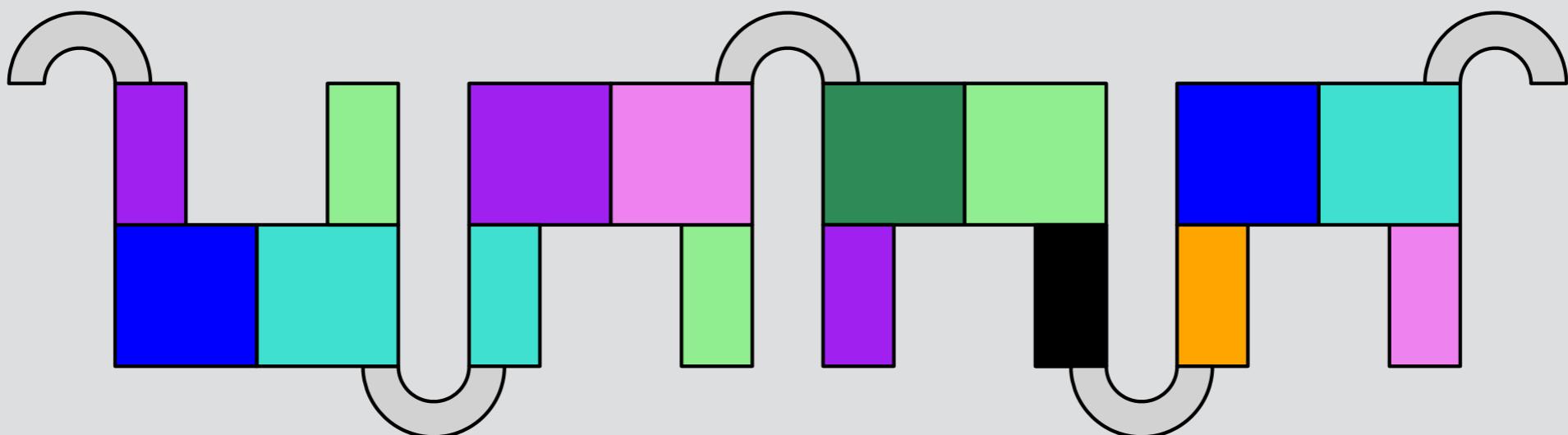


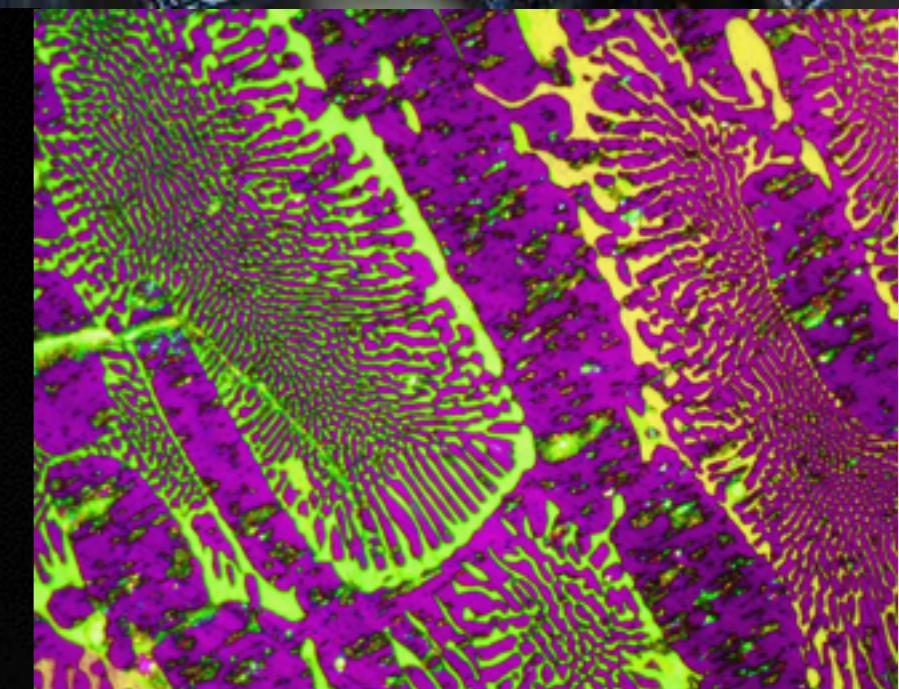
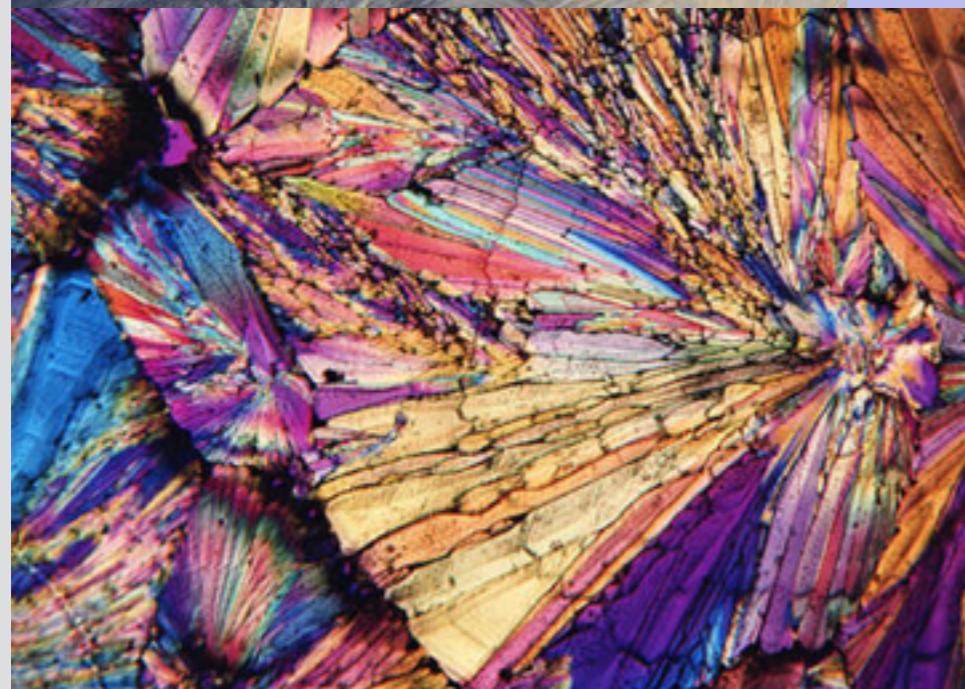
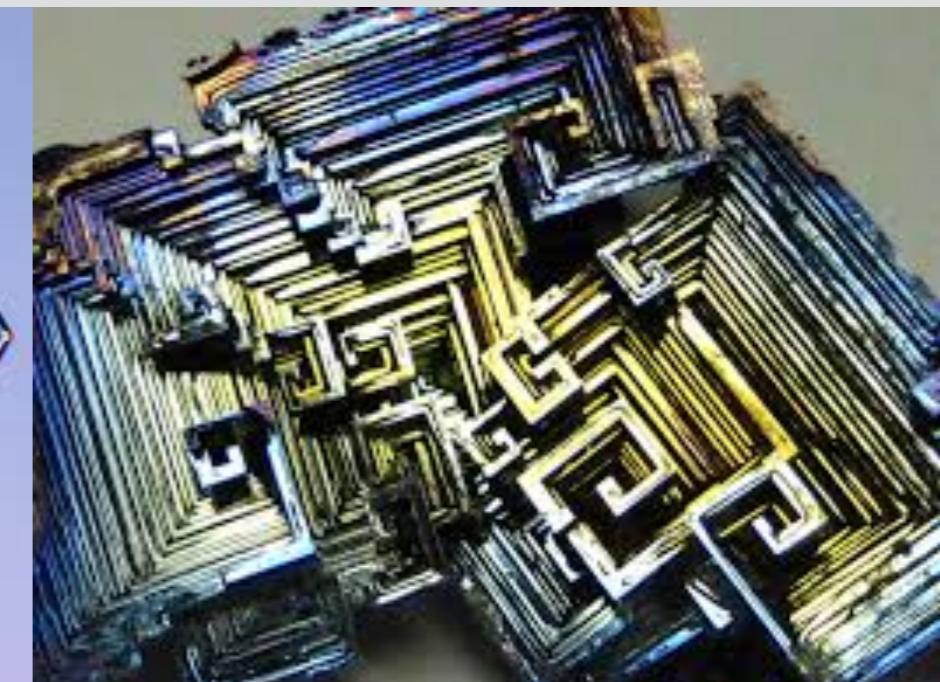
# The Limits of a Simple Model of Active Self-Assembly



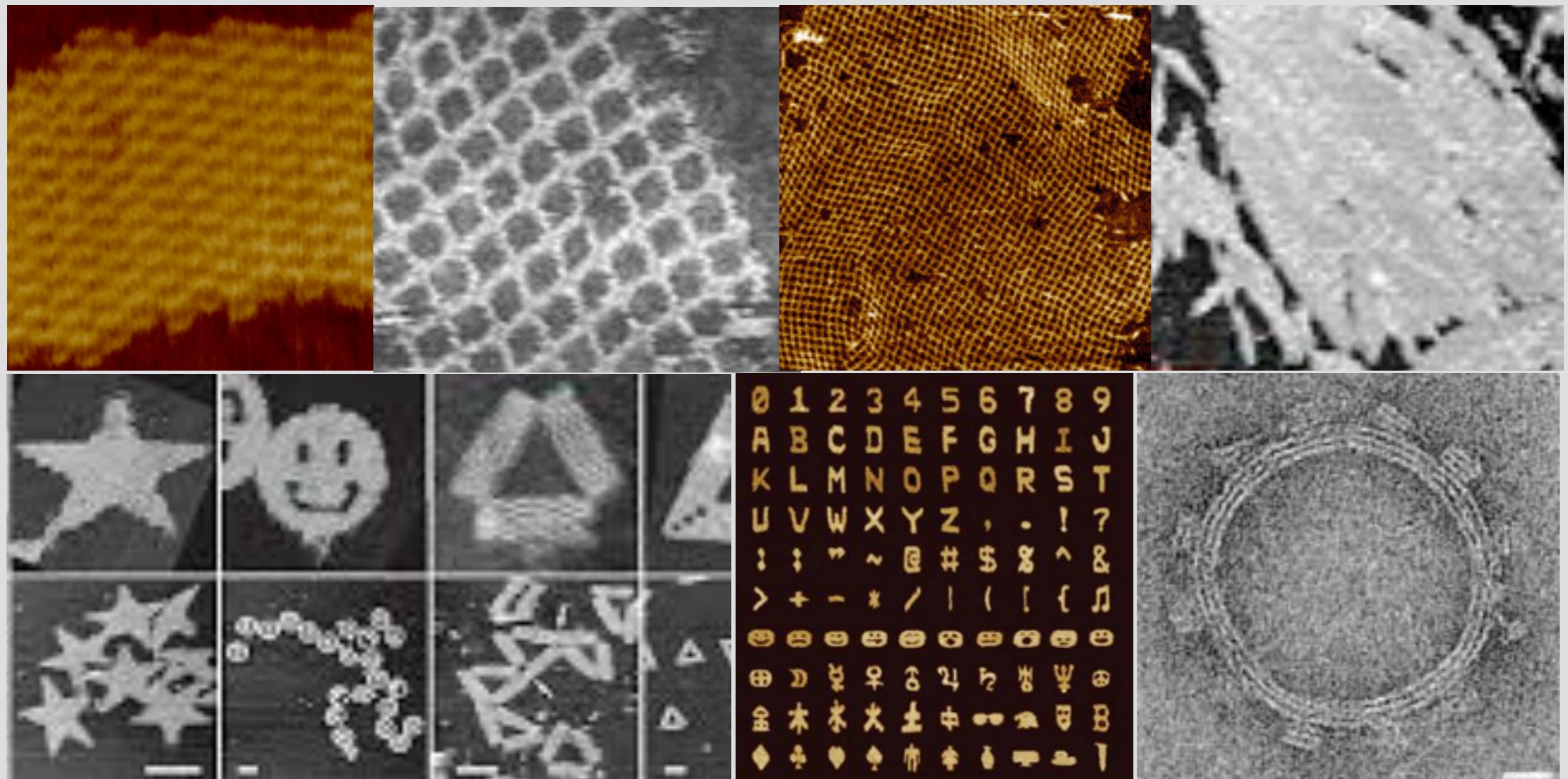
Andrew Winslow



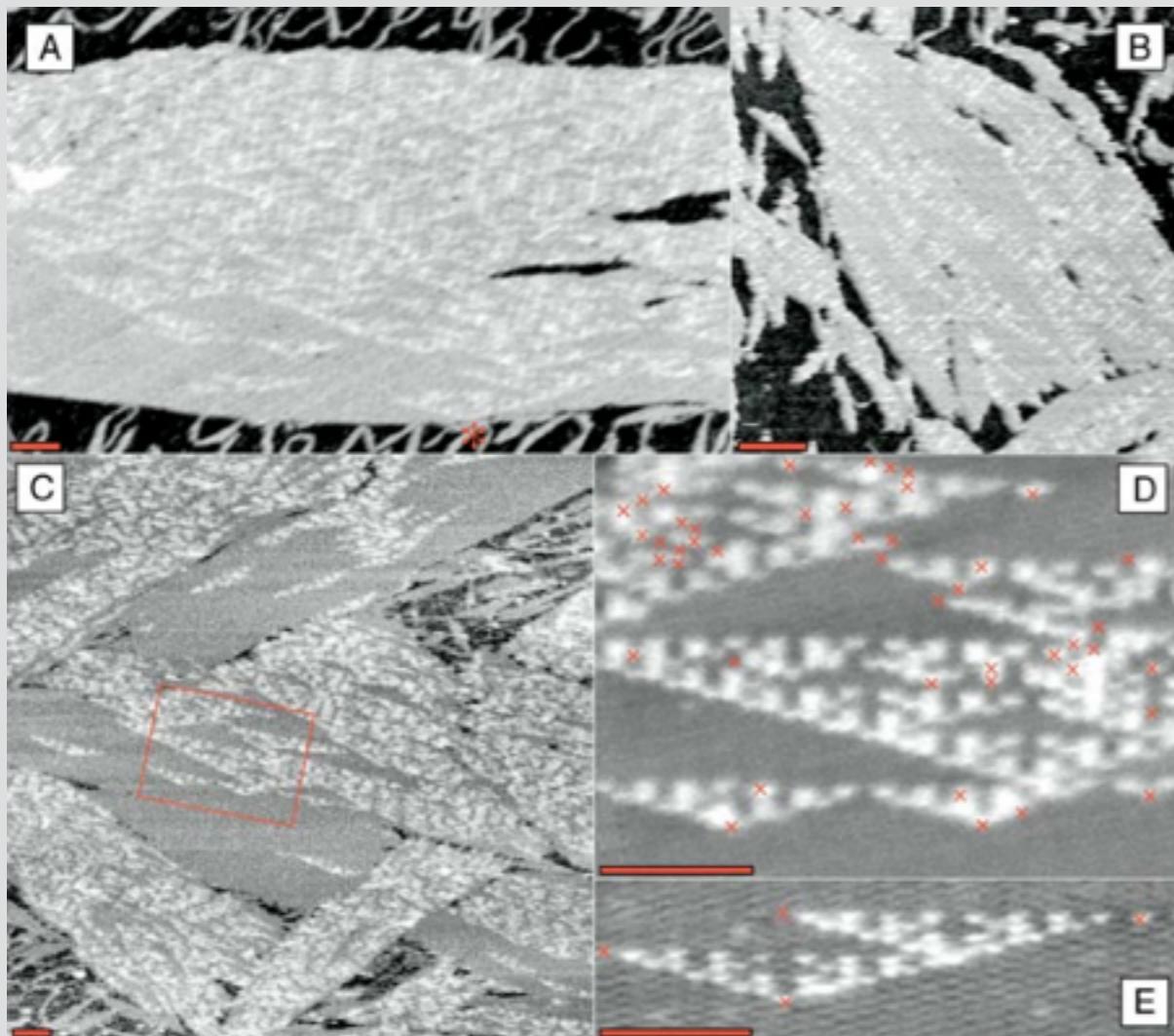
# Natural Nanoscale Self-Assembly



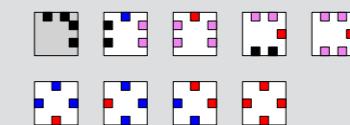
# Synthetic Nanoscale Self-Assembly



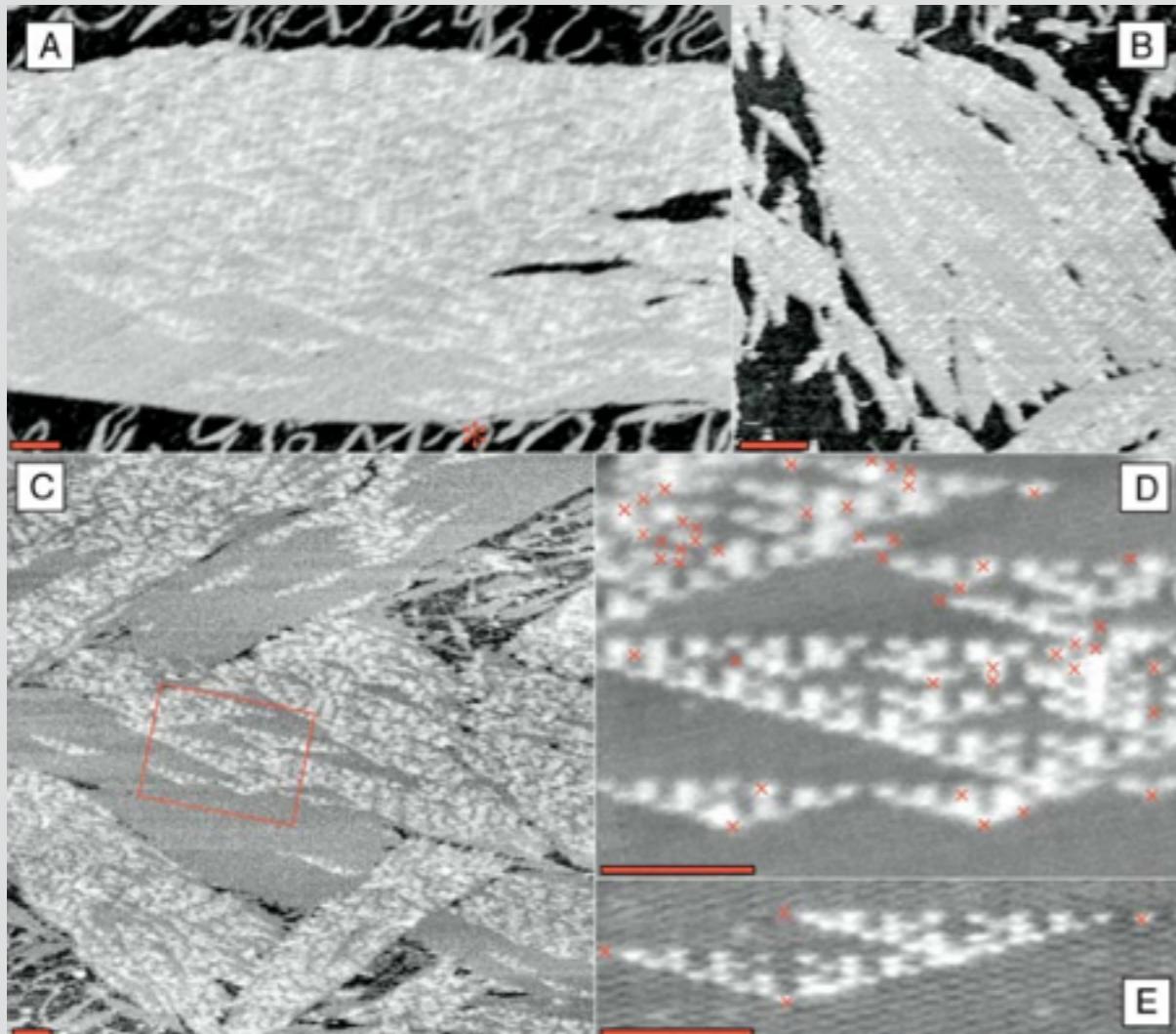
## Implementation (in DNA)



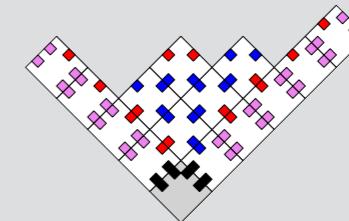
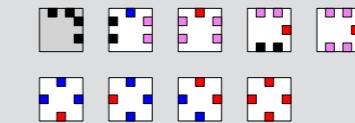
## Theoretical model (aTAM)



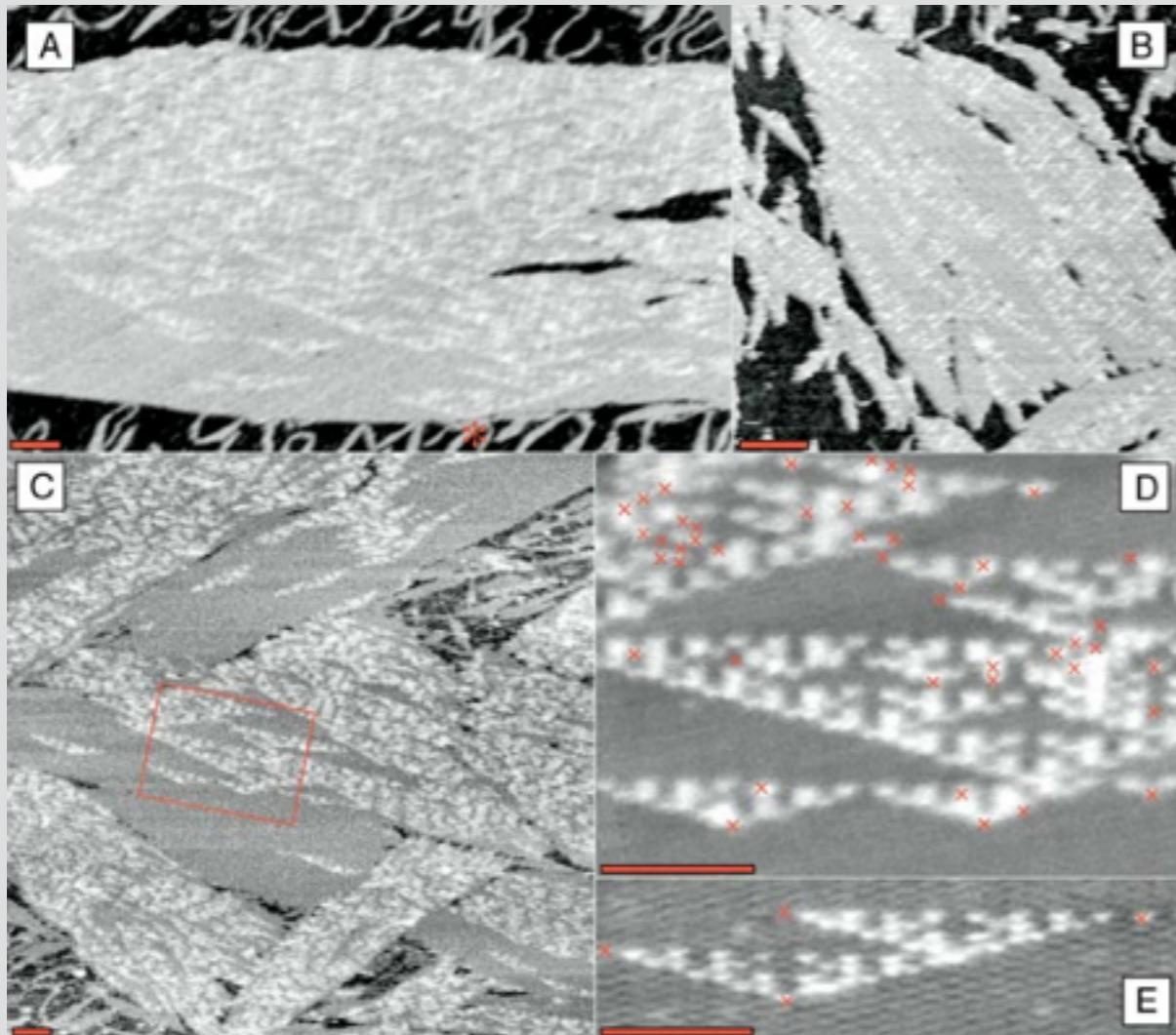
## Implementation (in DNA)



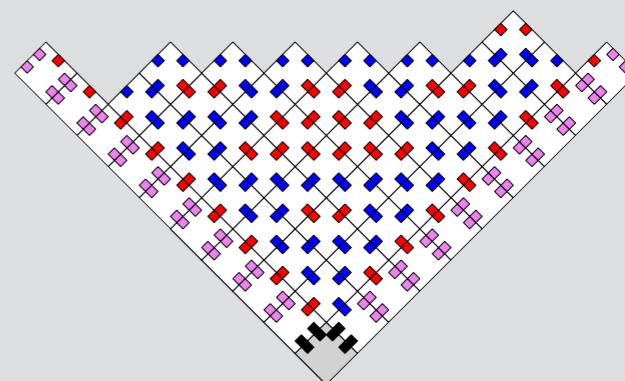
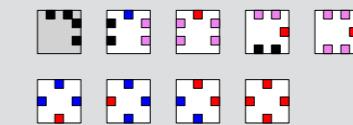
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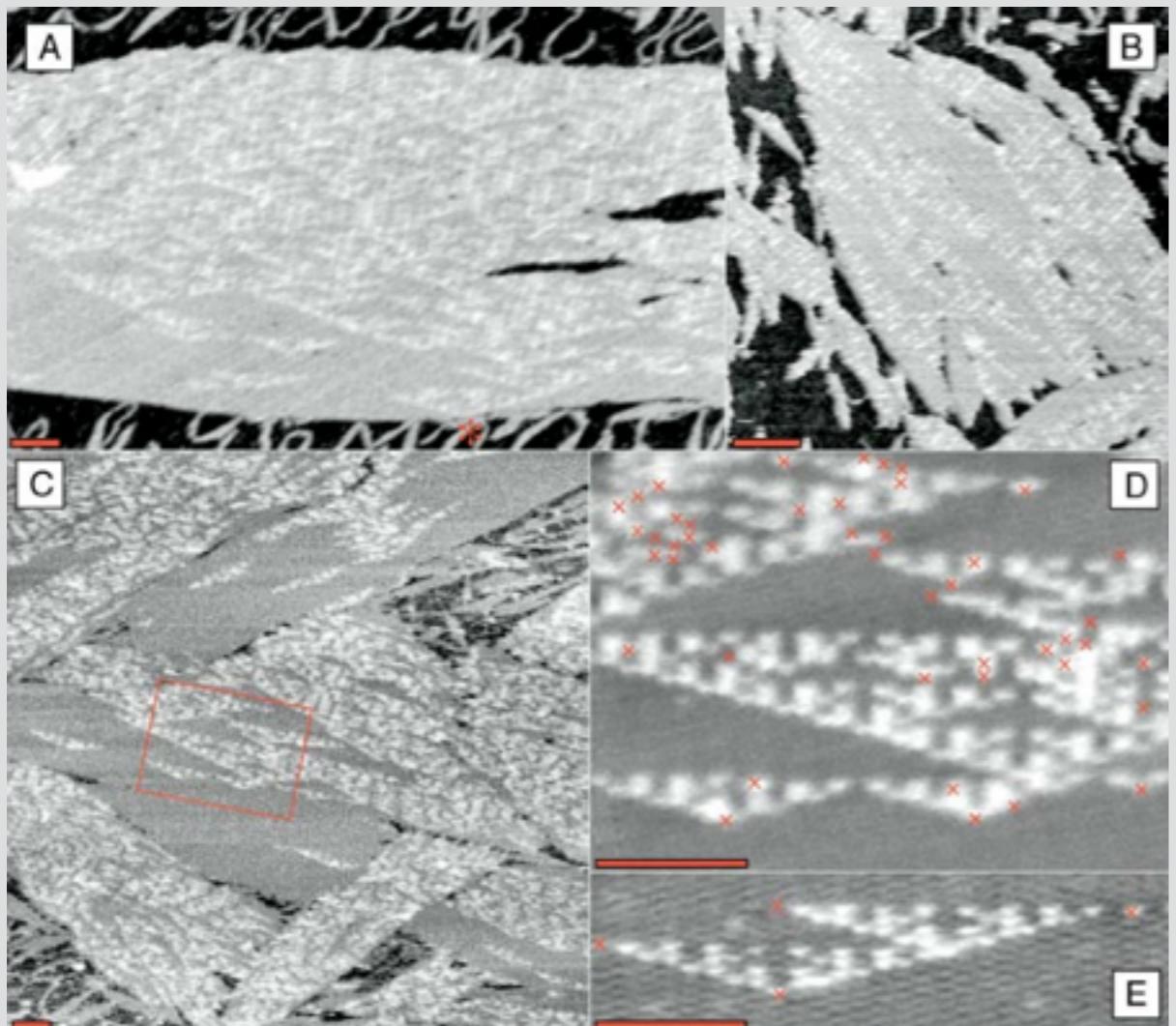
## Implementation (in DNA)



