dots 0]_1 + Q_1$$

$$[\dots]_1 + 1_1 \longleftrightarrow [\dots 1]_1 + Q_1$$

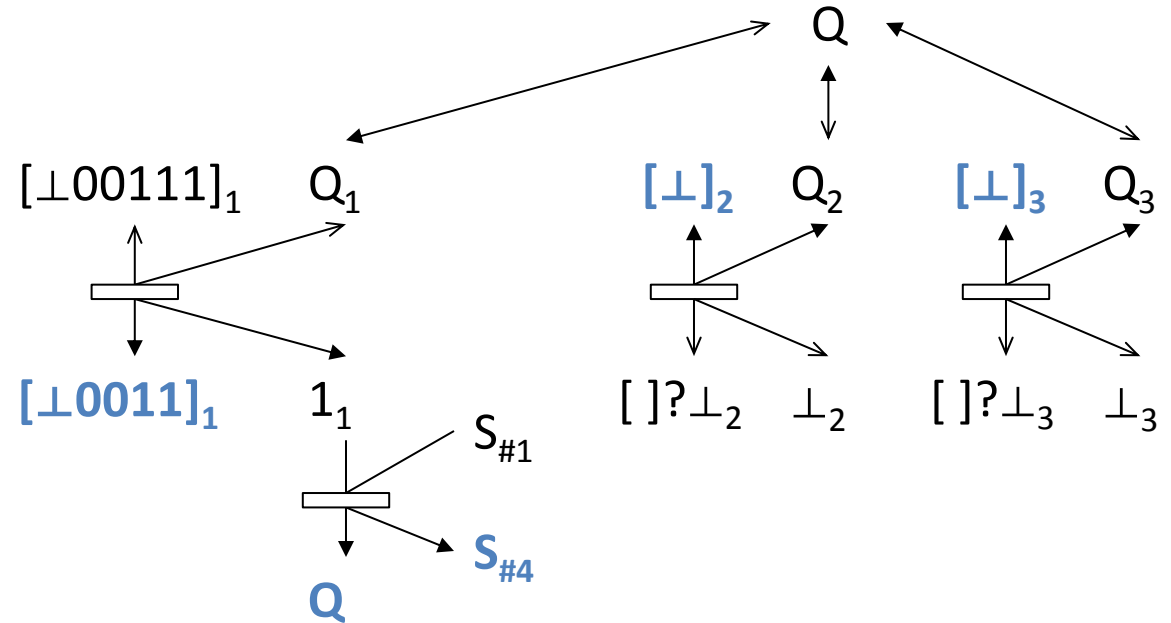
$$[\dots]_2 + 0_2 \longleftrightarrow [\dots 0]_2 + Q_2$$

$$[\dots]_2 + 1_2 \longleftrightarrow [\dots 1]_2 + Q_2$$

$$[\dots]_3 + 0_3 \longleftrightarrow [\dots 0]_3 + Q_3$$

$$[\dots]_3 + 1_3 \longleftrightarrow [\dots 1]_3 + Q_3$$

1.  $S_{\#1}, Q, [\perp 00111]_1, [\perp]_2, [\perp]_3$



2.  $S_{\#4}, Q, [\perp 00111]_1, [\perp]_2, [\perp]_3$

# A DNA stack machine implementation

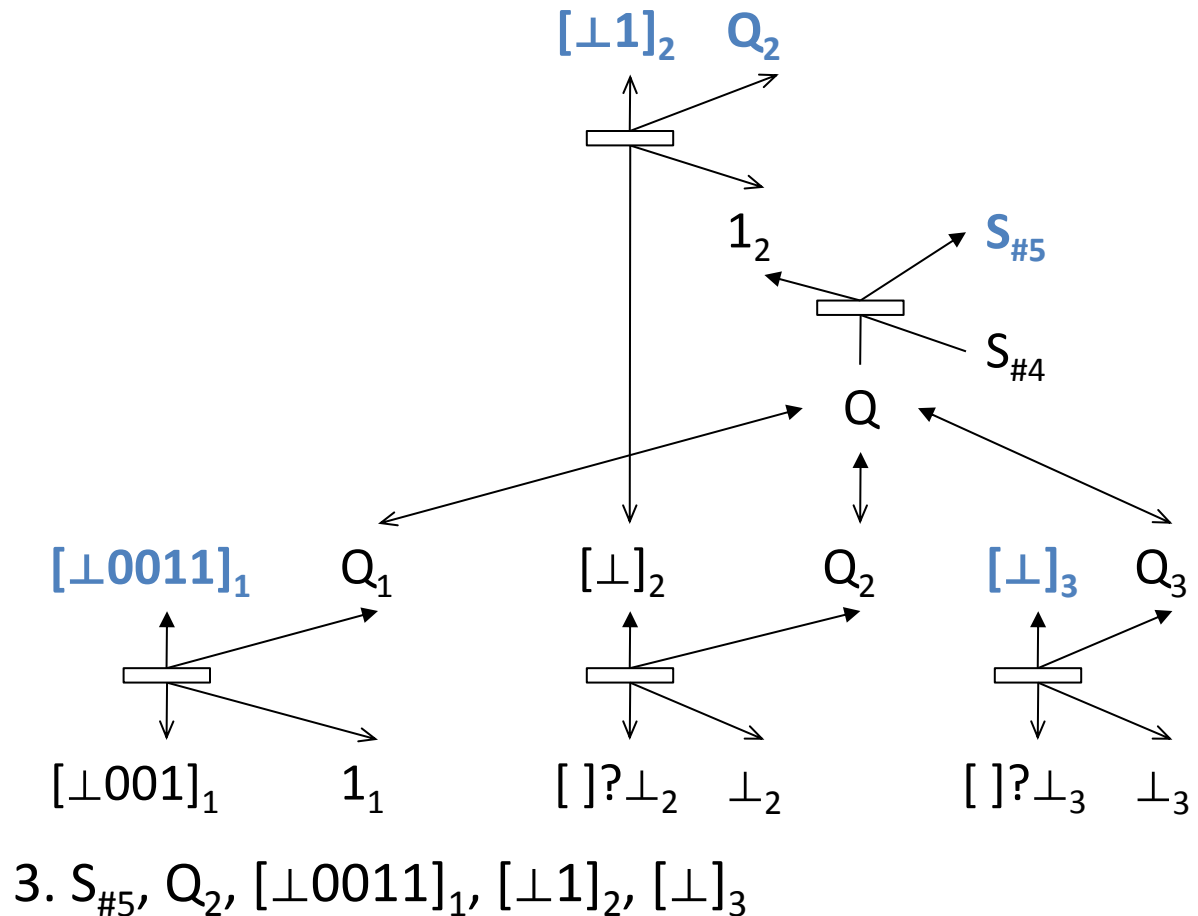
$$\begin{aligned}
 S_{\#1} + 0_1 &\longrightarrow S_{\#2} + Q \\
 S_{\#1} + 1_1 &\longrightarrow S_{\#4} + Q \\
 S_{\#1} + \perp_1 &\longrightarrow S_{\#6} + \perp_1 \\
 S_{\#2} + Q &\longrightarrow S_{\#3} + 0_2 \\
 S_{\#3} + Q &\longrightarrow S_{\#1} + 0_3 \\
 \boxed{S_{\#4} + Q &\longrightarrow S_{\#5} + 1_2} \\
 S_{\#5} + Q &\longrightarrow S_{\#1} + 1_3
 \end{aligned}$$

$$\begin{aligned}
 Q_1 &\longleftrightarrow Q \\
 Q_2 &\longleftrightarrow Q \\
 Q_3 &\longleftrightarrow Q
 \end{aligned}$$

$$\begin{aligned}
 [\dots]_1 + 0_1 &\longleftrightarrow [\dots 0]_1 + Q_1 \\
 [\dots]_1 + 1_1 &\longleftrightarrow [\dots 1]_1 + Q_1 \\
 [\dots]_2 + 0_2 &\longleftrightarrow [\dots 0]_2 + Q_2 \\
 [\dots]_2 + 1_2 &\longleftrightarrow [\dots 1]_2 + Q_2 \\
 [\dots]_3 + 0_3 &\longleftrightarrow [\dots 0]_3 + Q_3 \\
 [\dots]_3 + 1_3 &\longleftrightarrow [\dots 1]_3 + Q_3
 \end{aligned}$$

$$1. S_{\#1}, Q, [\perp 00111]_1, [\perp]_2, [\perp]_3$$

$$2. S_{\#4}, Q, [\perp 0011]_1, [\perp]_2, [\perp]_3$$



# A DNA stack machine implementation

$$\begin{aligned}
 S_{\#1} + 0_1 &\longrightarrow S_{\#2} + Q \\
 S_{\#1} + 1_1 &\longrightarrow S_{\#4} + Q \\
 S_{\#1} + \perp_1 &\longrightarrow S_{\#6} + \perp_1 \\
 S_{\#2} + Q &\longrightarrow S_{\#3} + 0_2 \\
 S_{\#3} + Q &\longrightarrow S_{\#1} + 0_3 \\
 S_{\#4} + Q &\longrightarrow S_{\#5} + 1_2 \\
 \boxed{S_{\#5} + Q &\longrightarrow S_{\#1} + 1_3}
 \end{aligned}$$

