

# On the Collective Behaviors of Bio-Inspired Distributed Systems



## Jury

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PhD defense

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- Distributed computing tasks in biological systems

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- The consensus problem with noisy communications
- The UNDECIDED-STATE and the 3-MAJORITY dynamics: contribution

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- Mentions to model and contribution

## 5. Conclusions & perspectives

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# Natural algorithms

Algorithms designed by **evolution** over millions of years [Chazelle, SODA 2009]

- migrating geese
- flocking cranes
- fish bait ball
- prey-predator systems
- synchronously flashing fireflies



flocking cranes



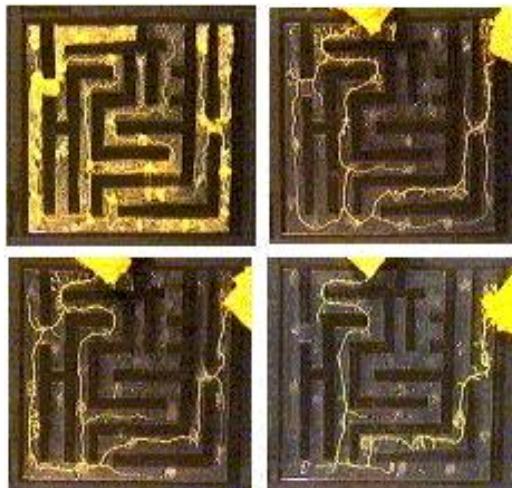
fish bait ball

"Georgia Aquarium Fish" by Mike Johnston

# Natural algorithms

The **computational lens** help catching **behavioral properties** of biological systems

- bird flocking convergence time [Chazelle, SODA 2009]
- slime mold computing shortest paths [Bonifaci et al., Journal of Theoretical Biology 2012]



**Slime mold**

John S. Macneil, Science 2000

5 - 1



**Flocking birds**

"Flocking birds" by davepatten, CC BY-NC-SA 2.0.

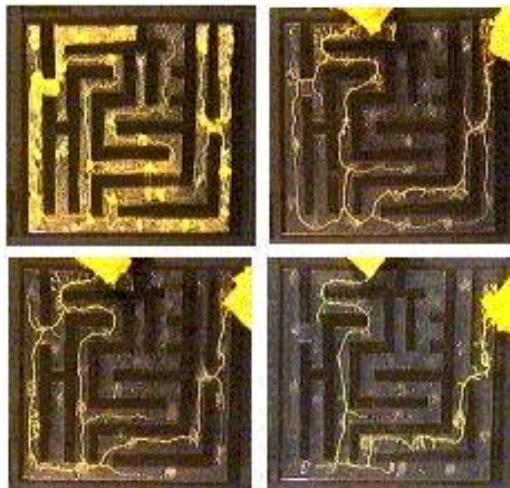
# Natural algorithms

The **computational lens** help catching **behavioral properties** of biological systems

- bird flocking convergence time [Chazelle, SODA 2009]
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On the other hand, biological systems help **designing** new **algorithms** for well-known problems

- distributed **maximal independent set** algorithm from the fly's nervous system [Afek et al., SCIENCE 2011]



Slime mold

John S. Macneil, Science 2000



Flocking birds

"Flocking birds" by davepatten, CC BY-NC-SA 2.0.

# Distributed computing tasks

Often, systems of **interacting agents** performing **collective tasks**

Other than MIS and bird flocking:

- Information spreading: schooling fish [Rosenthal et al., PNAS 2015]
- Reaching agreement: molecules [Carrol, Nature Immunology 2004], bacteria [Bassler, Cell 2002], social insects [Franks et al., 2002] (e.g. bees [Reina et al., Physical Review E 2017])
- Collective search: ants and bees [Feinerman et Korman, Distributed Computing 2017]



Foraging ants

"The Blueberry Hunters" by bob in swamp, CC BY 2.0.

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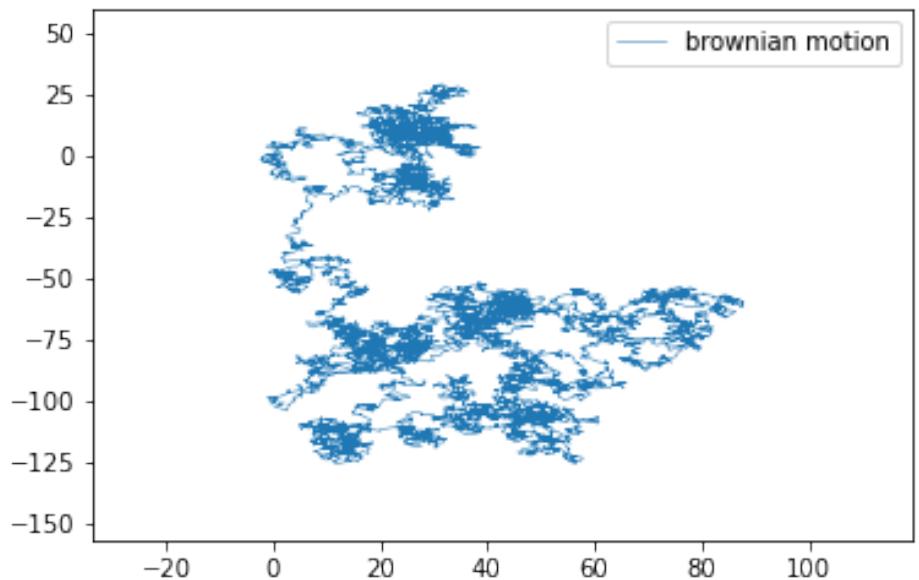
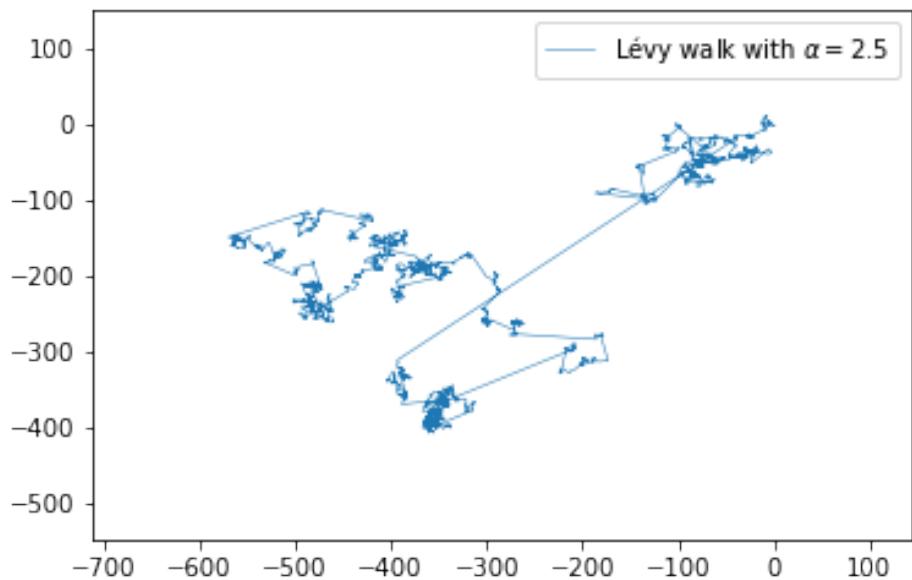
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# The Lévy walk



**Lévy walk (informal):**

*A Lévy walk is a random walk whose step-length density distribution is proportional to a power-law, namely, for each  $d \in \mathbb{R}^+$ ,  $f(d) \sim 1/d^\alpha$ , for some  $\alpha > 1$*

**Note:** the **speed** of the walk is **constant**

# Movement models and foraging theory

Lévy walks are used to model **movement patterns** [Reynolds, Biology Open 2018]

Examples:

- T cells within the brain
- swarming bacteria
- midge swarms
- termite broods
- schools of fish
- Australian desert ants
- a variety of molluscs



Desert ants

*Rhytidoponera mayri* workers. Credit: Associate Professor Heloise Gibb, La Trobe University

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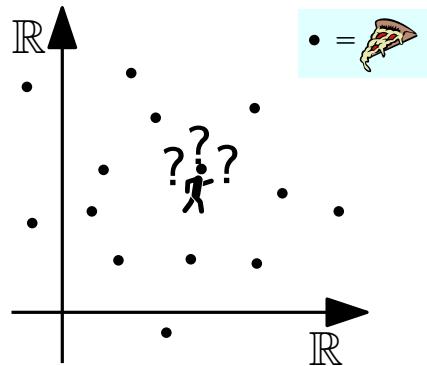
Rhytidoponera mayri workers. Credit: Associate Professor Heloise Gibb, La Trobe University

Widely employed in the **foraging theory**

- how animals search for food

# Lévy walk optimality

Foraging theory



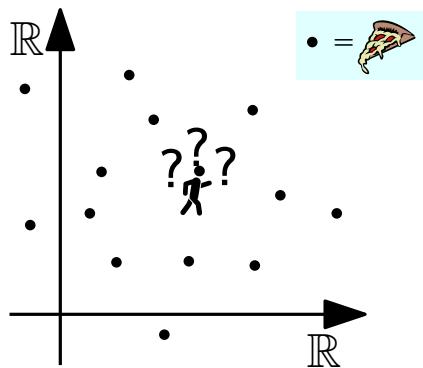
- distribution of food locations in  $\mathbb{R}^n$
- uninformed agent searching for food

*maximum expected food discovery rate*

[Viswanathan et al., Nature 1999]: Lévy walk with exponent  $\alpha = 2$  is optimal in any dimension, with some assumptions

# Lévy walk optimality

## Foraging theory

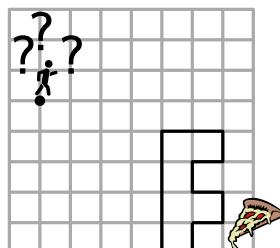


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*maximum expected food discovery rate*

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## Other search problems



- a target in the bidimensional torus  $\mathbb{T}$
- uninformed agent searching for it

*as fast as possible*

[Guinard et Korman, Sciences Advances 2021]: (truncated) Lévy walk with exponent  $\alpha = 2$  is optimal

# The Lévy flight foraging hypothesis

Formulation of an evolutionary hypothesis

**The Lévy flight foraging hypothesis** [Viswanathan et al., Physics of Life Reviews 2008]: since Lévy flights/walks **optimize random searches**, biological organisms must have therefore **evolved** to exploit Lévy flights/walks

- Special exponent  $\alpha = 2$

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We test **this hypothesis** by focusing on a **distributed search problem**:

- the **ANTS** (Ants Nearby Treasure Search) problem



# The ANTS problem

Introduced by [Feinerman et al., PODC 2012]:

Setting:

- $k$  (mutually) **independent agents** start moving on  $\mathbb{Z}^2$  from the origin
- time is **synchronous** and marked by a global clock
- one special node  $\mathcal{P} \in \mathbb{Z}^2$ , the **target**, placed by an **adversary** at unknown (Manhattan) distance  $\ell$  from the origin

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Task: find the target as fast as possible

Lower bound: for any  $k \geq 1$ , and for any search algorithm  $\mathcal{A}$ , the hitting time to find  $\mathcal{P}$  is  $\Omega(\ell^2/k + \ell)$  both with constant probability and in expectation

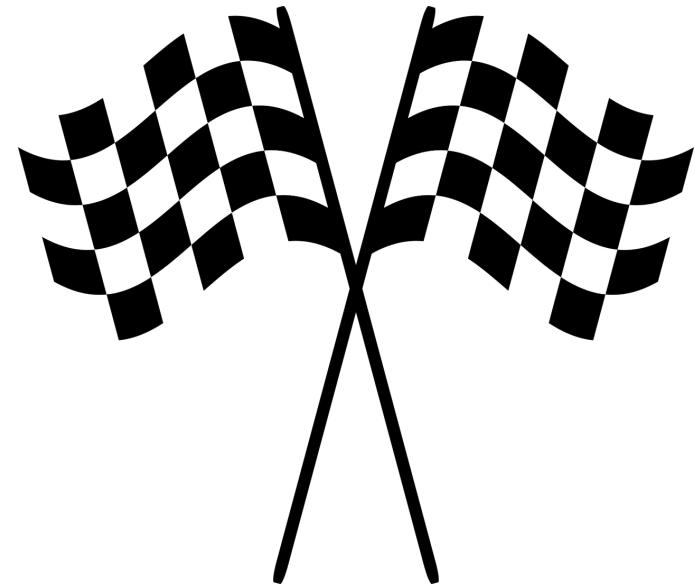


Image by OpenClipart-Vectors from Pixabay

# Our contributions

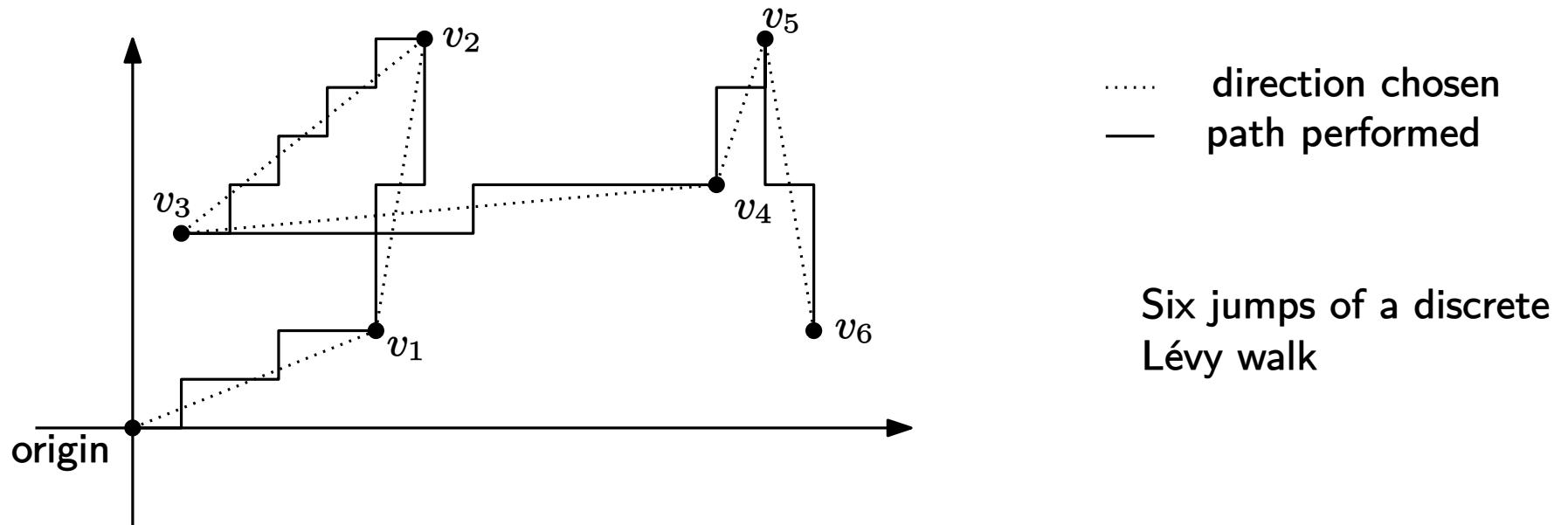
Based on the work [Clementi et al., PODC 2021]

- (i) first definition of Lévy walk in the discrete setting in  $\mathbb{Z}^2$ , time-homogeneous
- (ii) first analysis of the hitting time distribution of  $k$  parallel walks
- (iii) the Lévy walks can be employed to give a natural, almost-optimal solution to the ANTS problem (no advice, no communication)

# Our contributions

- (i) DEFINITION OF DISCRETE LÉVY WALK
- (ii) ANALYSIS OF THE PARALLEL HITTING TIME
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# Discrete Lévy walk