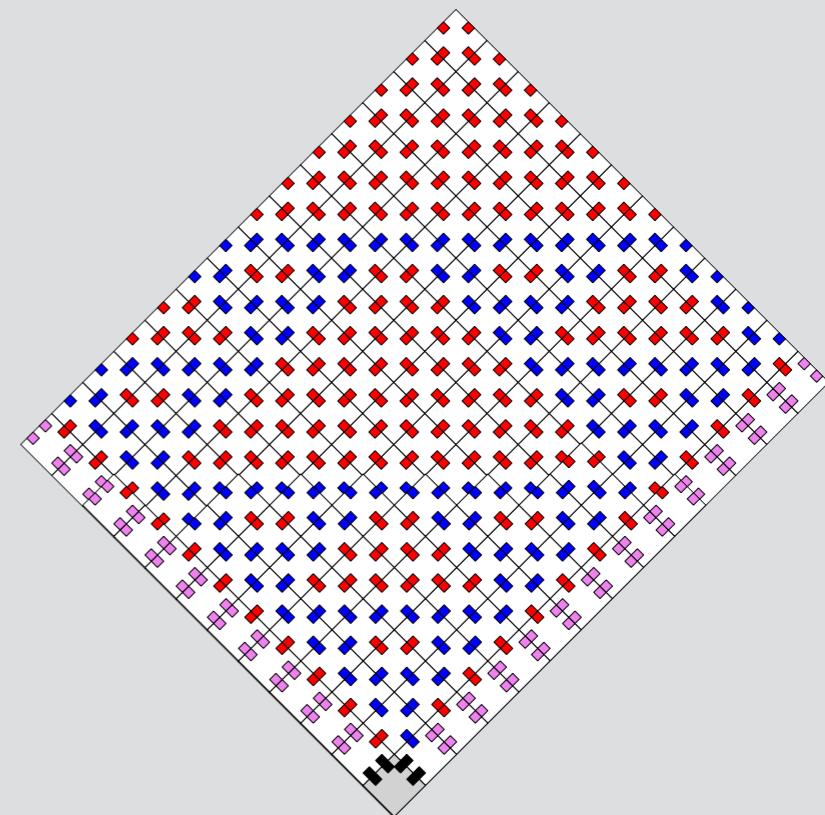
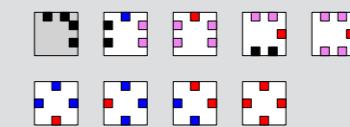
## Theoretical model (aTAM)



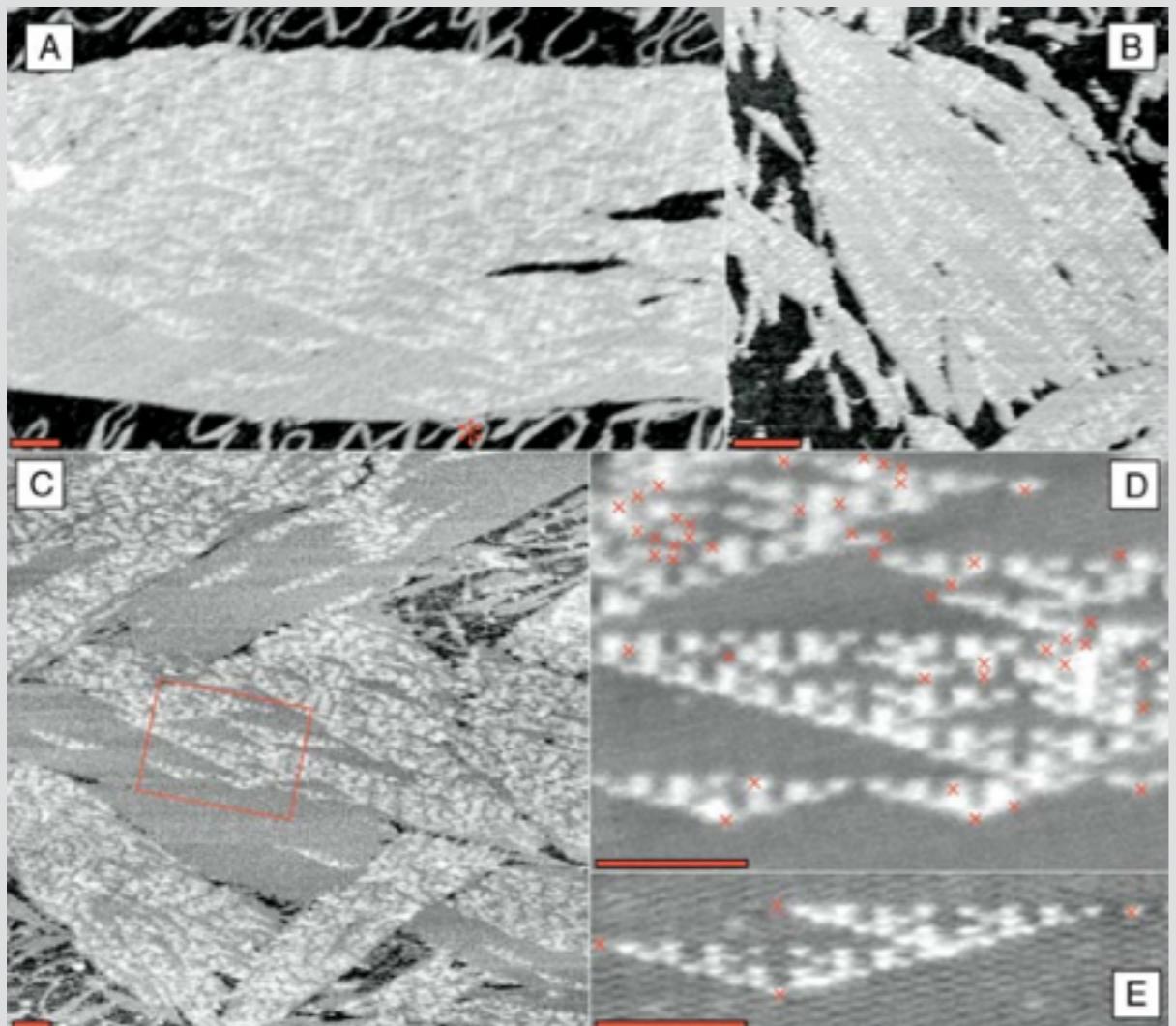
## Implementation (in DNA)



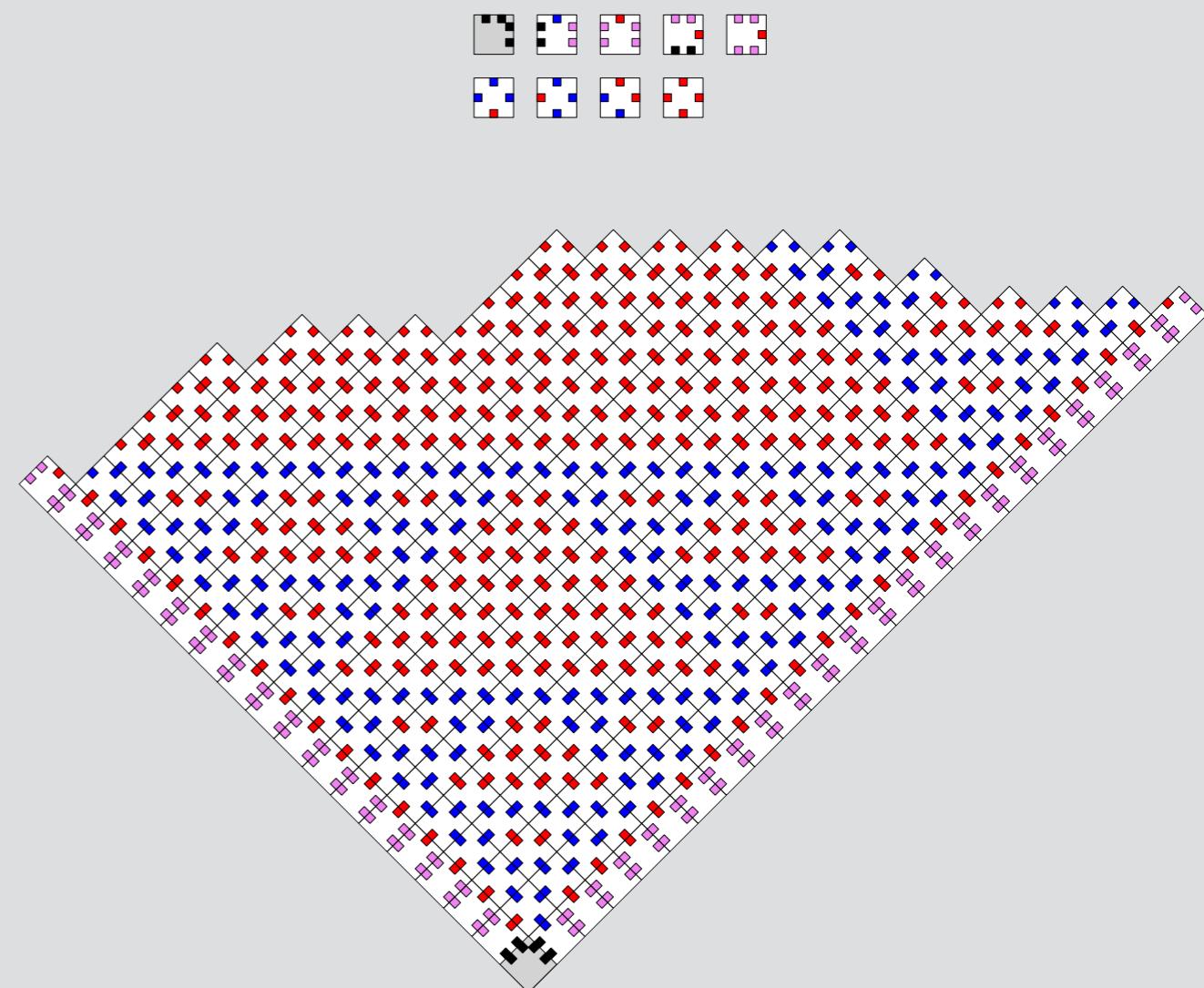
## Theoretical model (aTAM)



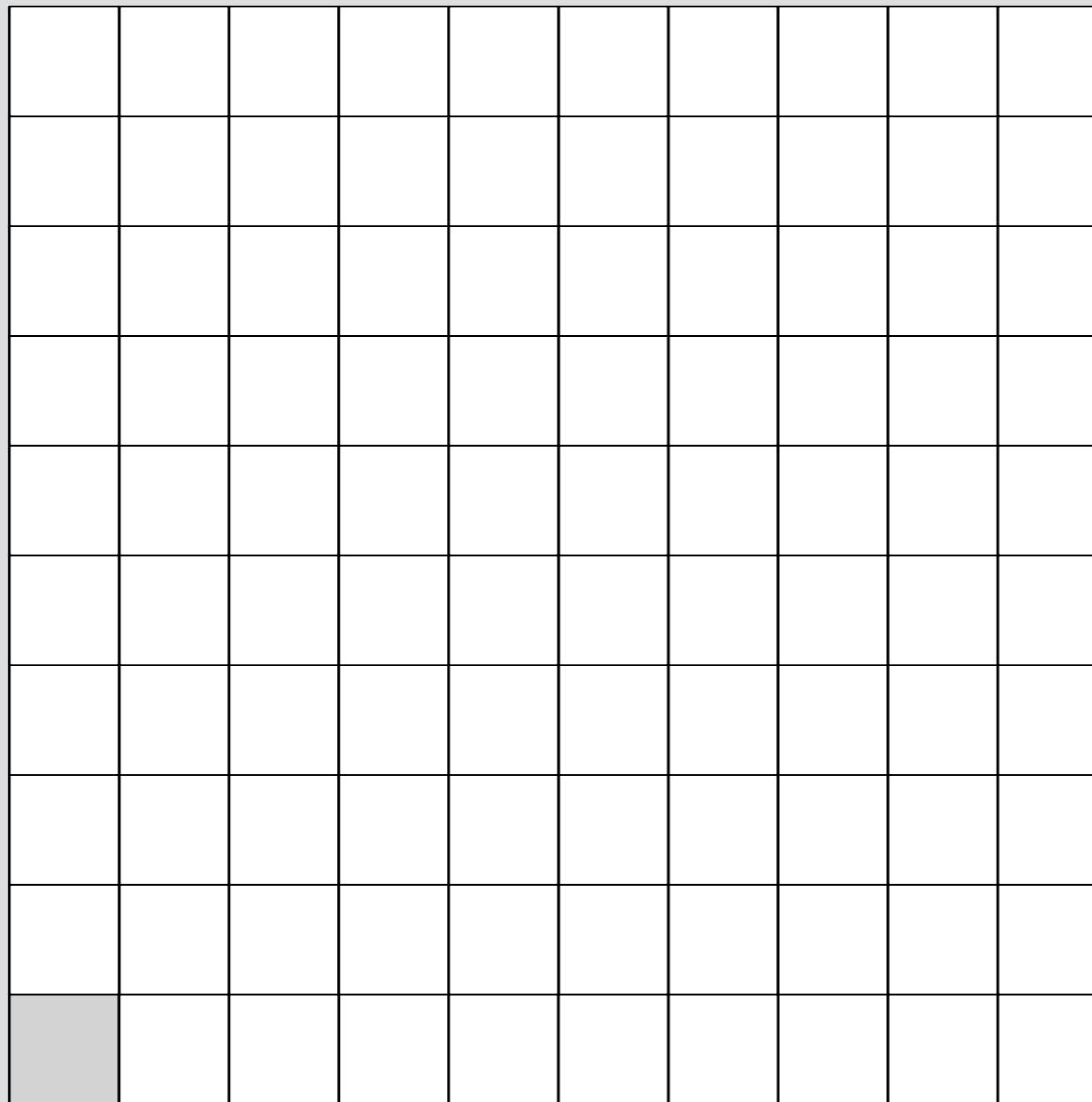
## Implementation (in DNA)



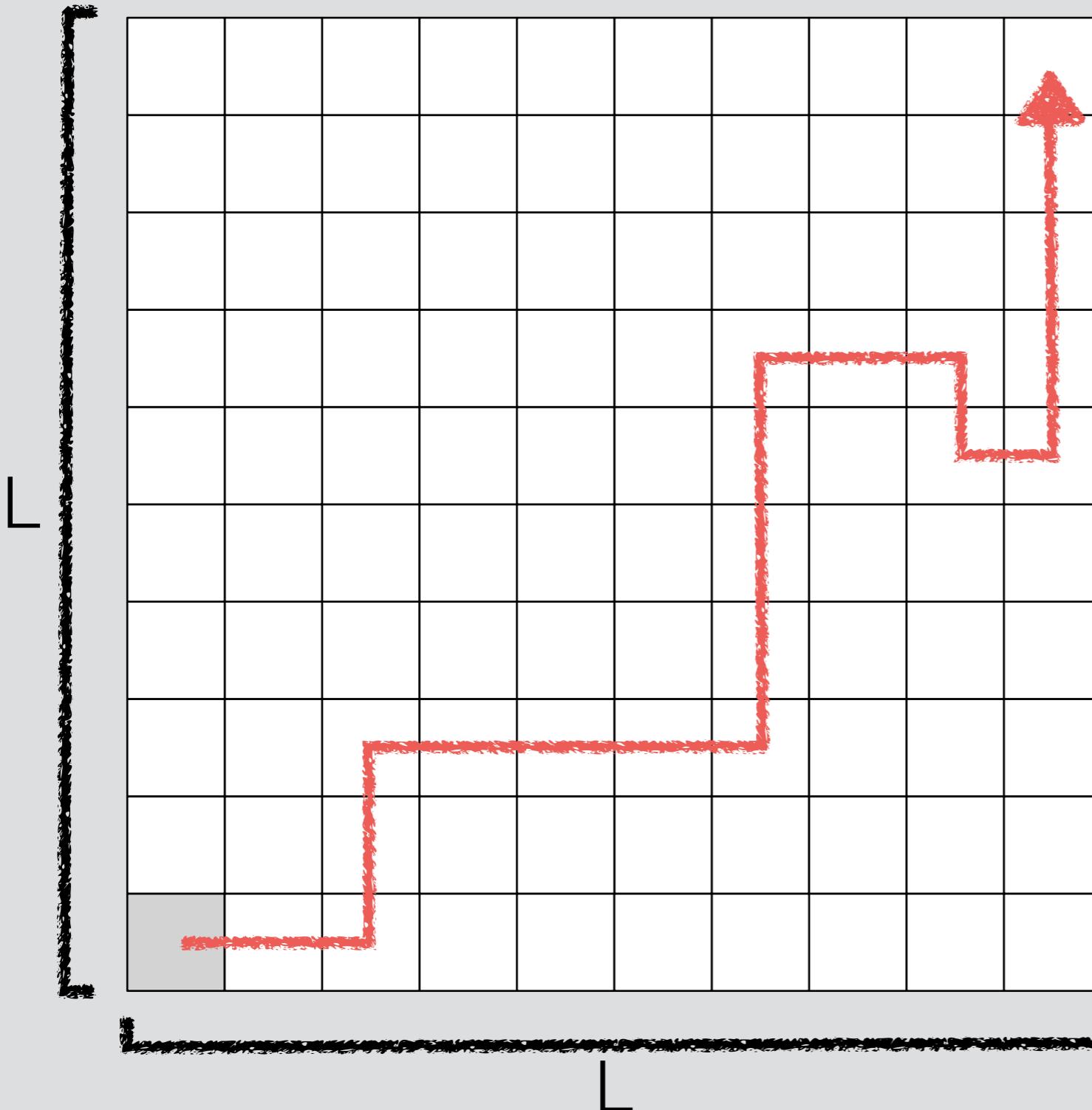
## Theoretical model (aTAM)



# A growth rate limitation

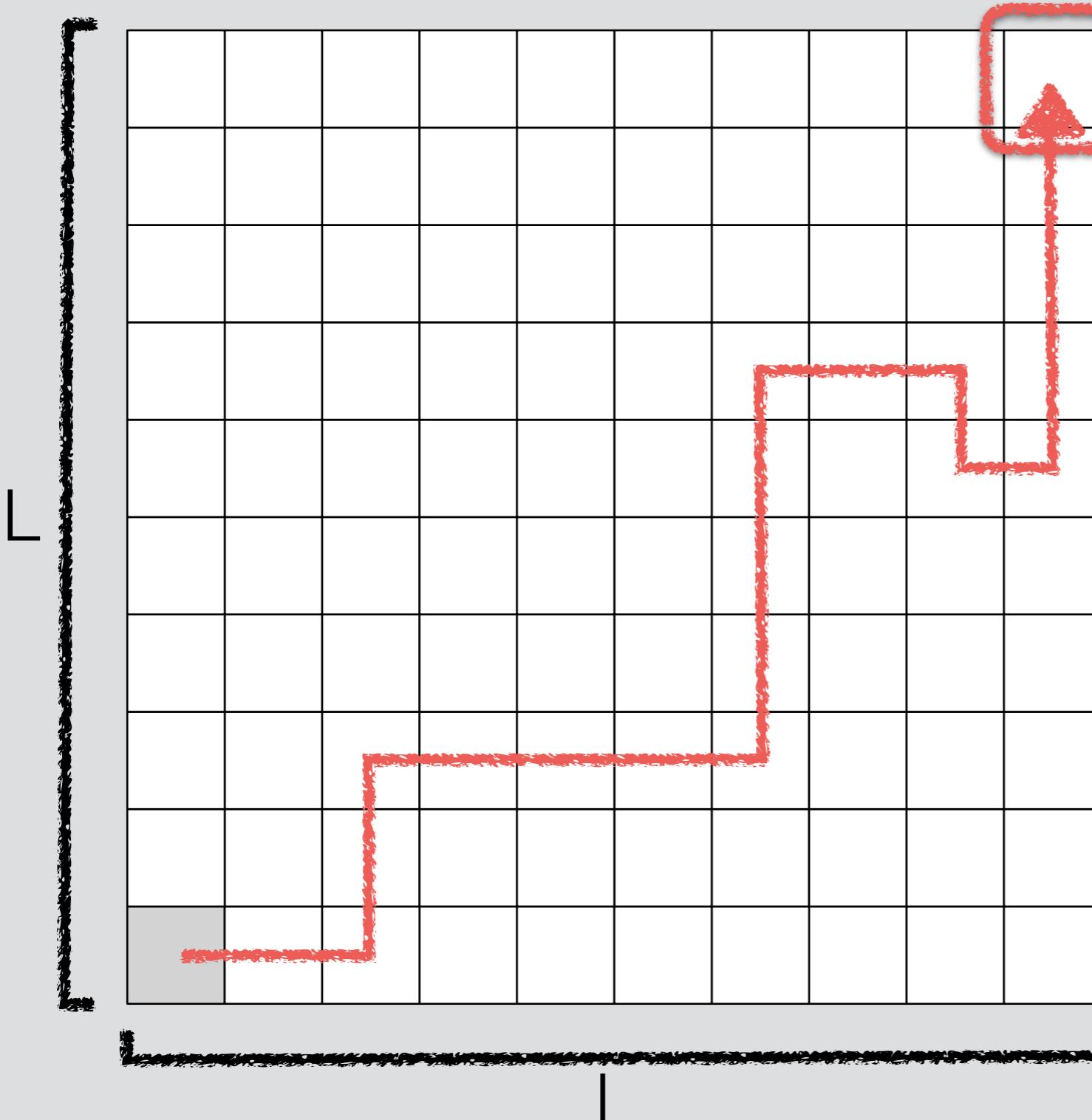


# A growth rate limitation



path length  $\geq L$

# A growth rate limitation



path length  $\geq L$

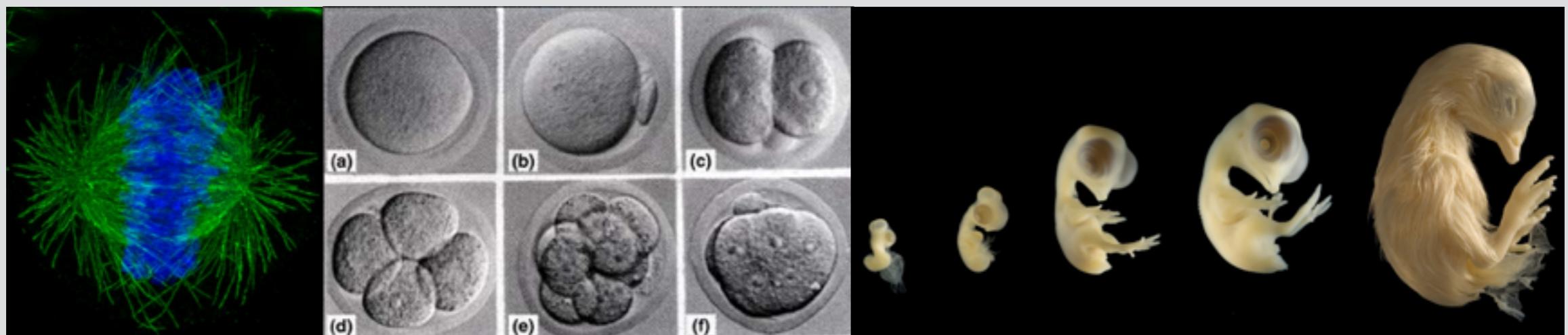
at least  $L$  tile prior  
tile placements

# Growth rates

- Takes  $\Omega(\text{diameter}) = \Omega(n^{0.5})$  expected time for  $n$  particles to assemble on a square lattice.
- So growth rate is  $O(n^2)$ .
- Similar polynomial bounds for other lattices.

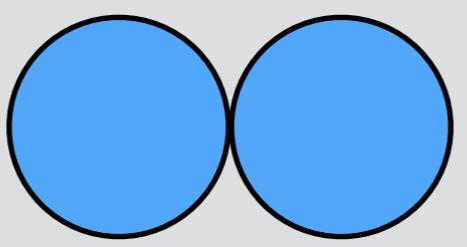
# Growth rates

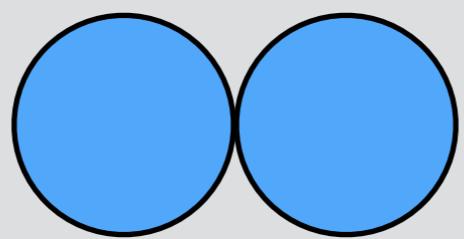
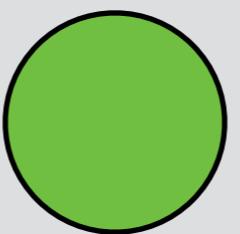
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- So growth rate is  $O(n^2)$ .
- Similar polynomial bounds for other lattices.
- But not all nanoscale growth is polynomial!

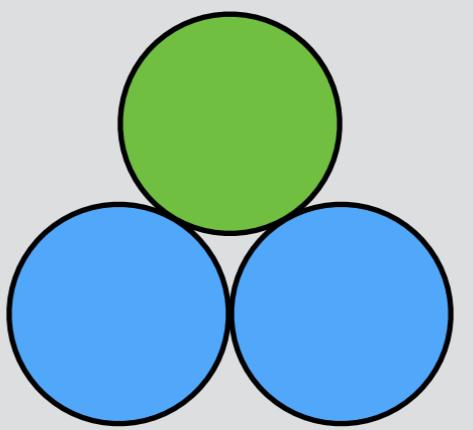


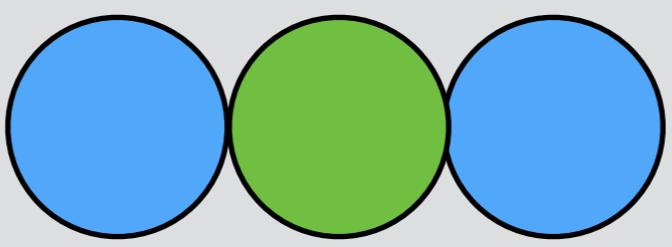
# Passive vs. Active Self-Assembly

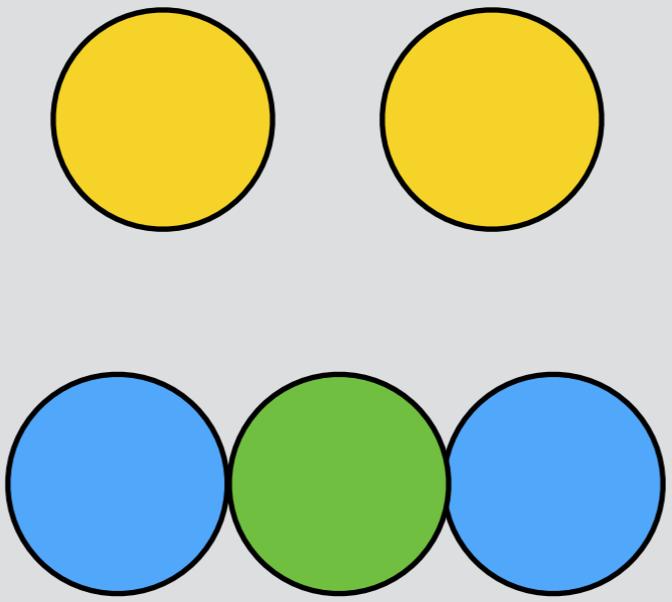
- Most current DNA self-assembly uses crystal-like **passive** growth: bonds and geometry do not change.
- Some natural systems use **active** growth: bonds and geometry change.
- Active growth enables exponential growth rates.