$$Q_1 \longleftrightarrow Q$$

$$Q_2 \longleftrightarrow Q$$

$$Q_3 \longleftrightarrow Q$$

$$[\dots]_1 + 0_1 \longleftrightarrow [\dots 0]_1 + Q_1$$

$$[\dots]_1 + 1_1 \longleftrightarrow [\dots 1]_1 + Q_1$$

$$[\dots]_2 + 0_2 \longleftrightarrow [\dots 0]_2 + Q_2$$

$$[\dots]_2 + 1_2 \longleftrightarrow [\dots 1]_2 + Q_2$$

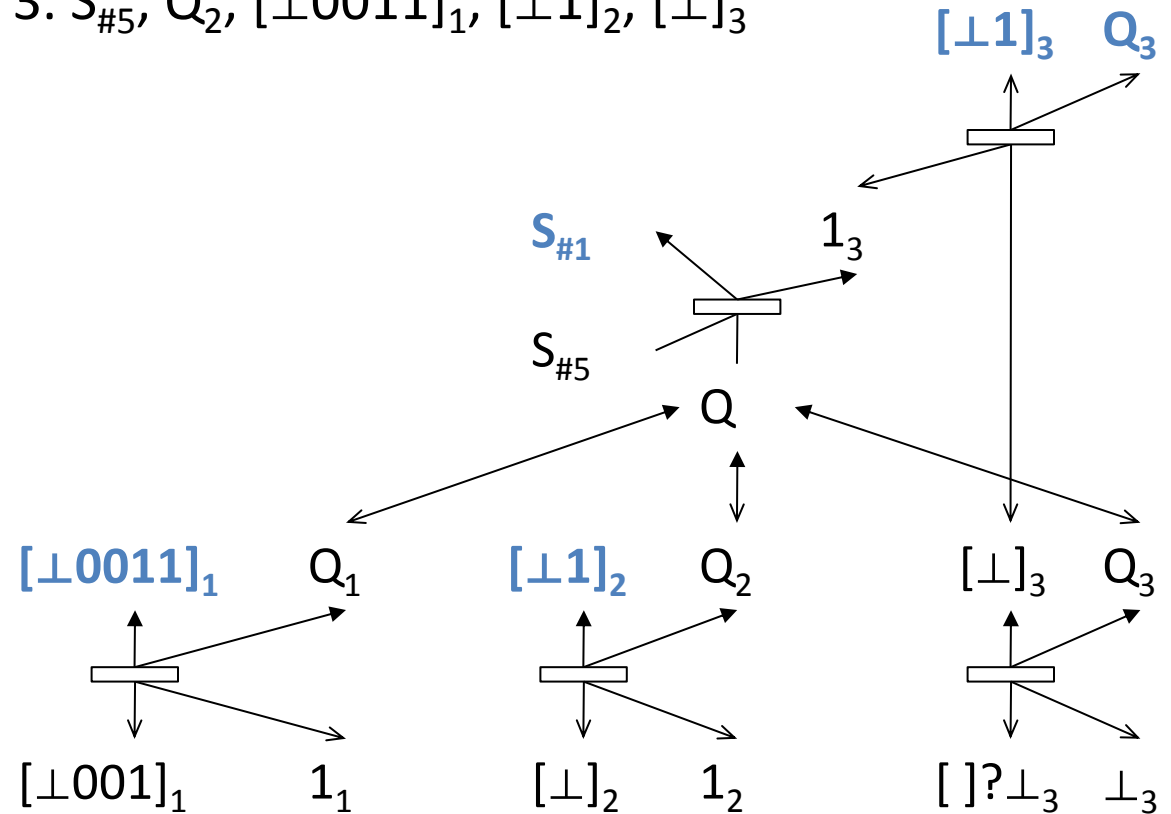
$$[\dots]_3 + 0_3 \longleftrightarrow [\dots 0]_3 + Q_3$$

$$[\dots]_3 + 1_3 \longleftrightarrow [\dots 1]_3 + Q_3$$

$$1. S_{\#1}, Q, [\perp 00111]_1, [\perp]_2, [\perp]_3$$

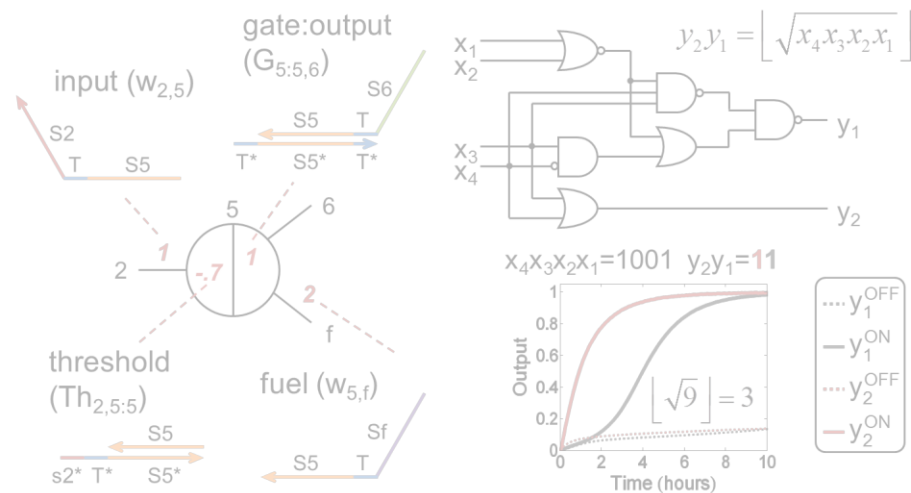
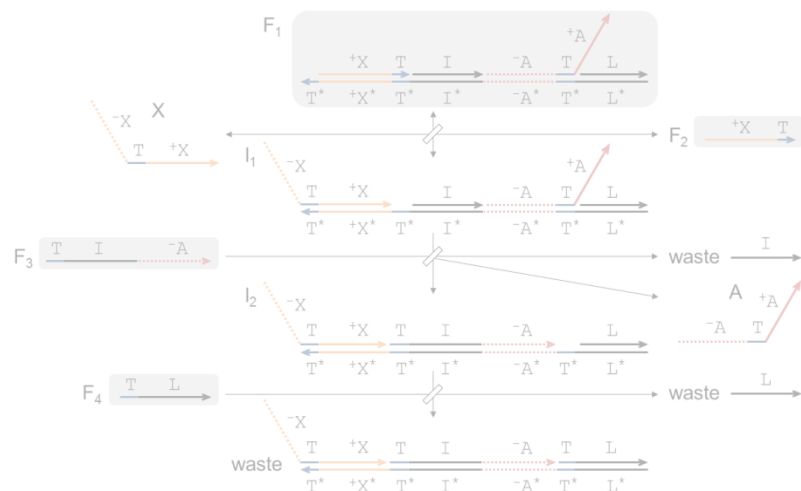
$$2. S_{\#4}, Q, [\perp 0011]_1, [\perp]_2, [\perp]_3$$