Let  $\alpha > 1$  be a real value

**Lévy walk:** the agent

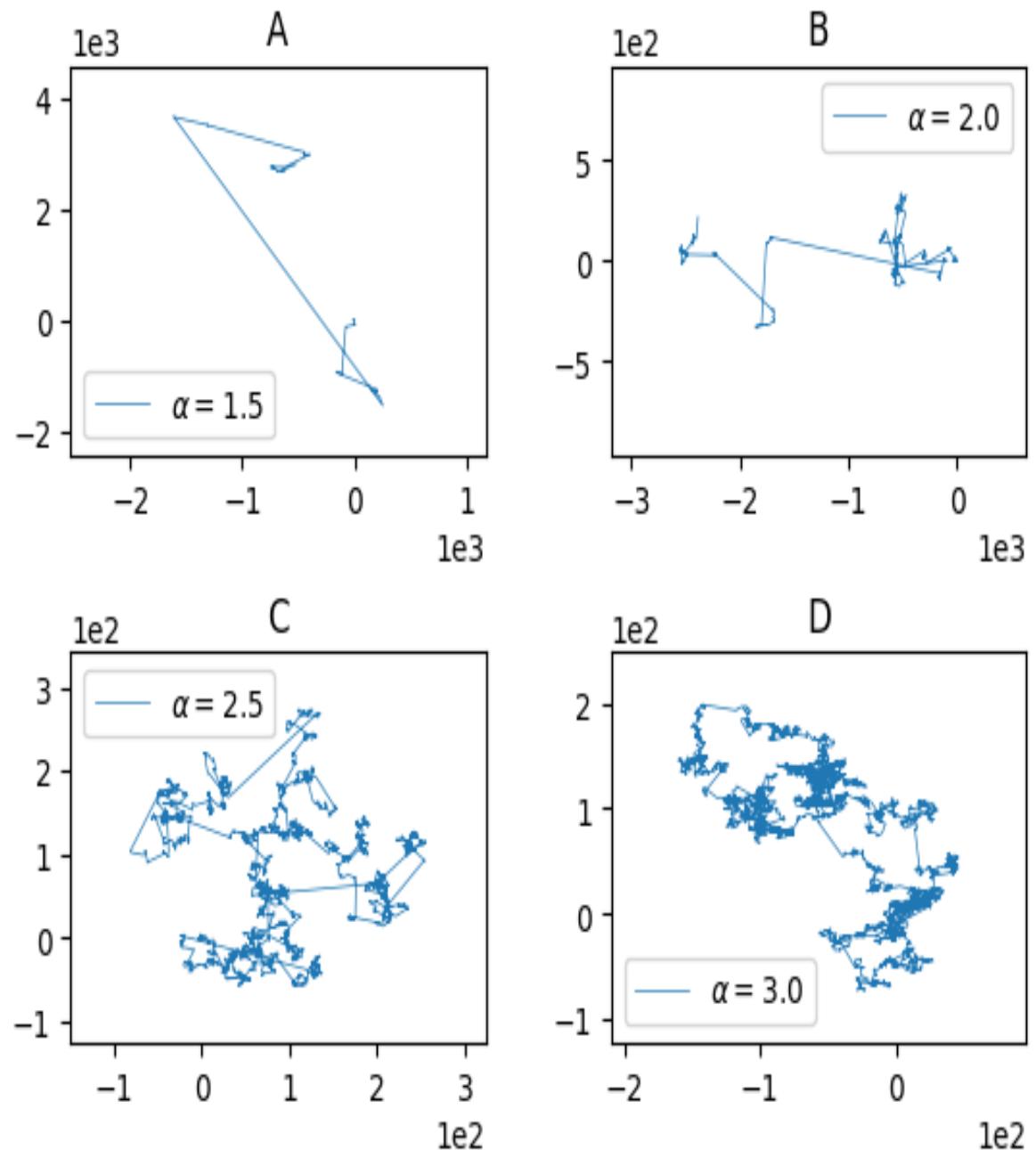
- a) chooses a **distance**  $d \in \mathbb{N}$  as follows:  $d = 0$  w.p.  $1/2$ , and  $d \geq 1$  w.p.  $c_\alpha/d^\alpha$
- b) chooses a **destination** u.a.r. among those at distance  $d$
- c) walks along an **approximating path** for  $d$  steps, one edge at a time, crossing  $d$  nodes
- d) **repeats** the procedure

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# $\alpha$ -behavior of Lévy walks

- **ballistic** diffusion: fig.s A and B
  - $1 < \alpha \leq 2$
- **super** diffusion: fig. C
  - $2 < \alpha < 3$
- **normal** diffusion: fig. D
  - $3 \leq \alpha$



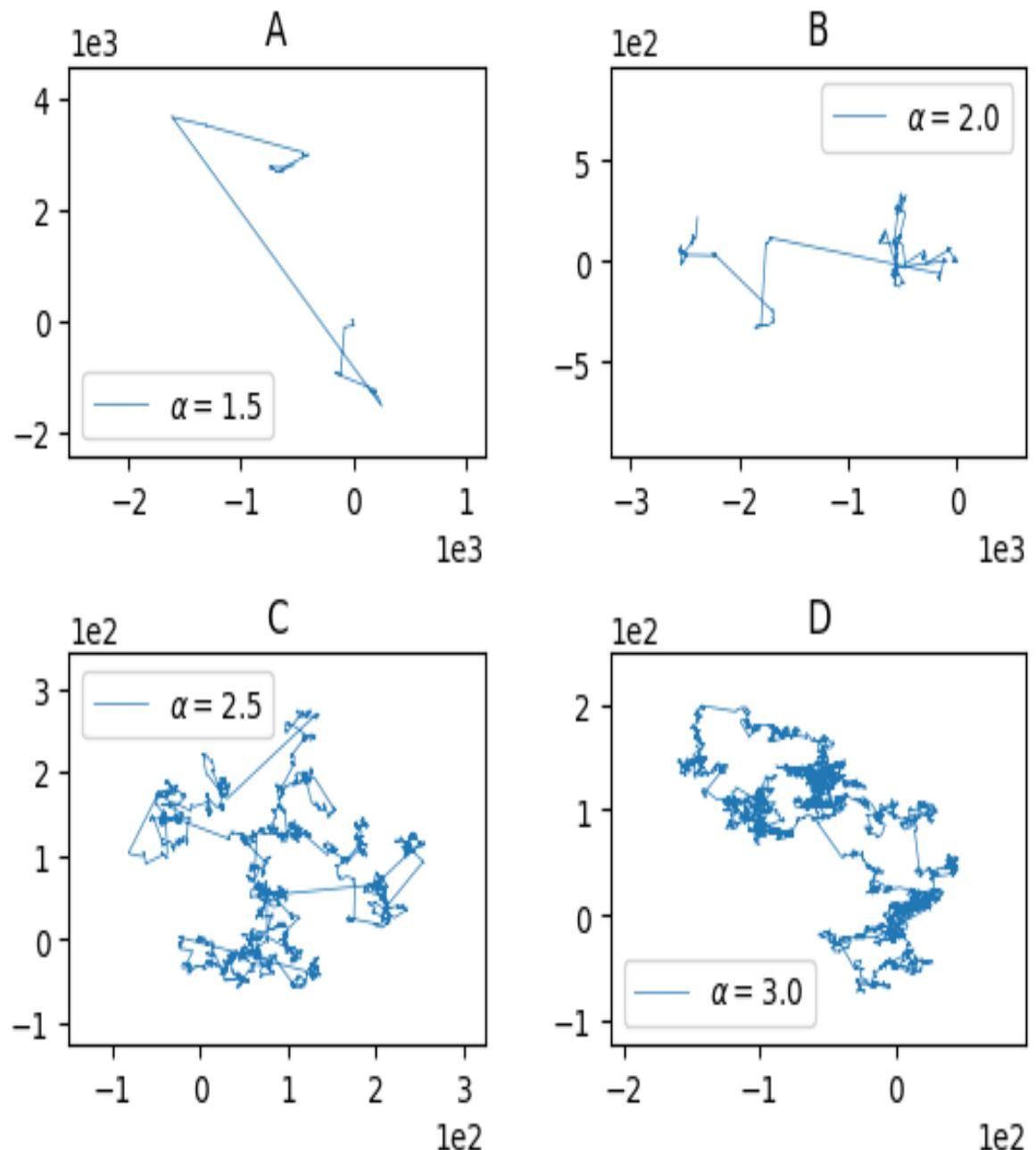
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## Expected jump-length

$$\int_1^\infty x^{-\alpha+1} dx$$

- $+\infty$  if  $\alpha \leq 2$
- $\frac{1}{\alpha-2}$  if  $\alpha > 2$



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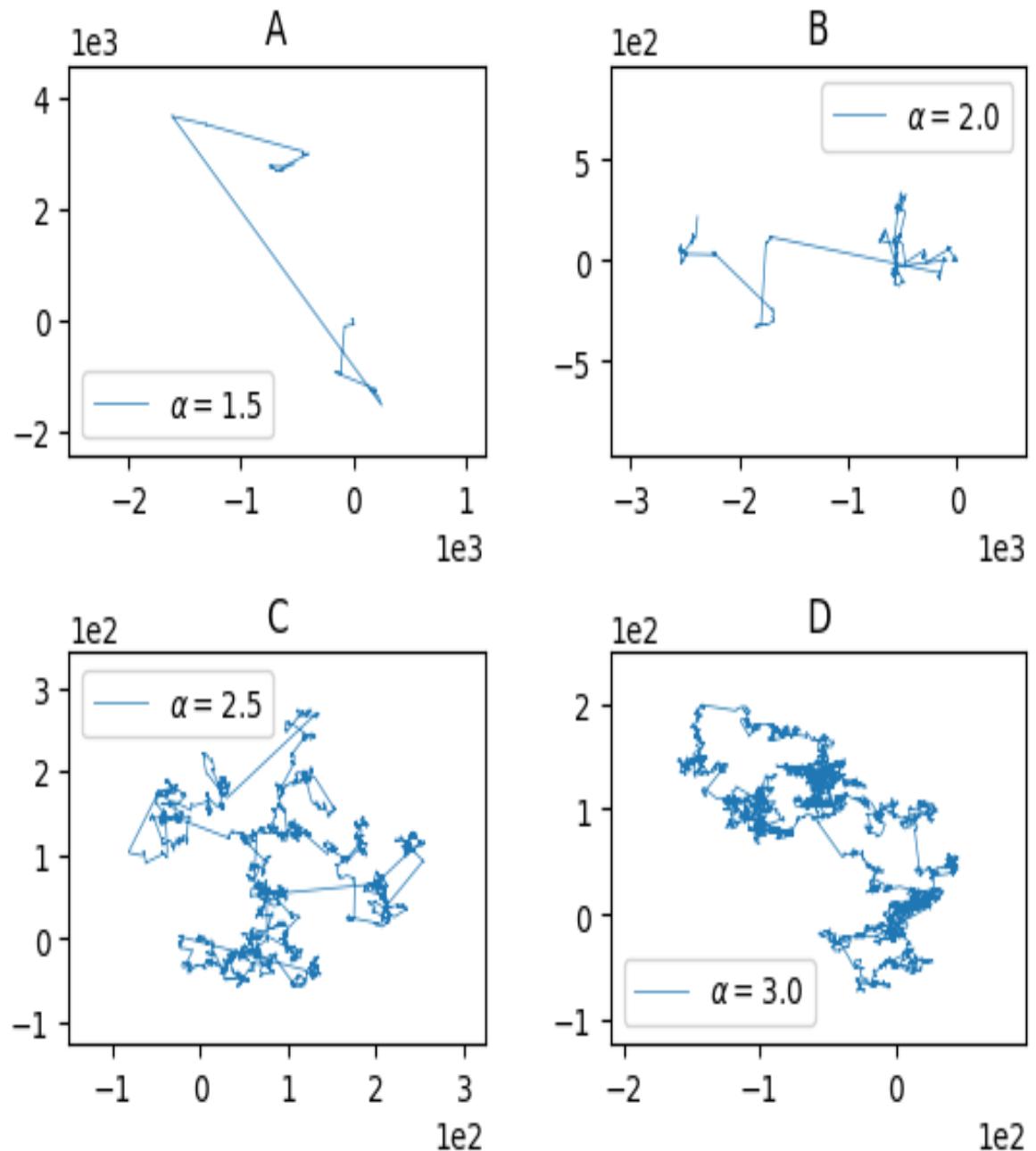
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## Jump-length second moment

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# Three ranges for $k$ and $\ell$

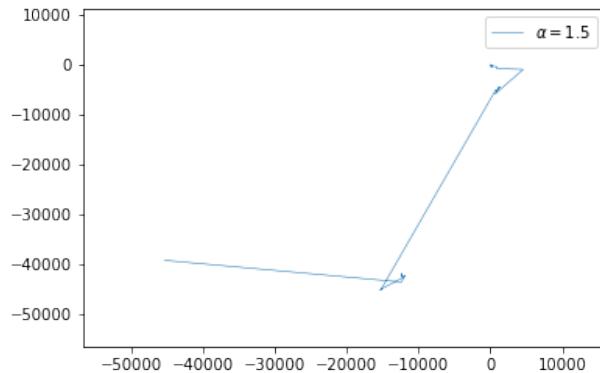
Recall:  $\ell$  target distance,  $k$  number of agents,  $\Omega(\ell^2/k + \ell)$  time **lower bound**

Three different possible settings:

1. **Close target:**  $\ell \leq k/\text{polylog}(k)$

Best strategy = **ballistic walks**

- any  $\alpha$  in  $(1, 2]$



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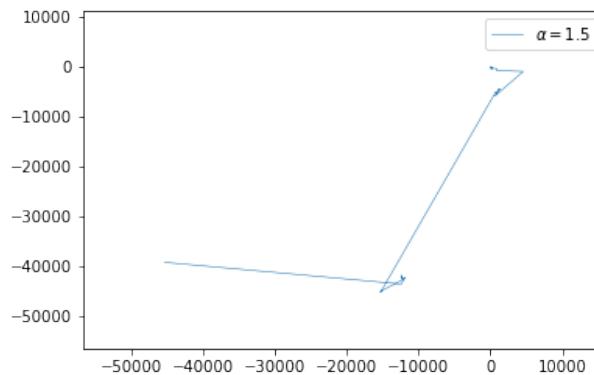
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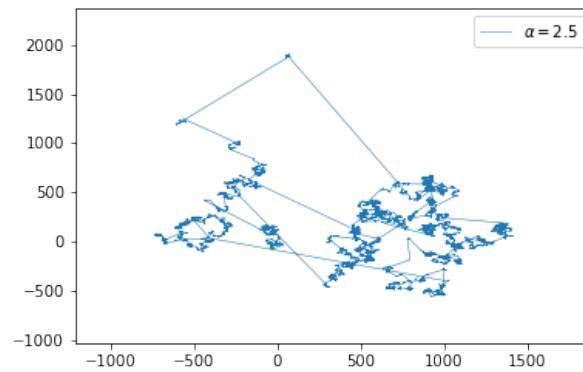
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2. **Far target:**  $k/\text{polylog}(k) \leq \ell \leq \exp(k^{\Theta(1)})$

Best strategy = **super-diffusive** range

- $\alpha = \underbrace{3 - \log k / \log \ell}_{\alpha^*} + \mathcal{O}\left(\frac{\log \log \ell}{\log \ell}\right)$



# Three ranges for $k$ and $\ell$

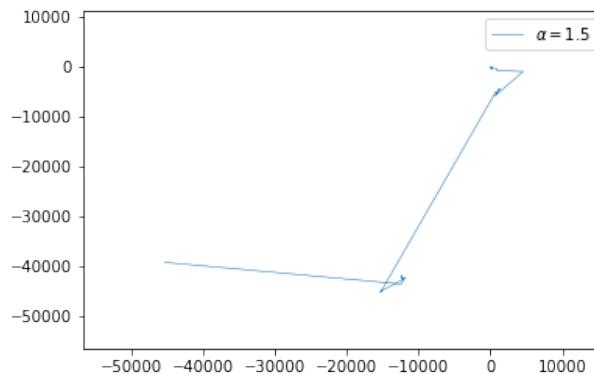
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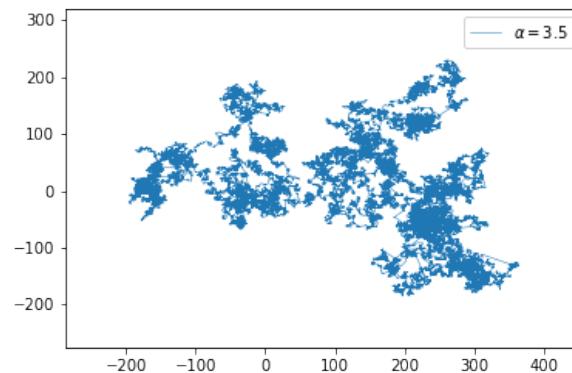
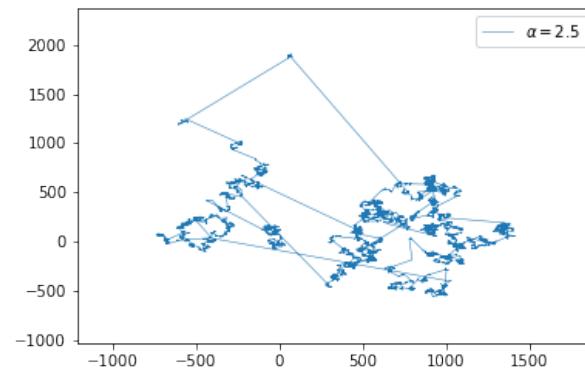
Best strategy = **super-diffusive range**

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3. **Very far target:**  $\exp(k^{\Theta(1)}) \leq \ell$

Best strategy = **diffusive walks**

- $[3, +\infty)$  (brownian-like behavior)



# Our contributions

(i) DEFINITION OF DISCRETE LÉVY WALK

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- proof's sketch

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# Far target case: proof's sketch

$\alpha^* = 3 - \log k / \log \ell$ : super-diffusive range  $\in (2, 3)$

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**Reverse engineering:**  $\alpha \in (2, 3)$

- analyze the hitting time of **a single Lévy flight** (Markov chain on  $\mathbb{Z}^2$ )
- derive bounds for a single **Lévy walk** through a **coupling argument**
- exploit **independence** to get results for  $k$  walks