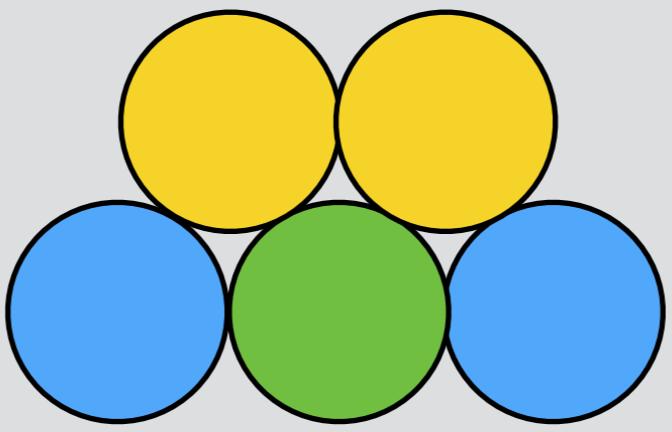


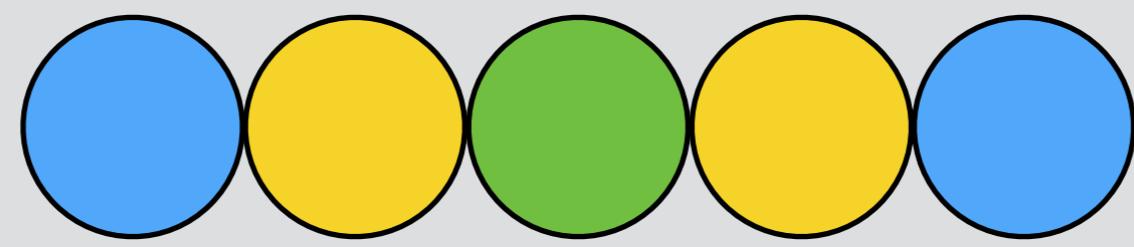


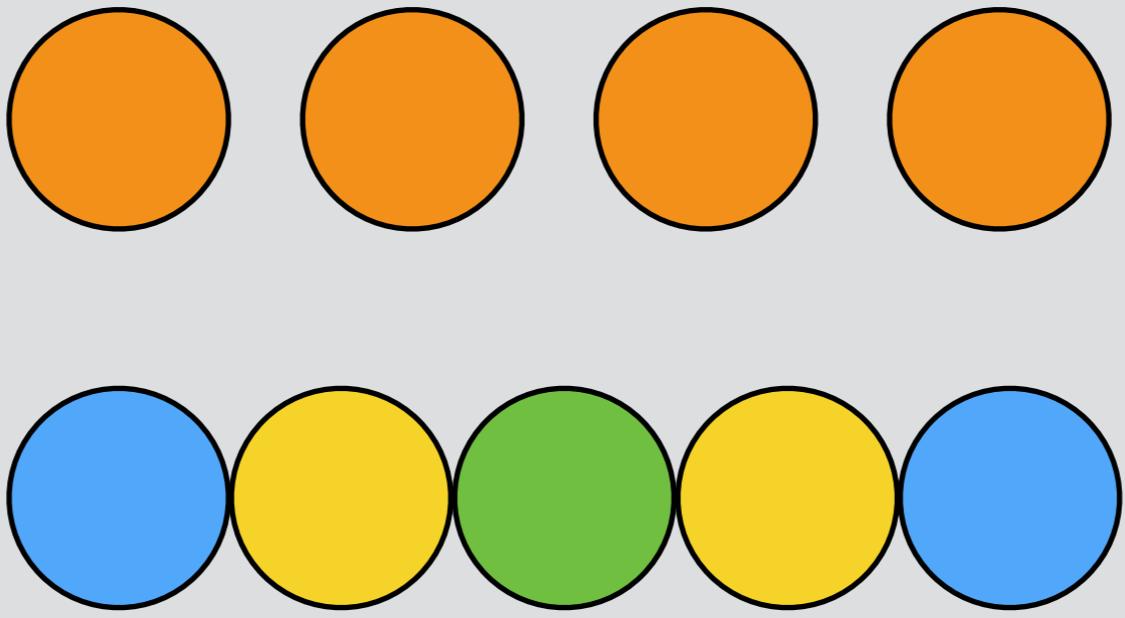


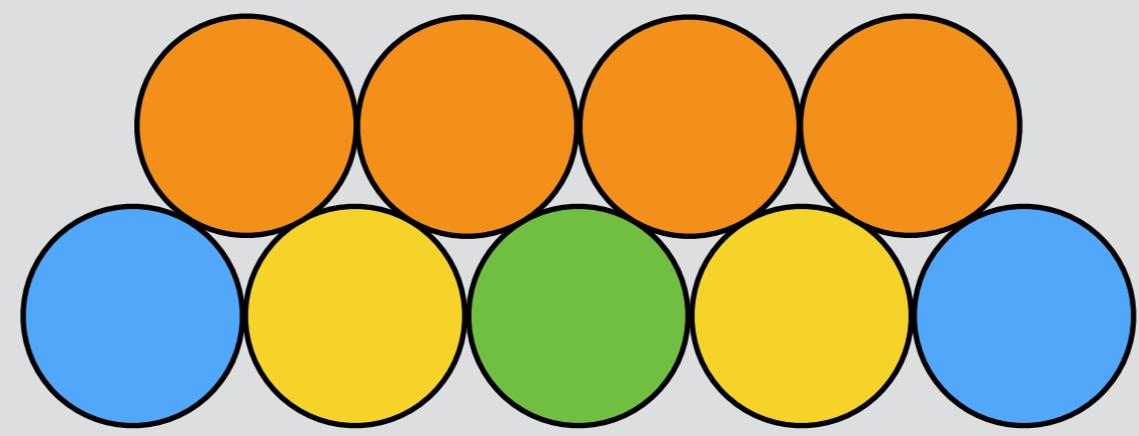


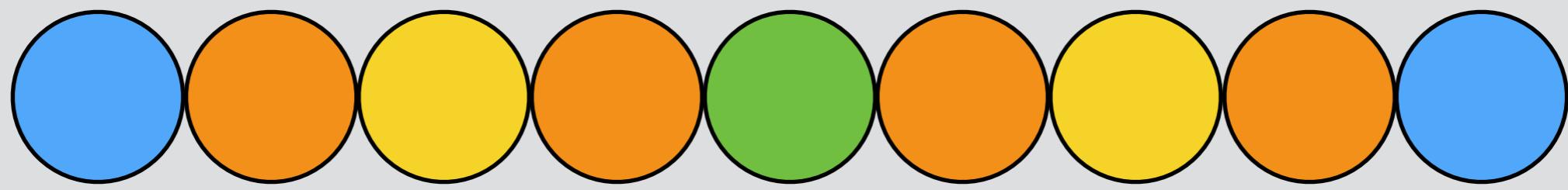


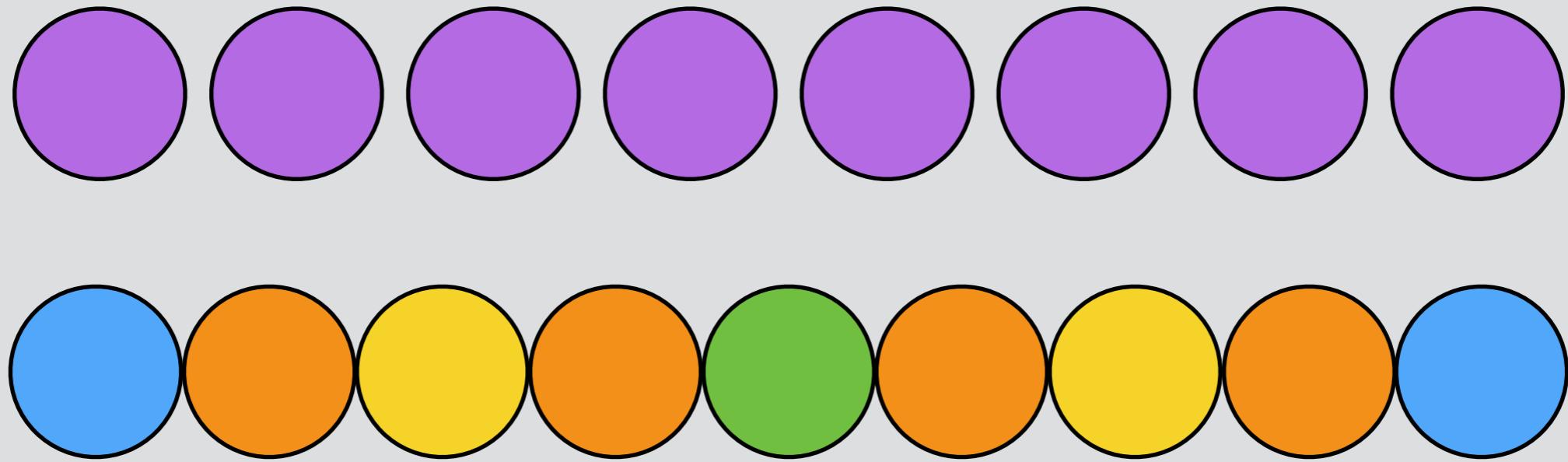


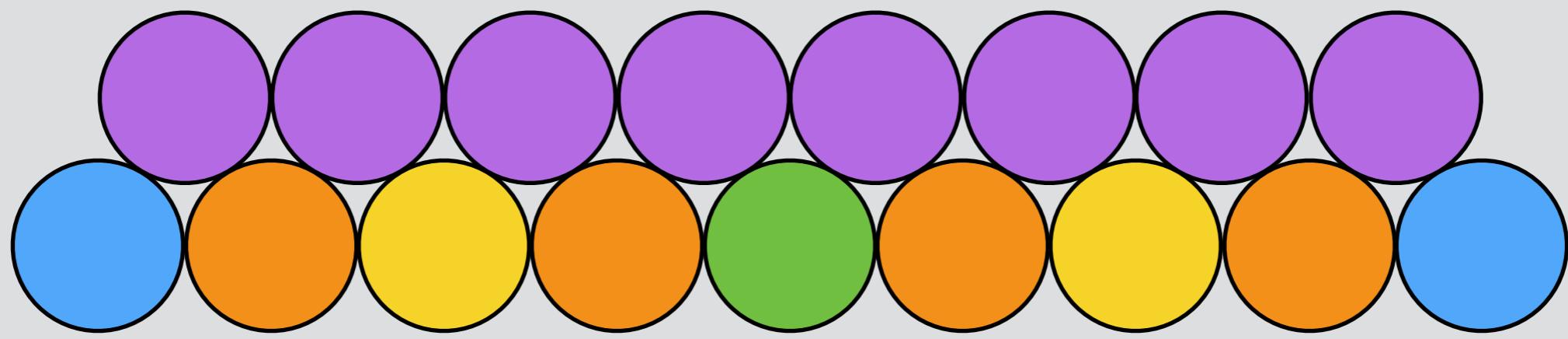


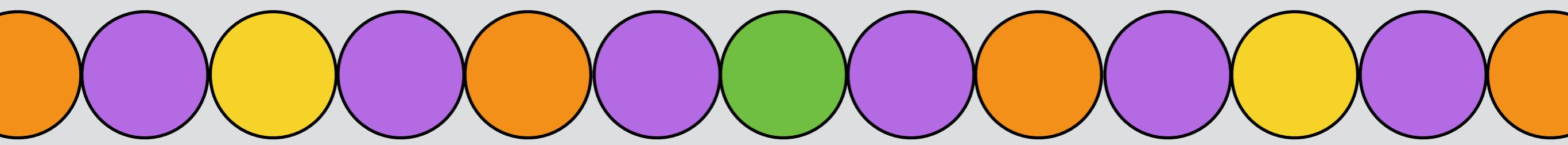




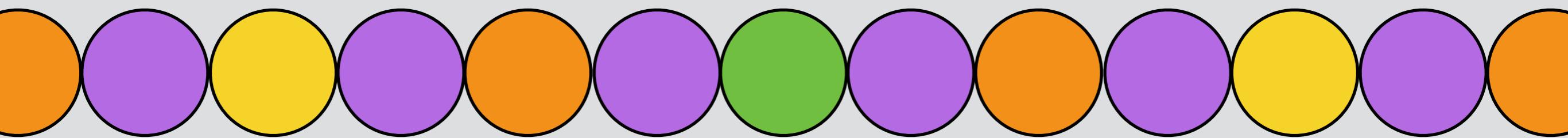






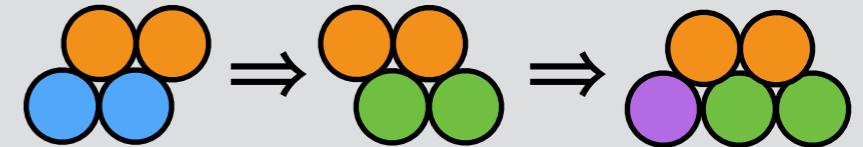


# Exponential growth!



# Active self-assembly models

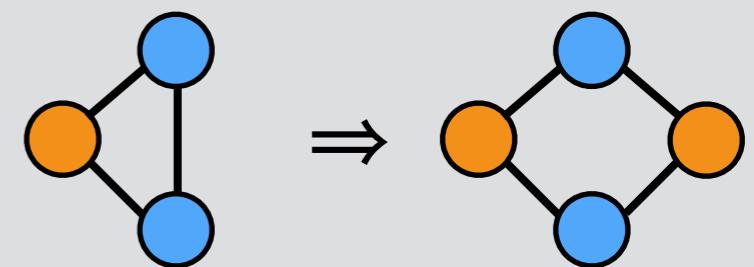
Nubots [Woods et al. ITCS 2012]:  
2D, flexible and rigid bonds, stateful particles.



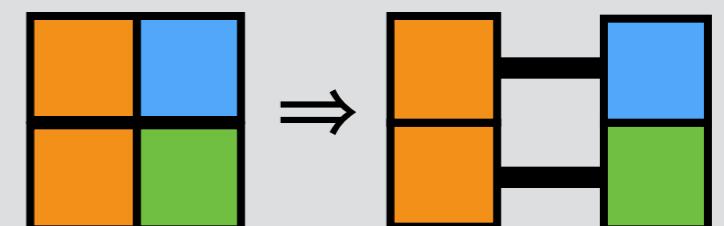
Insertion systems [Dabby, Chen SODA 2013]:  
1D, fixed shape, stateless particles.



Graph grammars [Klavins et al. ICRA 2004]:  
Geometry-less, stateless particles.



Crystalline robots [Rus, Vona ICRA 1999]:  
3D, stateful particles, global communication.



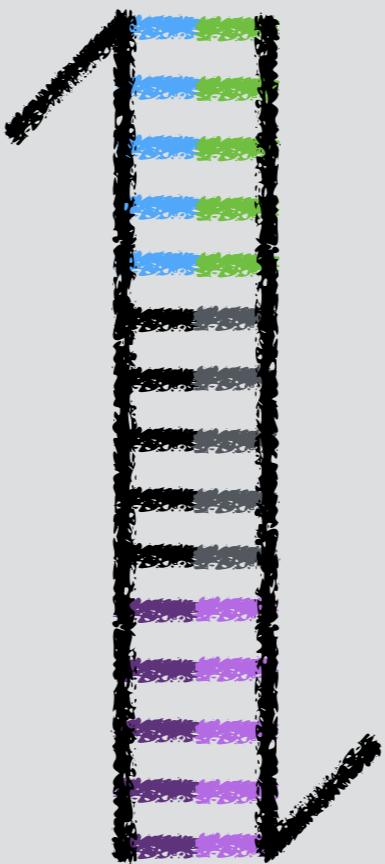
# Insertion systems

- Introduced by [Dabby, Chen 2013].
- A model of active self-assembly.
  - Implementable in DNA.
  - Capable of exponential growth.
  - Grows a linear structure by insertion of particles.

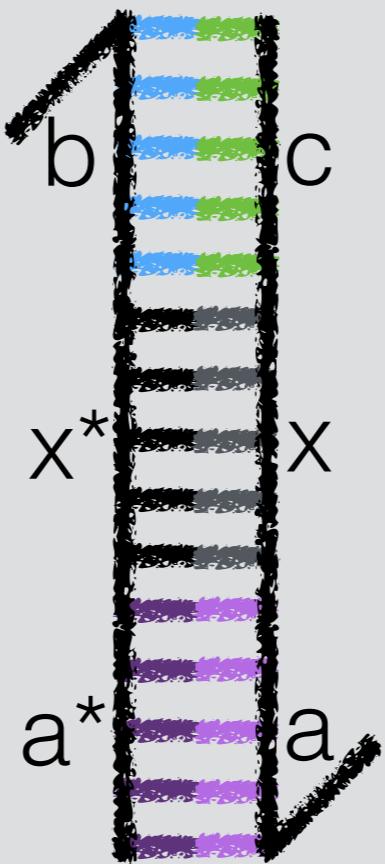
# Insertion systems

- Introduced by [Dabby, Chen 2013].
- A model of active self-assembly.
  - Implementable in DNA.
  - Capable of exponential growth.
  - Grows a linear structure by insertion of particles.
- Our work: bound capabilities of insertion systems.

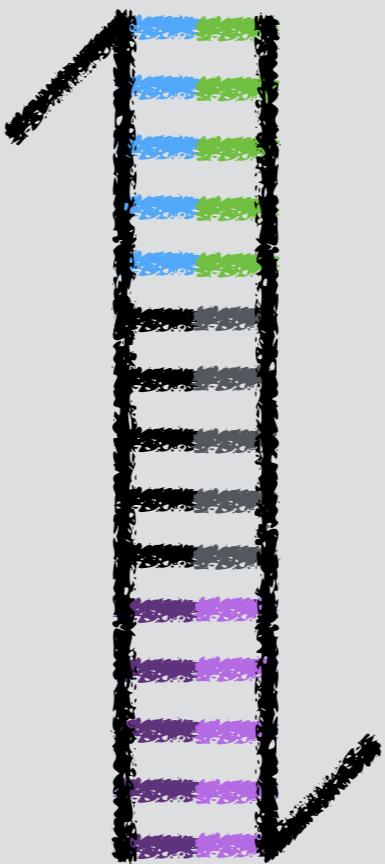
# DNA insertions



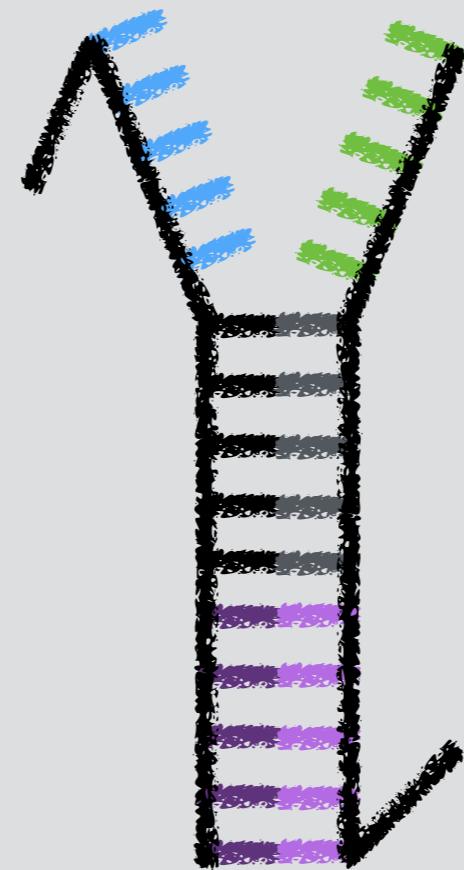
# DNA insertions



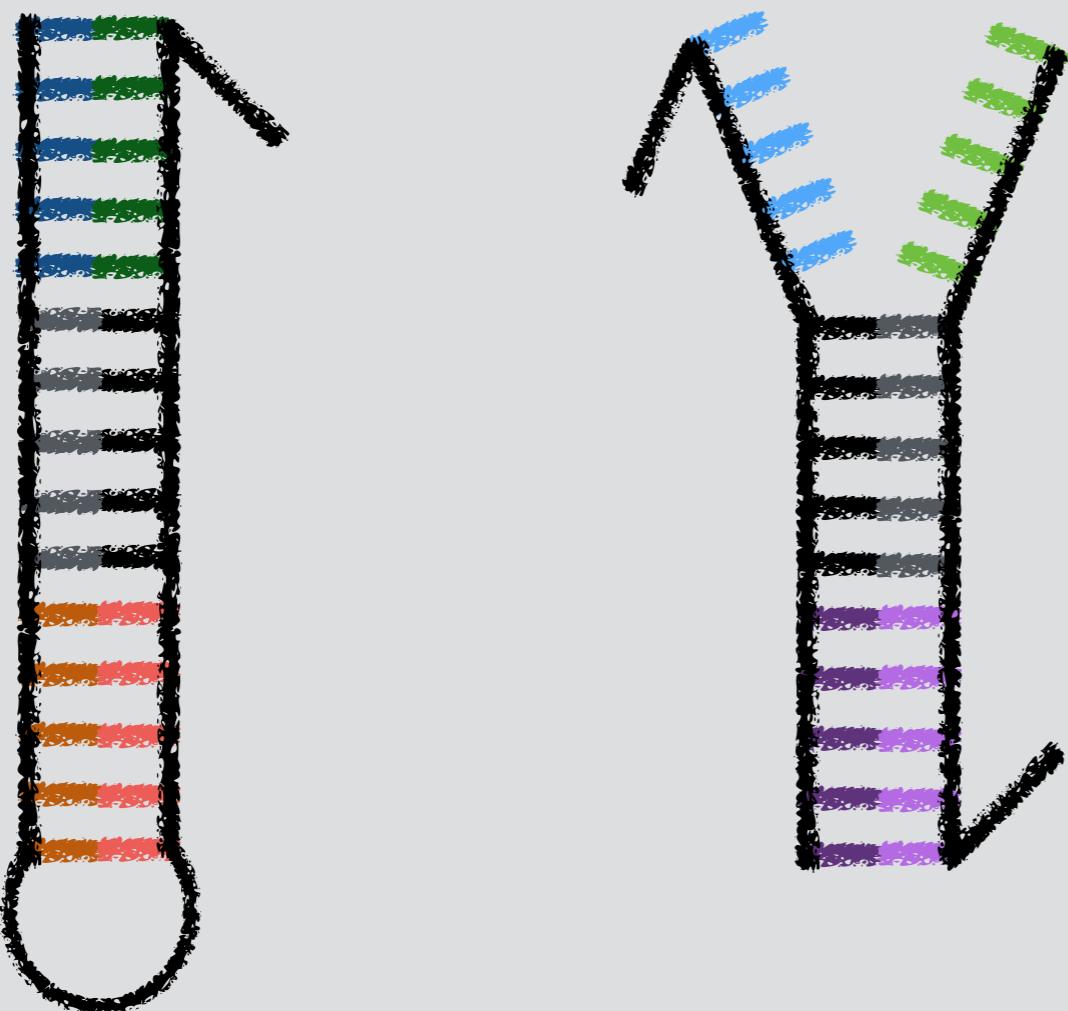
# DNA insertions



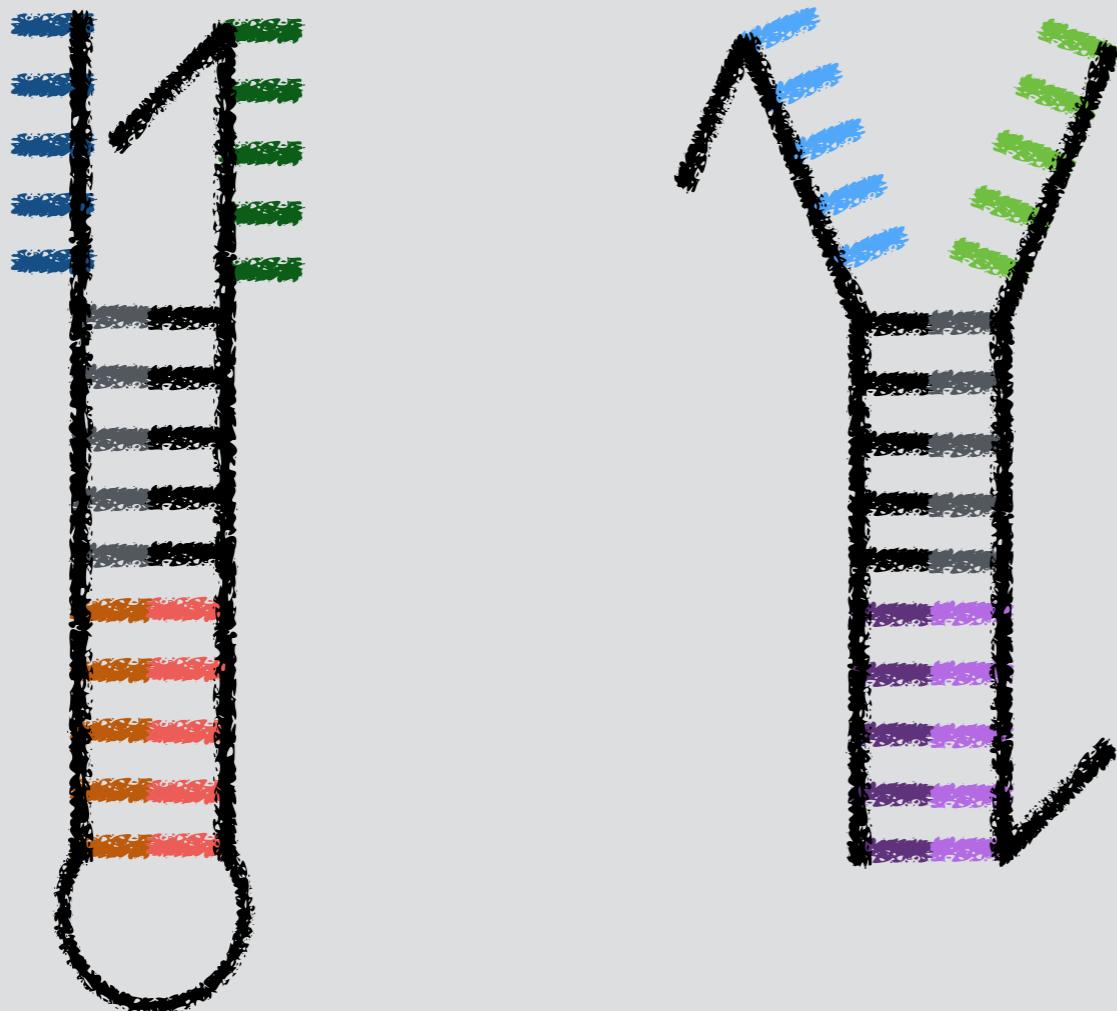
# DNA insertions



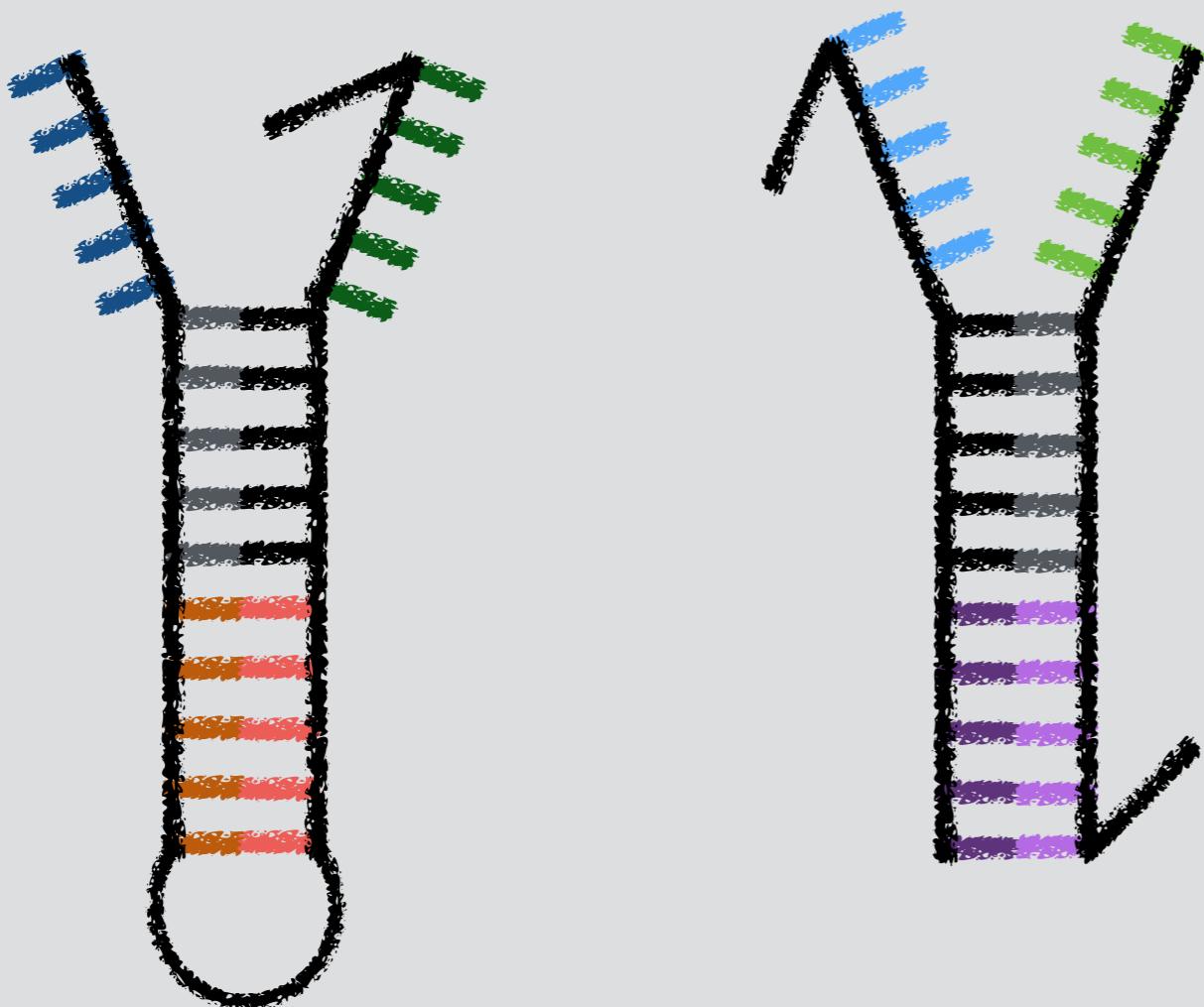
# DNA insertions



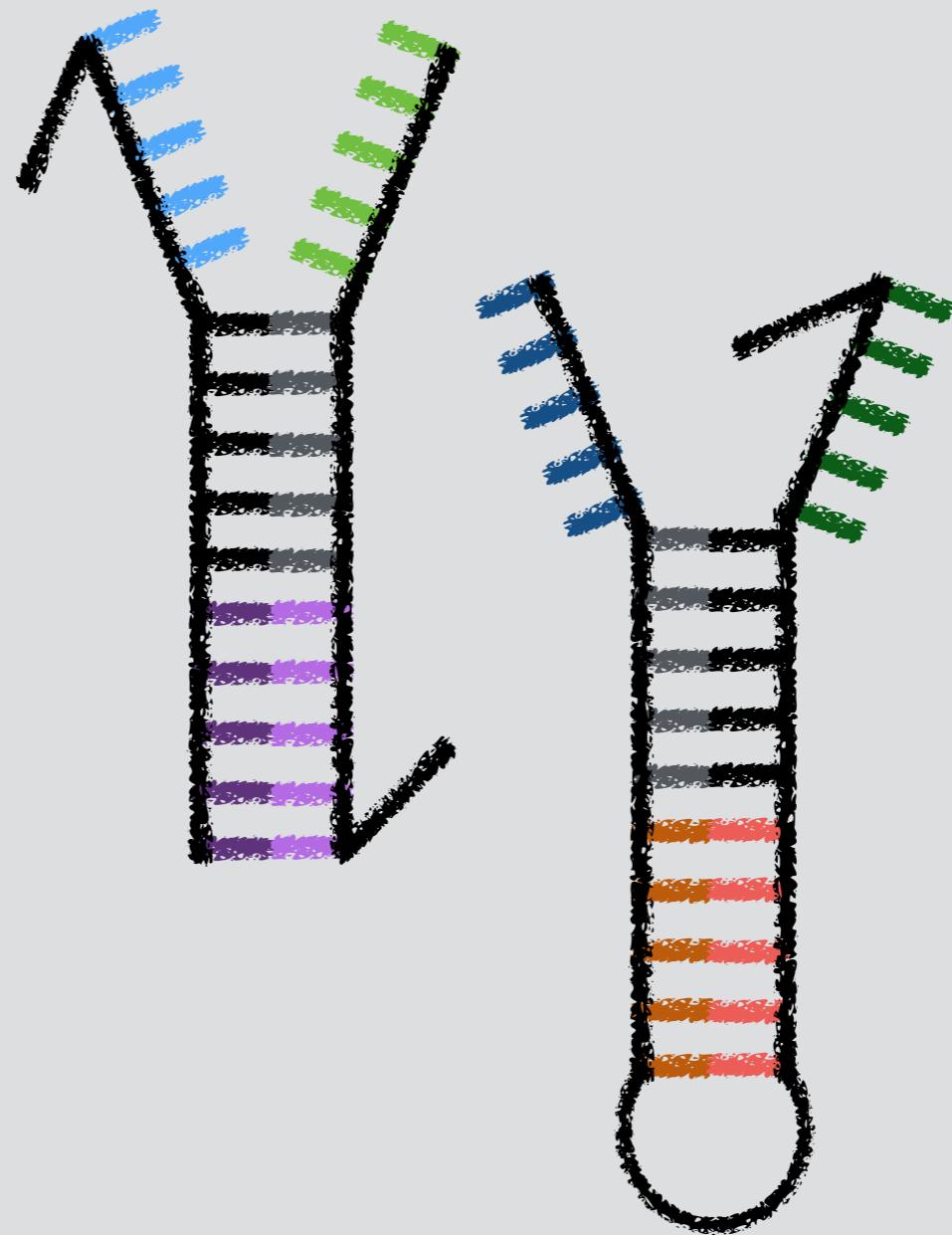
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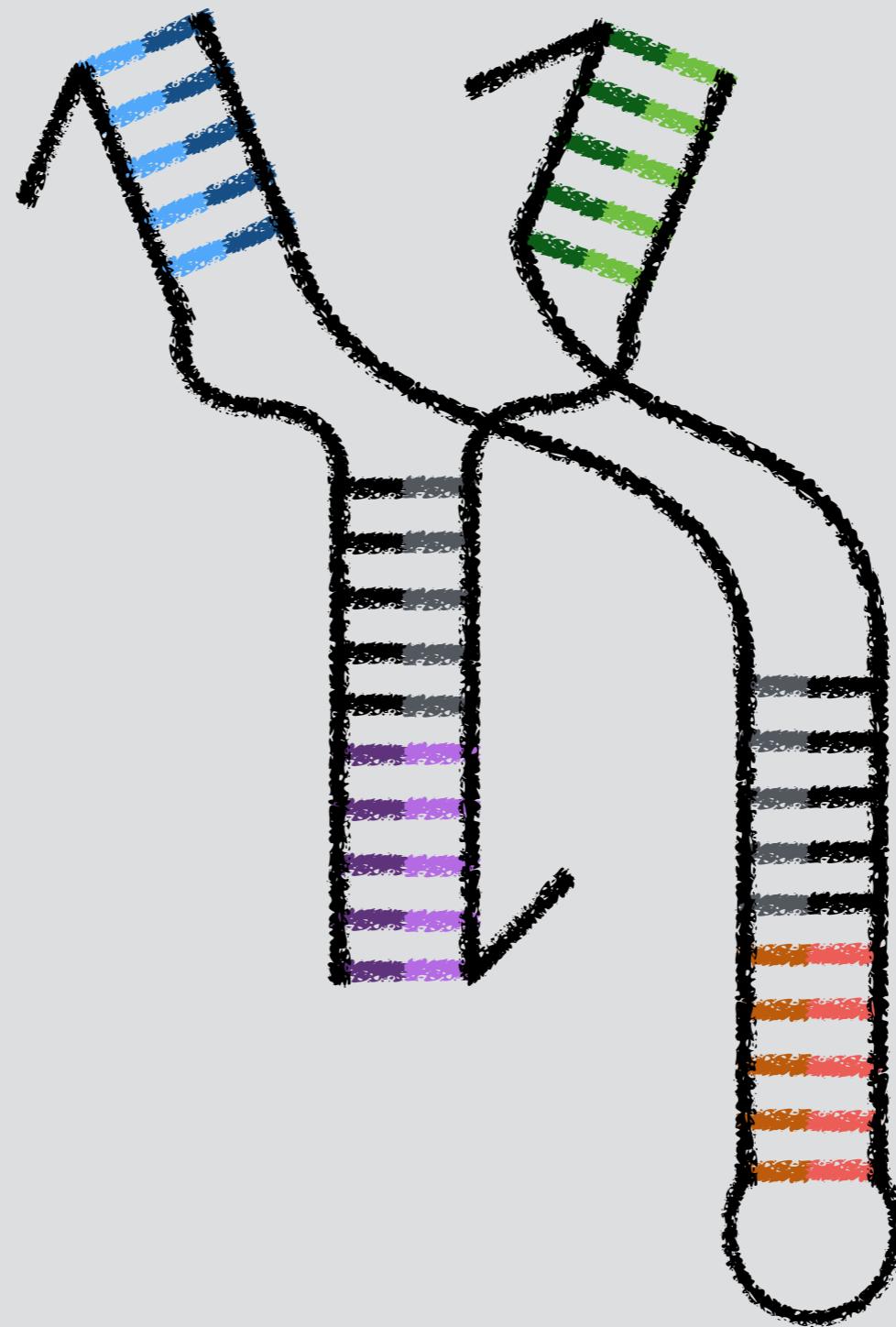
# DNA insertions