$$3. S_{\#5}, Q_2, [\perp 0011]_1, [\perp 1]_2, [\perp]_3$$

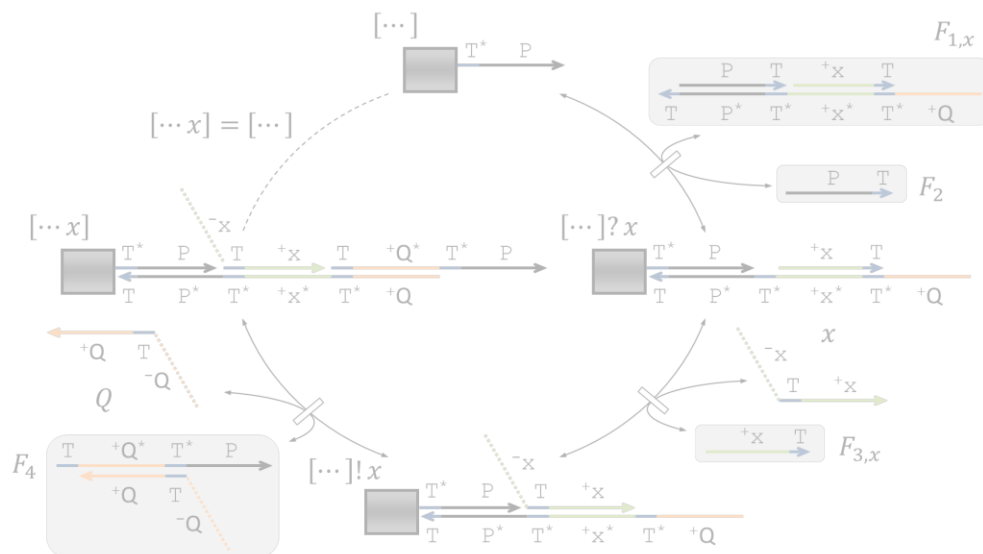


$$4. S_{\#1}, Q_3, [\perp 0011]_1, [\perp 1]_2, [\perp 1]_3$$

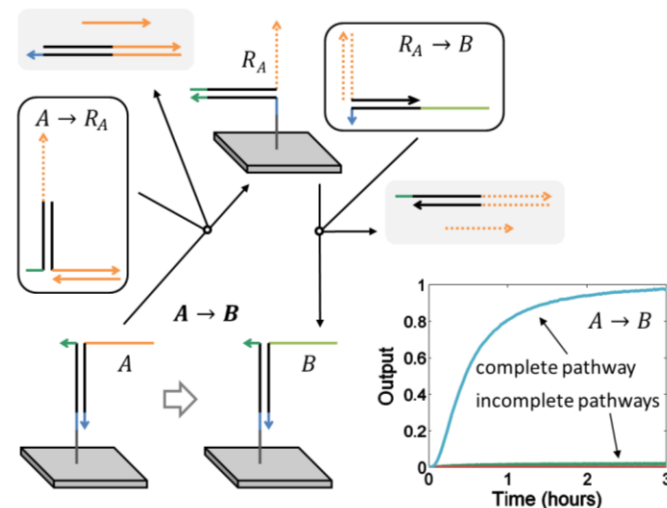
## Well-mixed CRNs



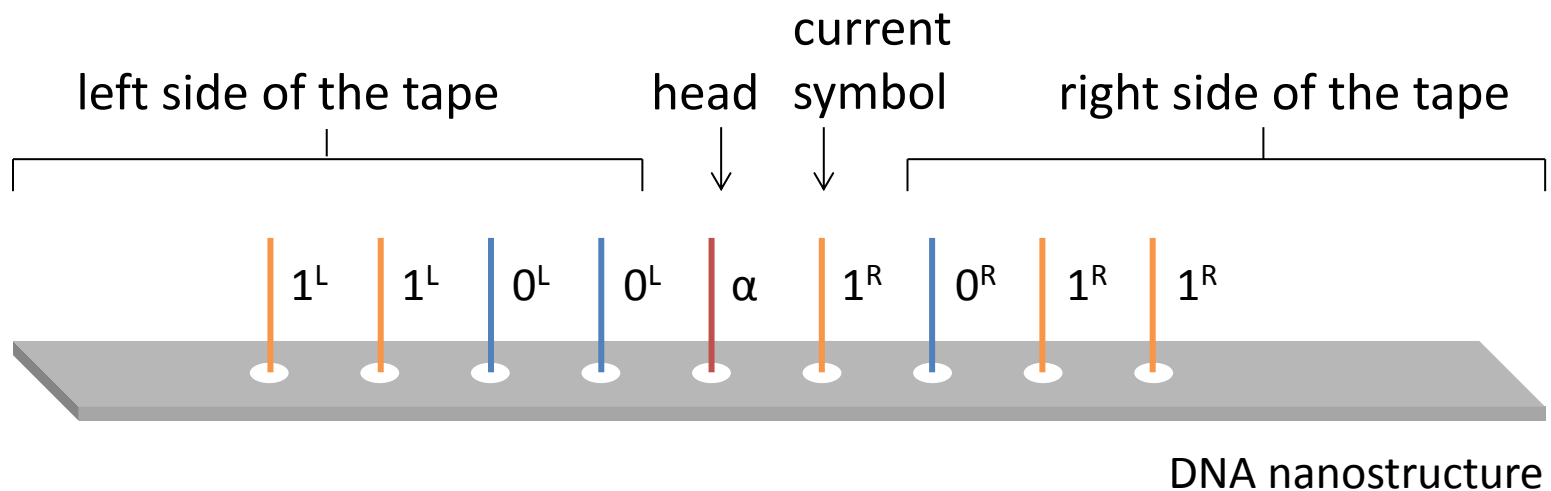
## Polymer CRNs



## Surface CRNs

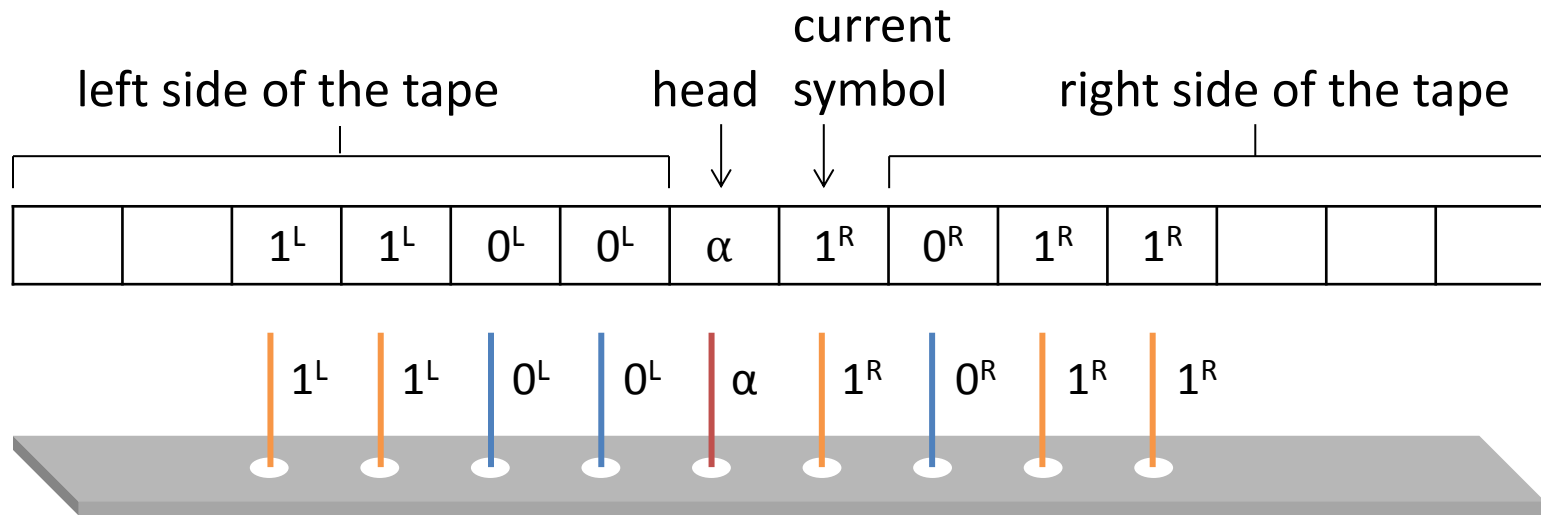


Can we use a DNA nanostructure to organize symbols and states represented as single-stranded DNA signals on a surface and serve as a tape for a molecular Turing machine?

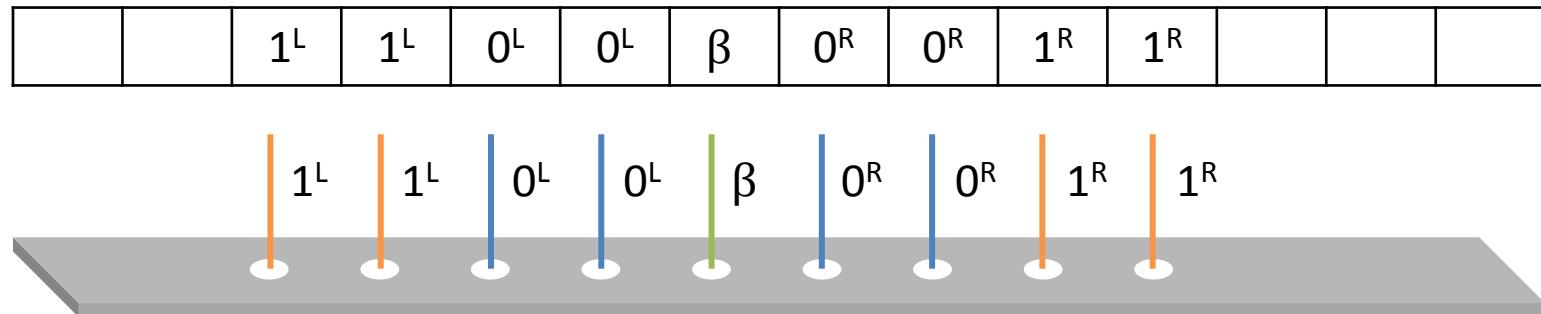




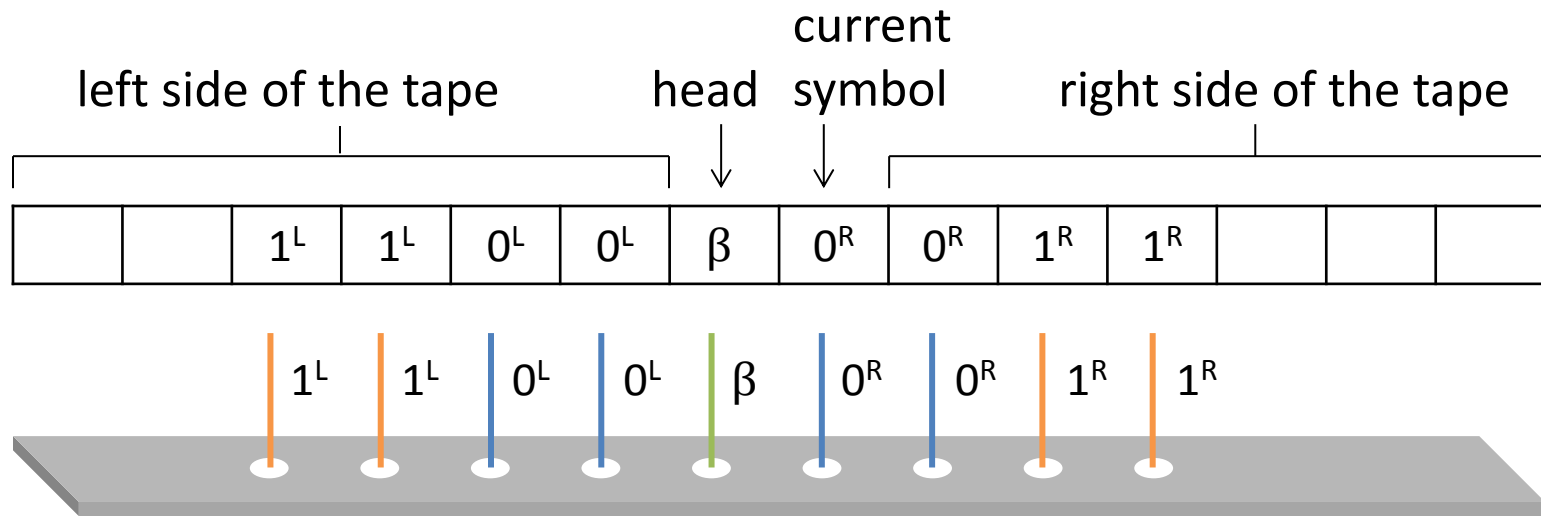
# Efficient molecular Turing machine



transition rule:  $\{\alpha, 1\} \rightarrow \{\beta, 0\}$   $\Downarrow$   $\alpha + 1^R \rightarrow \beta + 0^R$

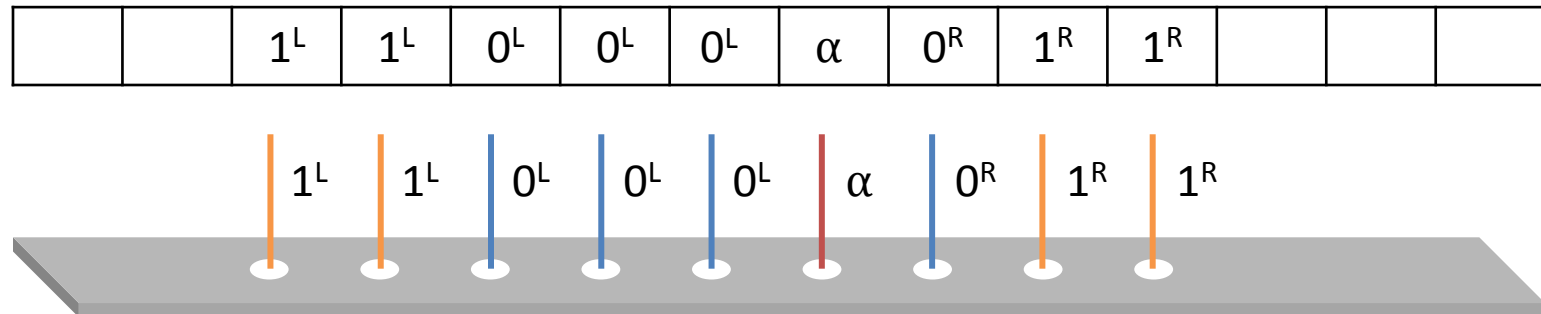


# Efficient molecular Turing machine

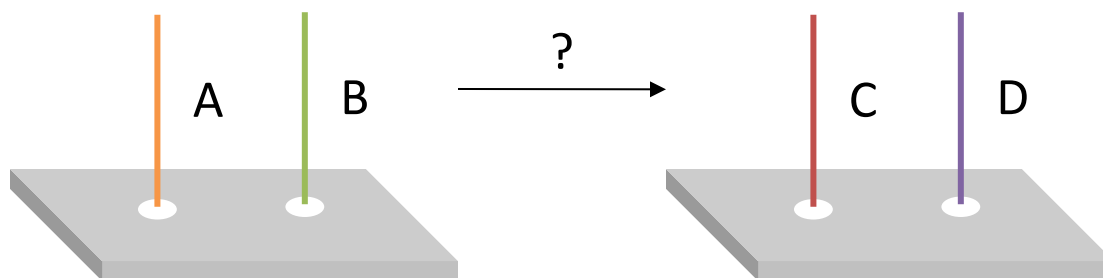
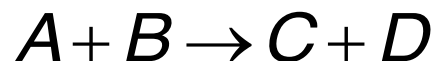


transition rule:  $\{\beta\} \rightarrow \{\alpha, +\}$

$$\begin{cases} \beta + 0^R \rightarrow 0^L + \alpha \\ \beta + 1^R \rightarrow 1^L + \alpha \end{cases}$$

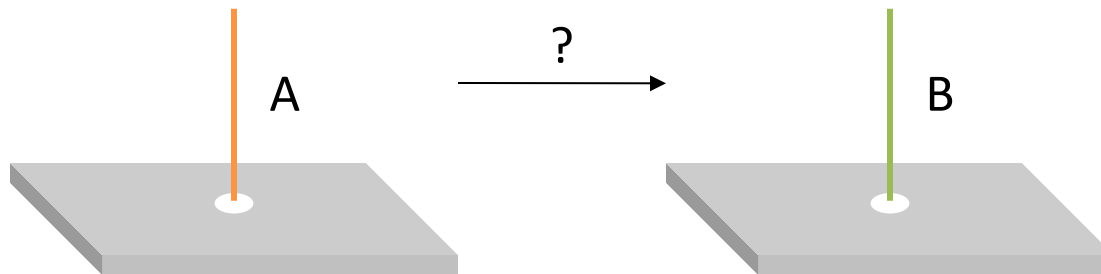
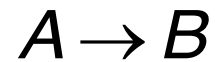


How do we cooperatively change two neighboring signals on a surface?