**Remark:** Here, we **DON'T** show why other values for  $\alpha$  are worse than  $\alpha^*$

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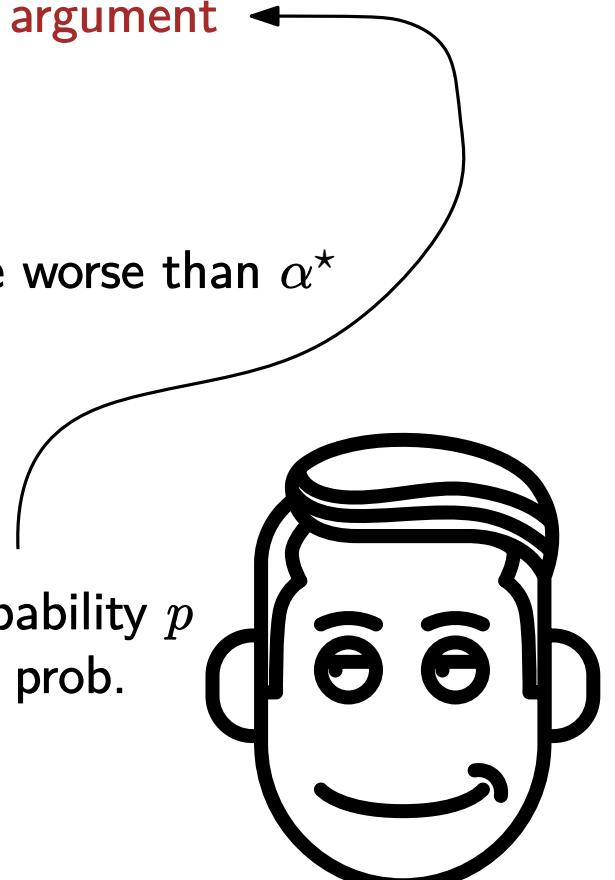
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**Remark:** Here, we **DON'T** show why other values for  $\alpha$  are worse than  $\alpha^*$

- $\mathcal{E}_t$  **event**: first  $t$  jumps have length  $\leq (t \log t)^{\frac{1}{\alpha-1}}$

**Coupling argument:** if a **Lévy flight** finds a node with probability  $p$  in time  $t$  (**conditional on  $\mathcal{E}_t$** ), then a **Lévy walk** finds it with prob.  $\approx p$  in time  $\mathcal{O}(t)$



# Single Lévy flight analysis

- $\mathcal{P}$  target node

Let

- $Z_u(t)$  = random variable of **number of visits** in  $u$  until time  $t$
- $\mathcal{E}_t$  = the **event** first  $t$  jumps have length  $\leq (t \log t)^{\frac{1}{\alpha-1}}$
- $a_t = \mathbb{E} [Z_{(0,0)}(t) \mid \mathcal{E}_t]$
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**Lemma:**  $p(t) = \mathbb{P}(Z_{\mathcal{P}}(t) > 0 \mid \mathcal{E}_t) \geq \mathbb{E}[Z_{\mathcal{P}}(t) \mid \mathcal{E}_t] / a_t$

Comes from two facts

- (i)  $\mathbb{E}[Z_{\mathcal{P}}(t) \mid \mathcal{E}_t] = \mathbb{E}[Z_{\mathcal{P}}(t) \mid Z_{\mathcal{P}}(t) > 0, \mathcal{E}_t] \cdot \mathbb{P}(Z_{\mathcal{P}}(t) > 0 \mid \mathcal{E}_t)$
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**Lemma:**  $a_t = \Theta(1)$

The process is **transient**

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We look for this expectation



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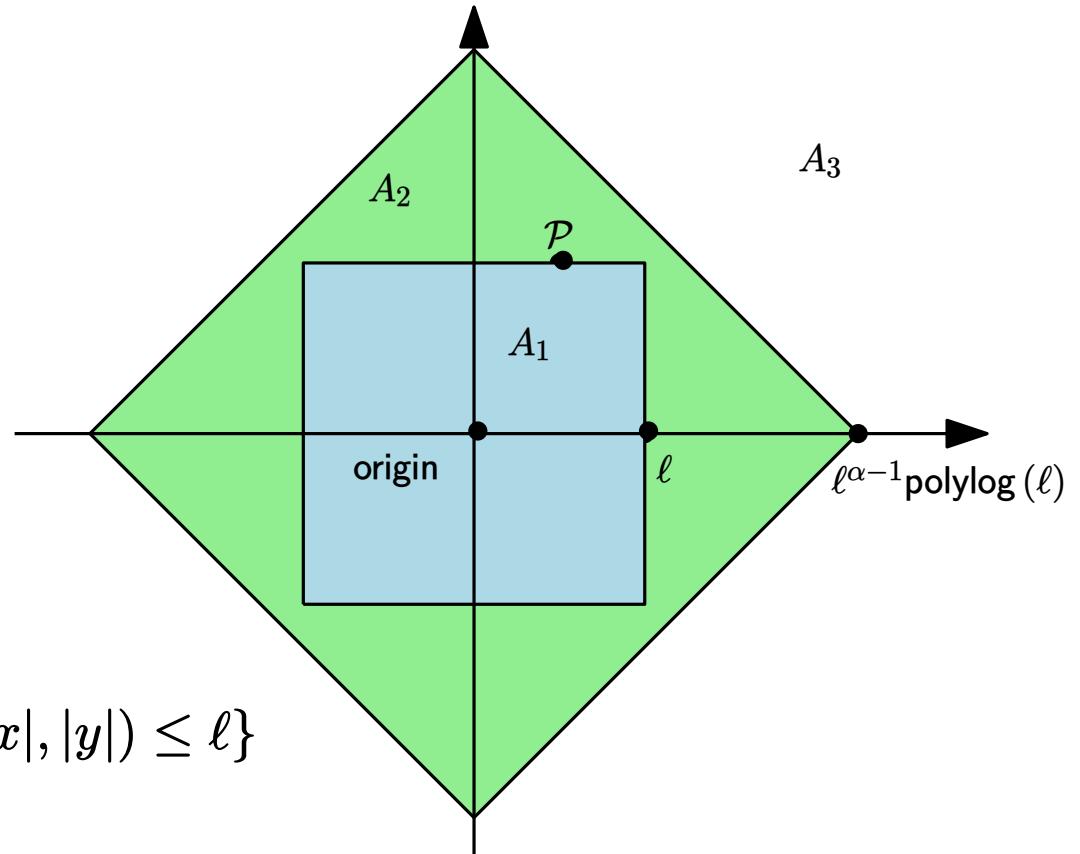
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# Partition of the space

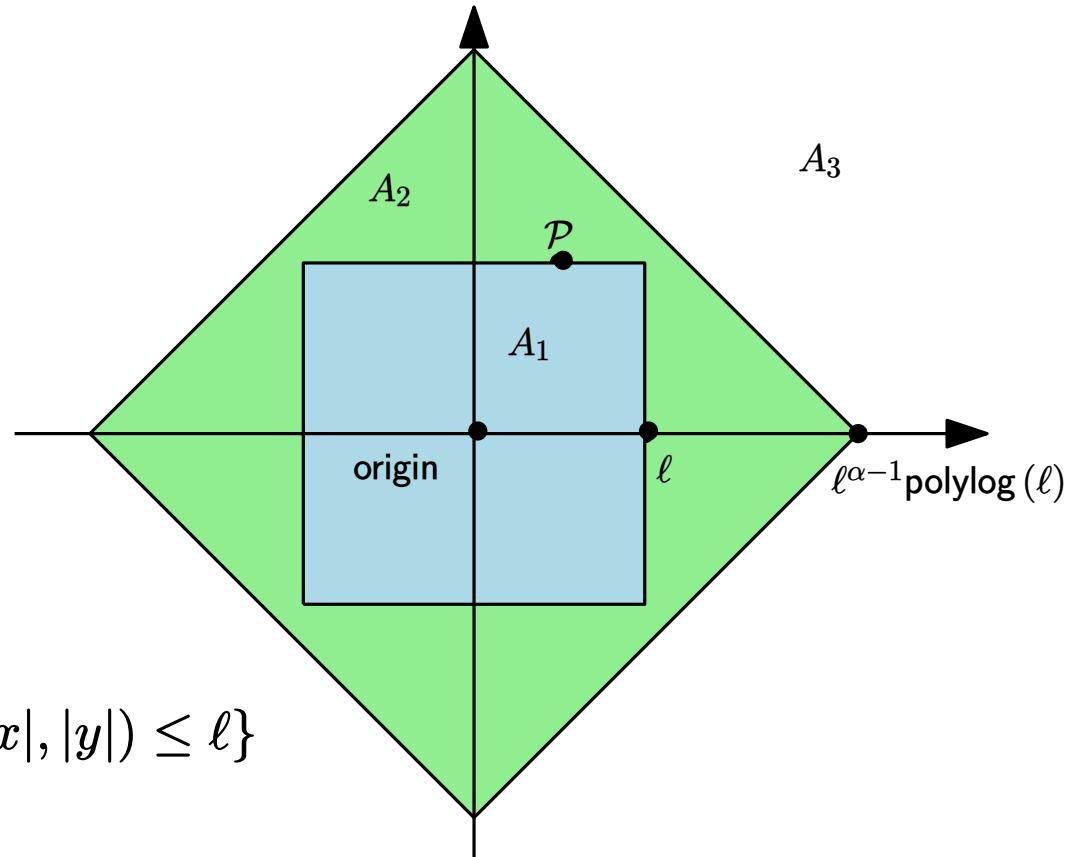
We **partition**  $\mathbb{Z}^2$  in three areas in the following way



- $A_1 = Q(\ell) = \{(x, y) \in \mathbb{Z}^2 : \max(|x|, |y|) \leq \ell\}$
- $A_2 = B_{\ell^{\alpha-1} \text{polylog}(\ell)}((0, 0)) \setminus A_1$
- $A_3 = \mathbb{Z}^2 \setminus (A_1 \cup A_2)$

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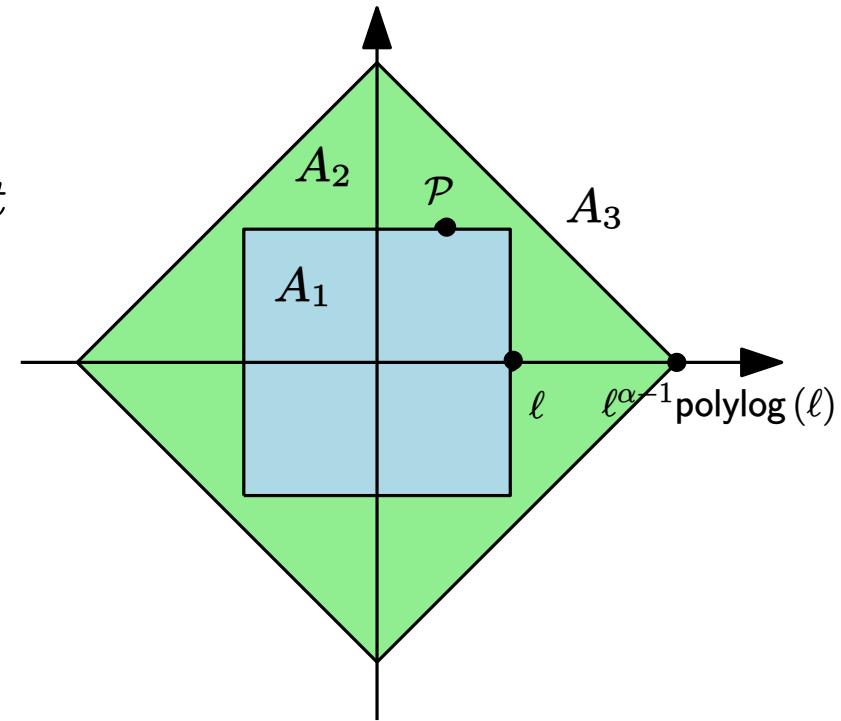
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**Why:** roughly  $\ell^{\alpha-1}$  jumps to **first jump** of length  $\geq \ell$

- Afterwards, the walk is **lost**

# Getting $\mathbb{E} [Z_{\mathcal{P}} (t) | \mathcal{E}_t]$

$Z_S (t)$ : total number of visits in the set  $S$  until time  $t$

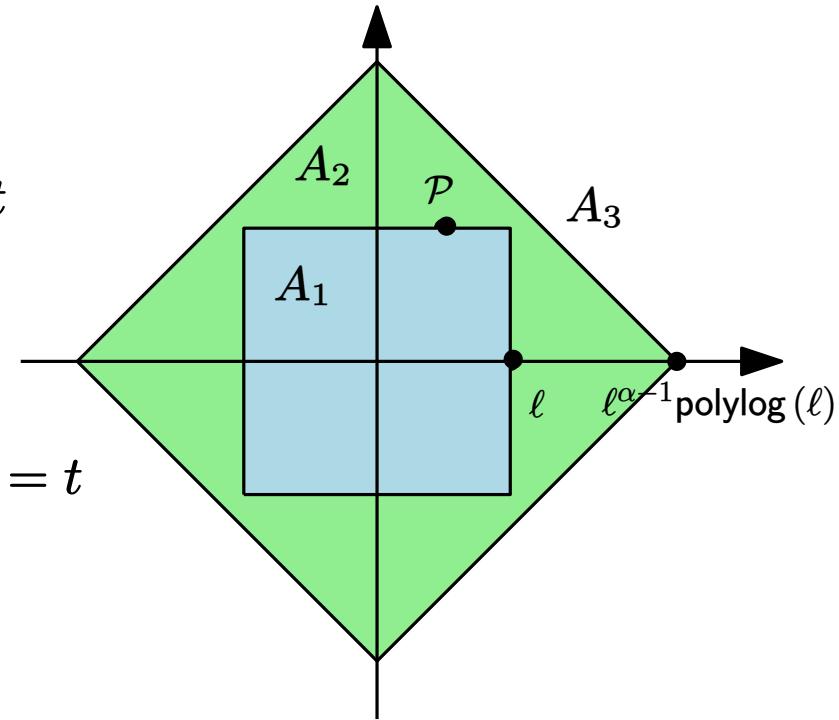


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As  $Z_{\mathbb{Z}^2} (t) = t$ , then

a)  $\mathbb{E} [Z_{A_1} (t) | \mathcal{E}_t] + \mathbb{E} [Z_{A_2} (t) | \mathcal{E}_t] + \mathbb{E} [Z_{A_3} (t) | \mathcal{E}_t] = t$



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For some  $t = \Theta(\ell^{\alpha-1})$ , we prove that:

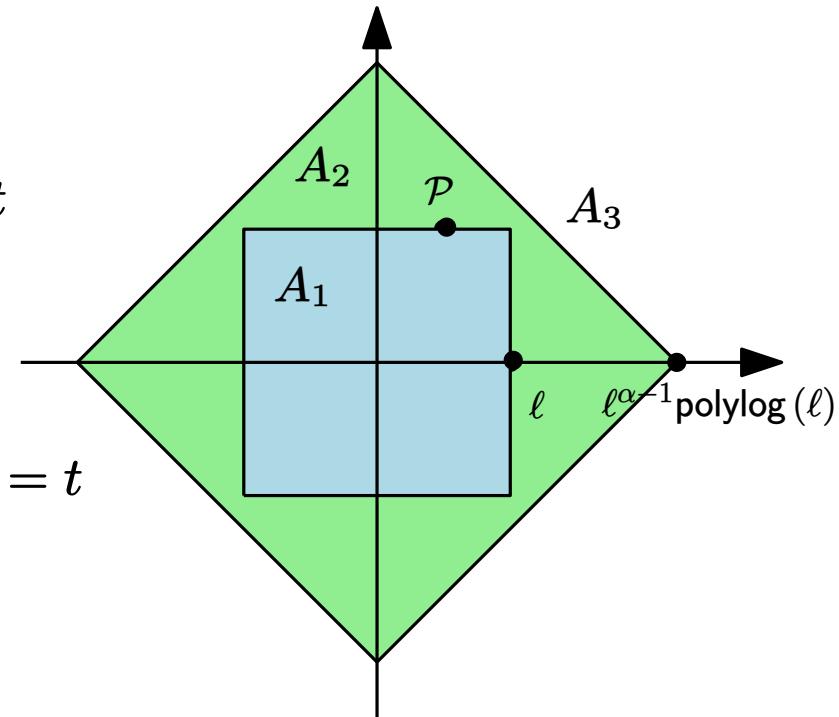
b)  $\mathbb{E} [Z_{A_1}(t) | \mathcal{E}_t] \leq \frac{3}{4}t$

$$|A_2|$$

c)  $\mathbb{E} [Z_{A_2}(t) | \mathcal{E}_t] \leq \mathbb{E} [Z_{\mathcal{P}}(t) | \mathcal{E}_t] \cdot \overbrace{\ell^2 \text{polylog}(\ell)}^{|\mathcal{P}|}$

Monotonicity property

d)  $\mathbb{E} [Z_{A_3}(t) | \mathcal{E}_t] = \mathcal{O}(t/\log t)$



# Getting $\mathbb{E}[Z_{\mathcal{P}}(t) | \mathcal{E}_t]$

$Z_S(t)$ : total number of visits in the set  $S$  until time  $t$

As  $Z_{\mathbb{Z}^2}(t) = t$ , then

a)  $\mathbb{E}[Z_{A_1}(t) | \mathcal{E}_t] + \mathbb{E}[Z_{A_2}(t) | \mathcal{E}_t] + \mathbb{E}[Z_{A_3}(t) | \mathcal{E}_t] = t$