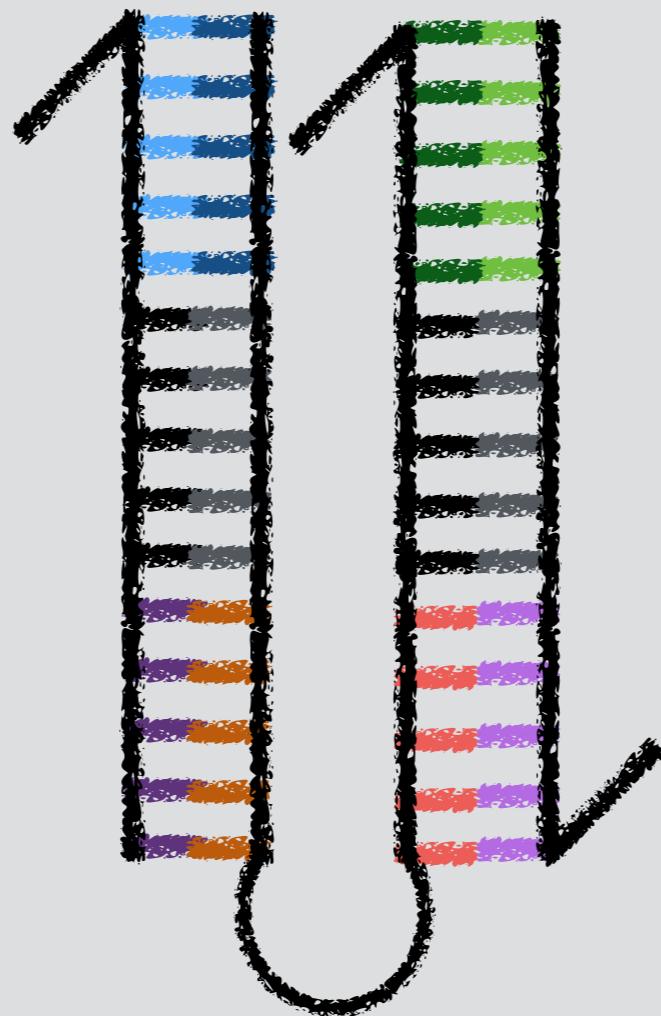
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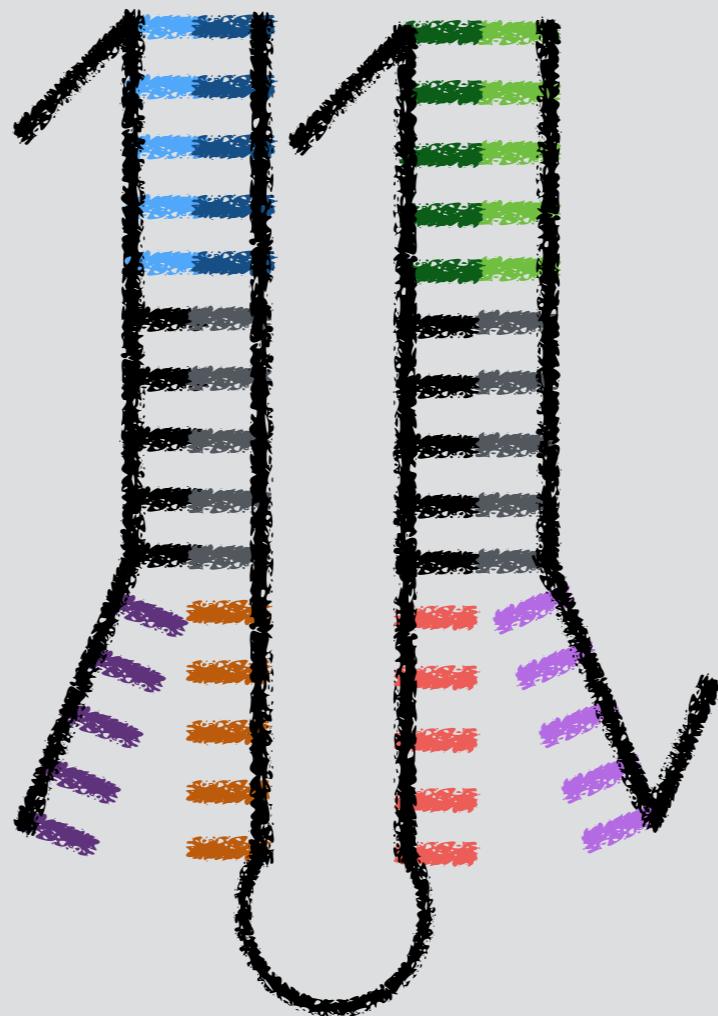
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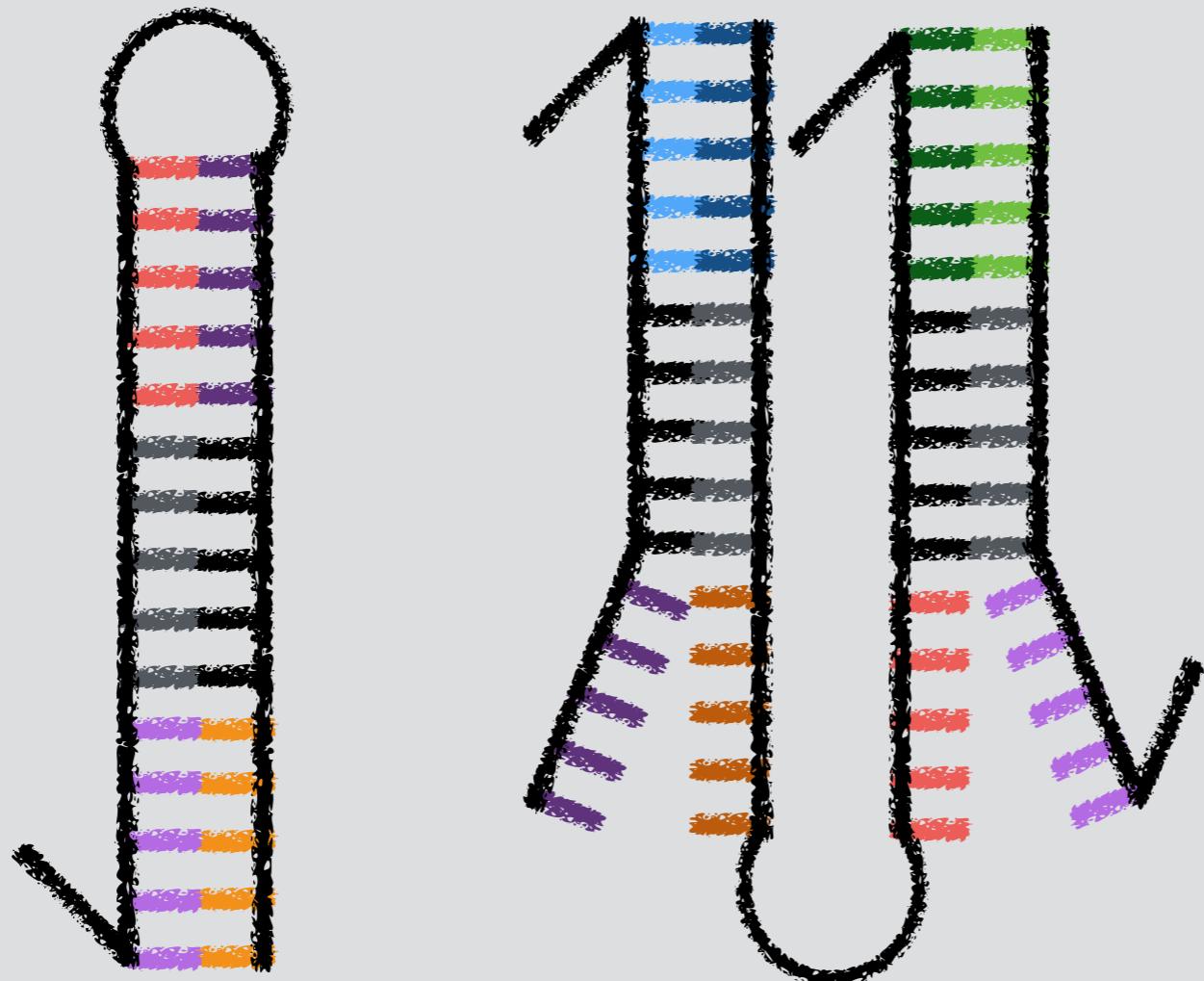
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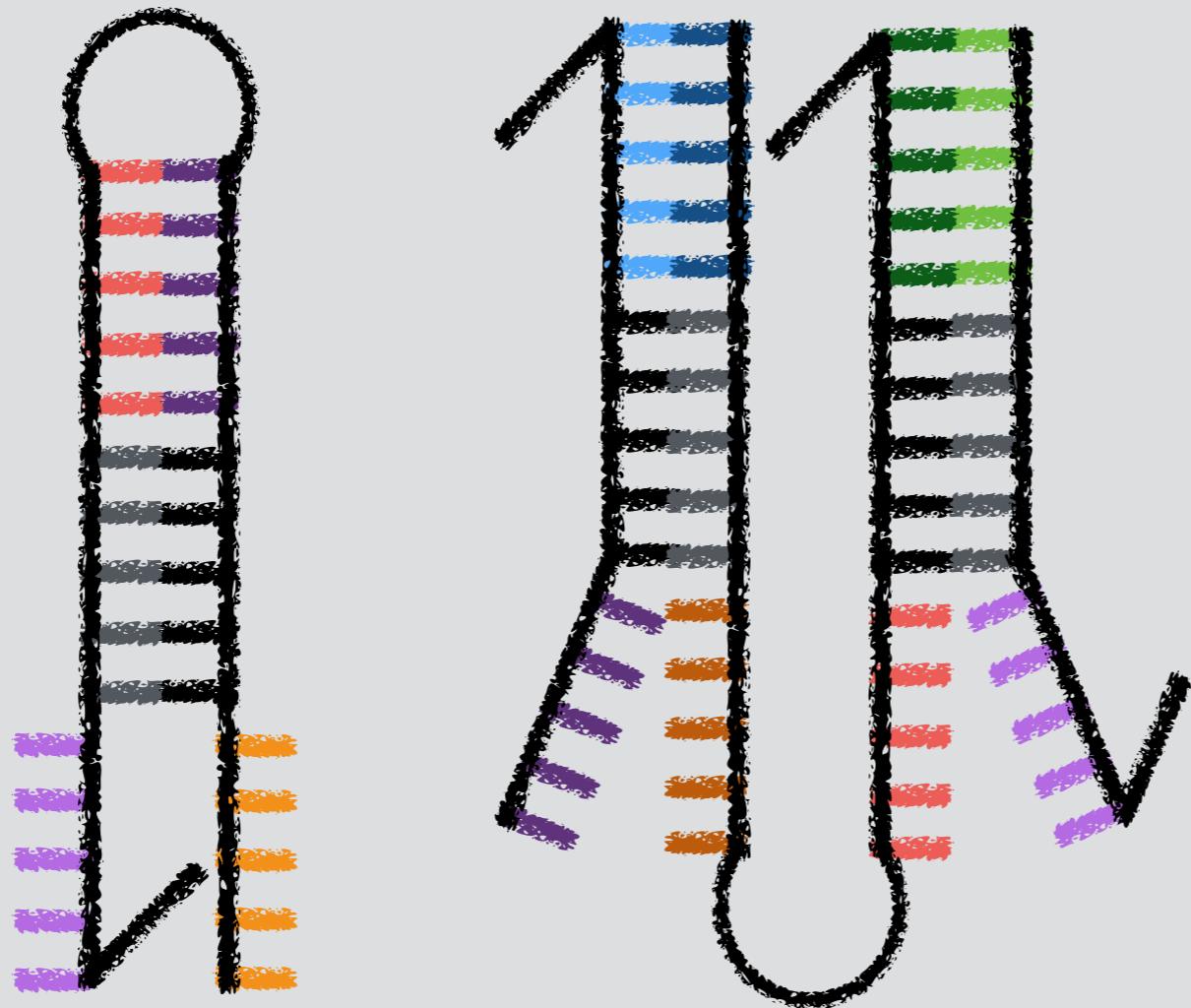
# DNA insertions



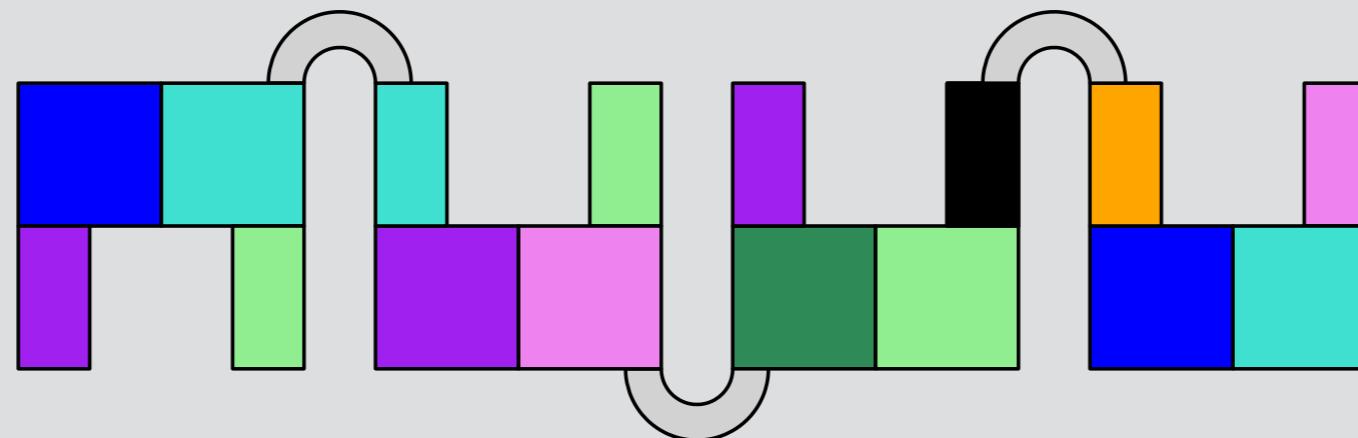
# DNA insertions



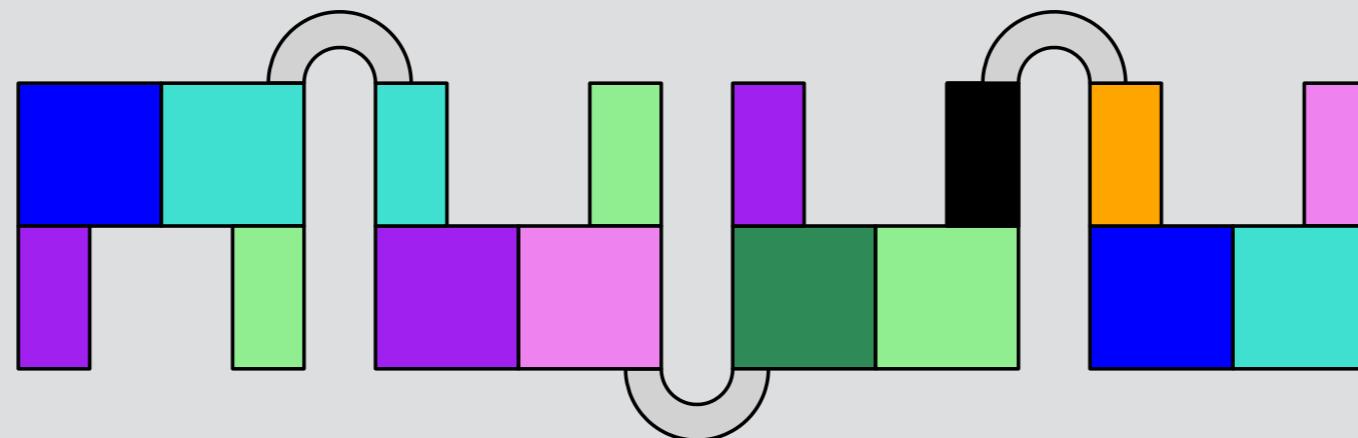
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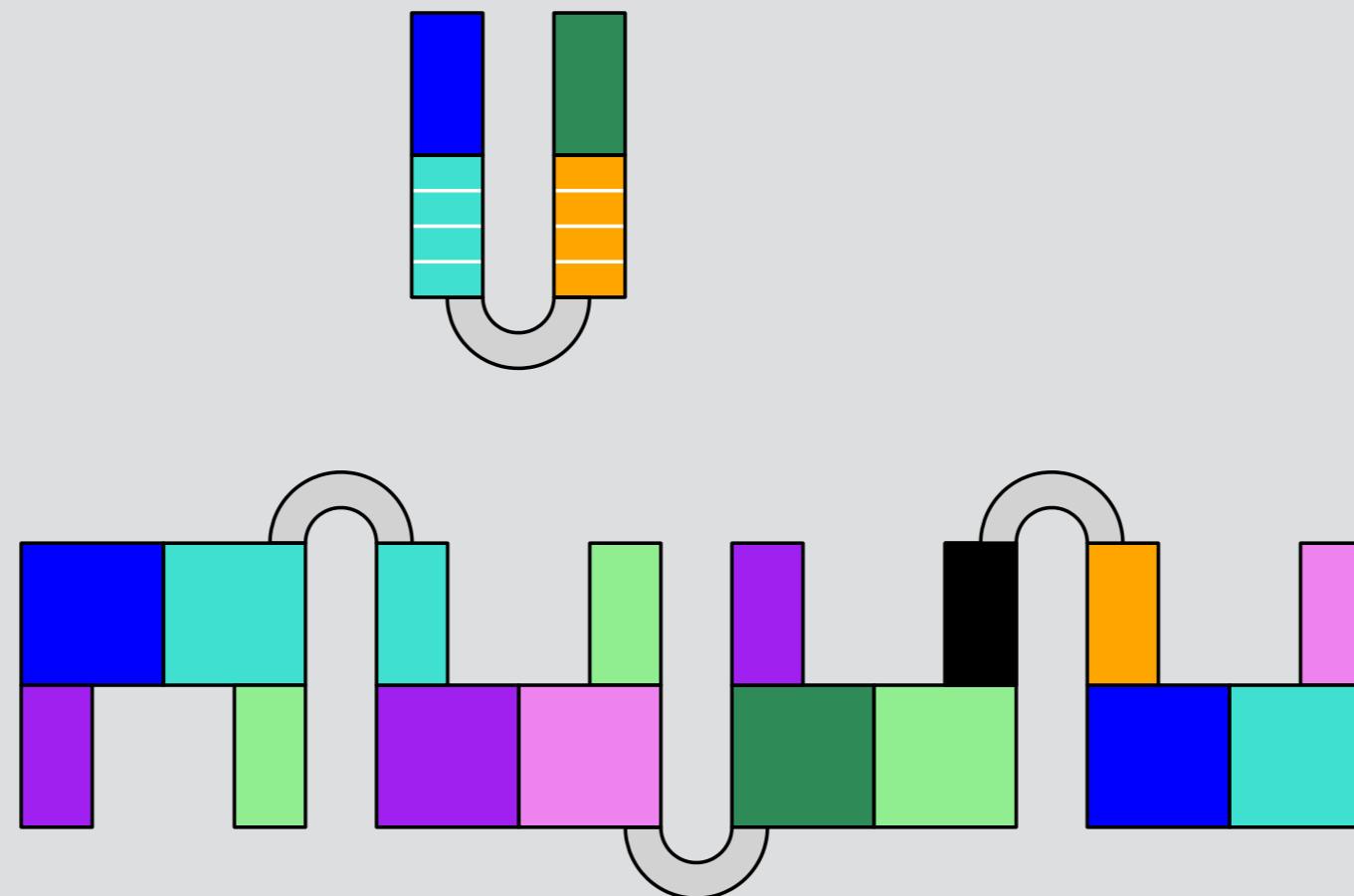
# Insertion systems



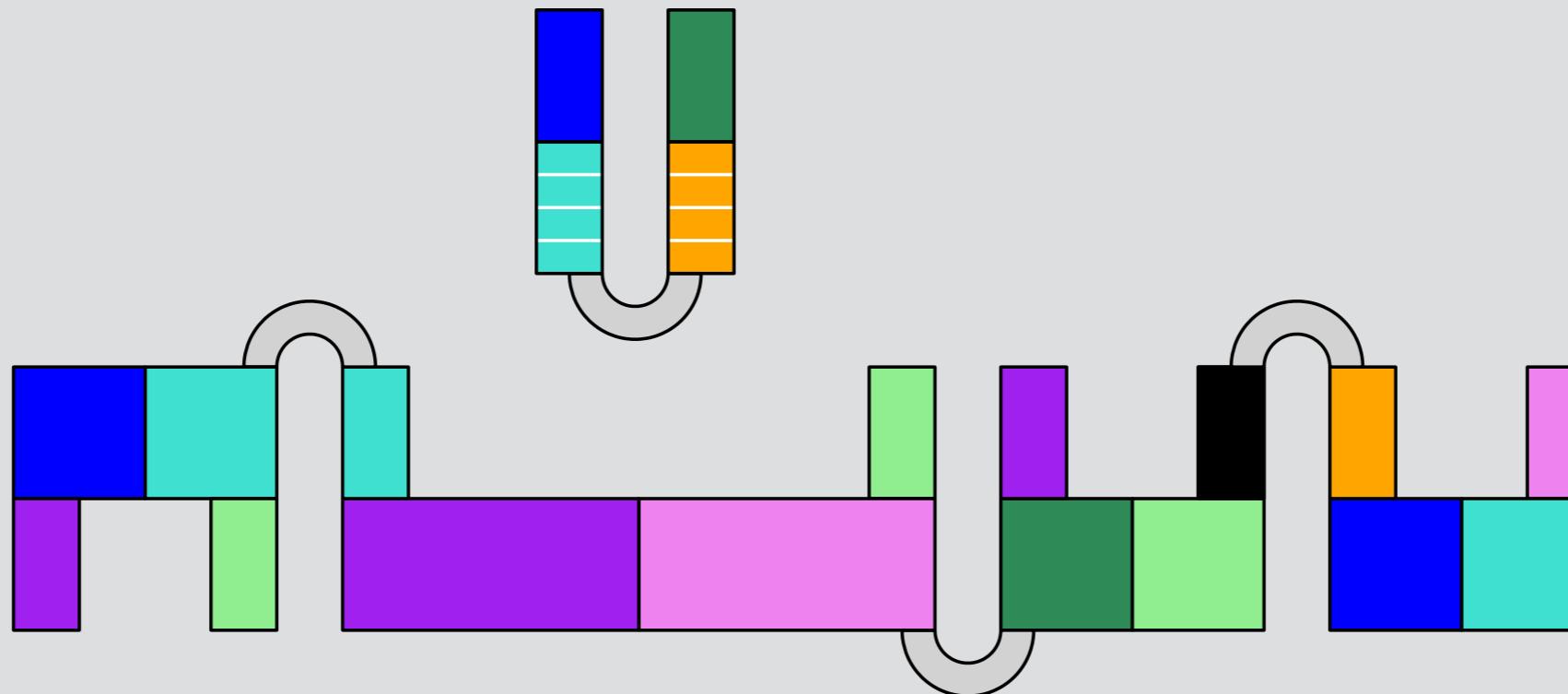
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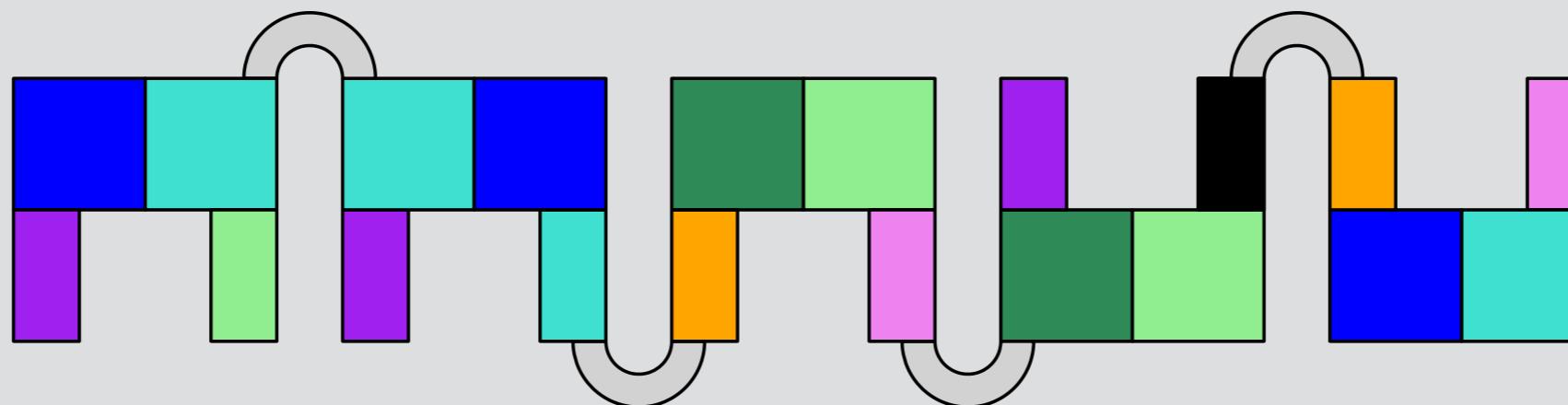
# Insertion systems



# Insertion systems

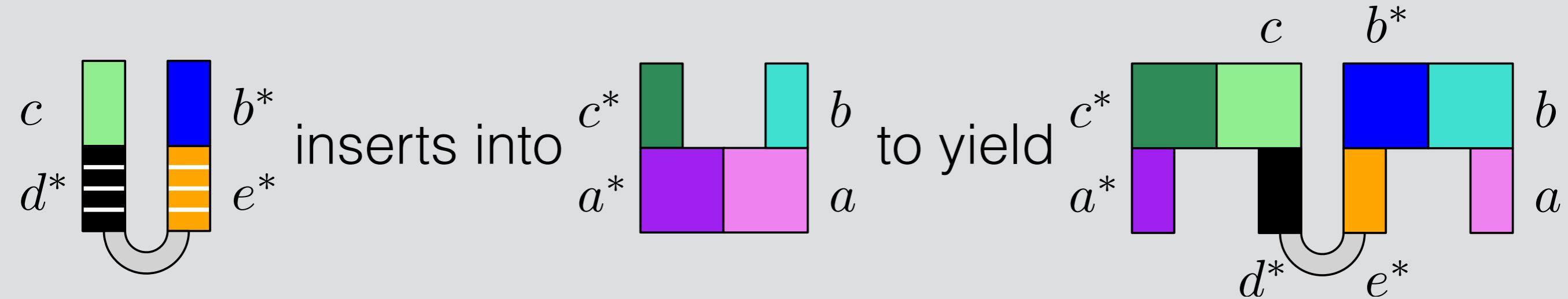


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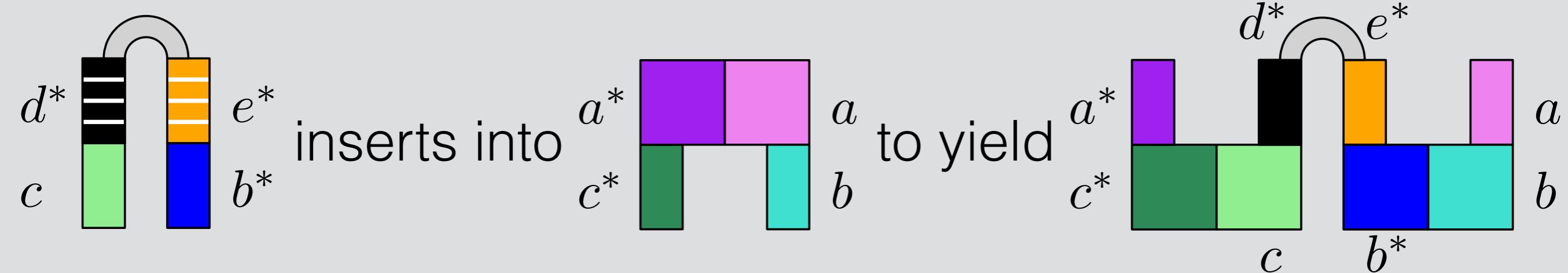