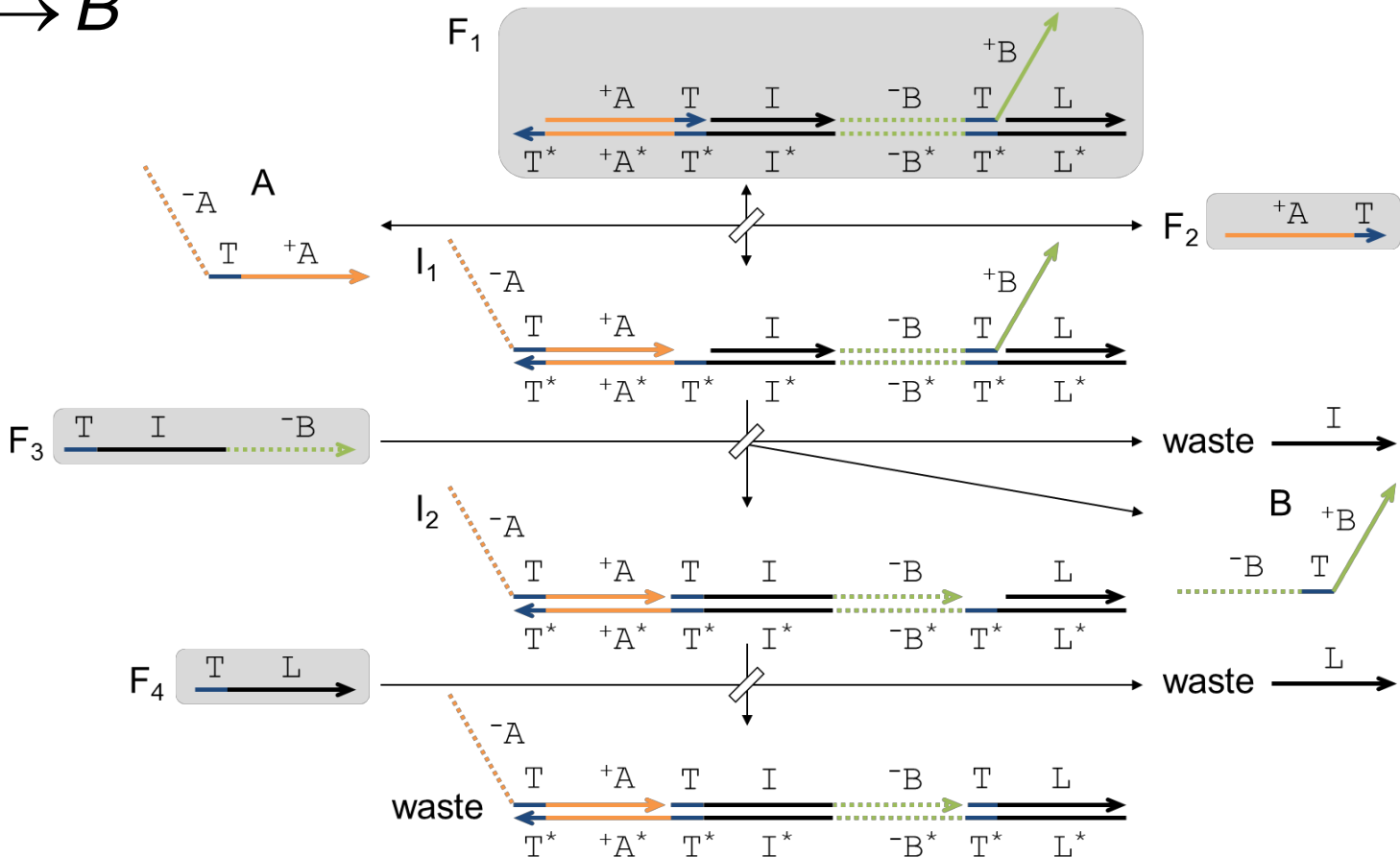
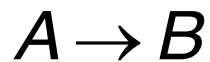
Surface-based formal bimolecular reaction

How do we change a signal on a surface from an original state to a new state?

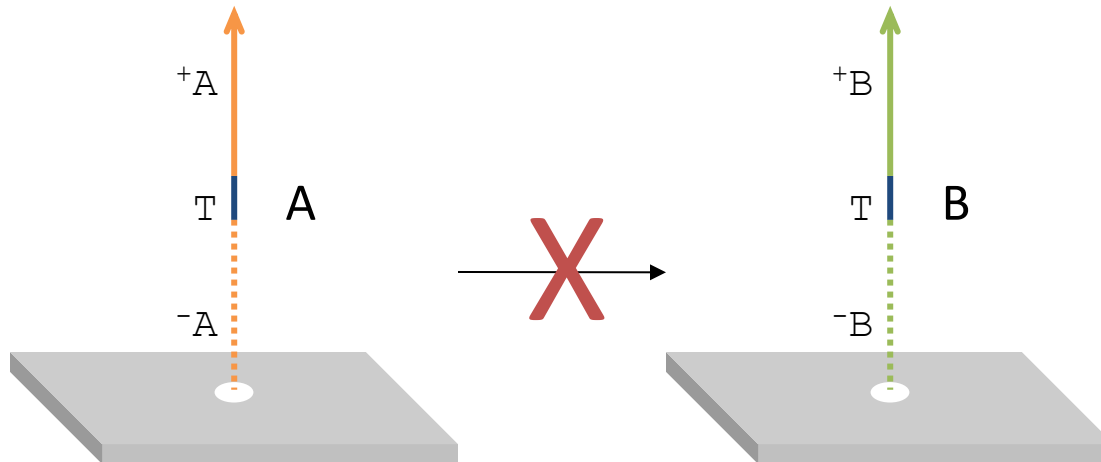
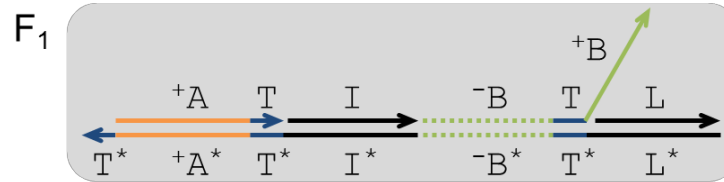
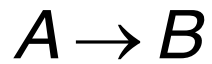


Surface-based formal unimolecular reaction

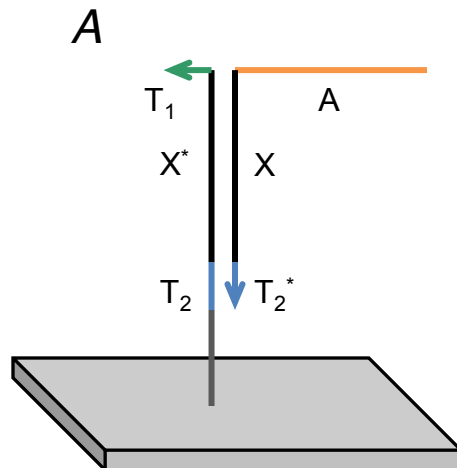
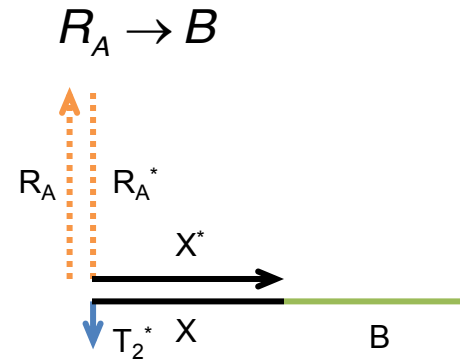
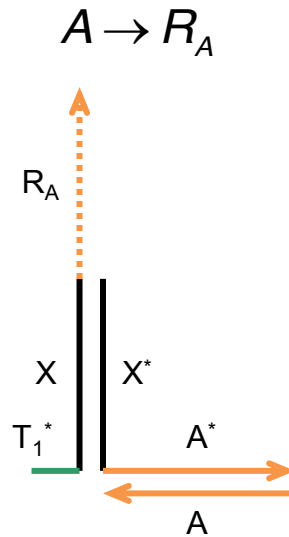
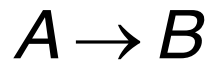
# Implementation of well-mixed formal unimolecular reaction



# Implementation of well-mixed formal unimolecular reaction

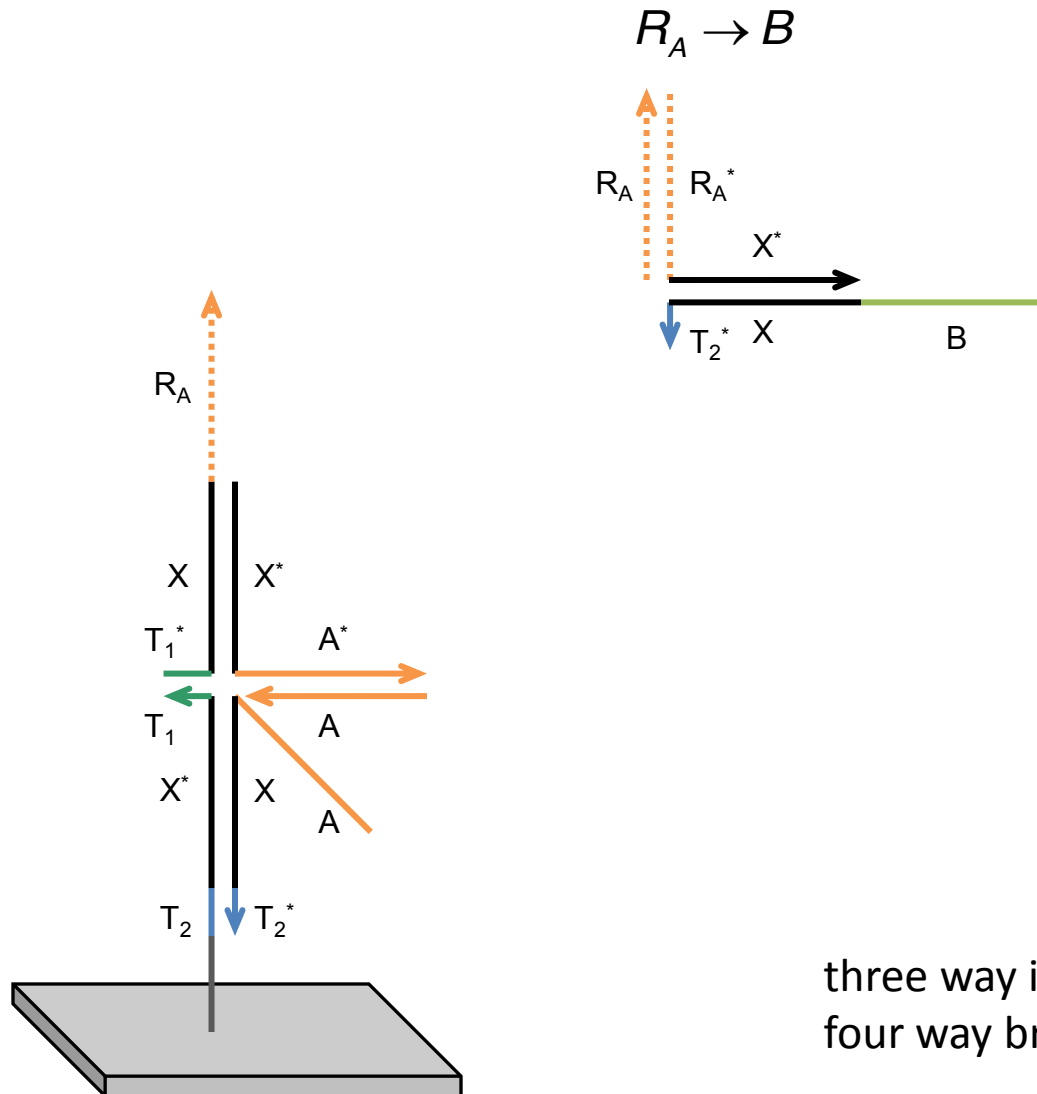
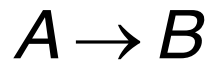