For some  $t = \Theta(\ell^{\alpha-1})$ , we prove that:

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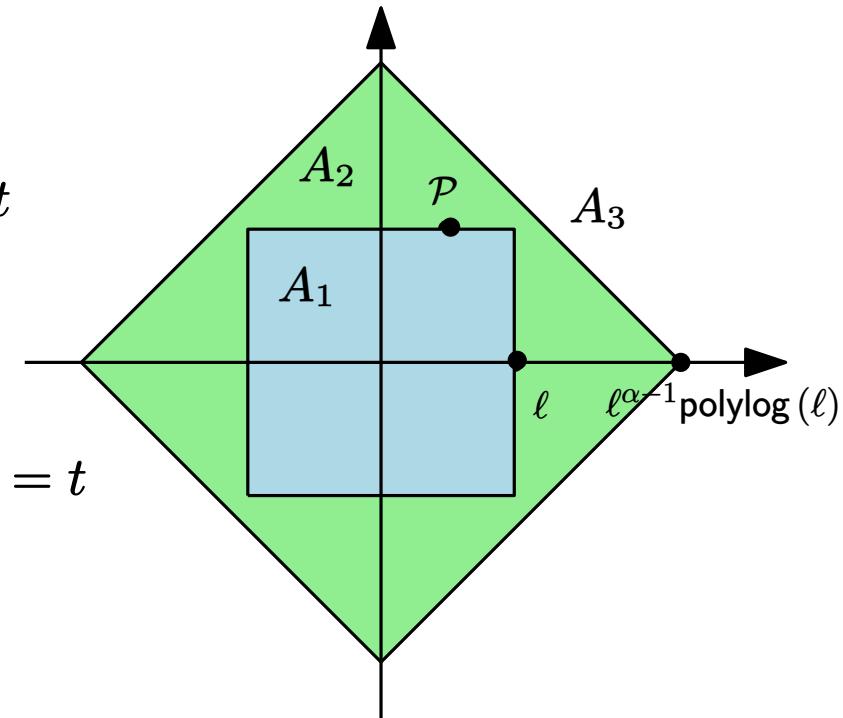
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Combine (a) with (b), (c), and (d) to get

$$\mathbb{E}[Z_{\mathcal{P}}(t) | \mathcal{E}_t] \geq 1 / (\ell^{3-\alpha} \text{polylog}(\ell))$$



# Getting $p(t)$ and $\alpha^*$

**Reminder:**  $p(t) = \mathbb{P}(Z_{\mathcal{P}}(t) > 0 \mid \mathcal{E}_t) \geq \mathbb{E}[Z_{\mathcal{P}}(t) \mid \mathcal{E}_t] / a_t$

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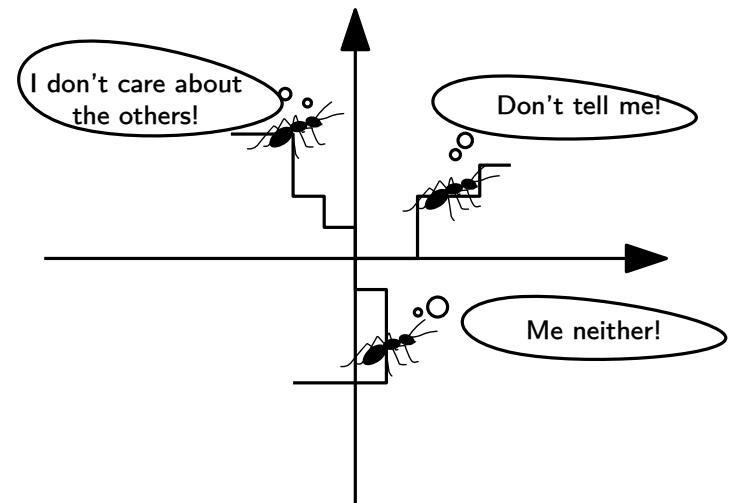
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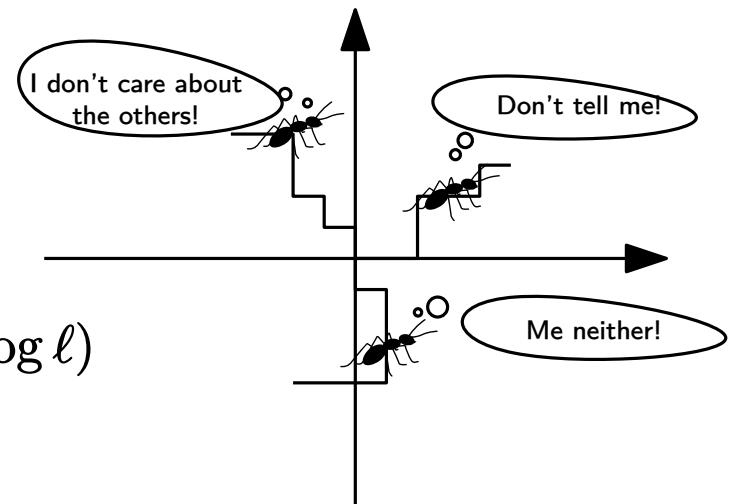
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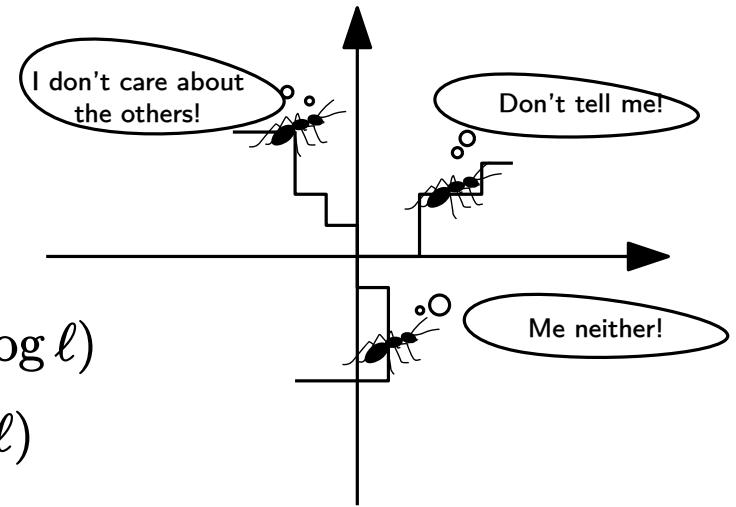
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$\alpha^*$

$$\alpha = 3 - \log k / \log \ell + \mathcal{O}(\log \log \ell / \log \ell)$$
$$t = \mathcal{O}(\ell^{\alpha-1}) \leq (\ell^2/k + \ell) \text{polylog}(\ell)$$



# Our contributions

- (i) DEFINITION OF DISCRETE LÉVY WALK
- (ii) ANALYSIS OF THE PARALLEL HITTING TIME
- (iii) ALGORITHM FOR THE ANTS PROBLEM

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**Algorithm:** each agent  $u$  *samples* u.a.r. a real number  $\alpha_u \in (2, 3)$ . Then, it *performs* a discrete *Lévy walk* with *exponent*  $\alpha_u$

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**Proof outline:**

Fix some  $\epsilon = \mathcal{O}(\log \log \ell / \log \ell)$

We use:  $\ell < \exp(k^{\Theta(1)})$  ( $\iff k \geq \text{polylog}(\ell)$ ) + Chernoff bound

$\implies$  at least  $\Theta(\epsilon k)$  agents sample an exponent in the range  $(\alpha^* - \epsilon, \alpha^* + \epsilon)$  w.h.p.

$\Theta(\epsilon k)$  agents are sufficient to ensure high probability to find the target fast enough

# Recap

We

- provide a **definition** of a discrete version of the **Lévy walk**
- analyze the **hitting time** of  $k$  parallel **Lévy walks**
- show that for any choices of  $k$  and  $\ell$  from a wide range, **Lévy walks** are an **almost-optimal search strategy** for the ANTS problem

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- show that for any choices of  $k$  and  $\ell$  from a wide range, **Lévy walks** are an **almost-optimal search strategy** for the ANTS problem
- mathematically corroborate the **Lévy flight foraging hypothesis**
- argue the non (universal) optimality of exponent  $\alpha = 2$

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## 3. Opinion dynamics

- The consensus problem with noisy communications
- The UNDECIDED-STATE and the 3-MAJORITY dynamics: contribution

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## 5. Conclusions & perspectives

# The consensus problem

**Input:** system of  $n$  agents supporting opinions, with a communication network

**Task:** designing a protocol which brings the system in finite time to a configuration such that

1. all agents support the same opinion (**AGREEMENT**)
2. the final opinion is among the initial ones (**VALIDITY**)
3. the agreement keeps on unless external events occur (**STABILITY**)

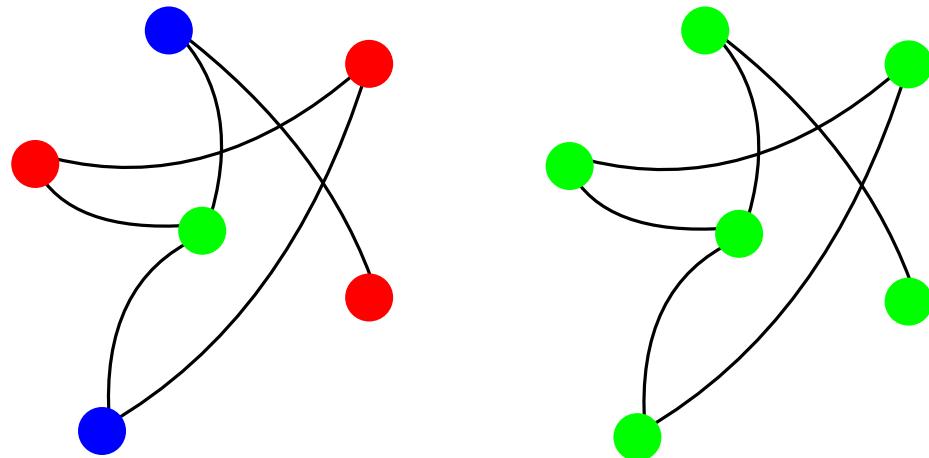
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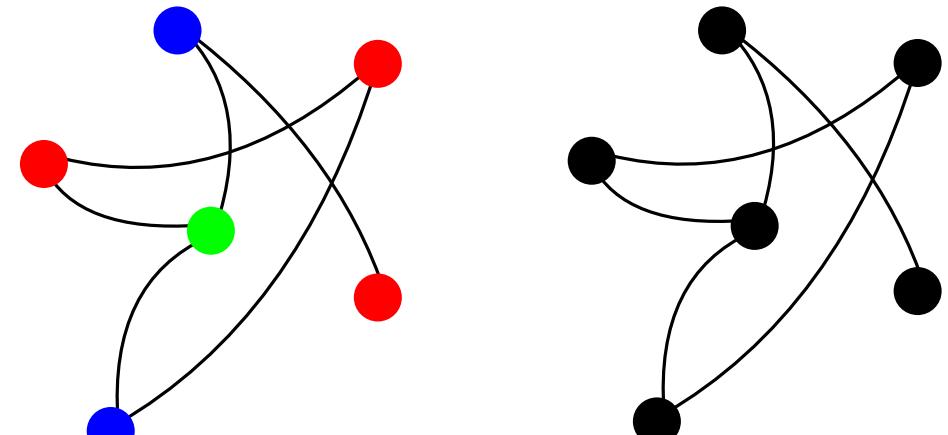
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time  $t = 0$   $\longrightarrow$  time  $t > 0$



valid consensus

time  $t = 0$   $\longrightarrow$  time  $t > 0$



non valid consensus

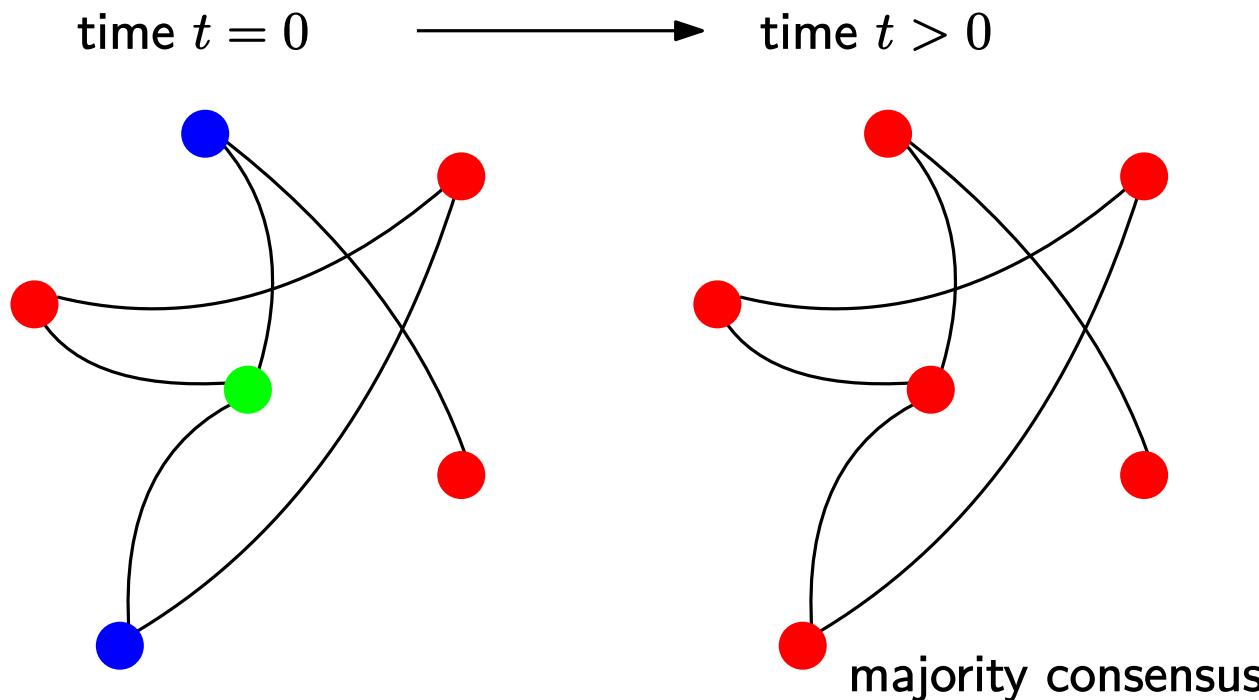
# The majority consensus problem

- 1. AGREEMENT
- 2. VALIDITY
- 3. STABILITY



- 1. AGREEMENT
- 2. MAJORITY
- 3. STABILITY

2. MAJORITY property: the **final opinion** is the **initial majority** one



# Opinion dynamics

Opinion dynamics: class of simple, lightweight parallel protocols for distributed tasks (e.g., consensus, clustering, etc.)