# Definitions and examples

# Insertions



$(c, d^*, e^*, b^*)^+$  inserts into  $(a^*, c^*)(b, a)$  to yield  $(a^*, c^*)(c, d^*, e^*, b^*)(b, a)$



$(d^*, c, b^*, e^*)^-$  inserts into  $(c^*, a^*)(a, b)$  to yield  $(c^*, a^*)(d^*, c, b^*, e^*)(a, b)$

## **Insertion system:**

Monomer types:  $(1,x,2,b)^+$   $(x,2^*,a,3^*)^-$   $(3,x,4,b)^+$

Initiator:  $(a,1^*)(b^*,a^*)$

---

## **Insertion system:**

Monomer types:  $(1,x,2,b)^+$   $(x,2^*,a,3^*)^-$   $(3,x,4,b)^+$

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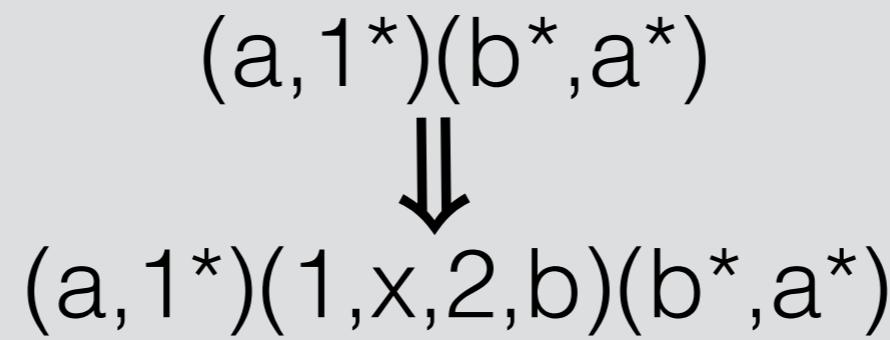
---

$(a,1^*)(b^*,a^*)$

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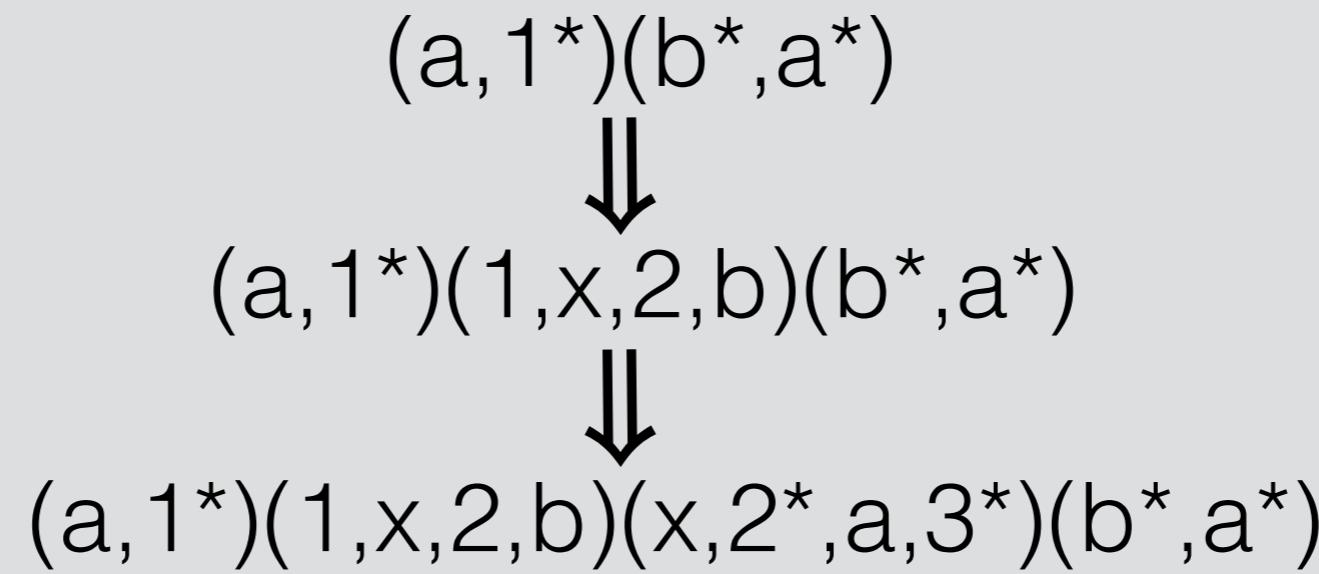


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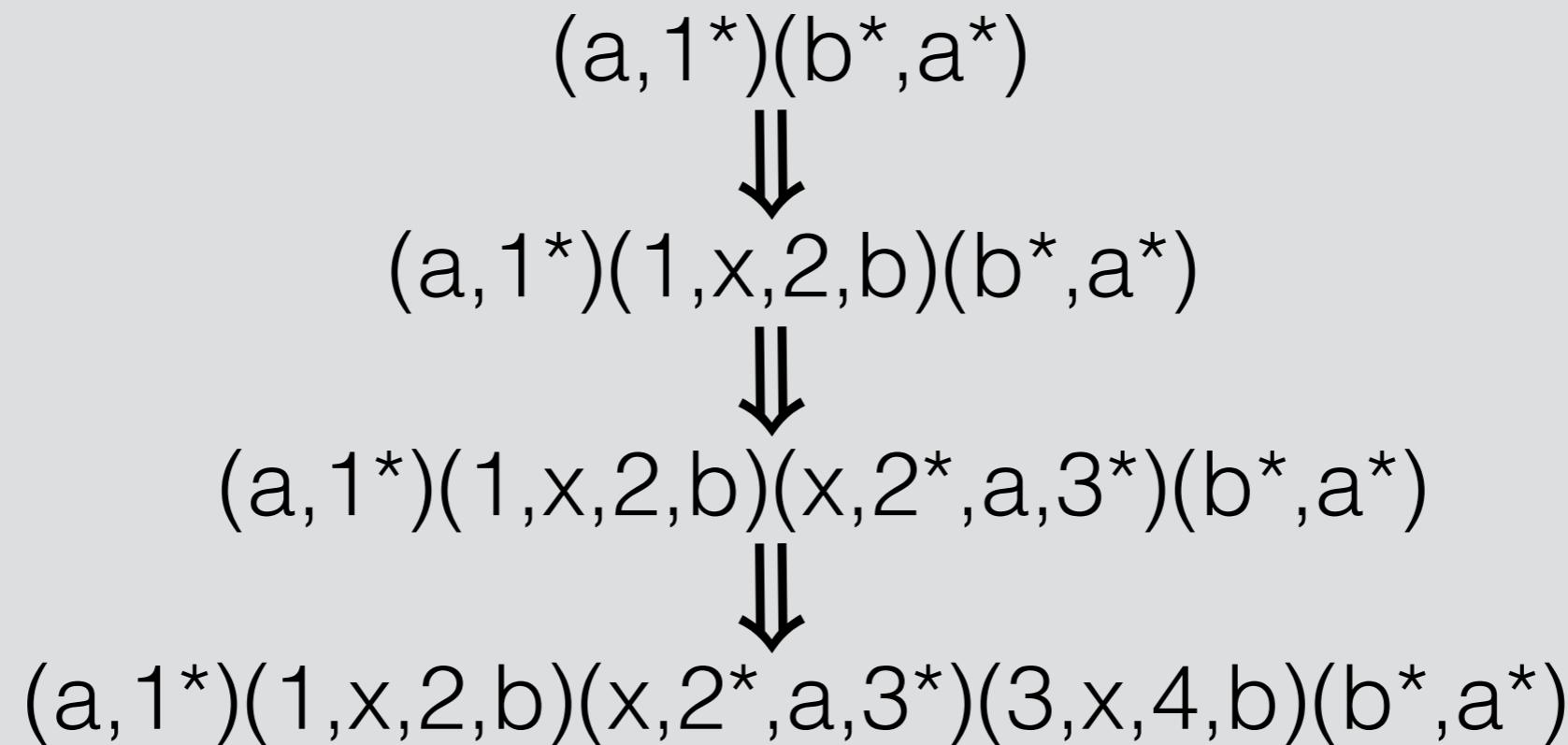


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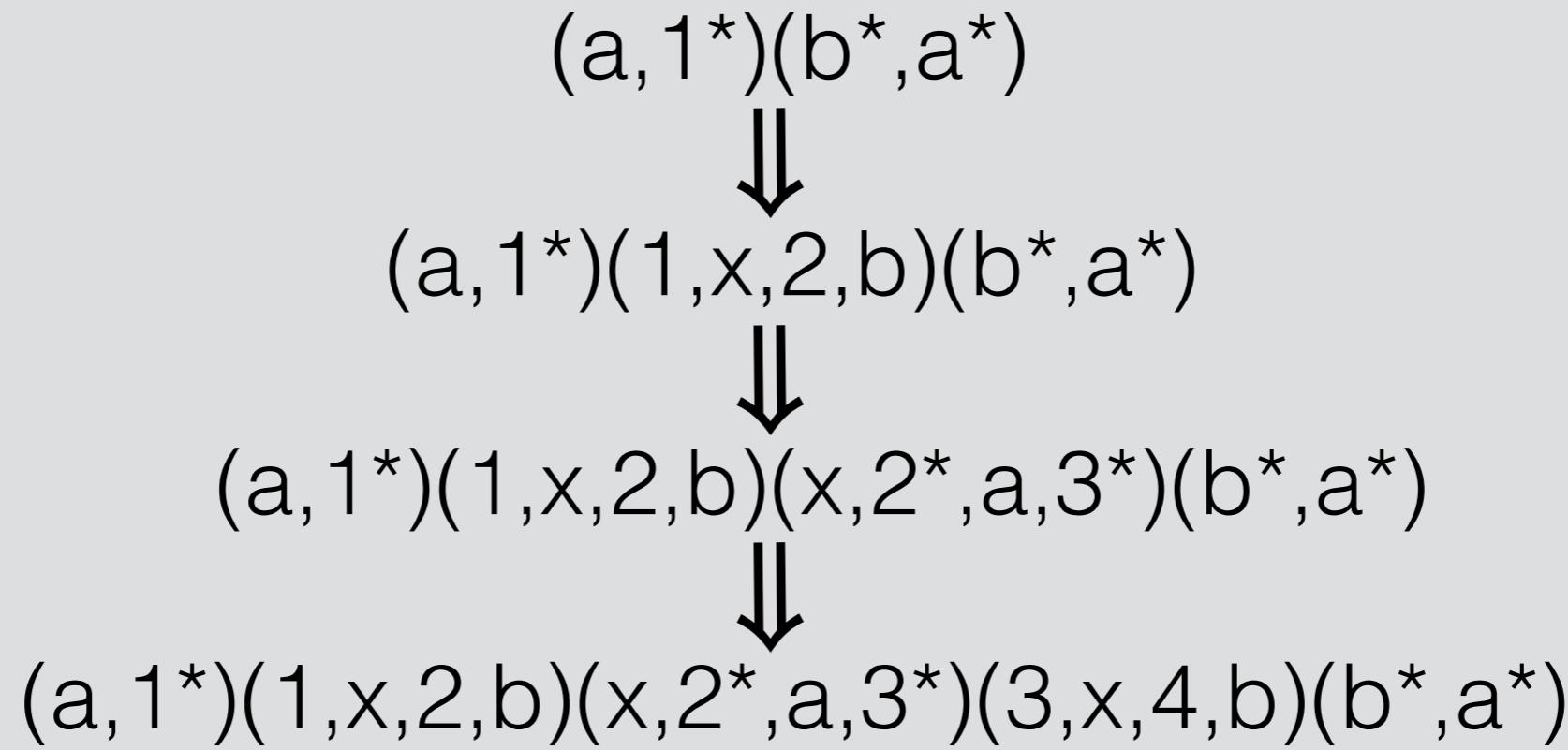
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Monomer types:  $(1,x,2,b)^+$   $(x,2^*,a,3^*)^-$   $(3,x,4,b)^+$

Initiator:  $(a,1^*)(b^*,a^*)$



*Terminal polymer of length 5*

# Insertion time

- Each monomer type has a concentration in  $[0,1]$ .
- Concentrations of all types in a system must sum to  $\leq 1$ .
- An insertion occurs after time  $t$  with:
  - $t$  an exponential random variable with rate  $c$ .
  - $c$  is the total concentration of insertable monomers.

## **Insertion system:**

Monomer types:  $(1,x,2,b)^+$   $(x,2^*,a,3^*)^-$   $(3,x,4,b)^+$

Initiator:  $(a,1^*)(b^*,a^*)$

---

## **Insertion system:**

Monomer types:  $(1,x,2,b)^+$   $(x,2^*,a,3^*)^-$   $(3,x,4,b)^+$

Concentrations:      0.25                  0.25                  0.5

Initiator:  $(a,1^*)(b^*,a^*)$

---

## Insertion system:

Monomer types:  $(1,x,2,b)^+$   $(x,2^*,a,3^*)^-$   $(3,x,4,b)^+$

Concentrations: 0.25 0.25 0.5

Initiator:  $(a,1^*)(b^*,a^*)$

---

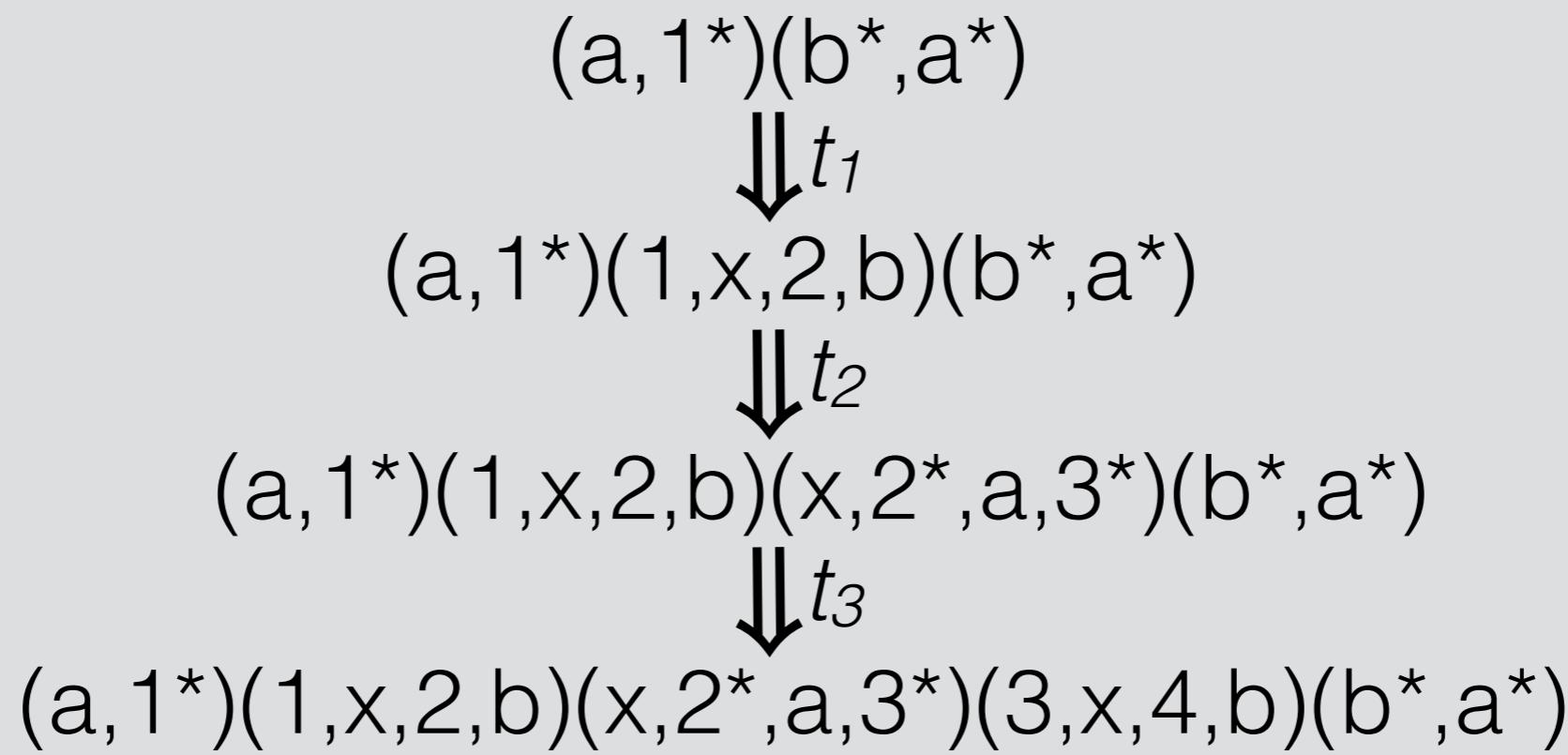
$(a,1^*)(b^*,a^*)$

## Insertion system:

Monomer types:  $(1,x,2,b)^+$   $(x,2^*,a,3^*)^-$   $(3,x,4,b)^+$

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Initiator:  $(a,1^*)(b^*,a^*)$



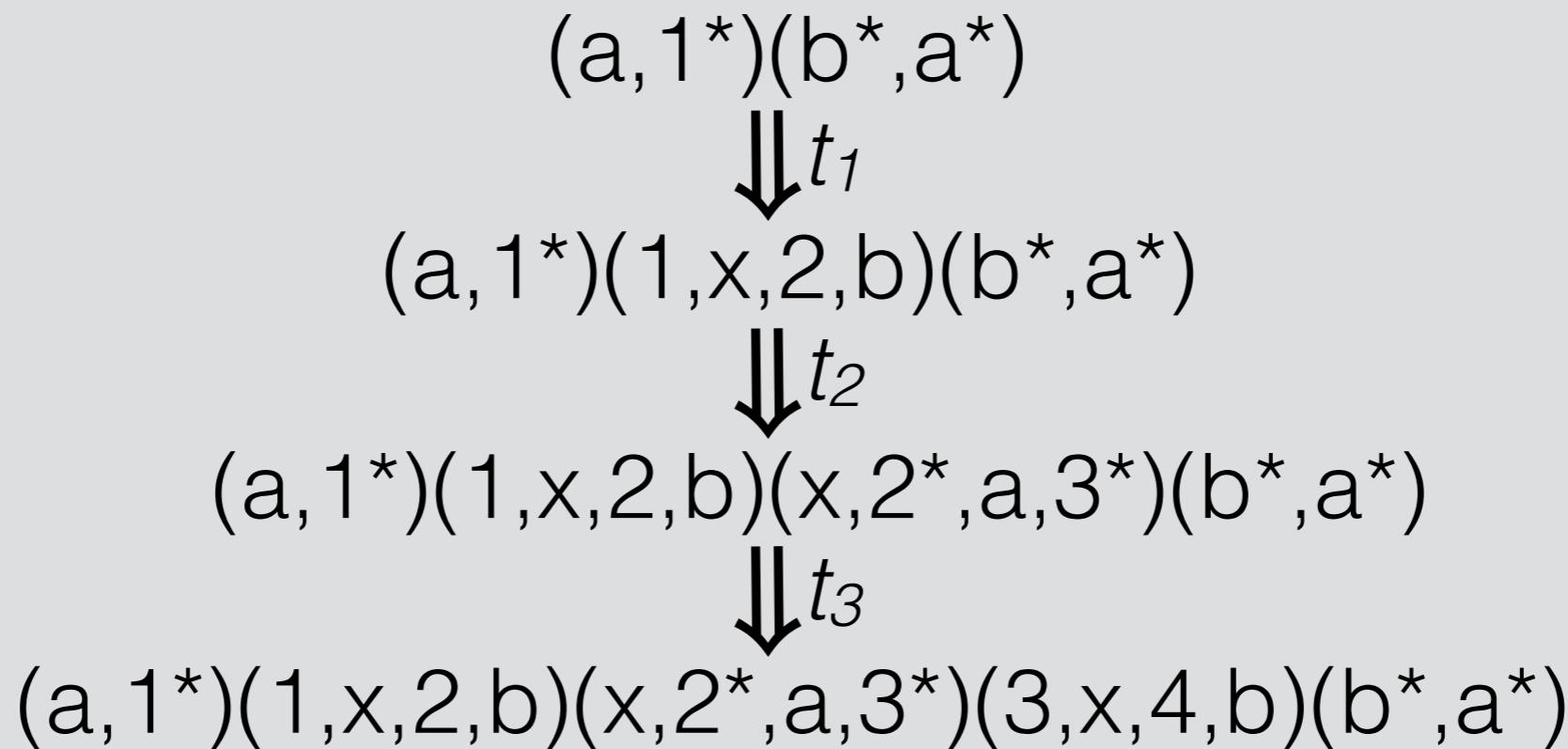
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Monomer types:  $(1,x,2,b)^+$   $(x,2^*,a,3^*)^-$   $(3,x,4,b)^+$

Concentrations: 0.25 0.25 0.5

Initiator:  $(a,1^*)(b^*,a^*)$



*Terminal polymer of length 5*

*Expected time:  $t_1 + t_2 + t_3$ , with*

$$E[t_1] = E[t_2] = 4, E[t_3] = 2.$$

$$4 + 4 + 2 = 12$$

## Insertion system:

Monomer types:  $(1^*, 2, 2, 1^*)^+$     $(x, 0^*, 2^*, x)^-$     $(x, 2^*, 0, x)^-$

Concentrations:      0.5                          0.1                          0.4

Initiator:  $(0, 1)(1, 0^*)$

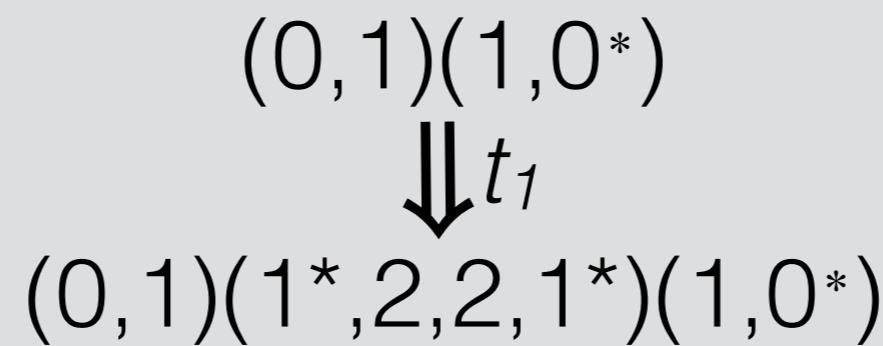
$(0, 1)(1, 0^*)$

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Monomer types:  $(1^*, 2, 2, 1^*)^+$   $(x, 0^*, 2^*, x)^-$   $(x, 2^*, 0, x)^-$

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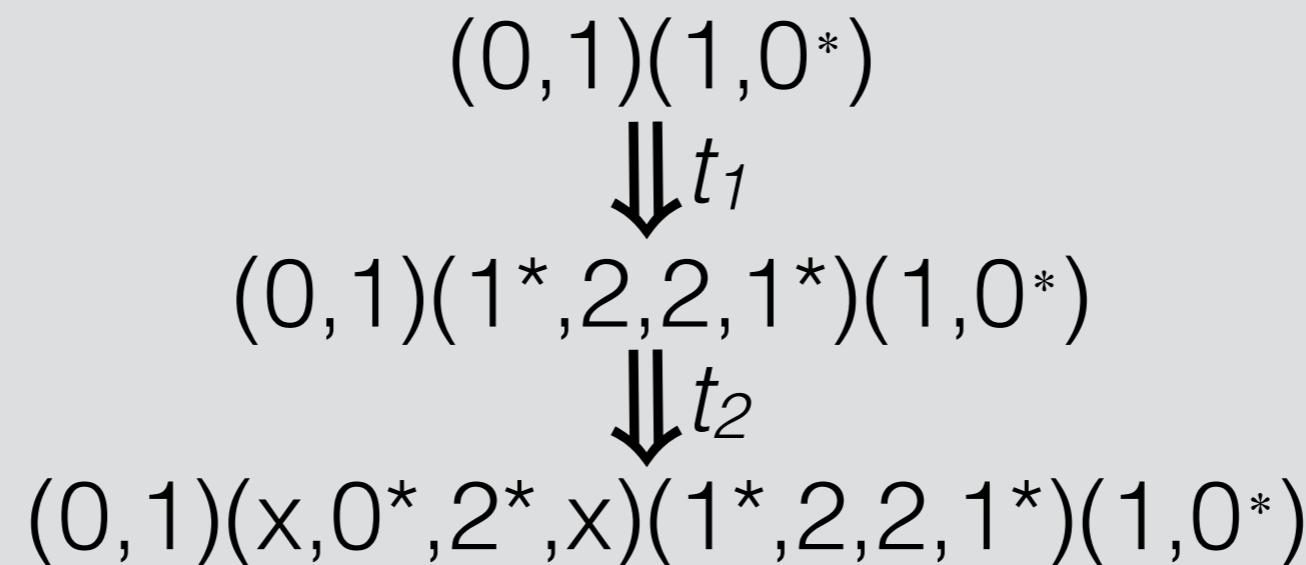


## Insertion system:

Monomer types:  $(1^*, 2, 2, 1^*)^+$     $(x, 0^*, 2^*, x)^-$     $(x, 2^*, 0, x)^-$

Concentrations:      0.5                          0.1                          0.4

Initiator:  $(0, 1)(1, 0^*)$

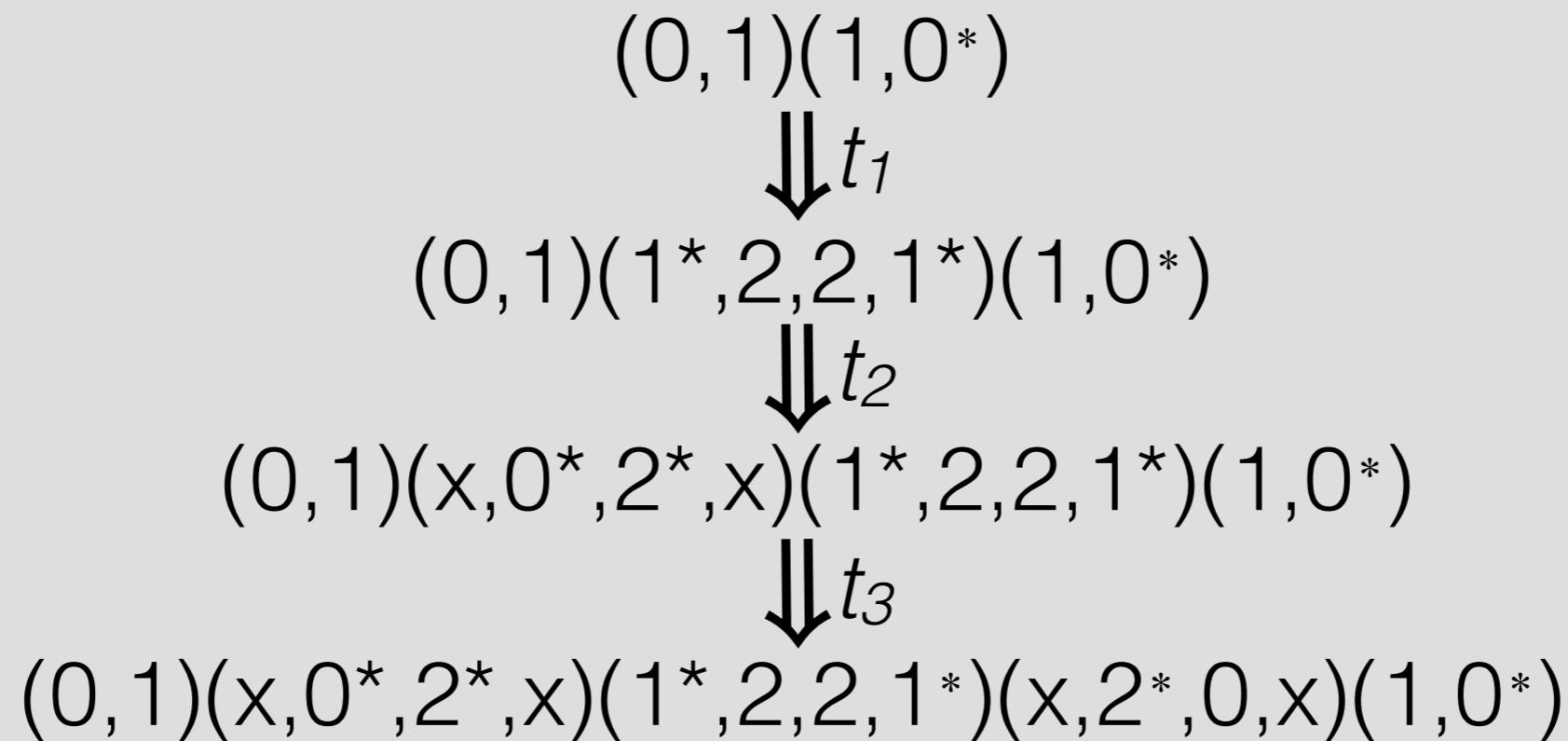


# Insertion system:

Monomer types:  $(1^*, 2, 2, 1^*)^+$     $(x, 0^*, 2^*, x)^-$     $(x, 2^*, 0, x)^-$