# Implementation of surface-based formal unimolecular reaction



three way initiated  
four way branch migration

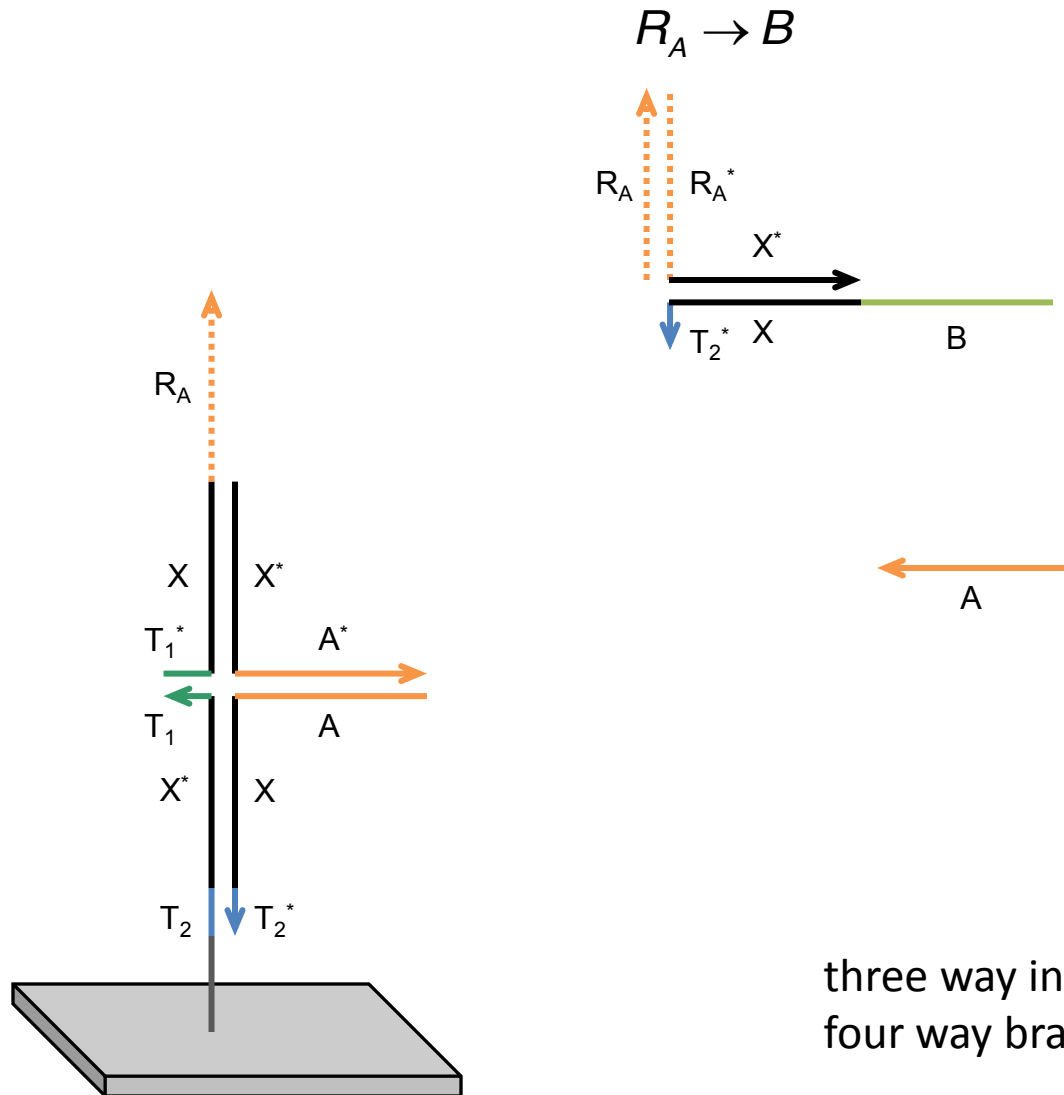
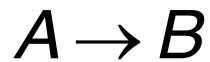
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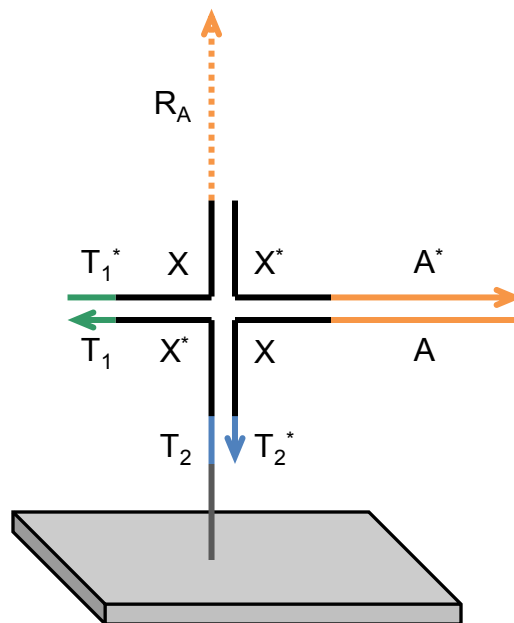
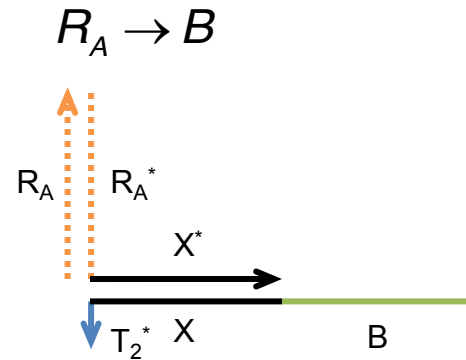
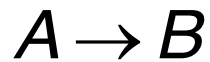


# Implementation of surface-based formal unimolecular reaction



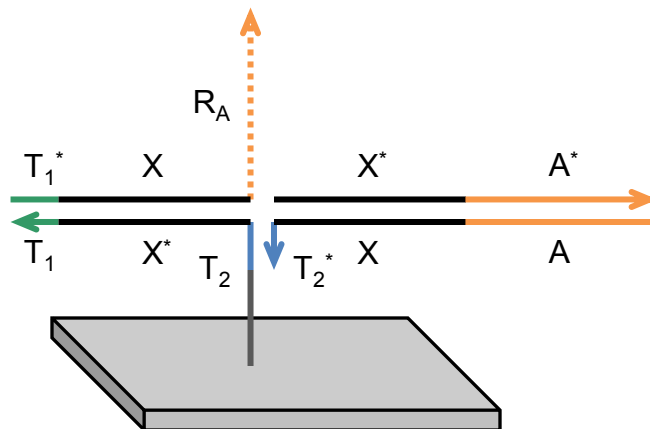
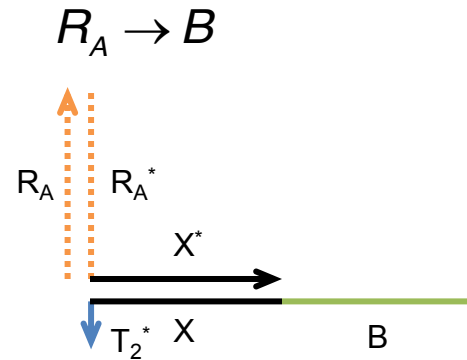
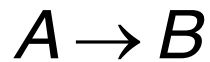
three way initiated  
four way branch migration