Many have been investigated, including:

- Voter Model [Hassin and Peleg, Inf. Comput. 2001] → linear dynamics
- Averaging dynamics [Becchetti et al., SODA 2017]
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Often, settings with adversarial Byzantine failures are investigated

Not realistic in biological scenarios; rather, uniform noise [Feinerman et al., PODC 2014]

# Uniform communication noise

Inspired by [Feinerman et al., PODC 2014], [Fraigniaud and Natale, PODC 2016]

$\Sigma$  set of  $k$  opinions,  $p \in [0, 1]$  constant

When  $u$  looks at  $v$ 's **opinion**  $x$

- a) with **probability**  $1 - p$ ,  $u$  **sees**  $x$
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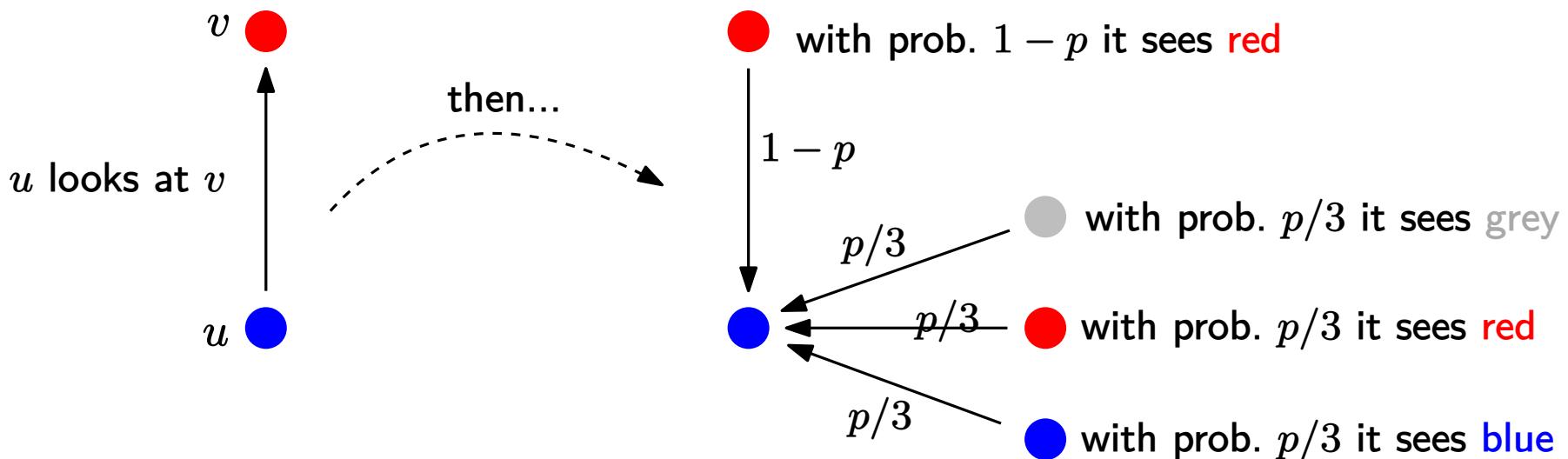
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Example:  $k = 3$



# 3-MAJORITY and UNDECIDED-STATE

**3-Majority** dynamics: each node  $u$

1. samples 3 **neighbors** u.a.r.
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**Undecided-State** dynamics: each node  $u$

1. samples a **neighbor**  $v$  u.a.r.
2. **pulls**  $v$ 's **opinion**
3. **updates** its opinion according to the following **table**

$u \setminus v$	opinion $i$	opinion $j$	undecided
opinion $i$	$i$	undecided	$i$
opinion $j$	undecided	$j$	$j$
undecided	$i$	$j$	undecided

# Our contribution

Based on [d'Amore et al., SIROCCO 2020], [d'Amore et Ziccardi, SIROCCO 2022], [d'Amore et al., Swarm Intelligence 2022]

We study the **Undecided-State** dynamics and the **3-Majority** dynamics with  $k = 2$  opinions in the presence of uniform noise in the complete graph

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For the **3-Majority** dynamics: phase-transition

- $p < 1/3$ :
- a value  $\bar{s} = \Theta(n)$  exists such that the bias of the system reaches the interval  $I_\varepsilon = [(1 - \varepsilon)\bar{s}, (1 + \varepsilon)\bar{s}]$  in time  $\mathcal{O}(\log n)$  w.h.p., and keeps in  $I_\varepsilon$  for time  $\text{poly}(n)$  w.h.p.  $\longrightarrow$  **almost-consensus**
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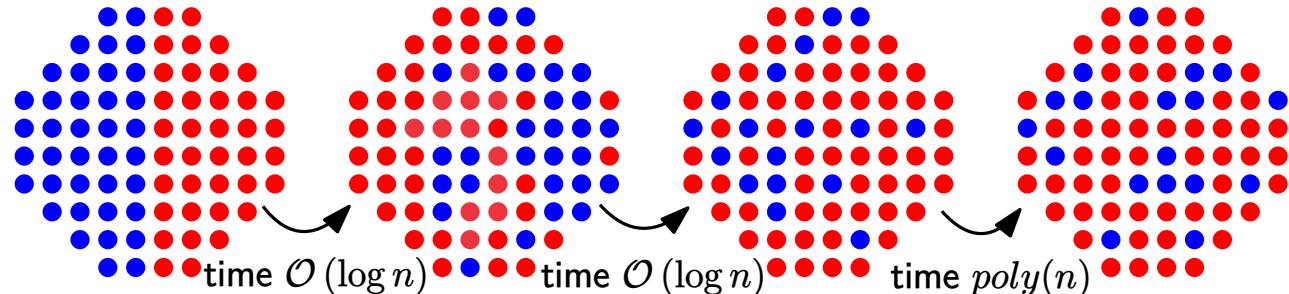
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For the **Undecided-State**: similar behavior

- phase-transition at  $p = 1/2$
- less characterized, more complex

# Recap

- **Undecided-State** dynamics and **3-Majority** dynamics **are not** implemented by biological systems
  - despite the **bio-inspiration**, **highly abstract model**
  - aiming to **capture** fundamental **phenomena** that (**very loosely**) relates to many biological systems

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  - despite the **bio-inspiration**, **highly abstract model**
  - aiming to **capture** fundamental **phenomena** that (**very loosely**) relates to many biological systems
- **Undecided-State** dynamics **more resilient** to **noise** than **3-Majority** dynamics
  - **phase-transitions**:  $p = 1/2$  vs  $p = 1/3$ 
    - $p = 1/2$  (USD) means **half** of the **communications** are **non-noisy** on average
    - $p = 1/3$  (3-Maj) means 2 out of 3 **pulled opinions** are **non-noisy** on average

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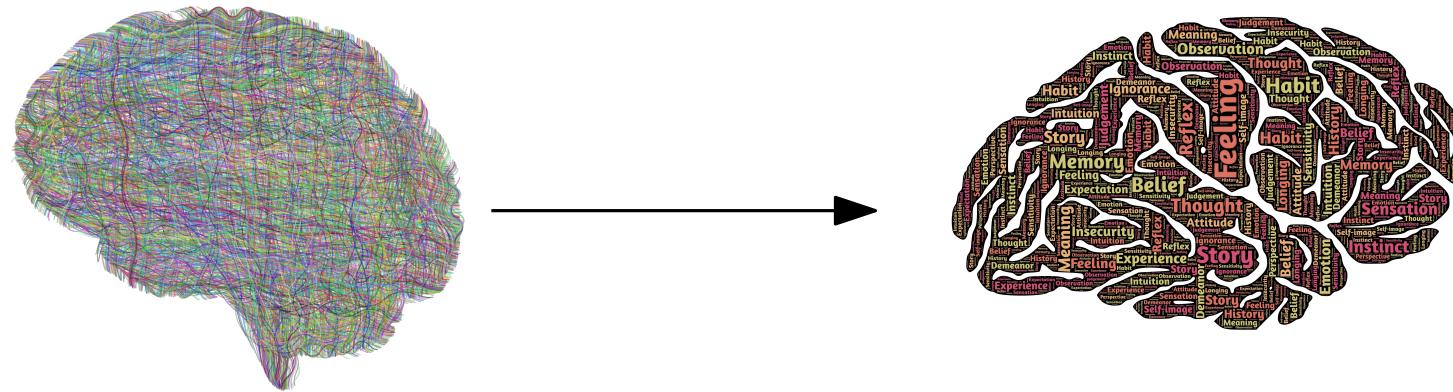
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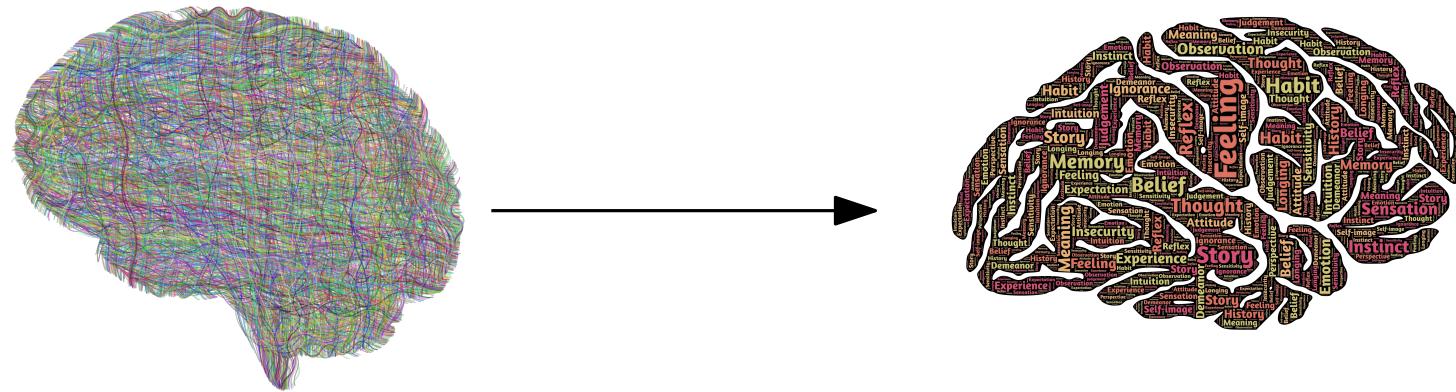
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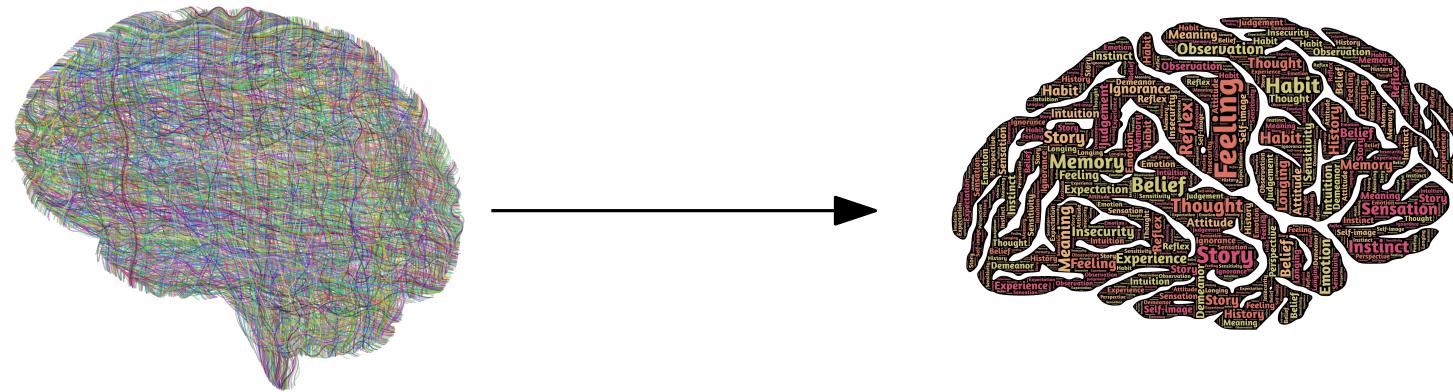


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**Assembly Calculus:** recently proposed **formal computational system** [Papadimitriou et al., PNAS 2020]

- explicit **purpose** of fitting Axel's logic
- bridging through **computation** the **gap** between **neurons** and **intelligence**

# Assemblies

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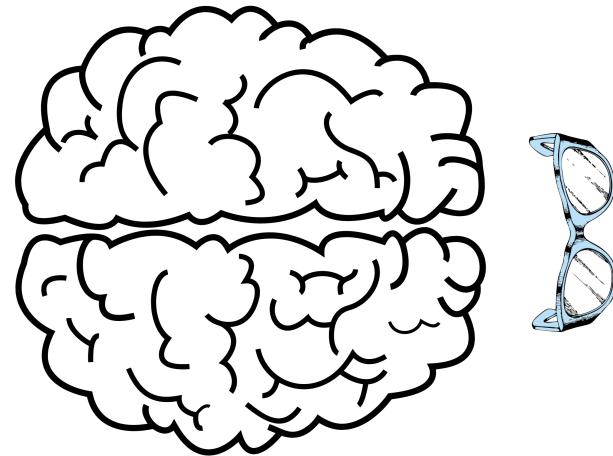
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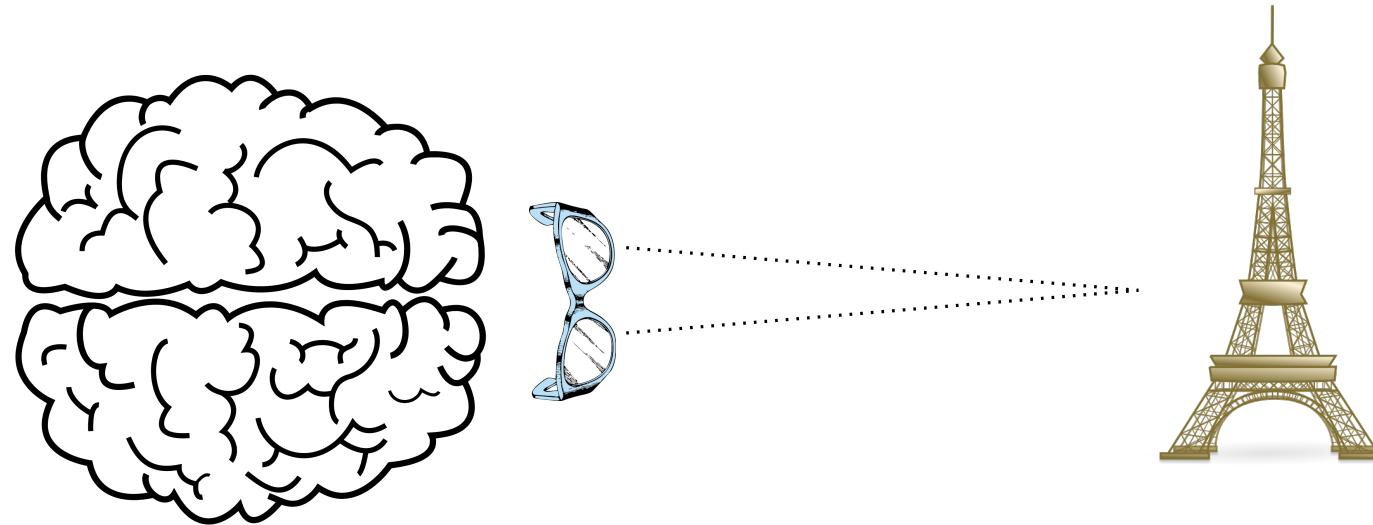
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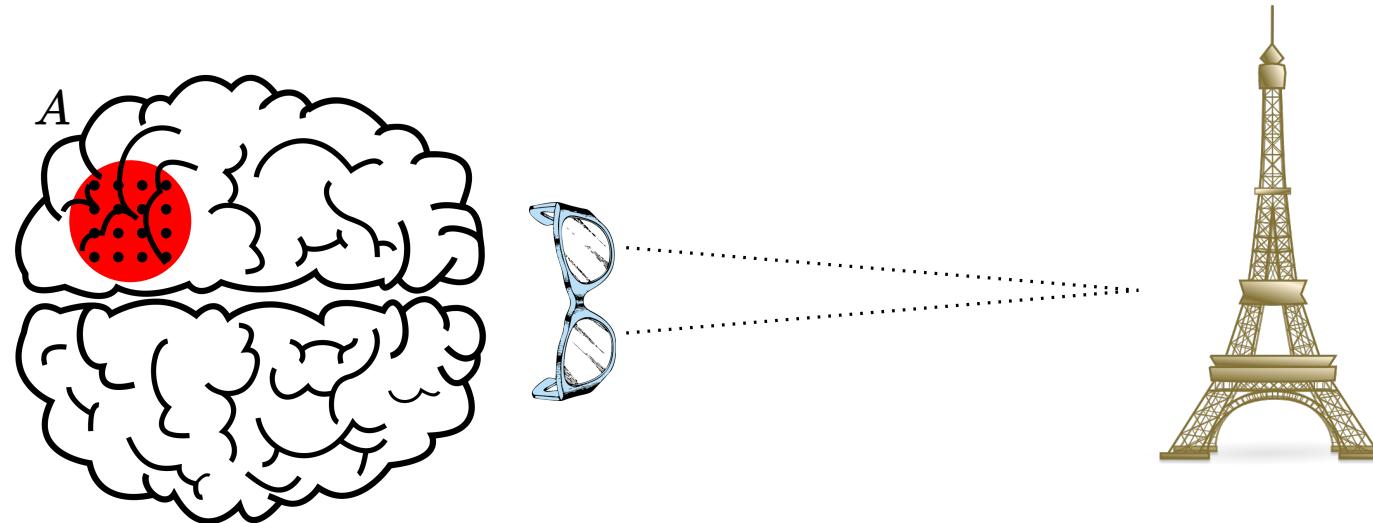
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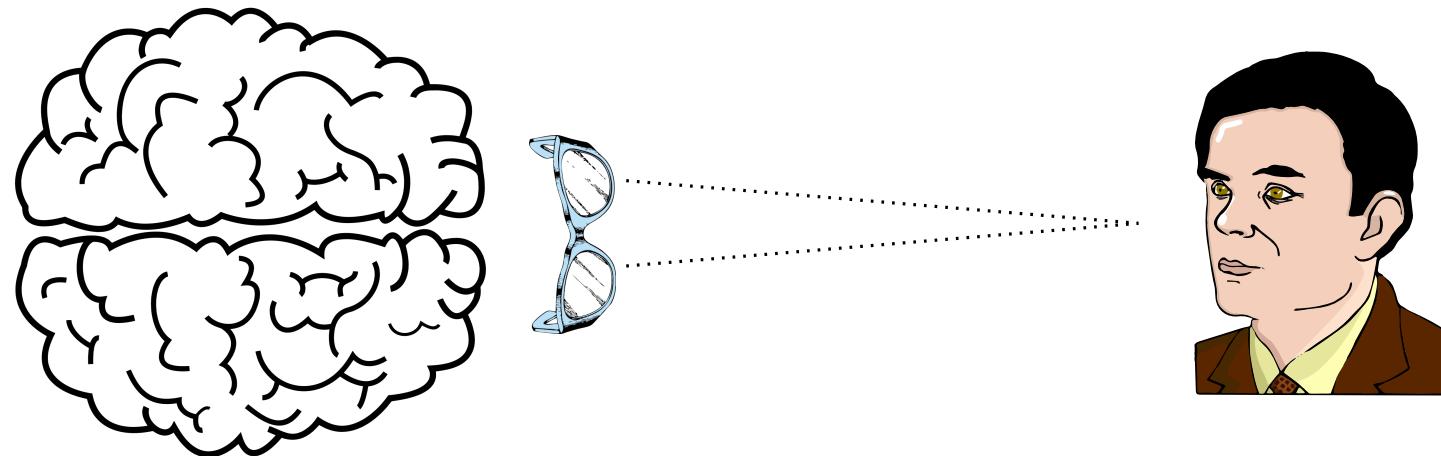
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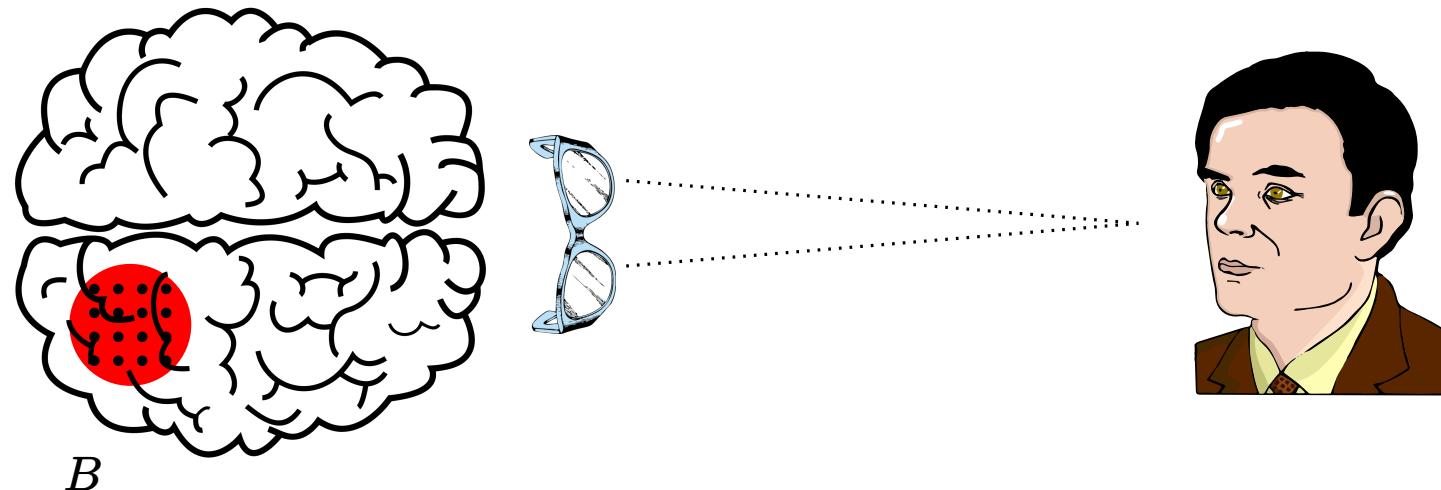
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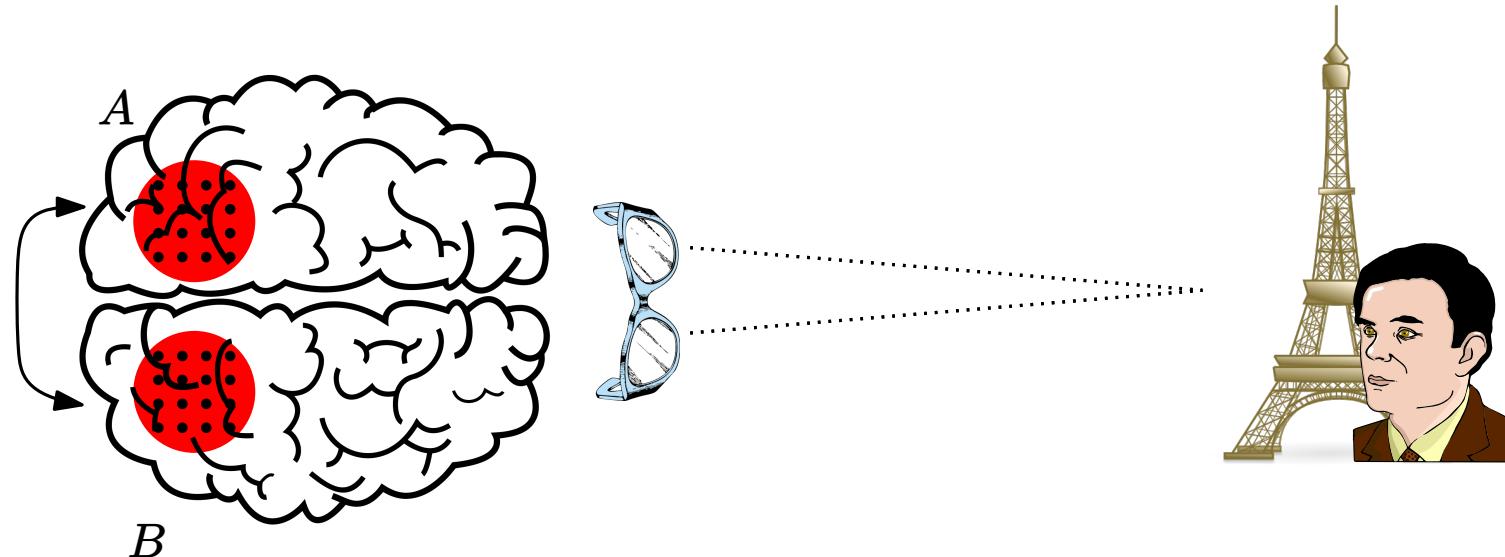
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**Association:** experiment by [Ison et al., Neuron 2015]