Concentrations:      0.5                          0.1                          0.4

Initiator:  $(0, 1)(1, 0^*)$

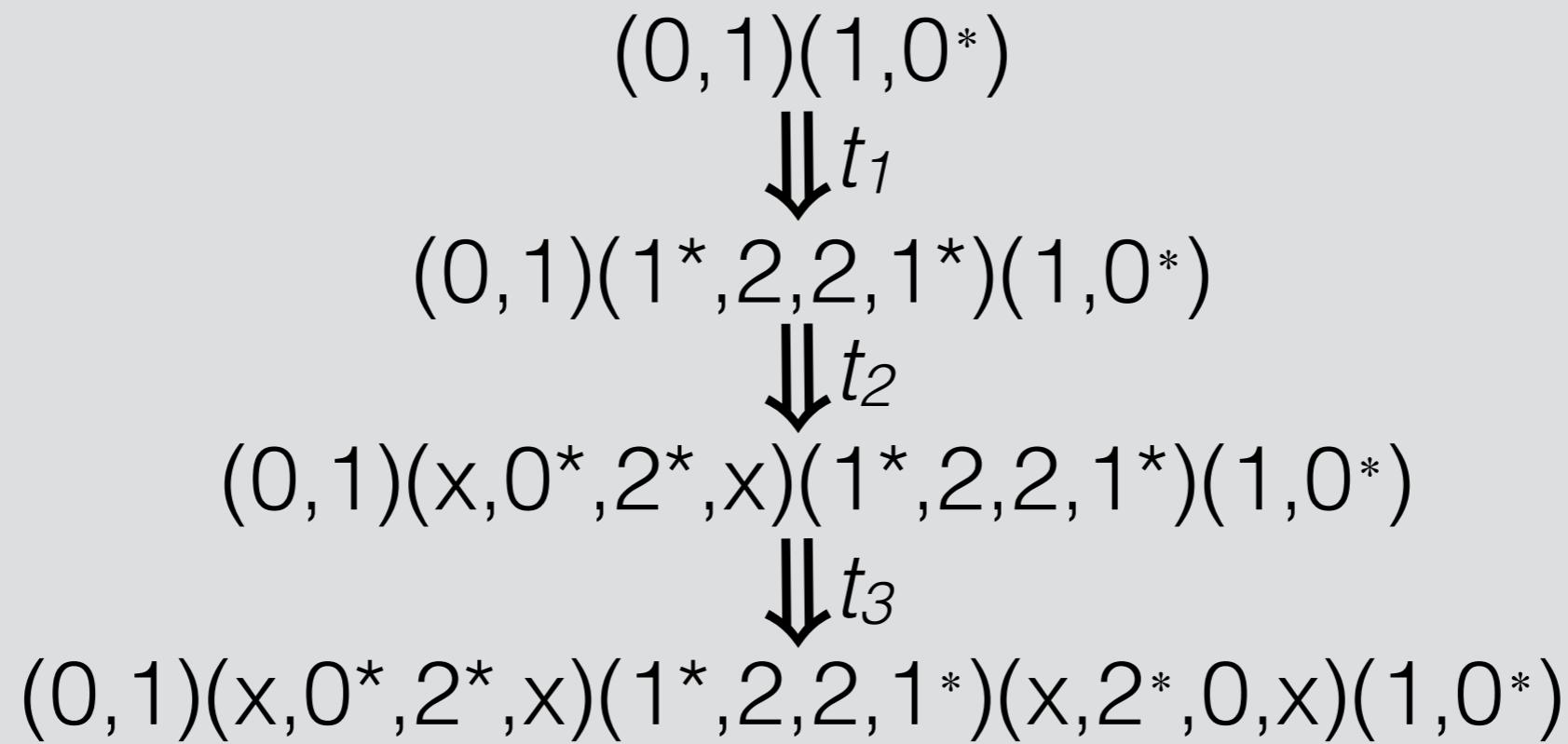


# Insertion system:

Monomer types:  $(1^*, 2, 2, 1^*)^+$     $(x, 0^*, 2^*, x)^-$     $(x, 2^*, 0, x)^-$

Concentrations:      0.5                          0.1                          0.4

Initiator:  $(0, 1)(1, 0^*)$



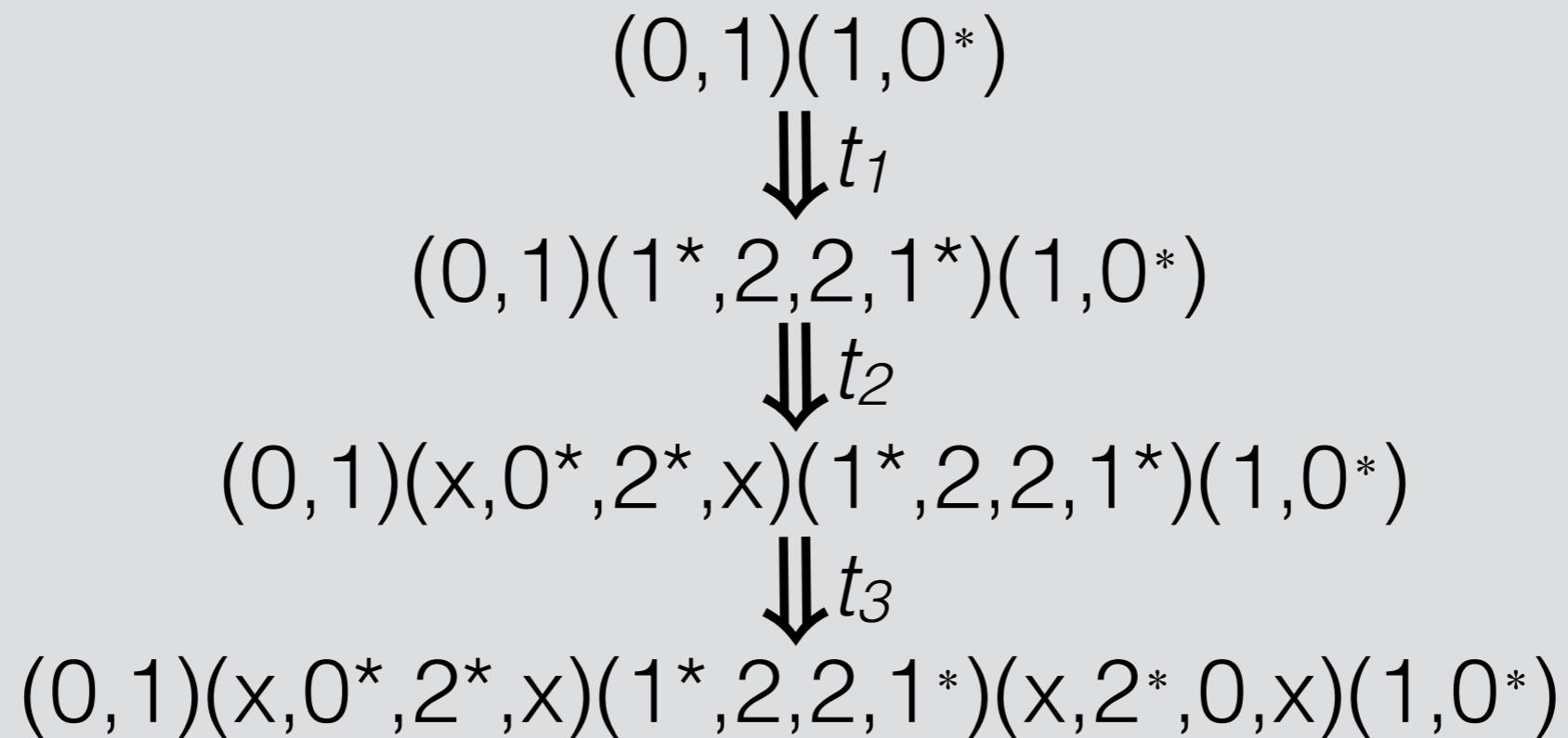
*Terminal polymer of length 5*

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Concentrations:      0.5                          0.1                          0.4

Initiator:  $(0, 1)(1, 0^*)$



*Terminal polymer of length 5*

*Expected time:  $t_1 + \max(t_2, t_3)$ , with*

$$E[t_1] = 2, E[t_2] = 10, E[t_3] = 2.5.$$
$$2 + 10.5 = 12.5$$

## Insertion system:

Monomer types:  $(1,x,2,b)^+$   $(x,2^*,a,1^*)^-$   $(x,2^*,a,x)^-$

Concentrations:      0.5                  0.4                  0.1

Initiator:  $(a,1^*)(b^*,a^*)$

---

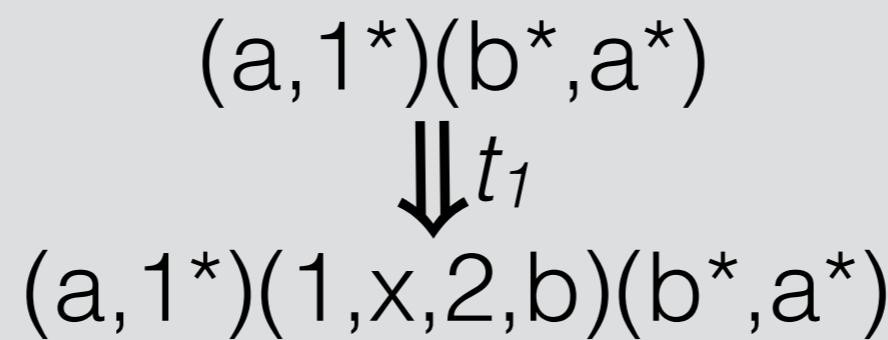
$(a,1^*)(b^*,a^*)$

## Insertion system:

Monomer types:  $(1,x,2,b)^+$   $(x,2^*,a,1^*)^-$   $(x,2^*,a,x)^-$

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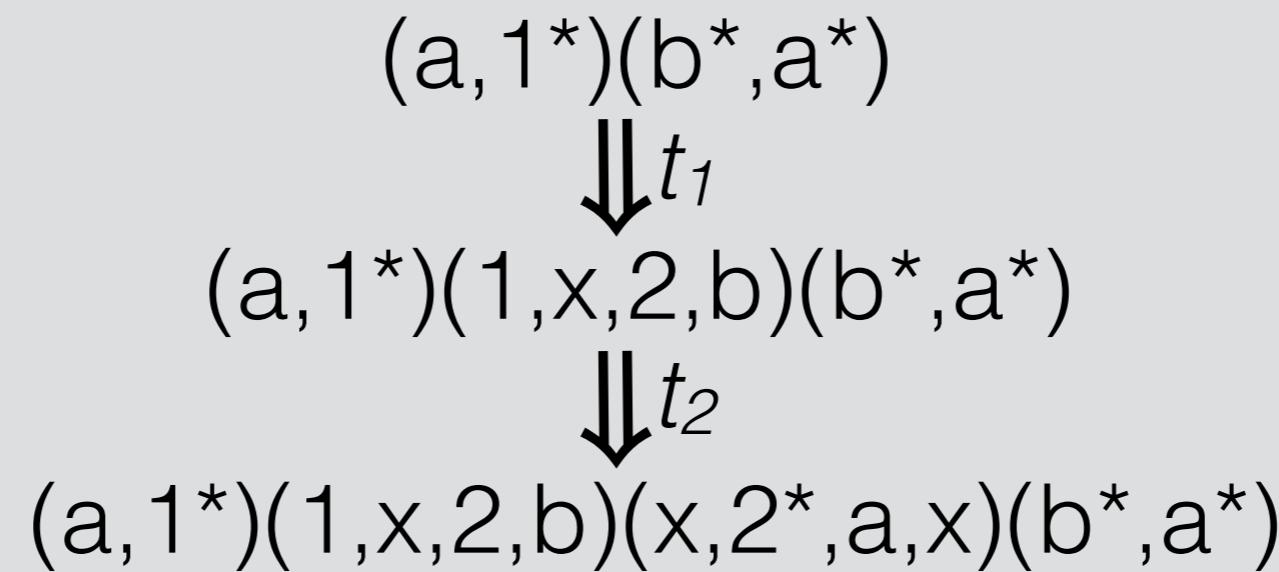


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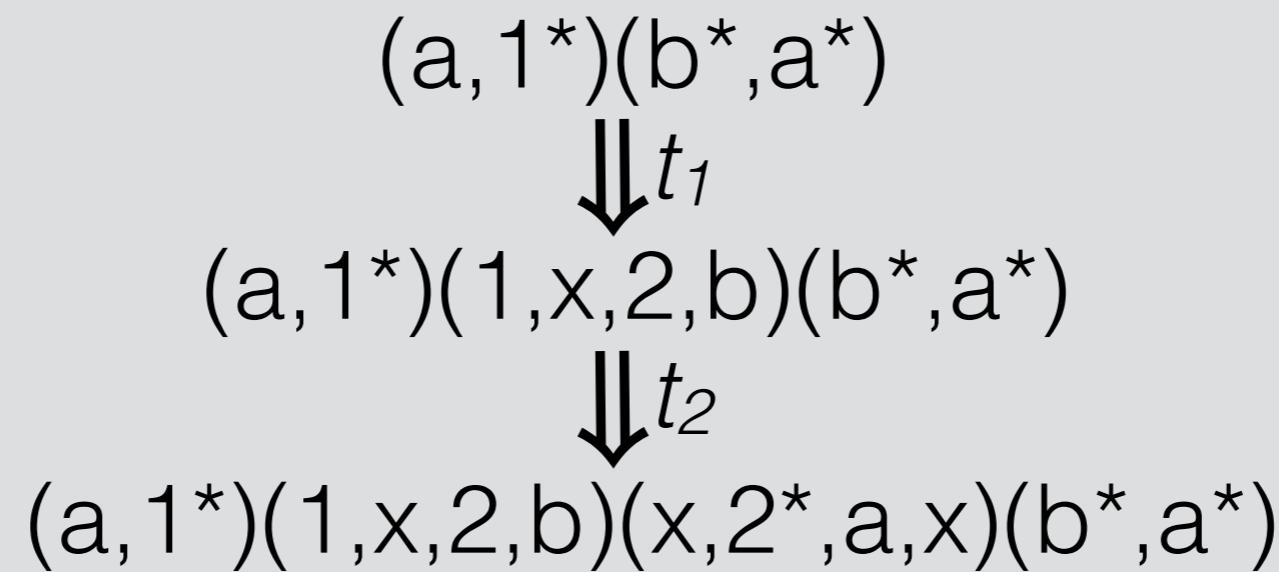


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Monomer types:  $(1,x,2,b)^+$   $(x,2^*,a,1^*)^-$   $(x,2^*,a,x)^-$

Concentrations: 0.5 0.4 0.1

Initiator:  $(a,1^*)(b^*,a^*)$



*Expected time:  $t_1 + t_2$ , with*

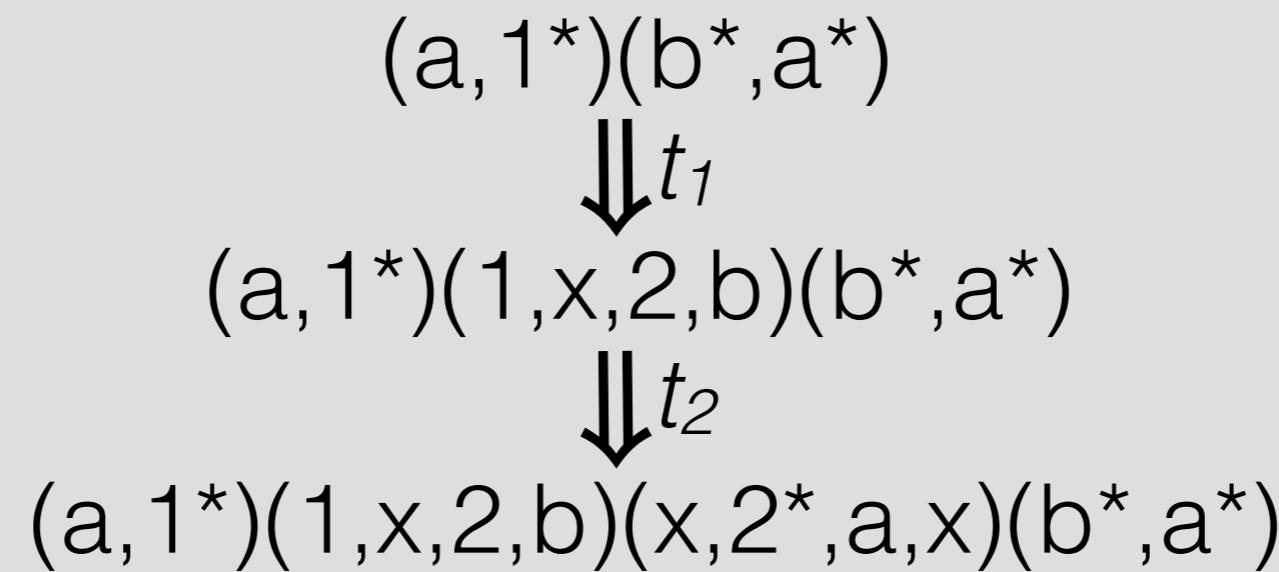
$$\begin{aligned} E[t_1] &= E[t_2] = 2. \\ 2 + 2 &= 4 \end{aligned}$$

## Insertion system:

Monomer types:  $(1,x,2,b)^+$   $(x,2^*,a,1^*)^-$   $(x,2^*,a,x)^-$

Concentrations: 0.5 0.4 0.1

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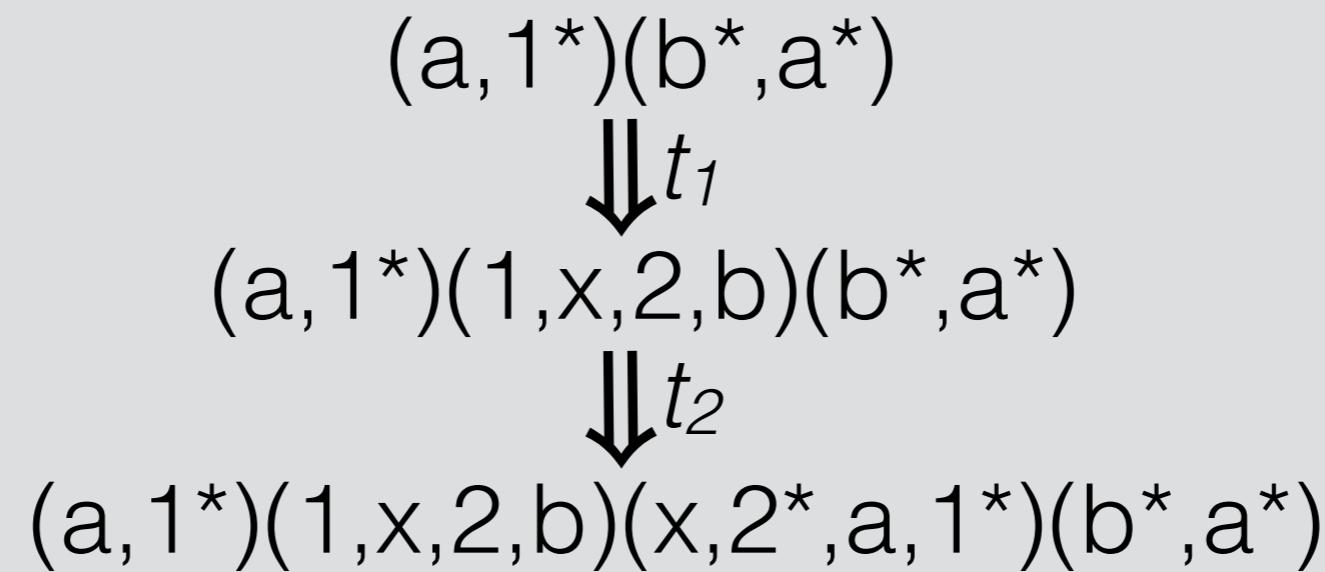


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Monomer types:  $(1,x,2,b)^+$   $(x,2^*,a,1^*)^-$   $(x,2^*,a,x)^-$

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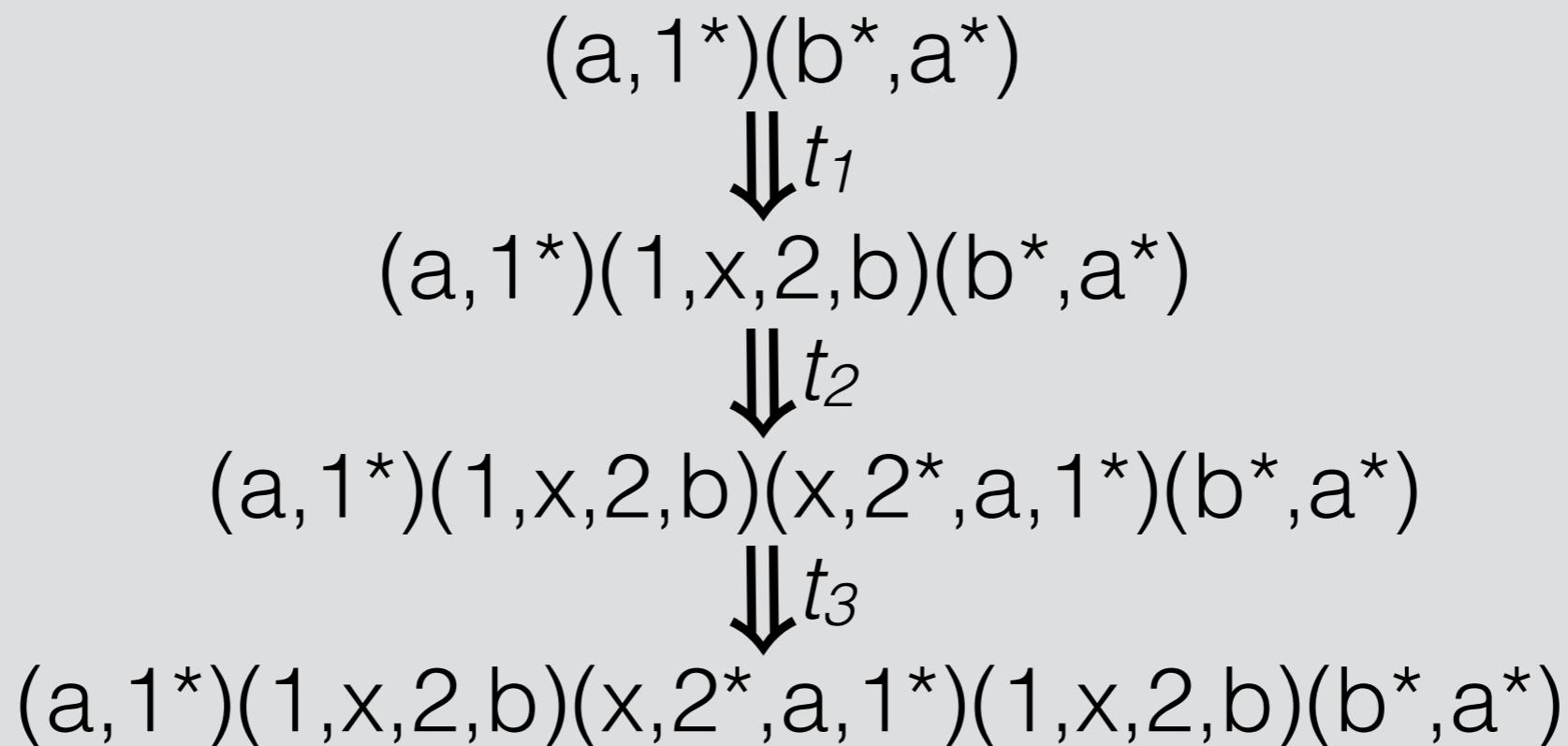


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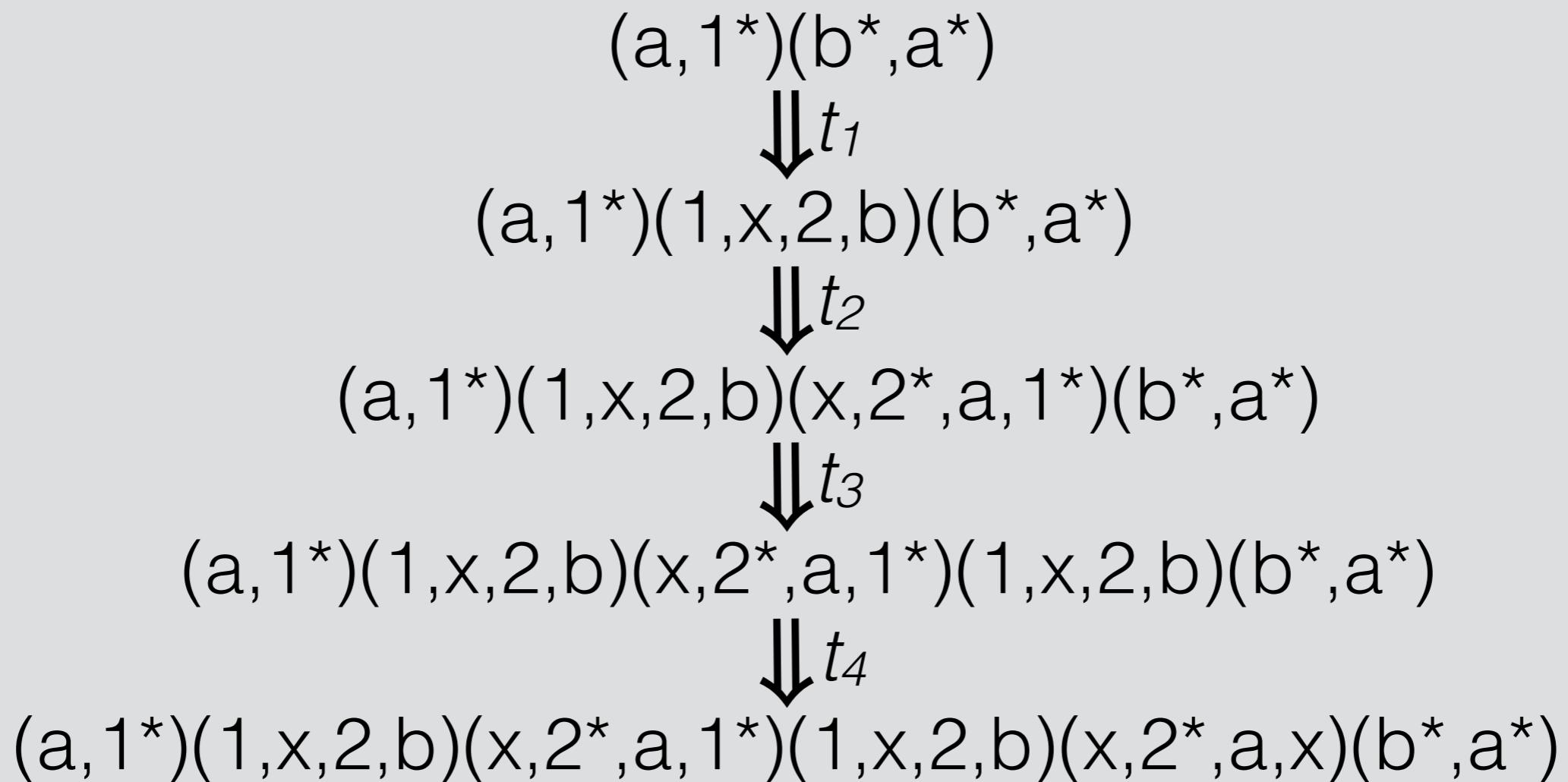


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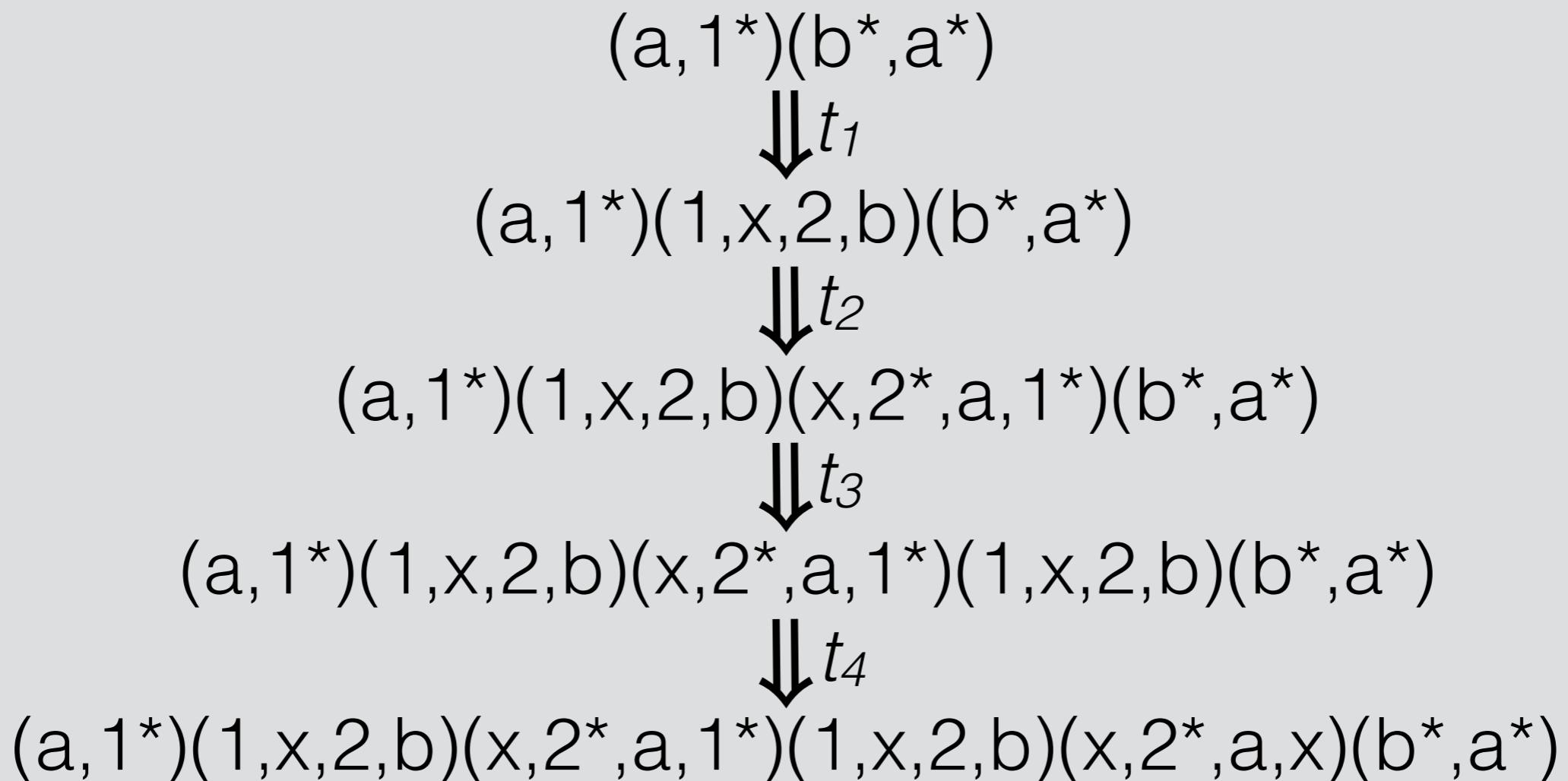


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Concentrations: 0.5 0.4 0.1

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*Expected time:  $t_1 + t_2 + t_3 + t_4$ , with*

$$E[t_1] = E[t_2] = E[t_3] = E[t_4] = 2.$$

$$2 + 2 + 2 + 2 = 8$$

# Expressive power

# Expressive Power

- Insertion systems: initiator + set of monomers = set of polymers, with terminal polymer subset.
- Context-free grammars: start symbol + set of rules = set of partial derivations, with string subset.
- Do insertion systems and context-free grammars have equal “expressive-ness”?