# Implementation of surface-based formal unimolecular reaction



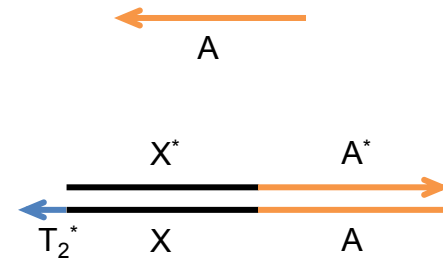
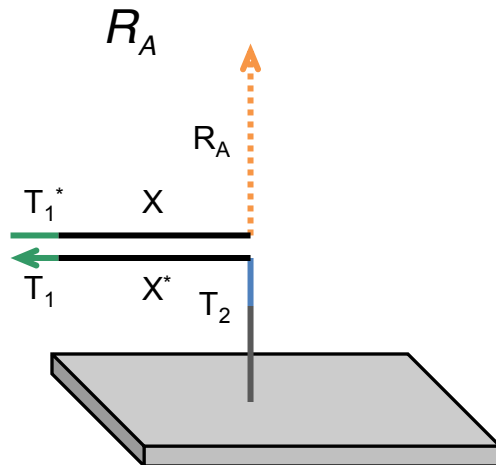
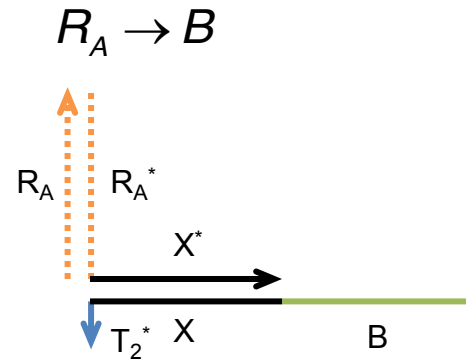
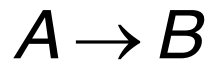
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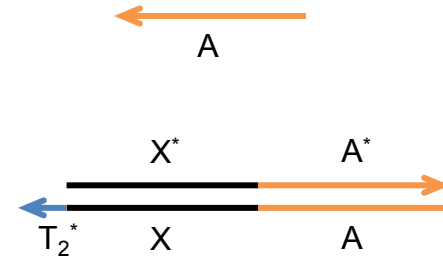
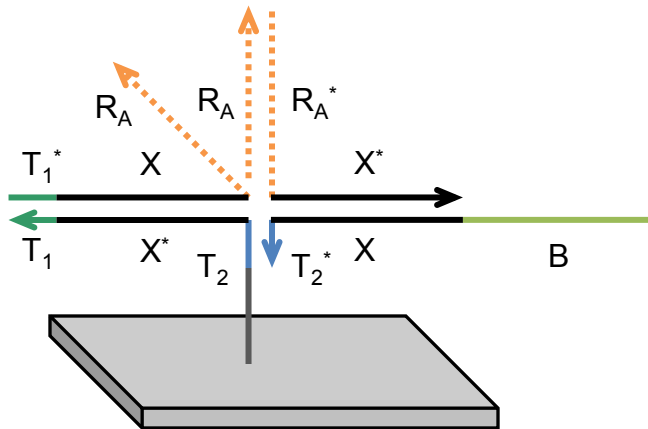
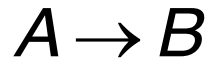
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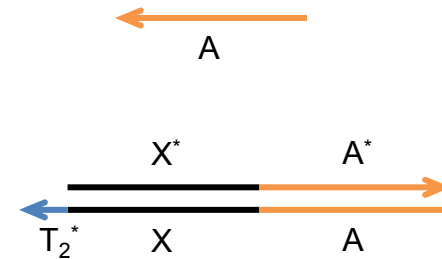
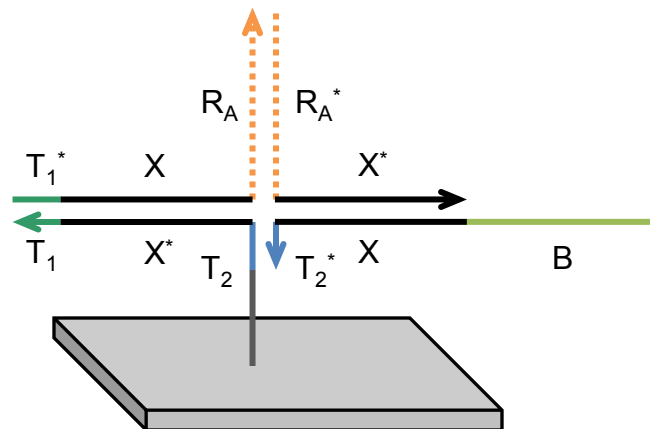
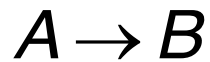
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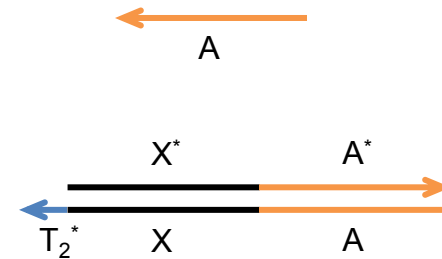
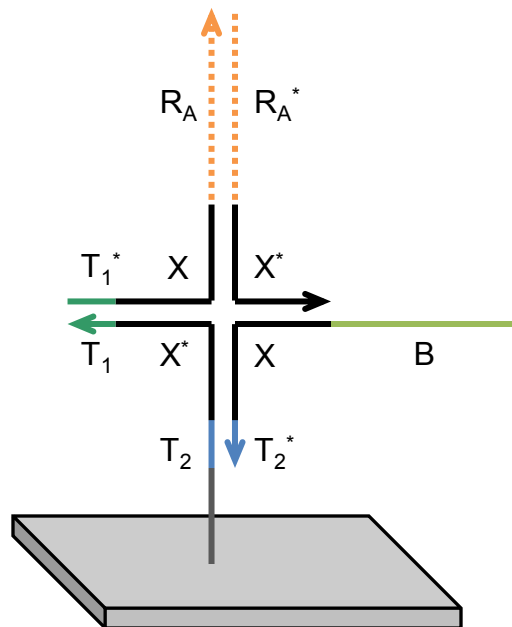
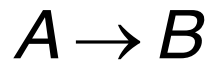
three way initiated  
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# Implementation of surface-based formal unimolecular reaction



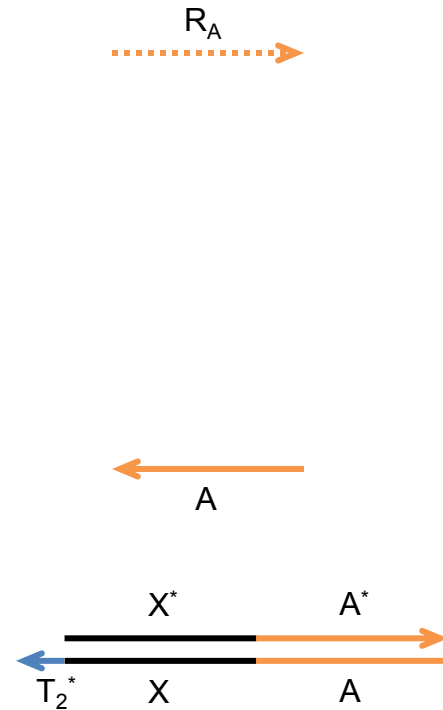
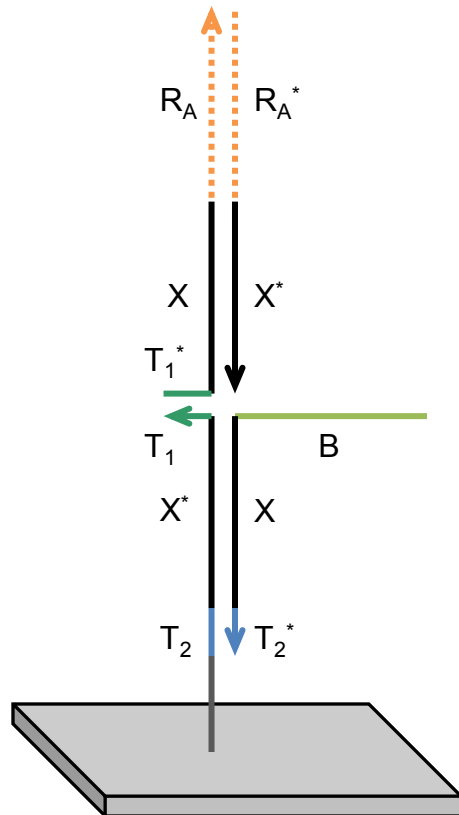
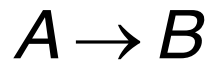
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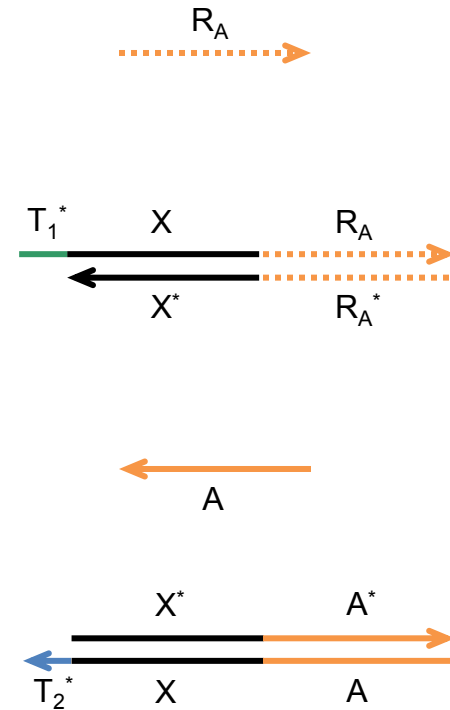
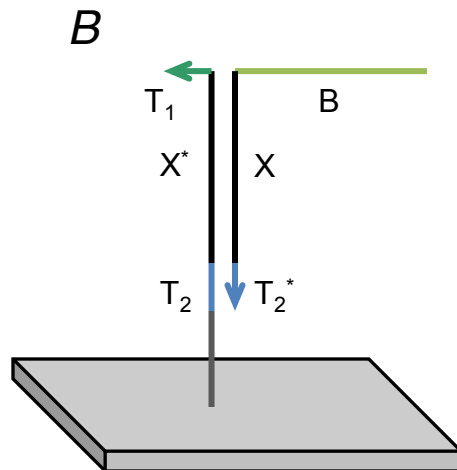
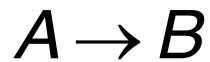
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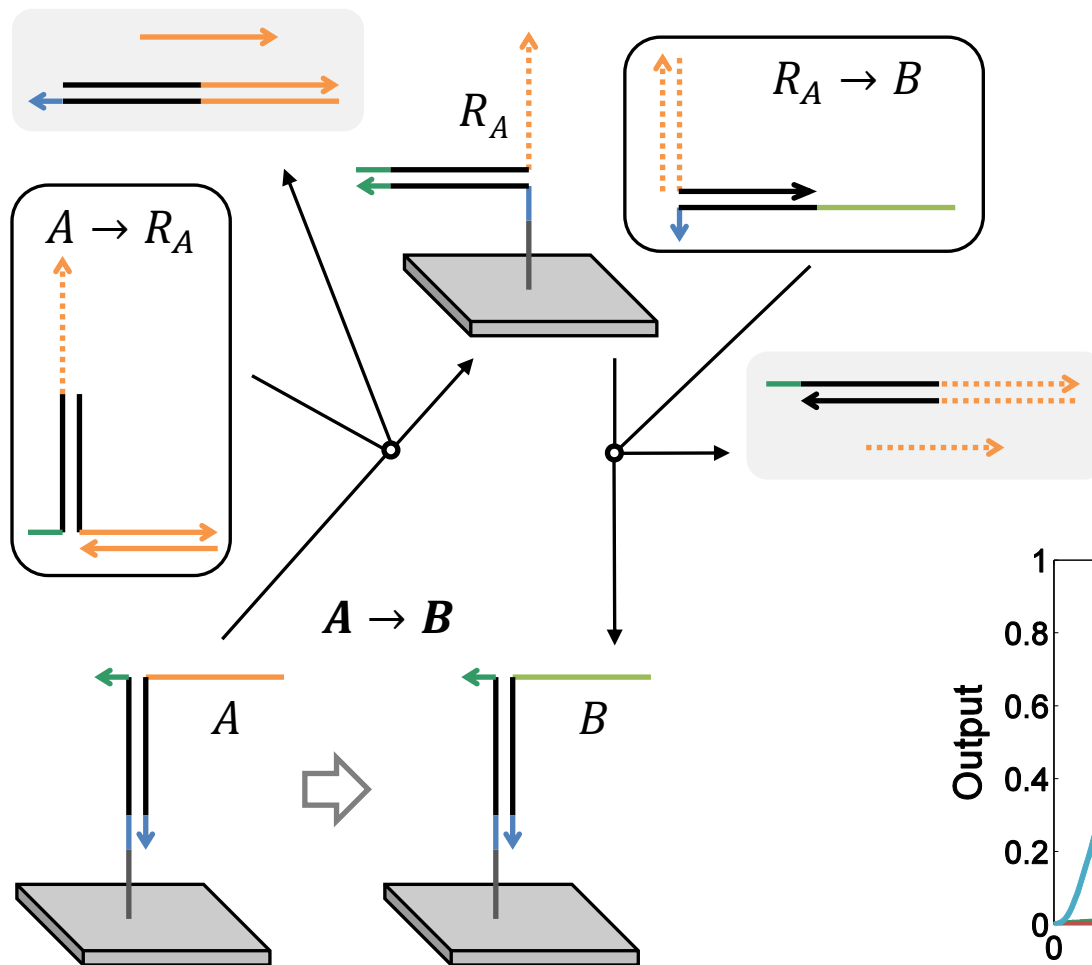