# Contribution

We empirically **demonstrate** that reasonably **large** and **complex** programs **run** correctly and reliably in the **assembly calculus** by implementing **planning strategies** in the **blocks world** (**Julia** programming language) [d'Amore et al., AAAI 2022]

**Possible moves:** • move a block from the **top** of a stack to the **table**

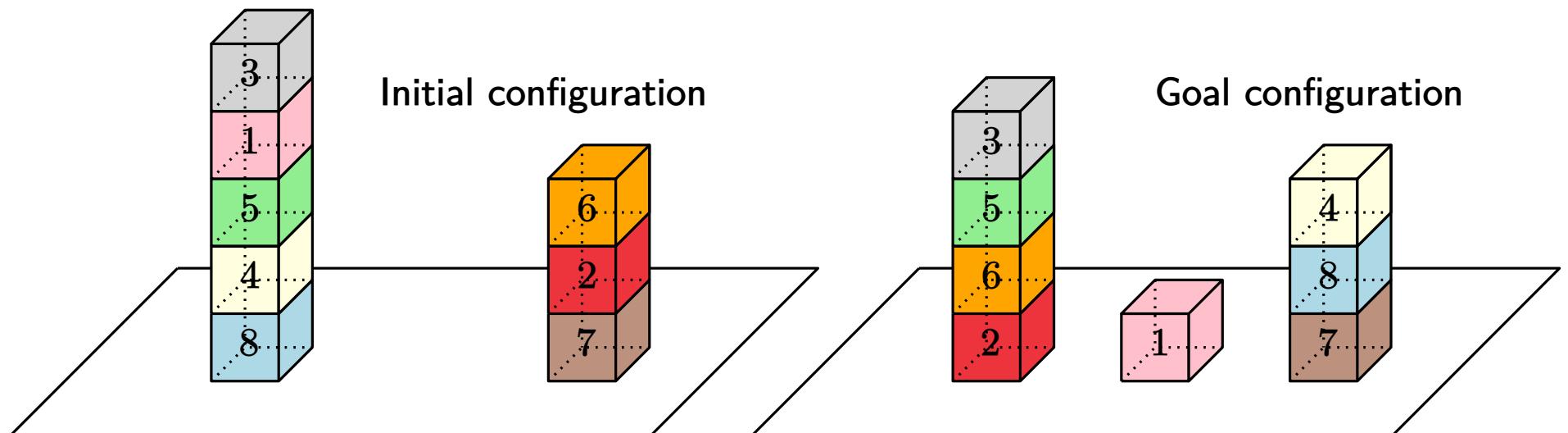
- move a block from the **top** of a stack to the **top** of another stack
- move a block from the **table** to the **top** of a stack

**Input:**

- set of **initial** stacks
- set of **goal** stacks

**Output:**

- the sequence of moves



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- Natural algorithms
- Distributed computing tasks in biological systems

## 2. Lévy walks

- Lévy walk's optimality
- Lévy flight foraging hypothesis
- The ANTS problem: contribution + proof's sketch

## 3. Opinion dynamics

- The consensus problem with noisy communications
- The UNDECIDED-STATE and the 3-MAJORITY dynamics: contribution

## 4. Assembly Calculus

- Mentions to model and contribution

## 5. Conclusions & perspectives

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Thank  
You!

# Appendix

1. Lévy walks
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# Defining the discrete Lévy walk

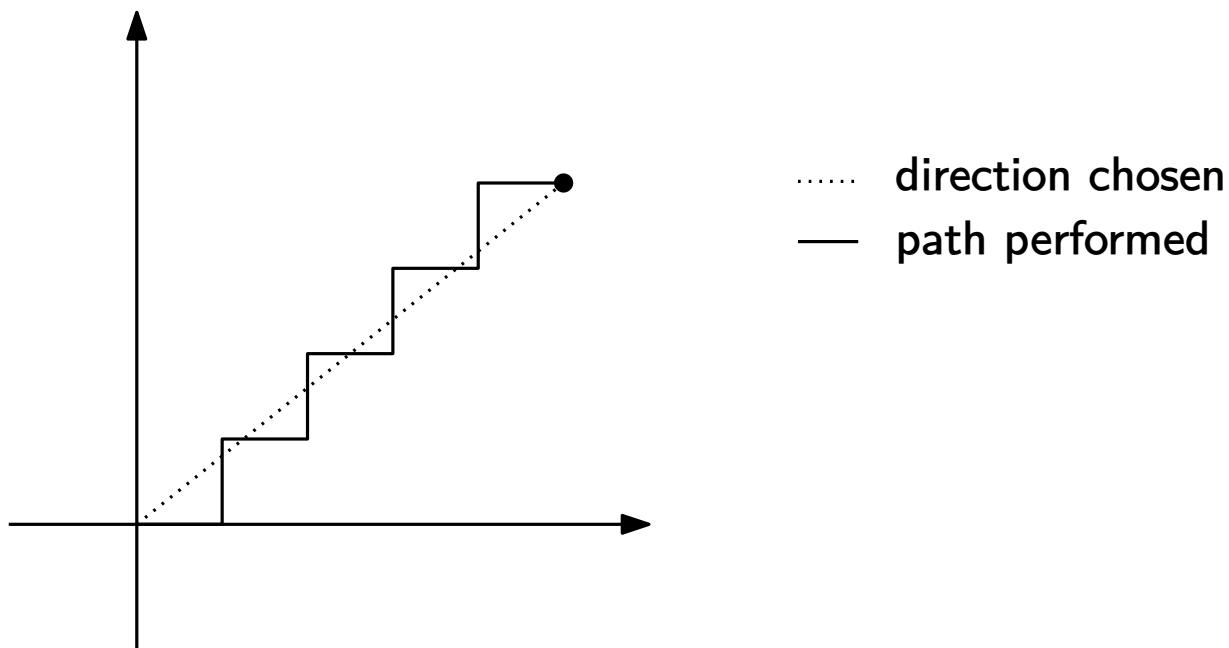
Two choices to make:

- define the **jump-length distribution**
- define a **notion of approximating** a **line-segment**

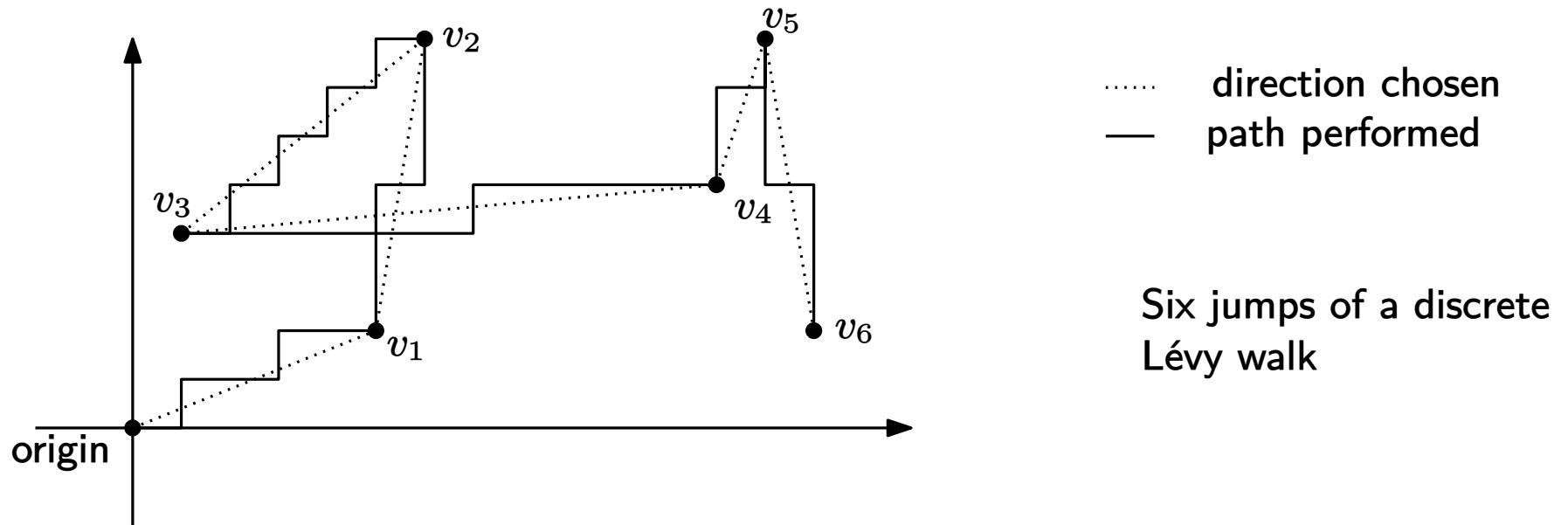
## Jump length distribution

- $d = 0$  with probability  $1/2$
- $d \geq 1$  with probability  $c_\alpha/d^\alpha$

## Approximation of a line-segment



# Discrete Lévy walk



Let  $\alpha > 1$  be a real value

**Lévy walk:** the agent

- a) chooses a **distance**  $d \in \mathbb{N}$  as follows:  $d = 0$  w.p.  $1/2$ , and  $d \geq 1$  w.p.  $c_\alpha/d^\alpha$
- b) chooses a **destination** u.a.r. among those at distance  $d$
- c) walks along an **approximating path** for  $d$  steps, one edge at a time, crossing  $d$  nodes
- d) **repeats** the procedure

# No advice, no communication

[Feinerman et Korman, DC 2017] proposes many solutions to the problem

Many considered settings, in which

- agents **exchange information** at the source node
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Their **best algorithm** in this case achieves **expected hitting time**

$$\mathcal{O} \left( (\ell^2/k + \ell) \log^{1+\epsilon} \ell \right),$$

for any fixed constant  $\epsilon > 0$

# No advice, no communication

Uniform algorithm proposed in [Feinerman et Korman, DC 2017]

(idea)

- i fix a subset of  $k_i$  **agents** to be moved
- ii fix a **ball** of some radius  $\ell_i$
- iii agents go to **random nodes** in the ball
- iv agents perform a **spiral search** of length  $d_i$  around the chosen nodes
- v agents **return** to the source node
- vi increase  $k_i$ ,  $\ell_i$ ,  $d_i$  and repeat (i)-(v)