# Expressive Power

- Insertion systems: initiator + set of monomers = set of polymers, with terminal polymer subset.
- Context-free grammars: start symbol + set of rules = set of partial derivations, with string subset.
- Do insertion systems and context-free grammars have equal “expressive-ness”? Yes.

# Expressive Power

Theorem: every insertion system can be expressed as a context-free grammar. [Dabby, Chen 2013]

Theorem: every context-free grammar can be expressed as an insertion system. [This work]

# Expressive Power

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Theorem: every context-free grammar can be expressed as an insertion system. [This work]

## **Insertion system:**

Monomer types:  $(1,x,2,b)^+$   $(x,2^*,a,3)^-$   $(3,x,4,b)^+$

Concentrations: 0.25 0.25 0.25

Initiator:  $(a,1)(b^*,a^*)$

## **Context-free grammar:**

Production rules:

$$S_{(a,1)(b^*,a^*)} \rightarrow S_{(a,1)(1,x)} S_{(2,b)(b^*,a^*)}$$

$$S_{(2,b)(b^*,a^*)} \rightarrow S_{(2,b)(x,2^*)} S_{(a,3)(b^*,a^*)}$$

$$S_{(a,3)(b^*,a^*)} \rightarrow S_{(a,3)(3,x)} S_{(4,b)(b^*,a^*)}$$

$$S_{(a,1)(1,x)} \rightarrow (a, 1)(1, x)$$

$$S_{(2,b)(x,2^*)} \rightarrow (2,b)(x,2^*)$$

$$S_{(a,3)(3,x)} \rightarrow (a,3)(3,x)$$

$$S_{(4,b)(x,4^*)} \rightarrow (4,b)(x,4^*)$$

Start symbol:  $S_{(a,1)(b^*,a^*)}$

# Expressive Power

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# Turning rules into monomers

Rule:  $A \rightarrow BC$

Derivation step:  $cdADee \Rightarrow cdBCDee$

Monomer type:  $(1,x,2,b)^+$

Insertion:  $(a,1)(b^*,a^*) \Rightarrow (a,1)(1,x,2,b)(b^*,a^*)$

*Rules completely replace non-terminals.*

*Insertions do not completely replace insertion sites.*

# Turning rules into monomers

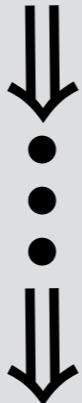
Rule:  $A_2 \rightarrow A_4 A_5$

Derivation step:  $A_1 A_2 A_3 \Rightarrow A_1 A_4 A_5 A_3$

(n non-terminals total)

Insertion site:  $u^*, s_a^*(s_d, u)$  with  $a+d \bmod n = 2$

Site modification:  $u^*, s_a^*(s_d, u)$



$u^*, s_a^*(s_b, u, \dots, u^*, s_c^*)(s_d, u)$

with  $a+b \bmod n = 4$ ,  $c+d \bmod n = 5$

Monomer types:  $\Theta(n)$  per rule.

# Polymer lengths

## Insertion system:

Monomer types:  $(1,x,2,b)^+$   $(x,2^*,a,1^*)^-$   $(x,2^*,a,x)^-$

Concentrations: 0.5 0.4 0.1

Initiator:  $(a,1^*)(b^*,a^*)$

---

$(a,1^*)(1,x,2,b)(x,2^*,a,x)(b^*,a^*)$

$(a,1^*)(1,x,2,b)(x,2^*,a,1^*)(1,x,2,b)(x,2^*,a,x)(b^*,a^*)$

$(a,1^*)(1,x,2,b)(x,2^*,a,1^*)(1,x,2,b)(x,2^*,a,1^*)(1,x,2,b)(x,2^*,a,x)(b^*,a^*)$

• • •

---

## Context-free grammar:

Production rules:  $S \rightarrow A$   $A \rightarrow aaA$   $A \rightarrow aa$

Start symbol: S

aa  
aaaa  
aaaaaaaa  
• • •

## Insertion system:

Monomer types:  $(1,x,2,b)^+$   $(x,2^*,a,1^*)^-$   $(x,2^*,a,x)^-$

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Initiator:  $(a,1^*)(b^*,a^*)$

$(a,1^*)(1,x,2,b)(x,2^*,a,x)(b^*,a^*)$

$(a,1^*)(1,x,2,b)(x,2^*,a,1^*)(1,x,2,b)(x,2^*,a,x)(b^*,a^*)$

Constructing arbitrarily long polymers is easy  
if infinite polymers allowed.

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Start symbol: S

aa  
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•••

# Constructing long polymers

Theorem: a system with  $k$  monomer types constructing a finite number of polymers can construct:

- polymers of length  $2^{\Theta(k^{1/2})}$  [Dabby, Chen 2013]
- polymers of length  $2^{\Theta(k^{3/2})}$  [This work]
- only polymers of length  $2^{O(k^{3/2})}$  [This work]

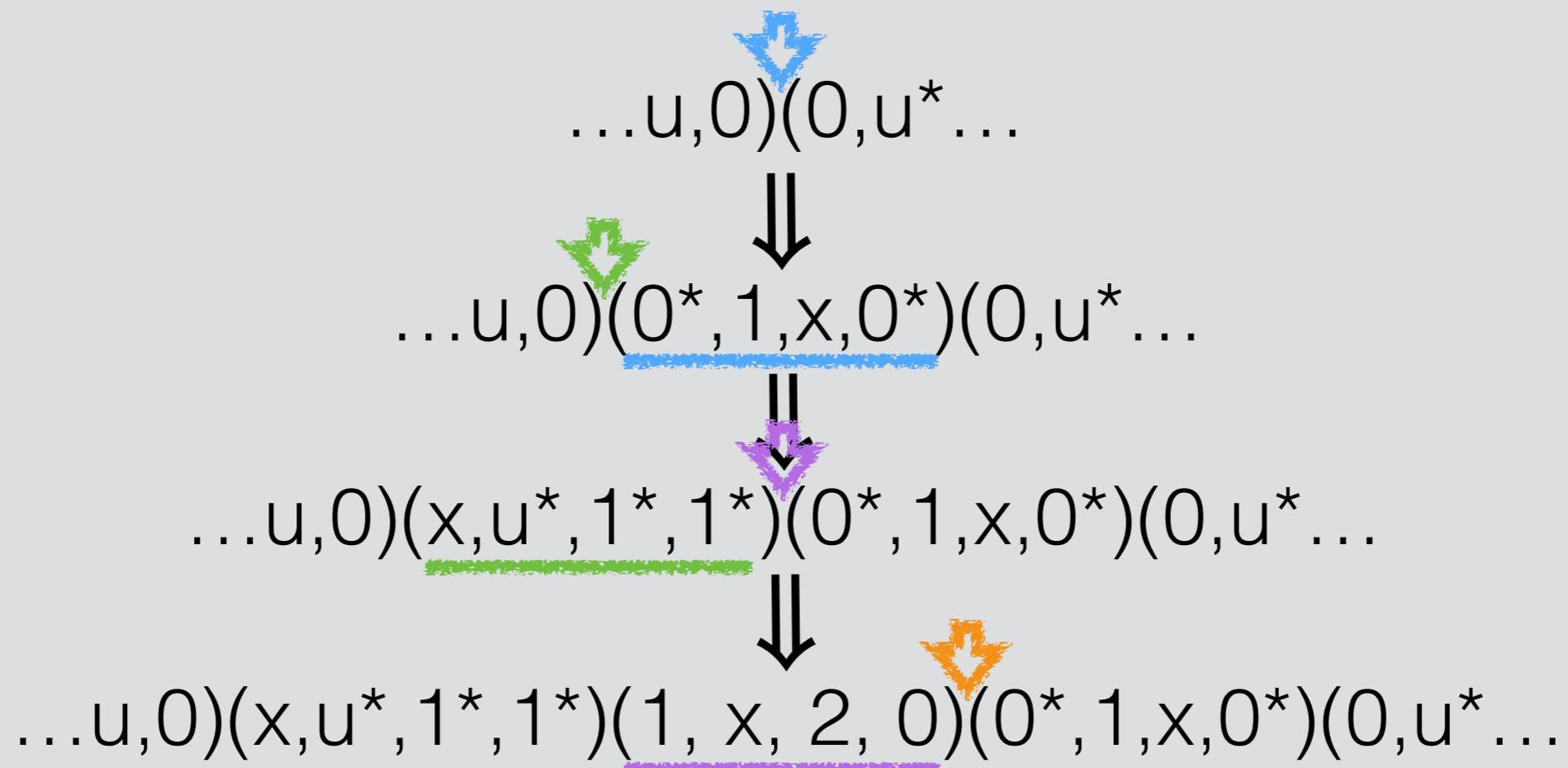
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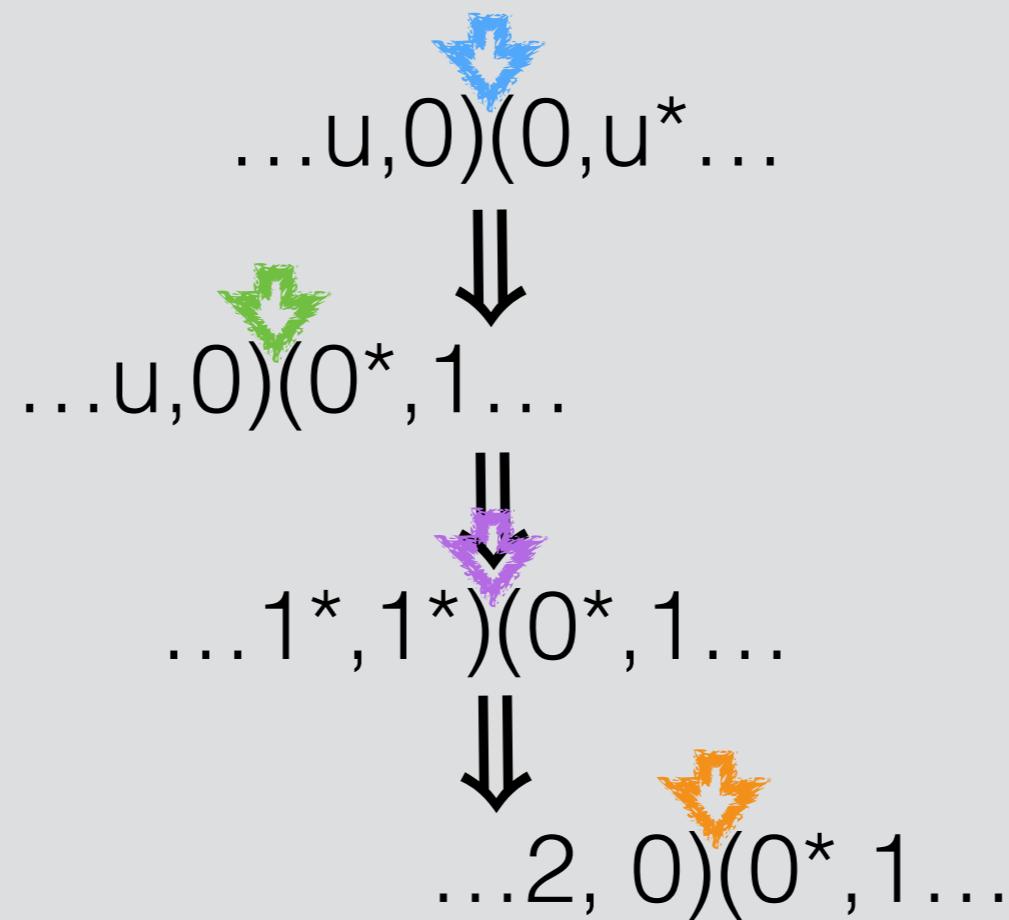
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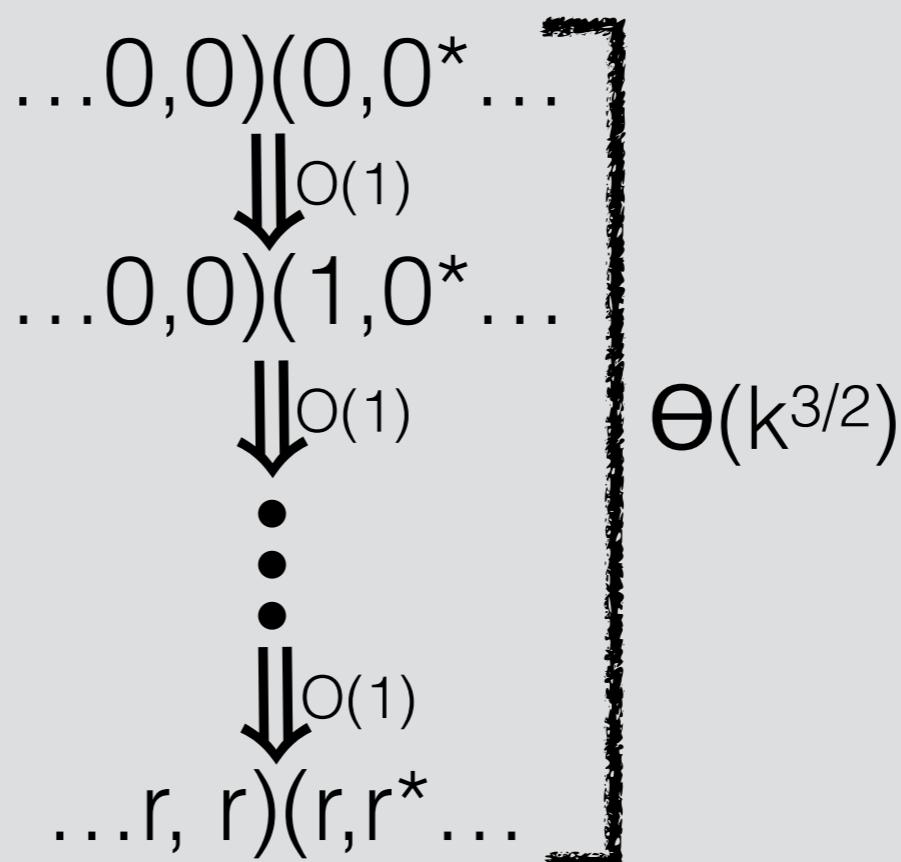
# Constructing long polymers

- Consider *insertion sequences*: repeated insertions into the site resulting from previous insertion.

$$\begin{array}{c} \dots u, 0)(0, u^* \dots \\ \Downarrow \\ \dots u, 0)(0^*, 1 \dots \\ \Downarrow \\ \dots 1^*, 1^*)(0^*, 1 \dots \\ \Downarrow \\ \dots 2, 0)(0^*, 1 \dots \end{array}$$

# Constructing long polymers

- Consider *insertion sequences*: repeated insertions into the site resulting from previous insertion.
- Ingredient 1: long insertion sequence with no repeated insertion sites.



# Constructing long polymers

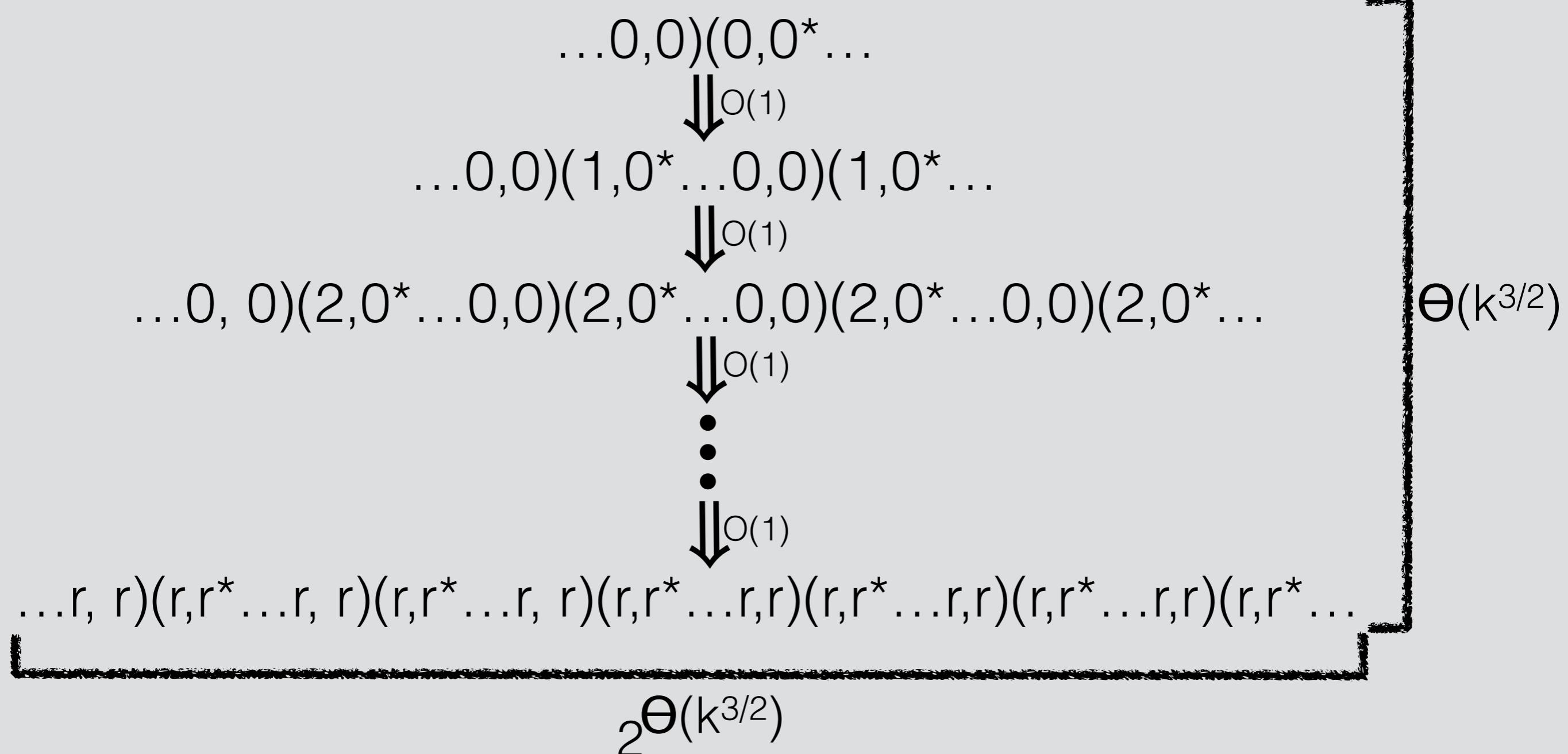
- Consider *insertion sequences*: repeated insertions into the site resulting from previous insertion.
- Ingredient 1: long insertion sequence with no repeated insertion sites.
- Ingredient 2: duplication of each site in sequence.

$$\begin{array}{c} \dots 0,0)(0,0^* \dots \\ \downarrow^{O(1)} \\ \dots 0,0)(1,0^* \dots 0,0)(1,0^* \dots \\ \downarrow^{O(1)} \\ \dots 0,0)(2,0^* \dots 0,0)(2,0^* \dots 0,0)(2,0^* \dots \end{array}$$

# Constructing long polymers

- Consider *insertion sequences*: repeated insertions into the site resulting from previous insertion.
- Ingredient 1: long insertion sequence with no repeated insertion sites.
- Ingredient 2: duplication of each site in sequence.
- Combine these for long polymers.

# Constructing long polymers



# Constructing long polymers

- Ingredient 1: long insertion sequence with no repeated insertion sites.

$S_a, S_b)(S_c, S_a^*$   
variables

Use  $r+1 = \Theta(k^{1/2})$  values of  $a, b, c \Rightarrow \Theta(r^3) = \Theta(k^{3/2})$  insertion sites.

Case:	$b < r$	$b = r, c < r$	$b = c = r, a < r$
Insertion sequence:	$S_a, S_b)(S_c, S_a^*$ $\Downarrow_{O(1)}$ $S_a, S_{b+1})(S_c, S_a^*$	$S_a, S_r)(S_c, S_a^*$ $\Downarrow_{O(1)}$ $S_a, S_0)(S_{c+1}, S_a^*$	$S_a, S_r)(S_r, S_a^*$ $\Downarrow_{O(1)}$ $S_{a+1}, S_0)(S_0, S_{a+1}^*$
Result:	$++b$	$b = 0, ++c$	$b = c = 0, ++a$

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# Constructing long polymers

$\dots S_a, S_r)(S_c, S_a^* \dots$

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$\dots S_a, S_0^*)(S_{c+1}^*, S_a^* \dots$

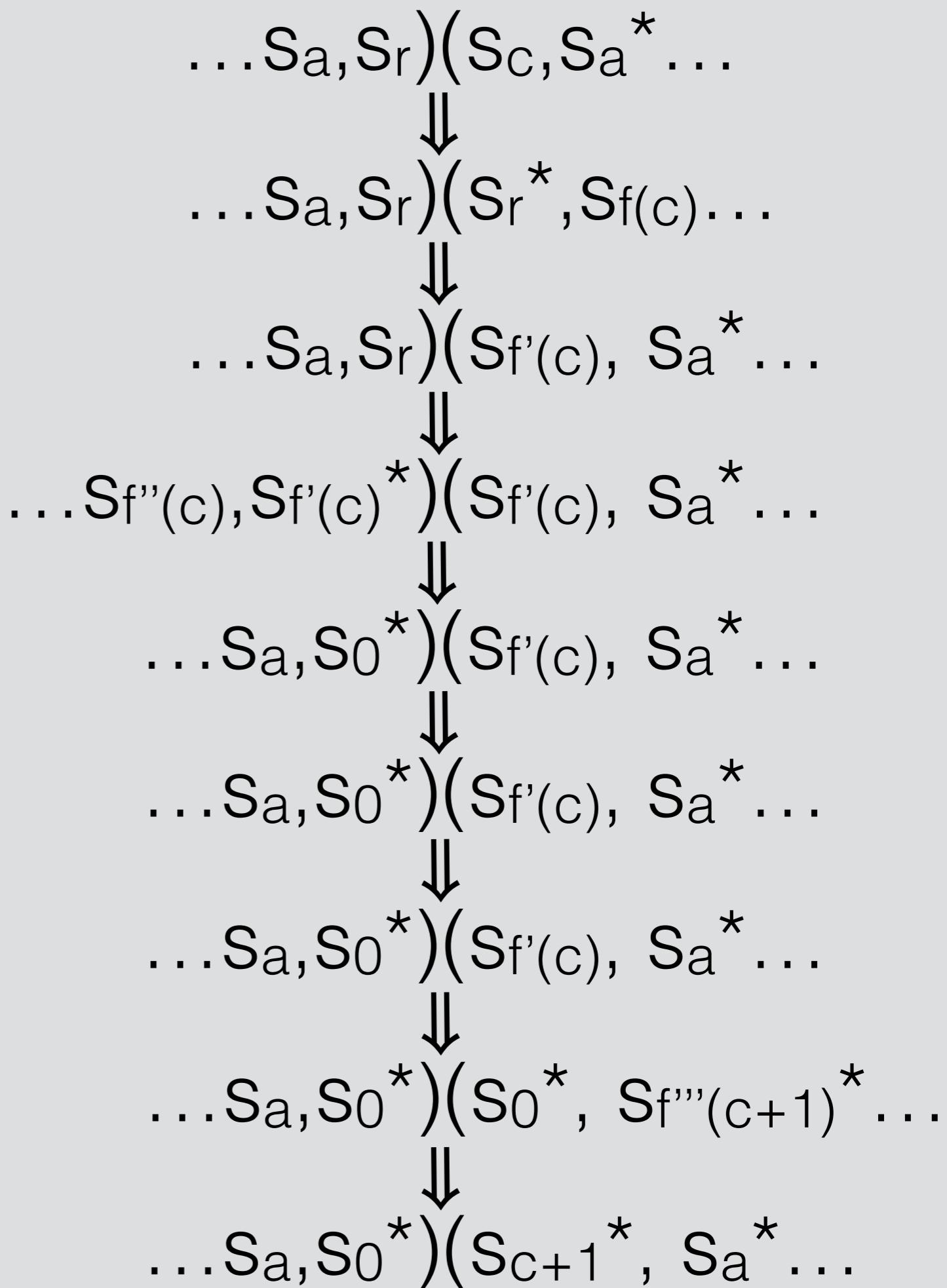
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# Constructing long polymers

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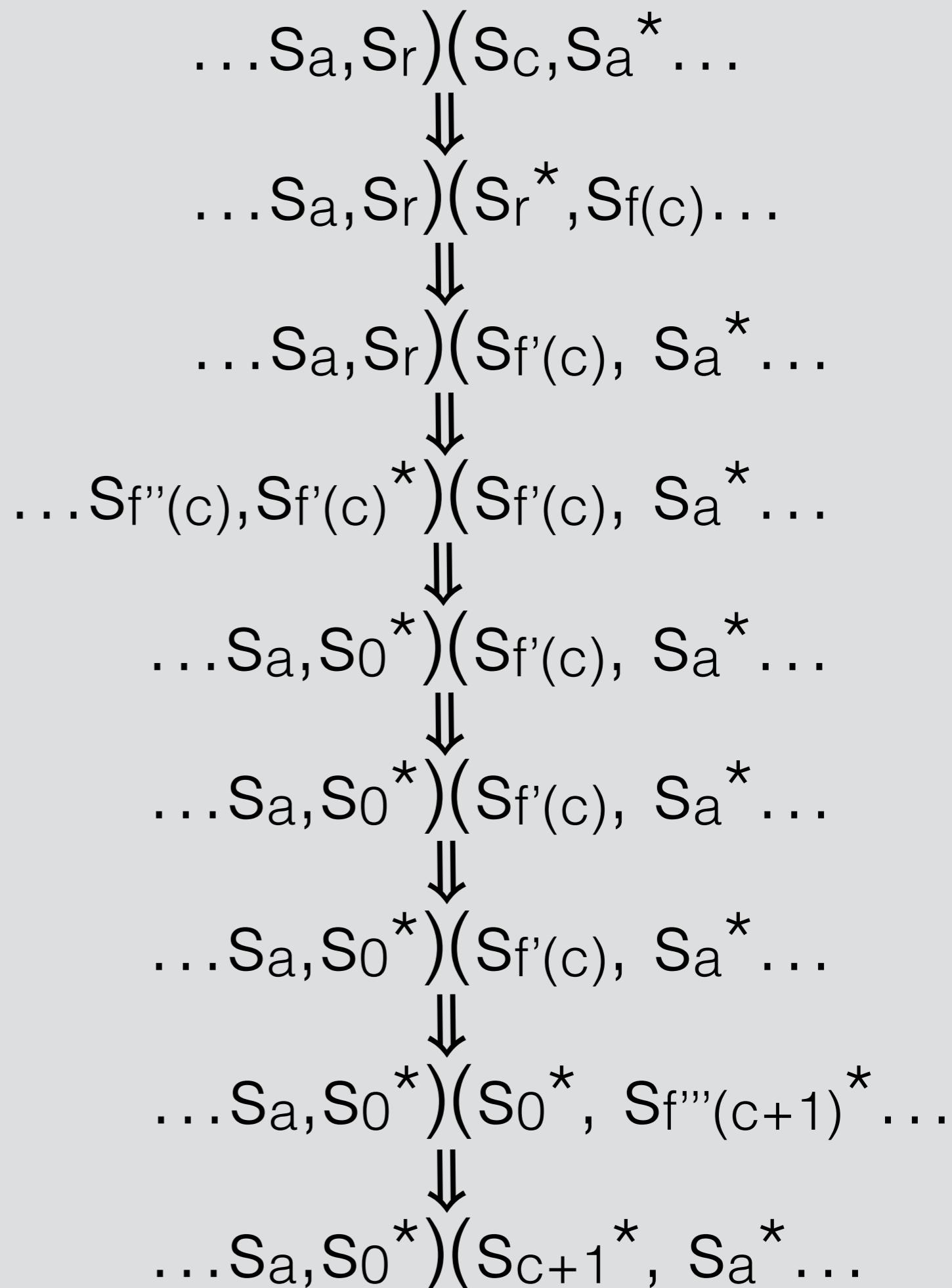
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Result:  $b = 0, ++c$

This is the easiest case.

A total of 23 monomer type families used.