# Implementation of surface-based formal unimolecular reaction

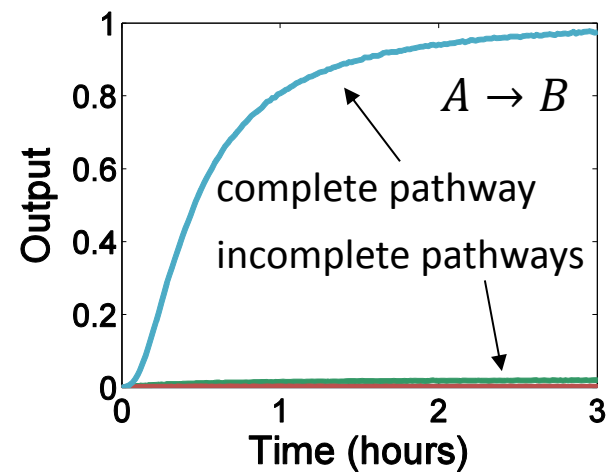


three way initiated  
four way branch migration

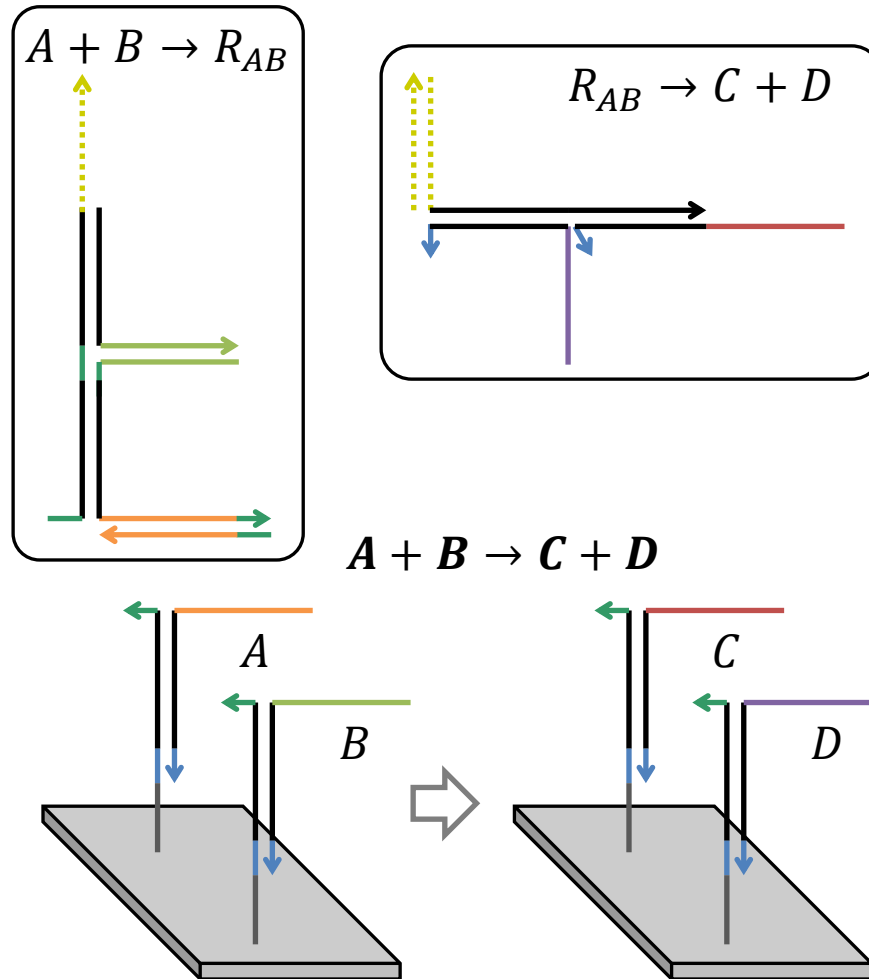
# Implementation of surface-based formal unimolecular reaction



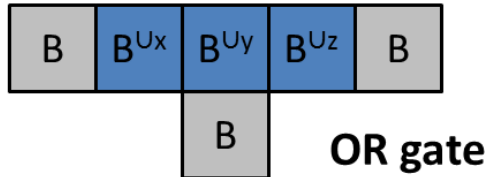
three way initiated  
four way branch migration



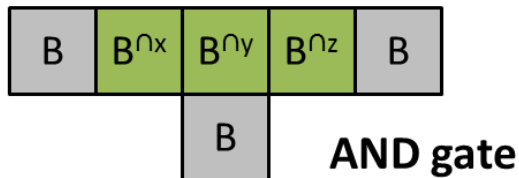
# Implementation of surface-based formal bimolecular reaction



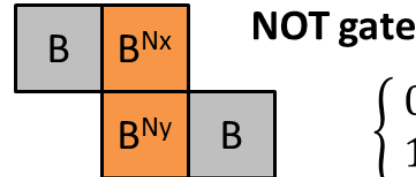
# Continuously active logic circuits with surface CRNs



$$\left\{ \begin{array}{l} 0^{Ux} + 0^{Uy} \rightarrow B^{Ux} + 0^{Uk} \\ 0^{Ux} + 1^{Uy} \rightarrow B^{Ux} + 1^{Uk} \\ 1^{Ux} + 0^{Uy} \rightarrow B^{Ux} + 1^{Uk} \\ 1^{Ux} + 1^{Uy} \rightarrow B^{Ux} + 1^{Uk} \\ 0^{Uk} + B^{Uz} \rightarrow B^{Uy} + 0^{Uz} \\ 1^{Uk} + B^{Uz} \rightarrow B^{Uy} + 1^{Uz} \end{array} \right.$$

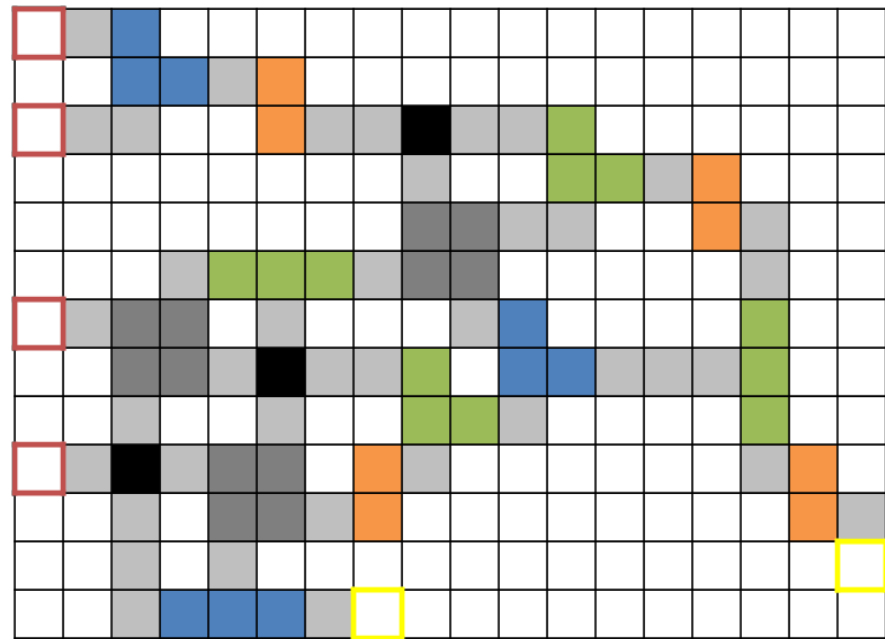


$$\left\{ \begin{array}{l} 0^{Nx} + 0^{Ny} \rightarrow B^{Nx} + 0^{Nk} \\ 0^{Nx} + 1^{Ny} \rightarrow B^{Nx} + 0^{Nk} \\ 1^{Nx} + 0^{Ny} \rightarrow B^{Nx} + 0^{Nk} \\ 1^{Nx} + 1^{Ny} \rightarrow B^{Nx} + 1^{Nk} \\ 0^{Nk} + B^{Nz} \rightarrow B^{Ny} + 0^{Nz} \\ 1^{Nk} + B^{Nz} \rightarrow B^{Ny} + 1^{Nz} \end{array} \right.$$



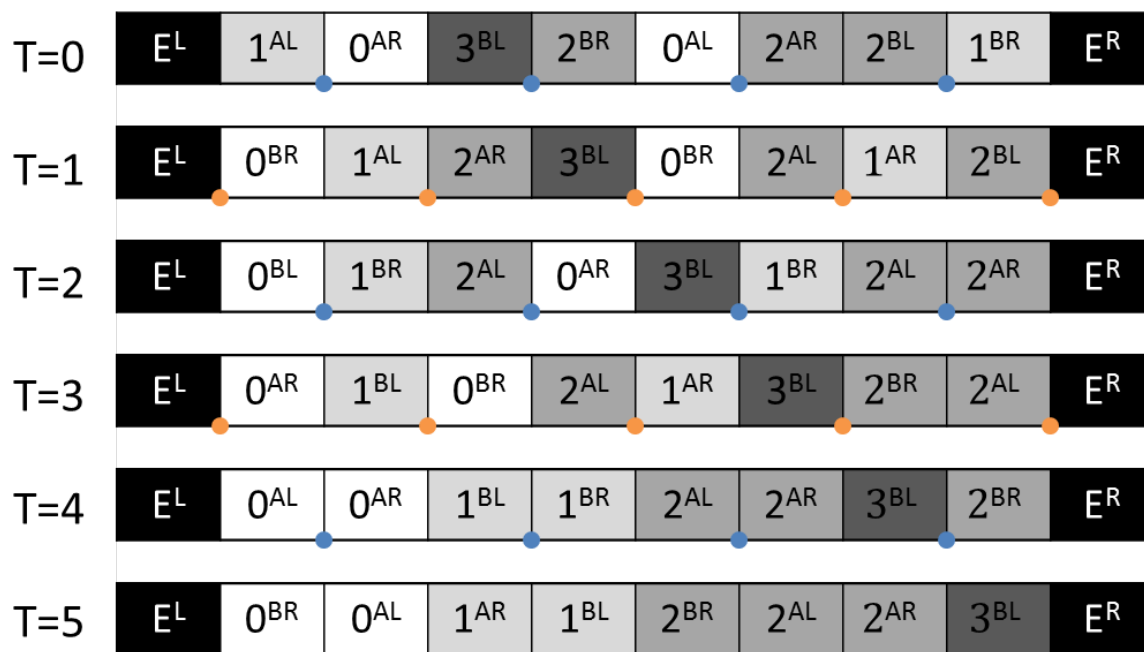
$$\left\{ \begin{array}{l} 0^{Nx} + B^{Ny} \rightarrow B^{Nx} + 1^{Ny} \\ 1^{Nx} + B^{Ny} \rightarrow B^{Nx} + 0^{Ny} \end{array} \right.$$