However, the above algorithm is not that **natural**

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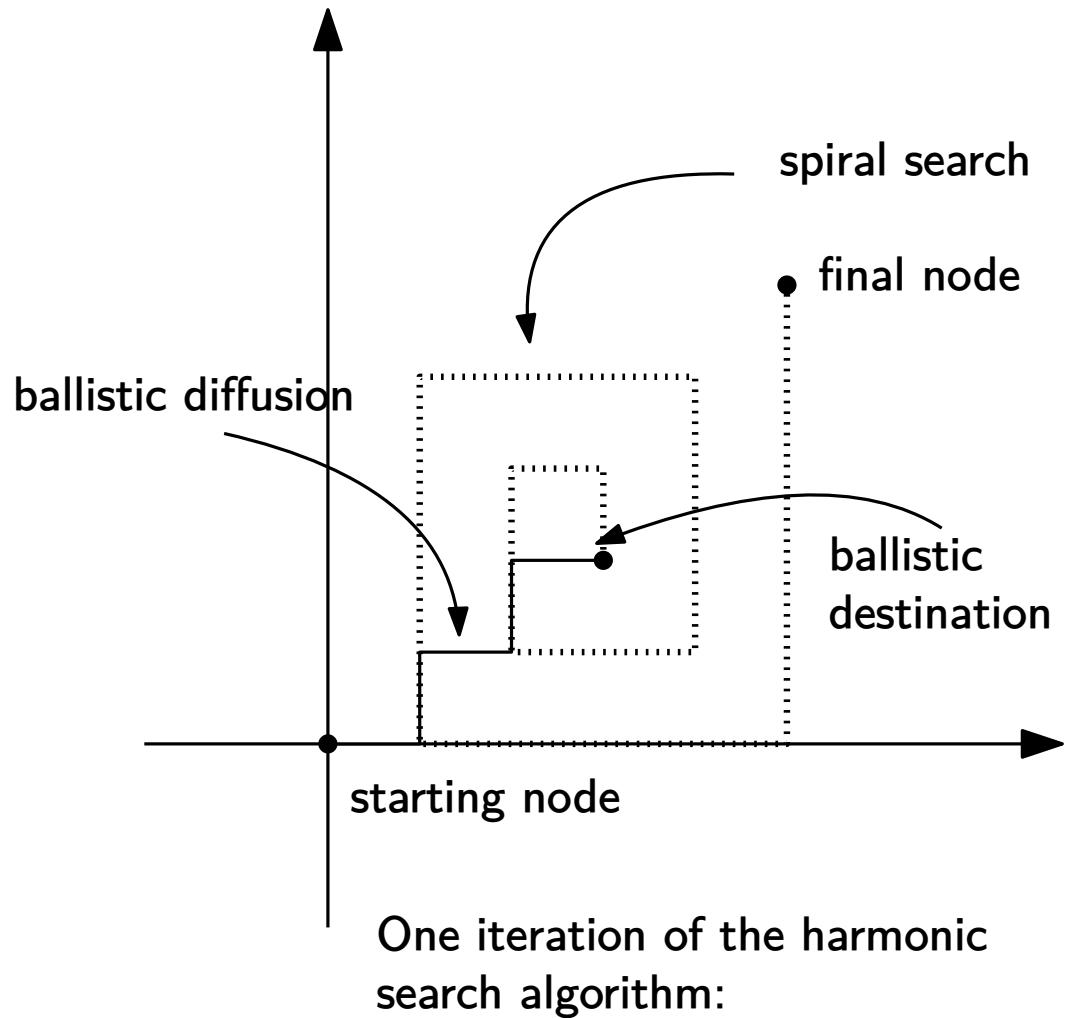
[Feinerman et Korman, DC 2017] proposes a more natural algorithm, the **Harmonic search algorithm (HSA)**

# The Harmonic search algorithm

HSA worsens performance, but increases probability: the hitting time is

$$O(\ell^{2+\delta}/k + \ell)$$

with probability  $1 - \epsilon$  for any fixed constants  $0 < \delta, \epsilon < 1$



# The Harmonic search algorithm

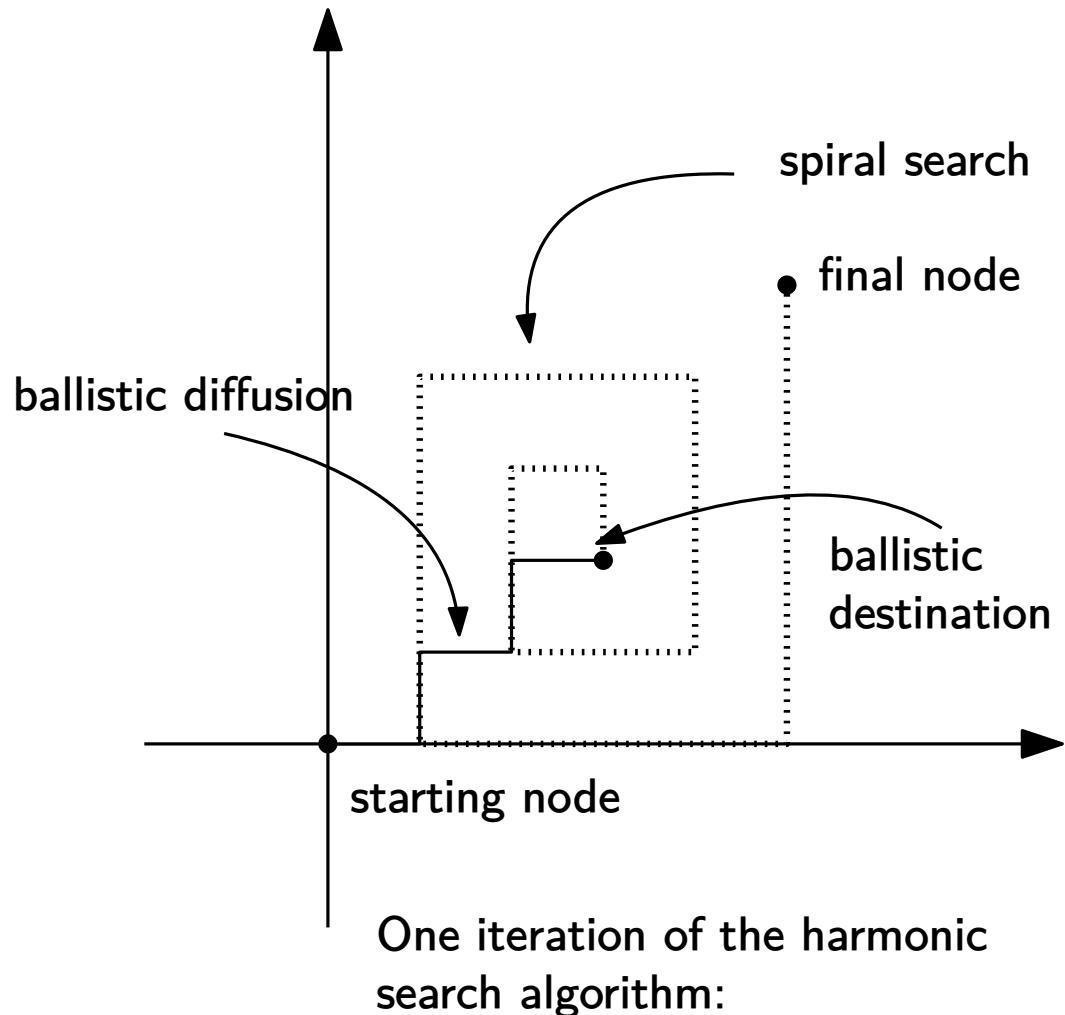
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**HSA:** each agent

- a) samples a **jump-length  $d$**  with a **power-law distribution** with exponent  $\alpha = 1 + \delta$  (small)
- b) (**ballistic diffusion**) moves to a destination at distance  $d$  chosen u.a.r.
- c) (**normal diffusion**) starts a spiral search for  $d^{\delta+2}$  steps
- d) **returns** in the origin and repeats



# Appendix

1. Lévy walks
2. Opinion dynamics
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# Techniques

For **consensus**:

- **symmetry breaking**: the bias has enough **standard deviation** to break symmetry (**drift analysis** results)
- applying **concentration inequalities** to construct a process  $M_t$  such that  $M_{t+1} \leq (1 - \delta)M_t$  w.h.p. as long as the **bias is outside**  $I_\varepsilon = [(1 - \varepsilon)\bar{s}, (1 + \varepsilon)\bar{s}]$  but at least  $\Omega(\sqrt{n \log n})$
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- constructing a super-martingale  $N_t$  such that  $\mathbb{E}[N_{t+1} | \mathcal{F}_t] \leq (1 - \delta)N_t$  involving the bias
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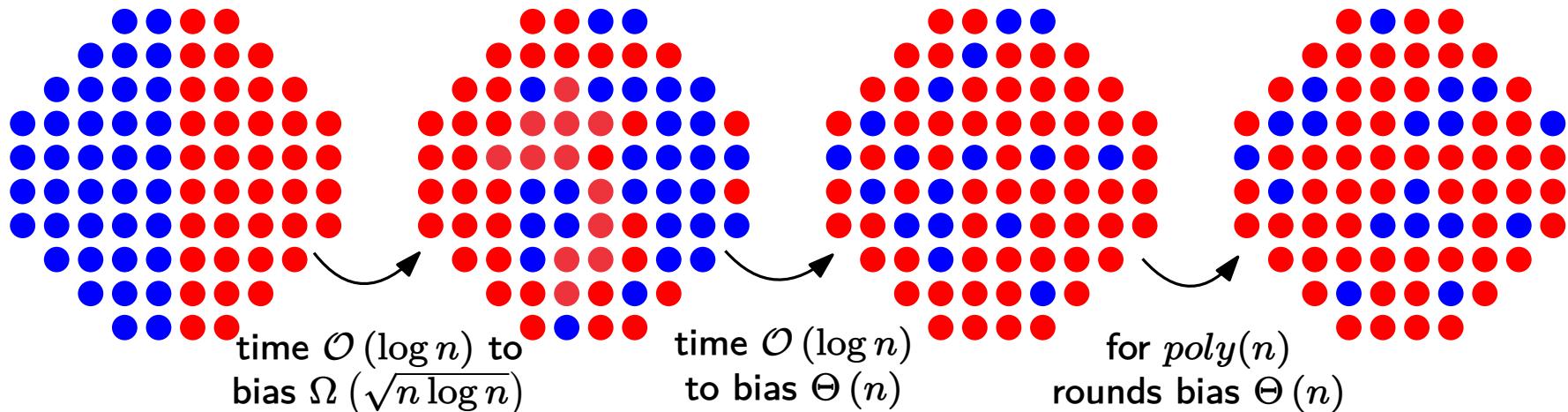
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Example: consensus



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# Assemblies

**Hypothesized** by [Hebb, 1949]: densely interconnected **sets of neurons** whose loosely synchronized **firing** in a **pattern** is **simultaneous** with the subject thinking of a particular **concept** or **idea**