# An upper bound for polymer length

- Every insertion sequence in polymer must consist of unique insertion sites  $a,b)(c,a^*$  that each accept a monomer.
- Prove there are  $O(k^{3/2})$  such sites.
- Proof idea: maximize  $|\{a\}|^*|\{b\}|^*|\{c\}|$  subject to
  - $|\{a\}|^*|\{b\}|, |\{a\}|^*|\{c\}| = O(k)$  since they are monomer right/left halves.
  - $|\{b\}|^*|\{c\}| = O(k)$  since site needs  $(b^*, \_, \_, c^*)^+$  inserted.
- $|\{a\}|^*|\{b\}|^*|\{c\}| = O(k^{3/2})$

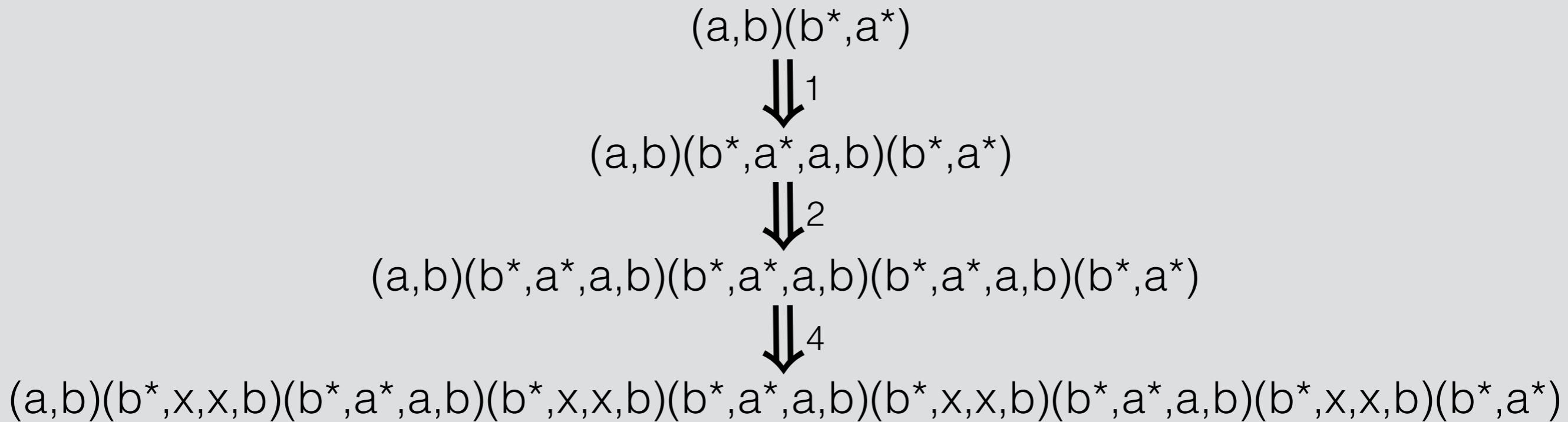
# Polymer growth speed

# Insertion system:

Monomer types:  $(b^*, a^*, a, b)^+$   $(b^*, x, x, b)^+$

Concentrations: 0.5 0.5

Initiator:  $(a, b)(b^*, a^*)$



*Each round of insertions takes  $O(1)$  expected time.*

*Construction of length  $n = 2^i - 1$  takes  $O(i)$  expected time.  
 $O(\log(n))$*

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$(a, b)(b^*, a^*)$   
 $\downarrow_1$   
Constructing polymers in  $O(\log(n))$   
expected time is easy  
if infinite polymers allowed.

$(a, b)(b^*, x, x, b)(b^*, a^*, a, b)(b^*, x, x, b)(b^*, a^*, a, b)(b^*, x, x, b)(b^*, a^*, a, b)(b^*, x, x, b)(b^*, a^*)$

*Each round of insertions takes  $O(1)$  expected time.*

*Construction of length  $n = 2^i - 1$  takes  $O(i)$  expected time.  
 $O(\log(n))$*

# Constructing polymers quickly

Theorem: a system constructing a finite number of polymers can **deterministically** construct a polymer of length  $n$  in:

- $O(\log^3(n))$  expected time [Dabby, Chen 2013]
- $O(\log^{5/3}(n))$  expected time [This work]
- only  $\Omega(\log^{5/3}(n))$  expected time [This work]

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# A lower bound for growth speed

- Consider longest insertion sequence carried out; must have length at least  $\log_2(n)$ .
- By polymer length upper bound, must insert  $\Omega(\log^{2/3}(n))$  different monomer types.
- Optimization problem: pick #insertions, concentrations of  $\log^{2/3}(n)$  monomer types into  $\log_2(n)$  sites to minimize total expected time.
- Assuming each site accepts 1 monomer type, minimum expected time is  $\Omega(\log^{5/3}(n))$ .

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System is *deterministic*.

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Theorem: a system constructing a finite number of polymers can **non-deterministically** construct a polymer of length  $n$  in:

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# Speedup by non-determinism

$$\dots S_1, S_0) (S_0^*, S_1^* \dots$$

$$\dots S_1, S_1) (S_1^*, S_1^* \dots$$

*Needs unique monomers.*

*Unique monomers  $\Rightarrow$  slow.*

# Speedup by non-determinism

$$\dots S_6, S_3) (S_3^*, S_6^* \dots$$

$$\dots S_6, S_4) (S_4^*, S_6^* \dots$$

*Needs unique monomers.*

*Unique monomers  $\Rightarrow$  slow.*

# Speedup by non-determinism

$\dots s_6, s_3)(s_3^*, s_6^* \dots$

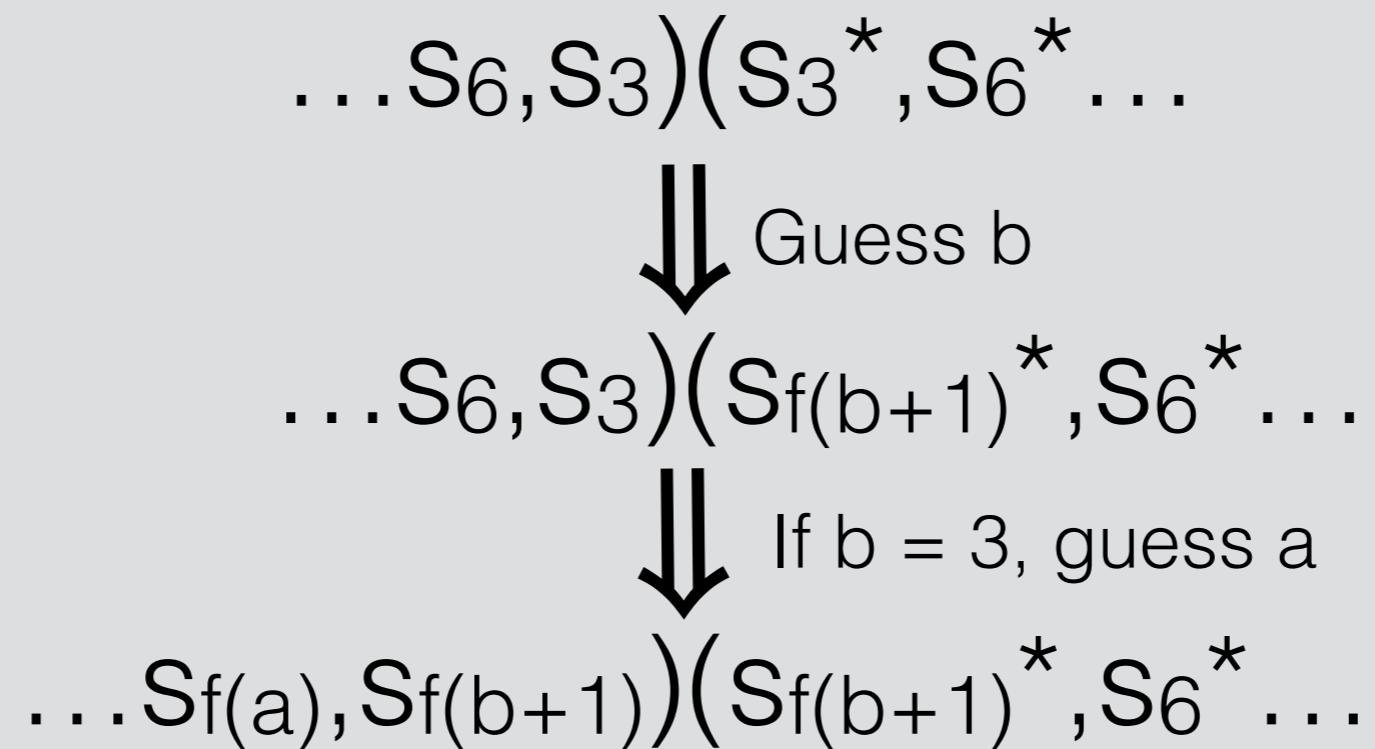
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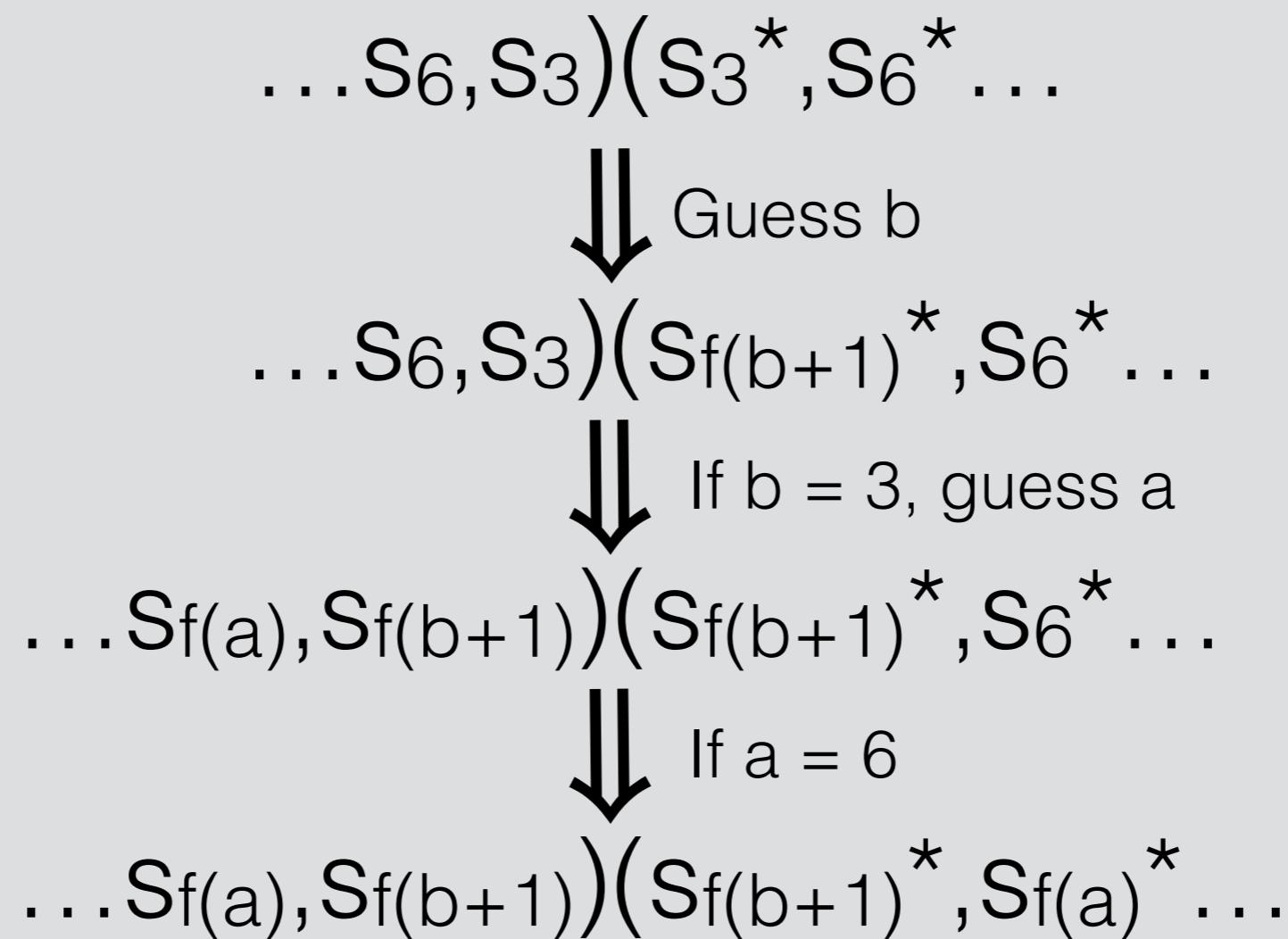
$\downarrow$  Guess b

$\dots s_6, s_3)(s_{f(b+1)}^*, s_6^* \dots$

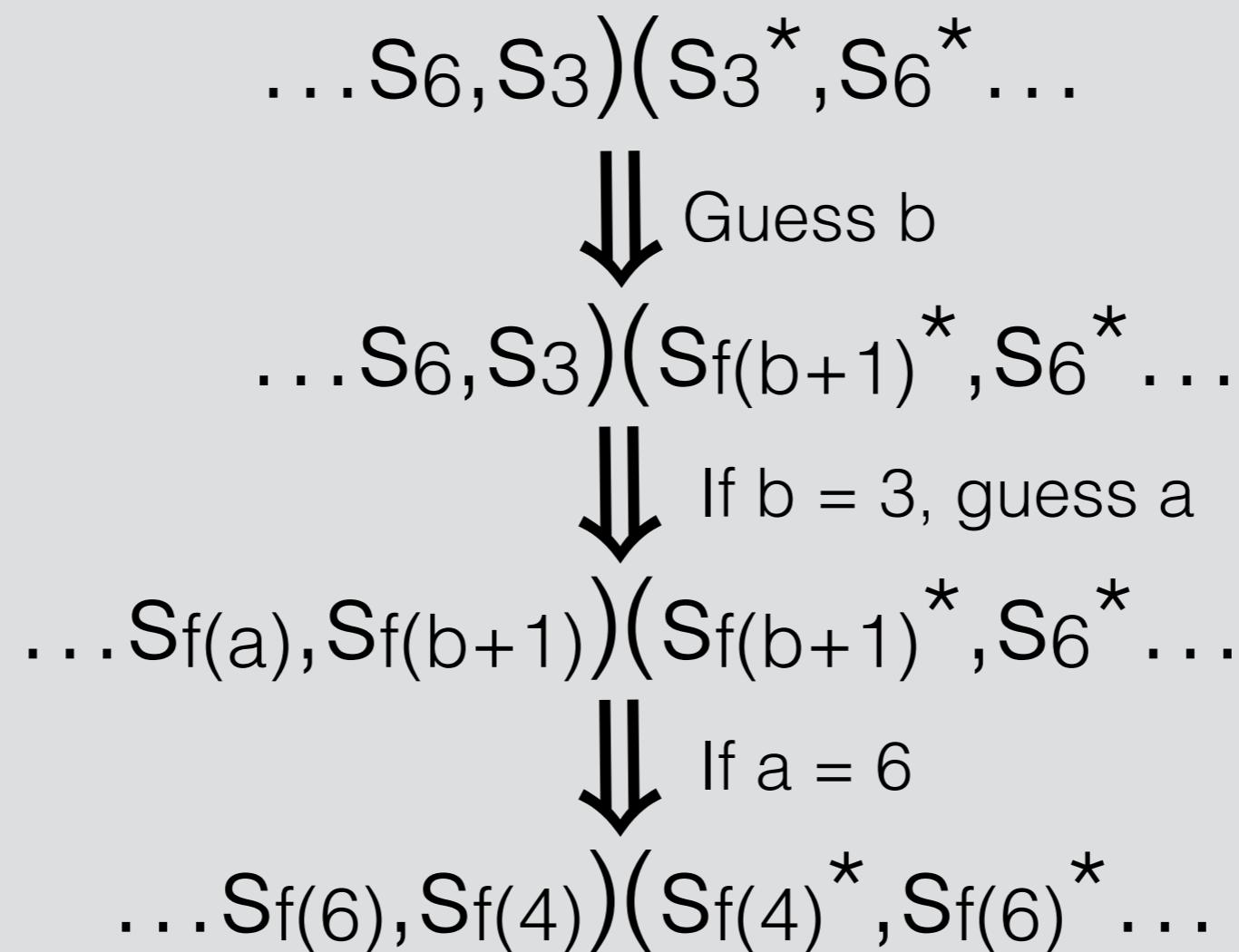
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$\downarrow$  Guess b

$\dots s_6, s_3)(s_{f(b+1)}^*, s_6^* \dots$

$\downarrow$  If  $b \neq 3$ , insertion sequence done.  
 $\times$

# Speedup by non-determinism

$\dots s_6, s_3)(s_3^*, s_6^* \dots$

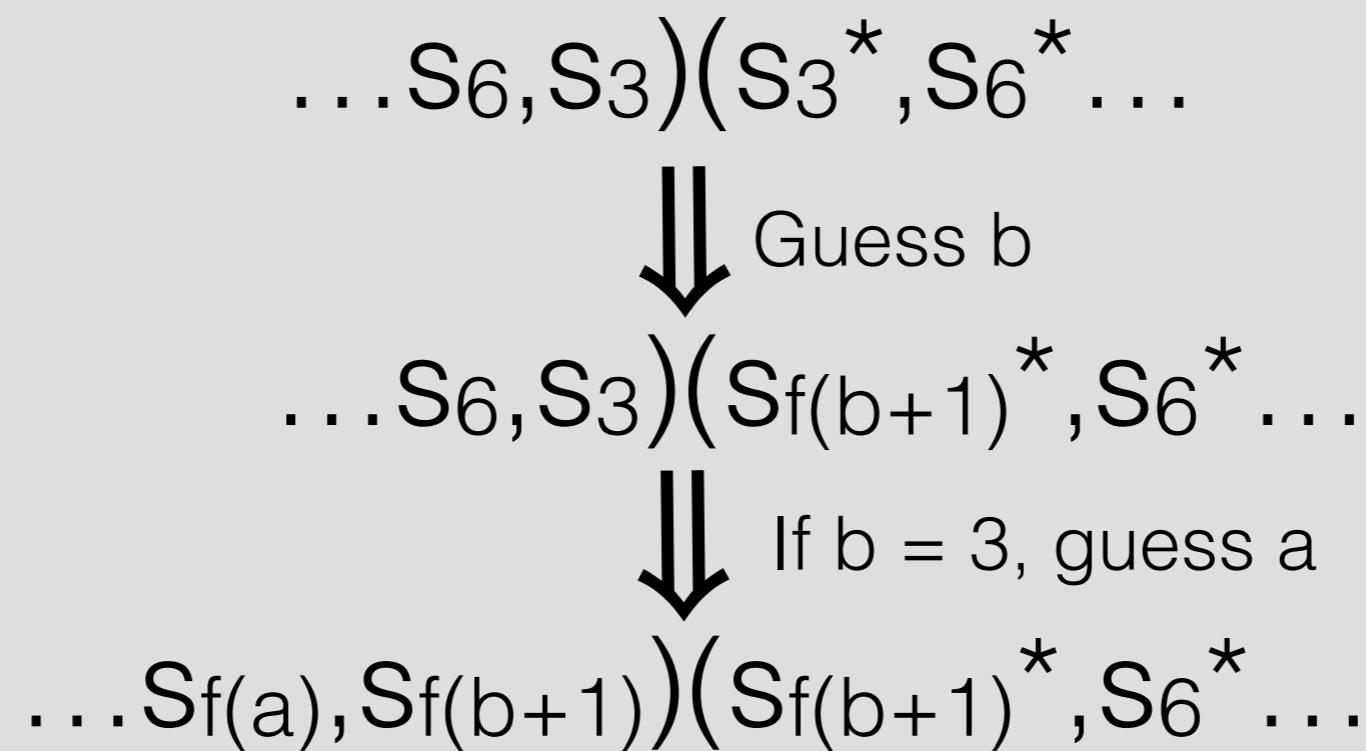
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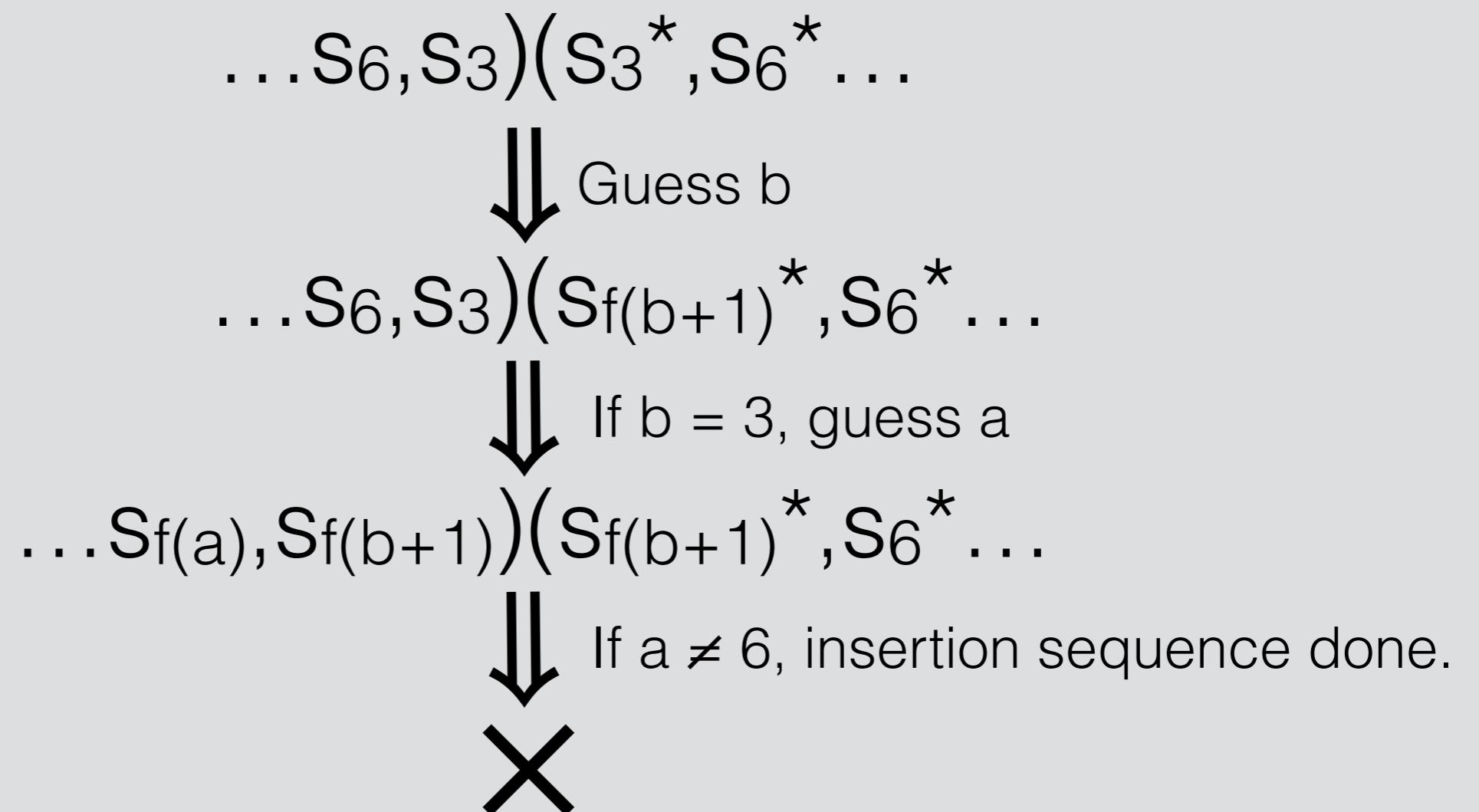
$\downarrow$  Guess b

$\dots s_6, s_3)(s_{f(b+1)}^*, s_6^* \dots$

# Speedup by non-determinism

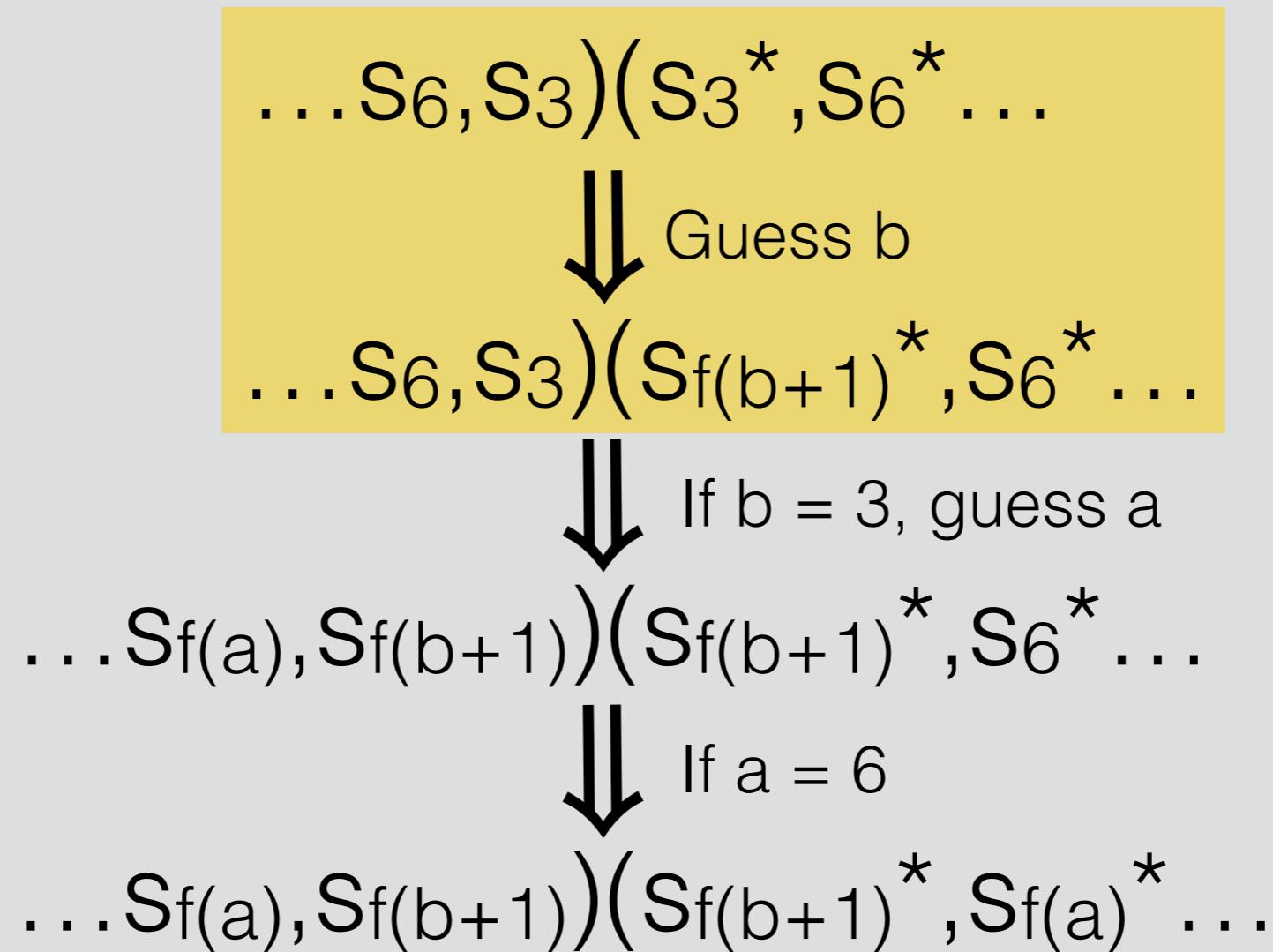


# Speedup by non-determinism



Why does guessing  
give a speed-up?

# Speedup by non-determinism



# Speedup by non-determinism

$$S_6, S_3)(S_3^*, S_6^*$$
  
 $\Downarrow \text{Guess } b$   
 $S_6, S_3)(S_{f(b+1)}^*, S_6^*$

$$S_6, S_5)(S_5^*, S_6^*$$
  
 $\Downarrow \text{Guess } b$   
 $S_6, S_5)(S_{f(b+1)}^*, S_6^*$

$$S_6, S_8)(S_8^*, S_6^*$$
  
 $\Downarrow \text{Guess } b$   
 $S_6, S_8)(S_{f(b+1)}^*, S_6^*$

$$S_6, S_{11})(S_{11}^*, S_6^*$$
  
 $\Downarrow \text{Guess } b$   
 $S_6, S_{11})(S_{f(b+1)}^*, S_6^*$

# Speedup by non-determinism

$S_6, S_3)(S_3^*, S_6^*$

↓ Guess b

$S_6, S_3)(S_{f(b+1)}^*, S_6^*$        $S_6, S_5)(S_5^*, S_6^*$   
The same set of monomers are  
insertable to *all* of these sites.

$S_6, S_8)(S_8^*, S_6^*$

↓ Guess b

$S_6, S_8)(S_{f(b+1)}^*, S_6^*$

$S_6, S_{11})(S_{11}^*, S_6^*$

↓ Guess b

$S_6, S_{11})(S_{f(b+1)}^*, S_6^*$

# Non-determinism speed-up

- Sites  $s_a, s_b)(s_b^*, s_a^*$  with  $\Theta(k^{1/2})$  values for a, b
  - So  $k = \Theta(\log(n))$
- Every site accepts  $\Theta(k^{1/2})$  monomers:  
 $\Omega(1/k^{1/2})$  total concentration.
- Expected insertion time is  $O(k^{1/2}) = O(\log^{1/2}(n))$ .
- Total expected construction time is  $O(\log^{3/2}(n))$ .

# Polymer length vs. growth speed

# Types vs Speed Tradeoff

Is it possible to construct a polymer using  $O(\log^{2/3}(n))$  monomer types in  $O(\log^{3/2}(n))$  expected time?

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Theorem: in a system constructing a finite number of polymers, constructing a polymer of length  $n$  using  $k$  monomer types takes  $\Omega(\log^2(n)/k^{1/2})$  expected time.

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$O(\log^{2/3}(n))$  types  $\Rightarrow \Omega(\log^{5/3}(n))$  expected time

$O(\log^{3/2}(n))$  expected time  $\Rightarrow \Omega(\log(n))$  types

# Conclusions

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- Insertion systems and context-free grammars are computationally equivalent.
- Growing finitely takes more monomer types and more time than infinite growth.
- Growing the longest finite polymers possible requires use of novel properties — cannot use CFG equivalence!
- Growing finite polymers fastest needs non-determinism and more monomer types.
- A smooth tradeoff between polymer length and growth rate.

# Thank you.

Joint work with Benjamin Hescott and Caleb Malchik.

Some results published in:

C. Malchik, A. Winslow, Tight bounds for active self-assembly with an insertion primitive, Proceedings of 22nd European Symposium on Algorithms, 677-688, 2014.