A square-root circuit on DNA origami:



blue sites: OR, green sites: AND, orange sites: NOT  
grey sites: wires, outlined sites: inputs or outputs

# Dynamically-updating cellular automata with Surface CRNs



each transition rule

$$\{x, y\} \rightarrow \{x^*, y^*\}$$

is implemented with:

$$\begin{cases} x^{AL} + y^{AR} \rightarrow x^{*BR} + y^{*AL} \\ x^{BL} + y^{BR} \rightarrow x^{*AR} + y^{*BL} \end{cases}$$

edge conditions for

each state  $x$  are

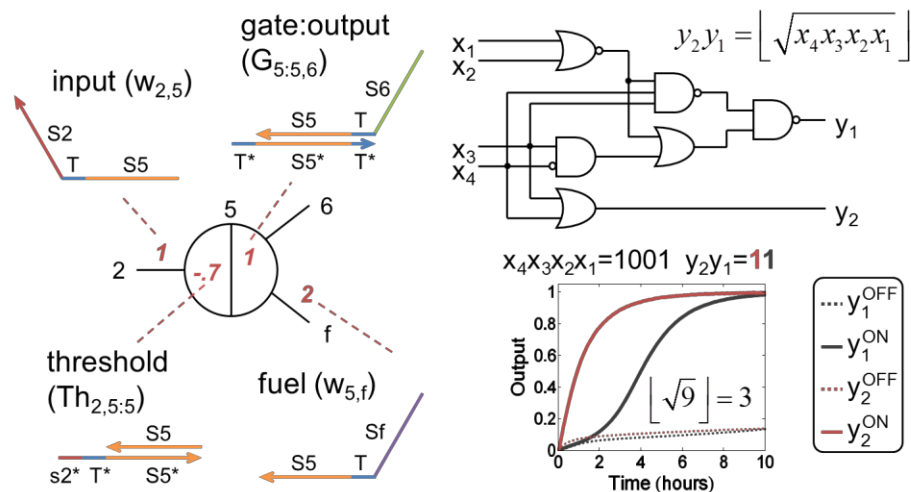
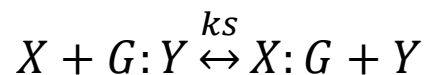
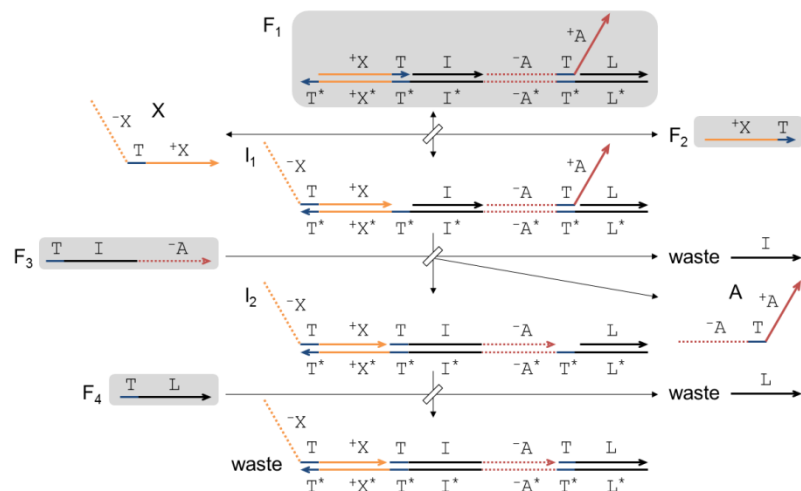
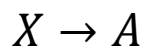
implemented with:

$$\begin{cases} E^L + x^{AR} \rightarrow E^L + x^{AL} \\ E^L + x^{BR} \rightarrow E^L + x^{BL} \\ x^{AL} + E^R \rightarrow x^{BR} + E^R \\ x^{BL} + E^R \rightarrow x^{AR} + E^R \end{cases}$$

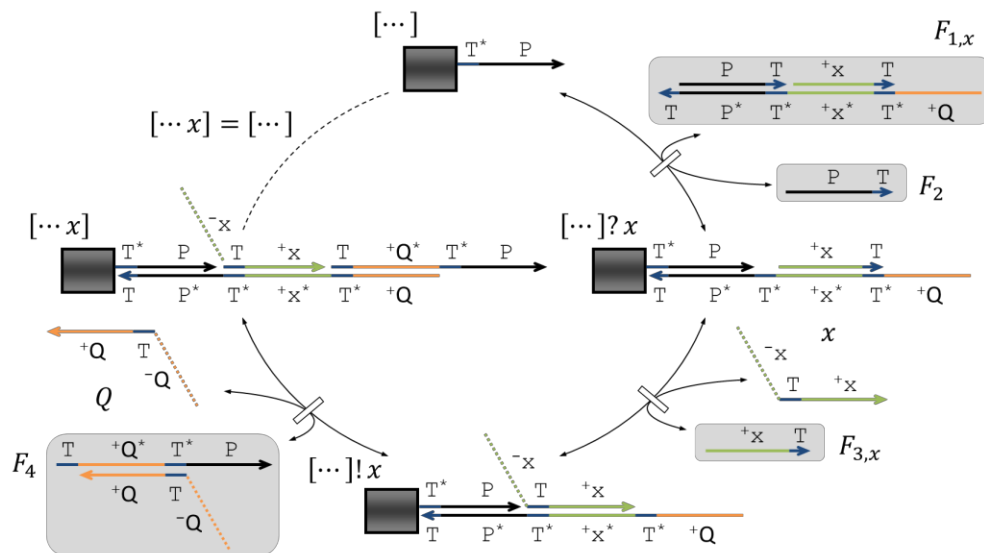
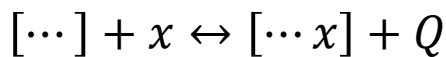
transition rules for sorting numbers between 0 to 3:

$$\begin{array}{llll} \{0, 0\} \rightarrow \{0, 0\} & \{0, 1\} \rightarrow \{0, 1\} & \{0, 2\} \rightarrow \{0, 2\} & \{0, 3\} \rightarrow \{0, 3\} \\ \{1, 0\} \rightarrow \{0, 1\} & \{1, 1\} \rightarrow \{1, 1\} & \{1, 2\} \rightarrow \{1, 2\} & \{1, 3\} \rightarrow \{1, 3\} \\ \{2, 0\} \rightarrow \{0, 2\} & \{2, 1\} \rightarrow \{1, 2\} & \{2, 2\} \rightarrow \{2, 2\} & \{2, 3\} \rightarrow \{2, 3\} \\ \{3, 0\} \rightarrow \{0, 3\} & \{3, 1\} \rightarrow \{1, 3\} & \{3, 2\} \rightarrow \{2, 3\} & \{3, 3\} \rightarrow \{3, 3\} \end{array}$$

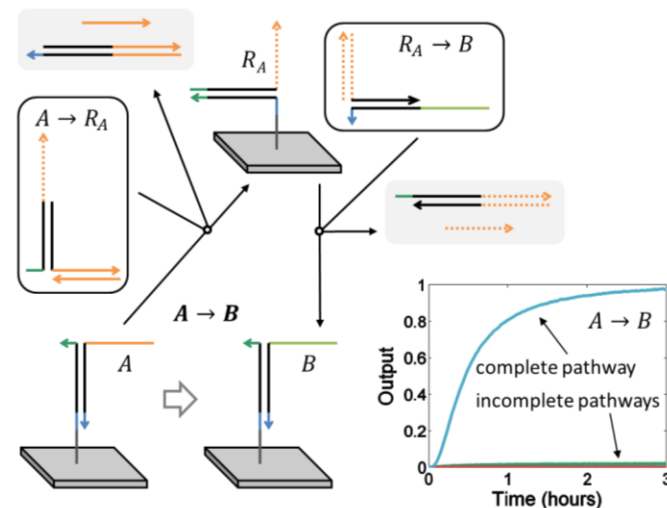
## Well-mixed CRNs



## Polymer CRNs



## Surface CRNs

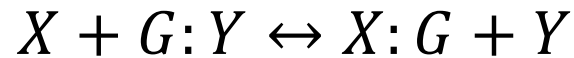


# Forward thinking and backward thinking

Given a computational task, how can we implement it with CRNs?

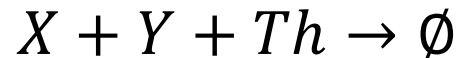
Given a few types of chemical reactions with experimentally successful implementations, what kind of interesting computational tasks can be performed?

1. What kind of interesting computational tasks can be performed with the following types of well-mixed reactions?



Feed-forward logic circuits: Qian et al, *Science* 2011

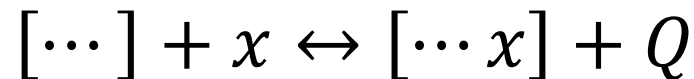
Neural networks (linear threshold circuits): Qian et al, *Nature* 2011



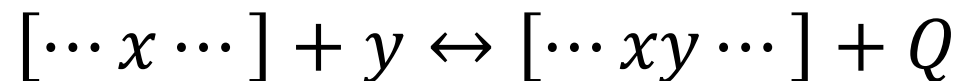
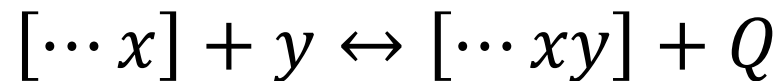
Linear I/O systems: Oishi et al, *IET Syst. Biol.* 2011



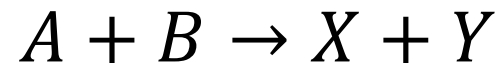
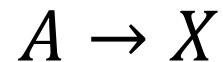
2. What kind of interesting computational tasks can be performed with the following types of polymer reactions?



Stack machines: Qian et al, *LNCS* 2011



3. What kind of interesting computational tasks can be performed with the following two types of surface reactions?



Sequential logic, Turing machines, cellular automata : Qian et al, *LNCS* 2014

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Career Award at Scientific Interface



Caltech

Biology and Biological Engineering