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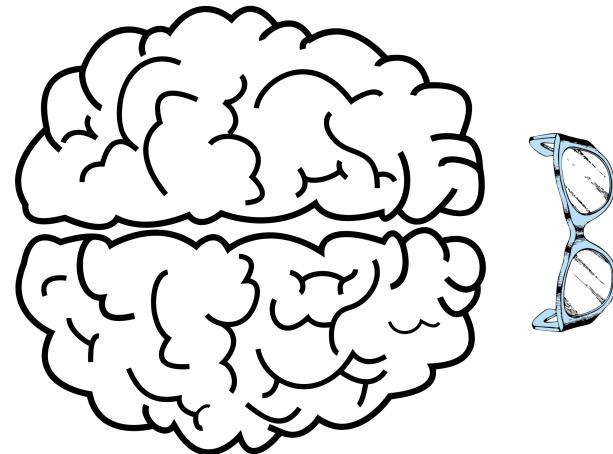
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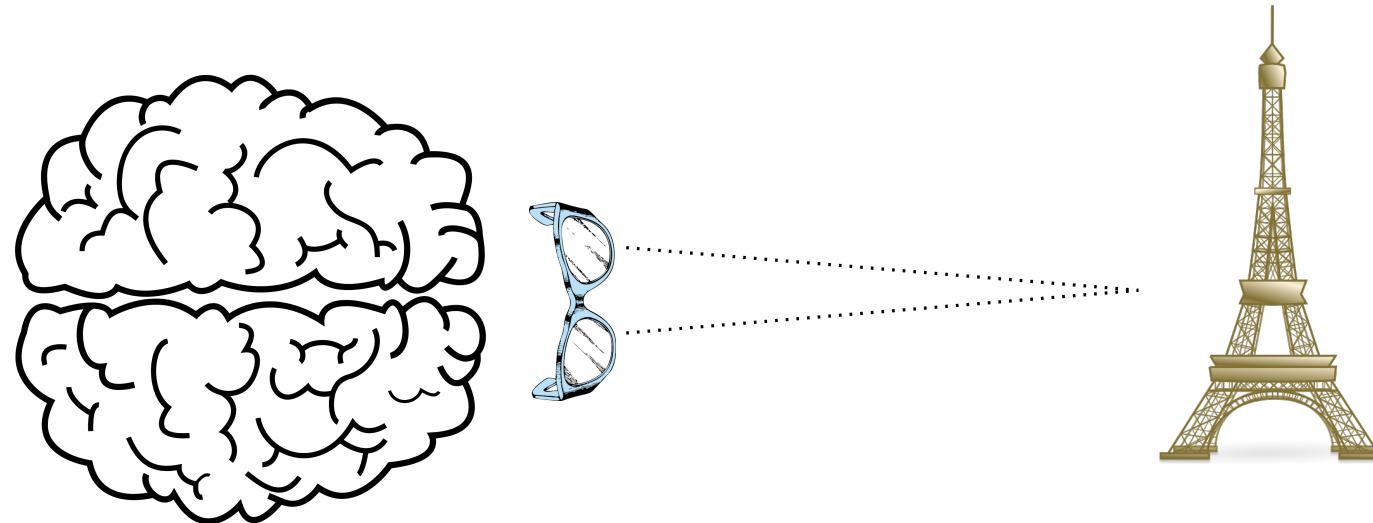
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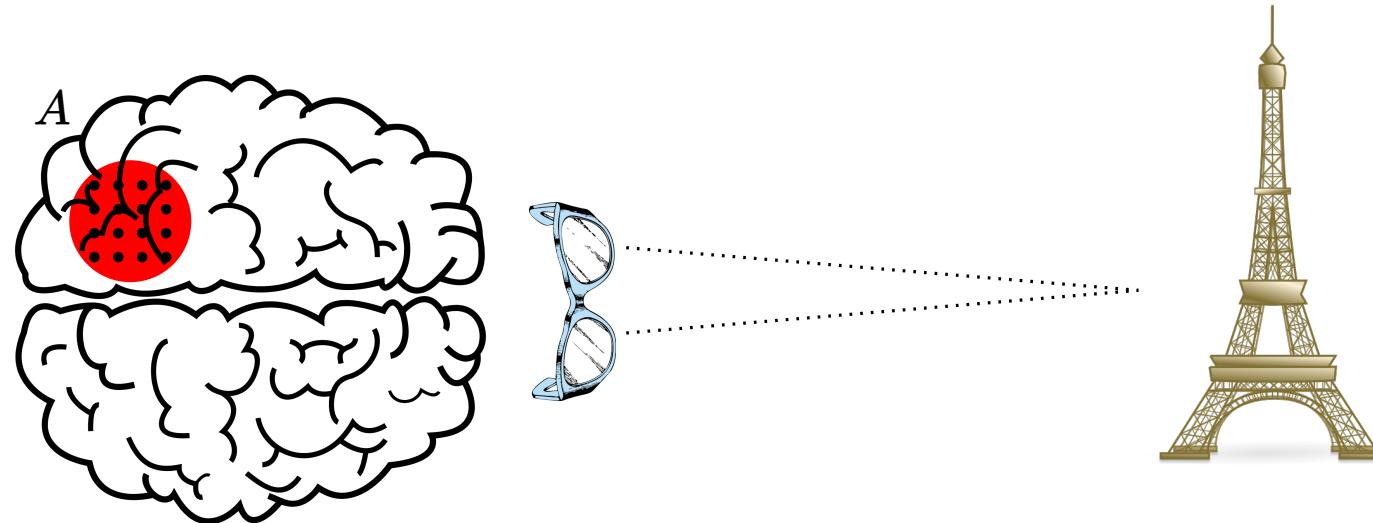
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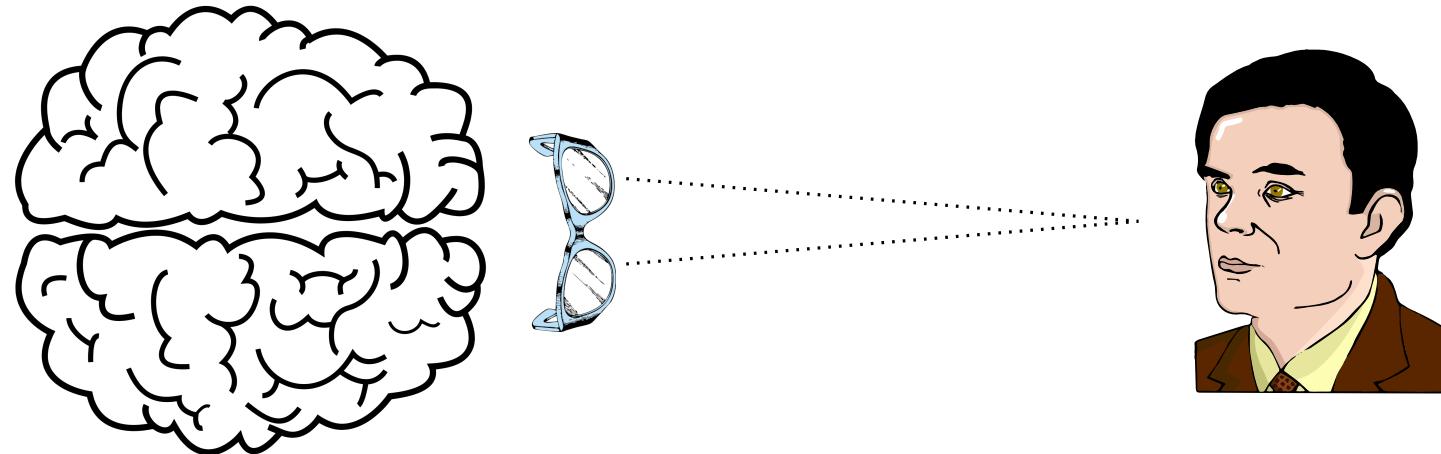
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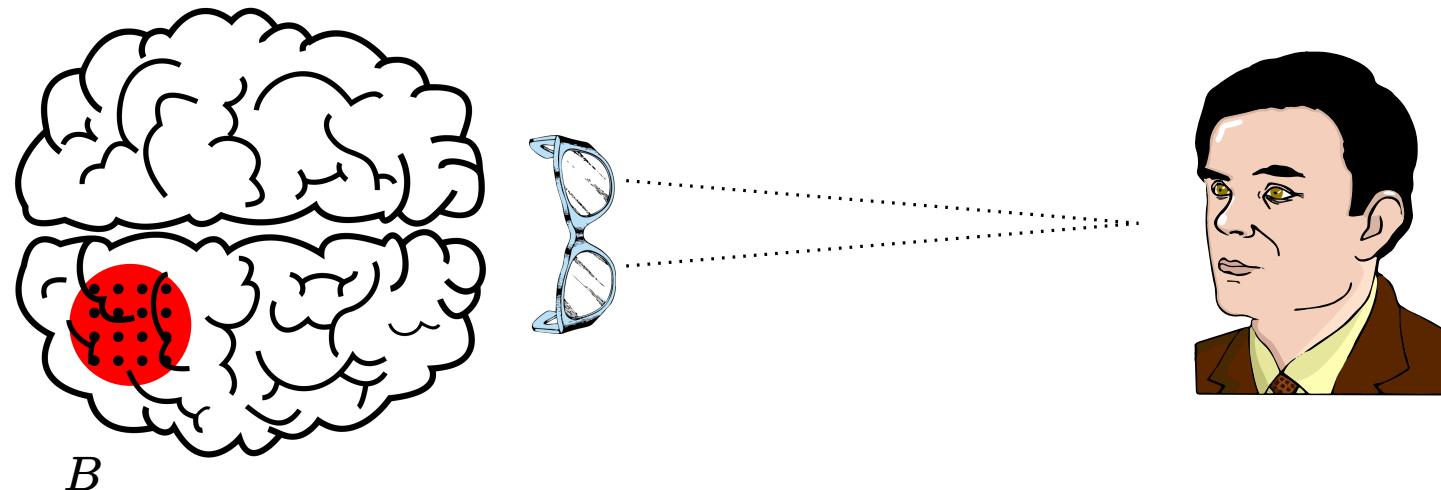
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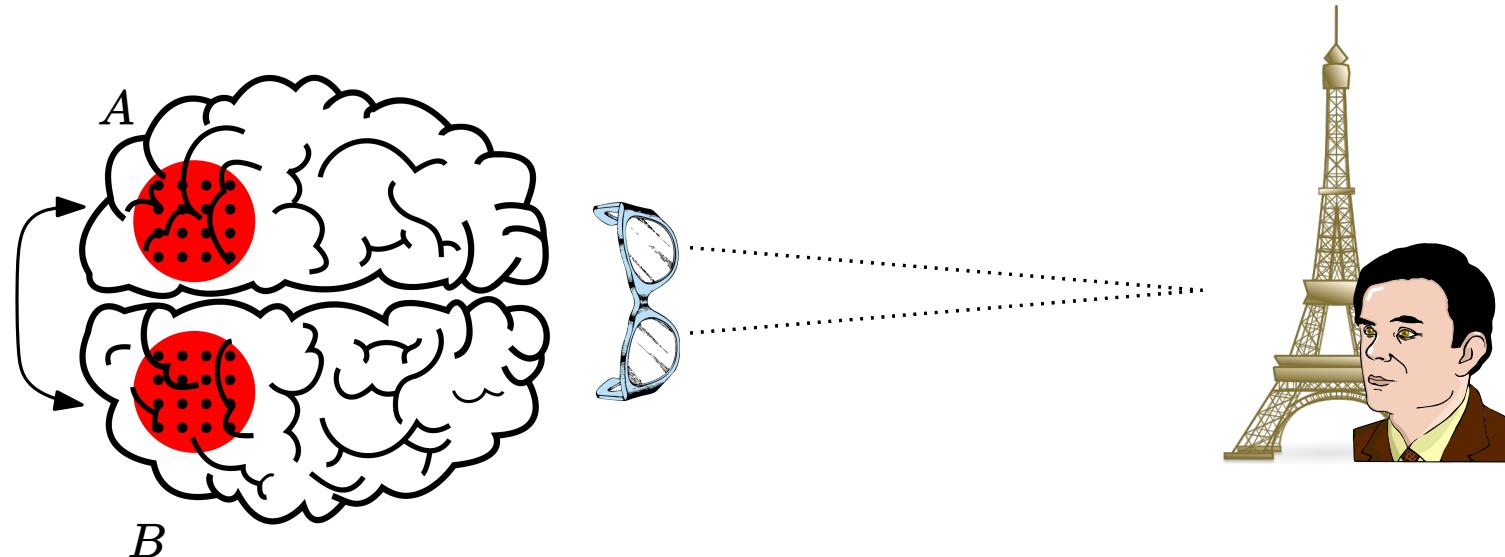
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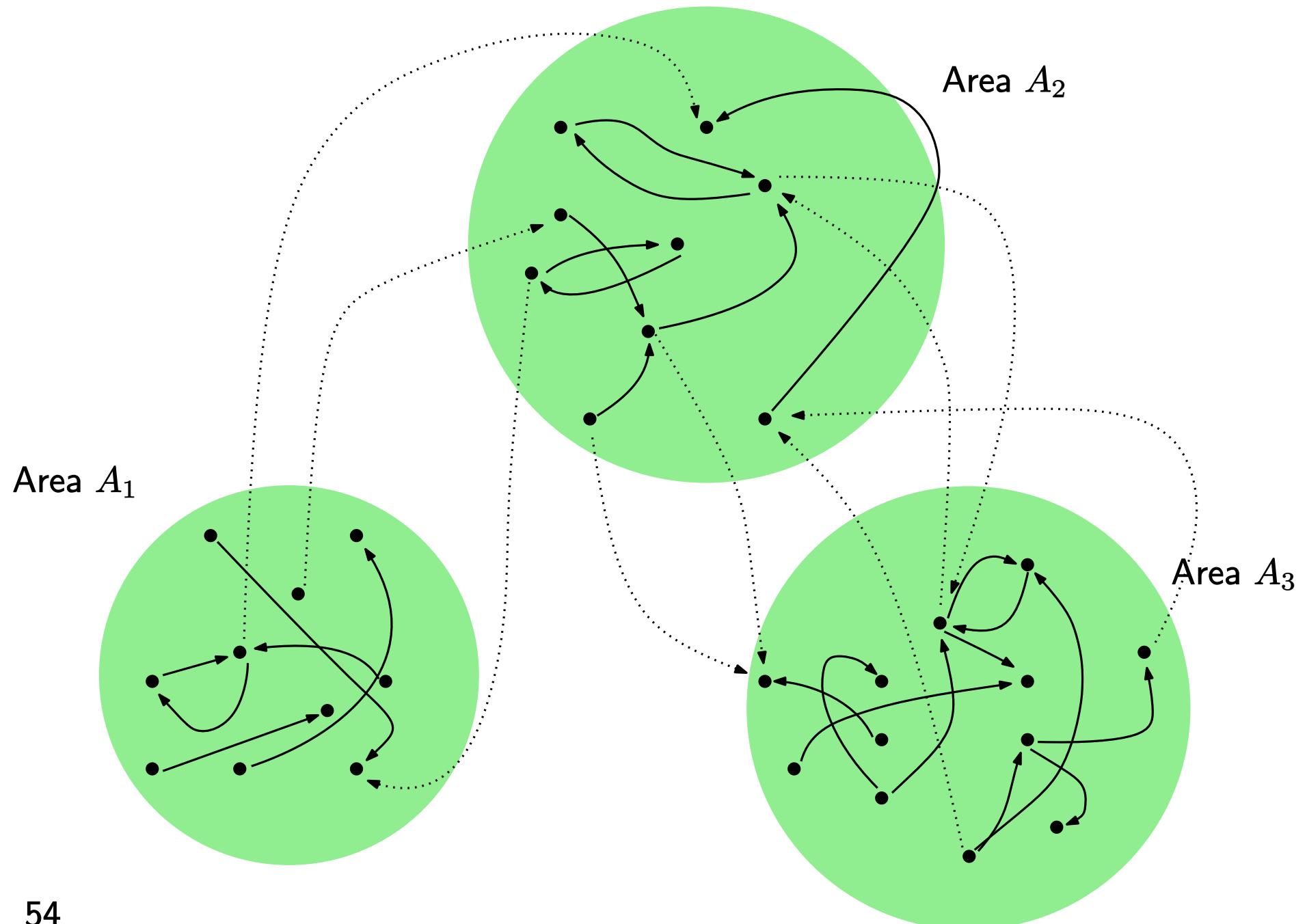
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- **Brain area:**  $A = G_{n_A, p_A}$  Erdös Rényi graph
- *Recall*  $G_{n,p} = (V, E)$ :
  - $|V| = n$
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- set of brain areas  $\mathcal{S} = \{A_1, A_2, \dots, A_m\}$
- $\mathcal{C} \subseteq \mathcal{S} \times \mathcal{S}$  set of ordered pairs of brain regions
- **The Brain:**  $\mathcal{B} = (V_{\mathcal{B}}, E_{\mathcal{B}})$  with
  - $V_{\mathcal{B}} = V_{A_1} \cup \dots \cup V_{A_m}$
  - $E_{\mathcal{B}} = E \cup E_{A_1} \cup \dots E_{A_m}$
  - $E$ :  $\forall (A, B) \in \mathcal{C}, \forall x \in A, y \in B, (x, y) \in E$  with probability  $p_{A,B}$

# Example of a Brain



# The AC dynamics

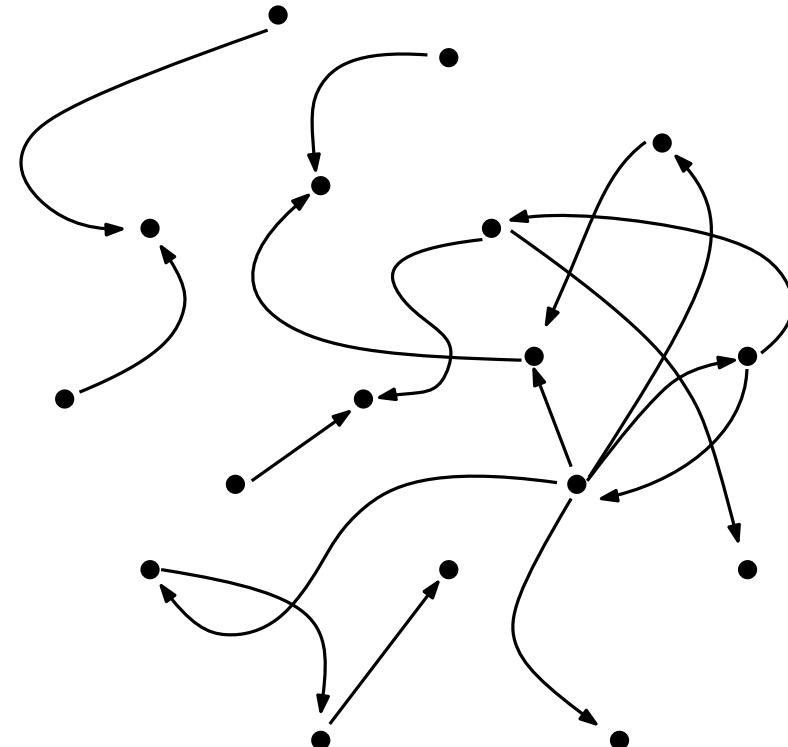
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All **fiber weights** initialized to be 1

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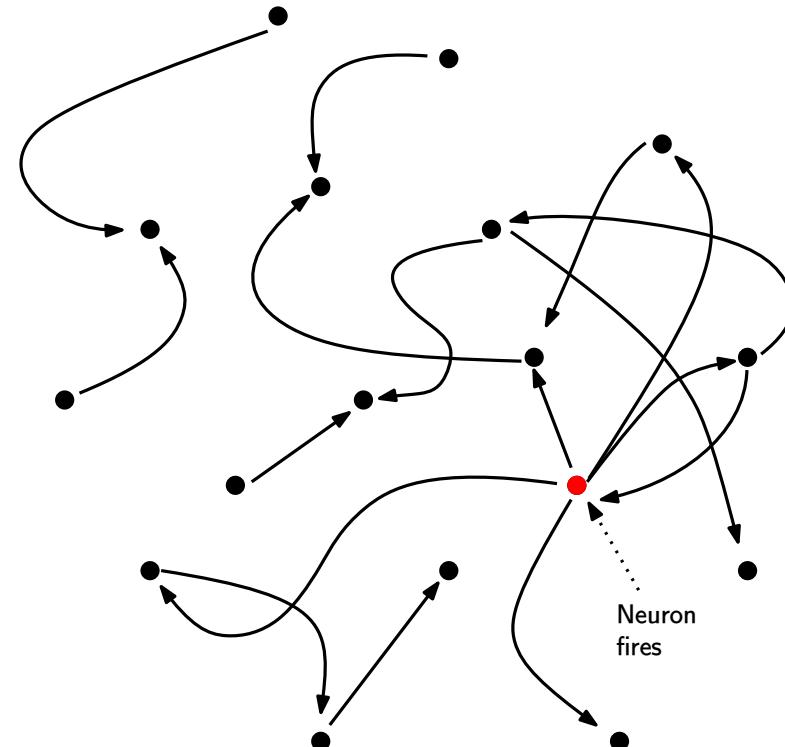
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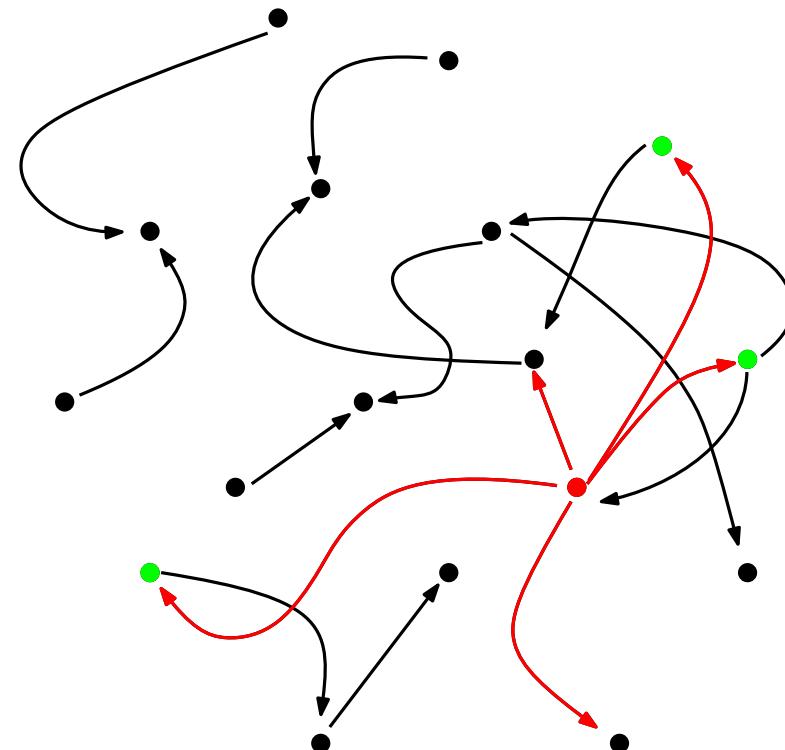
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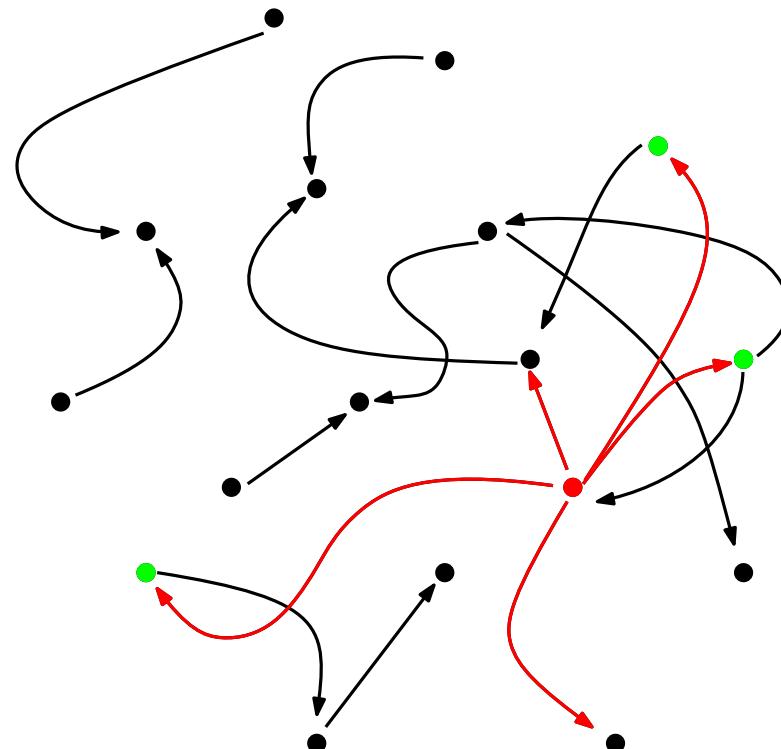
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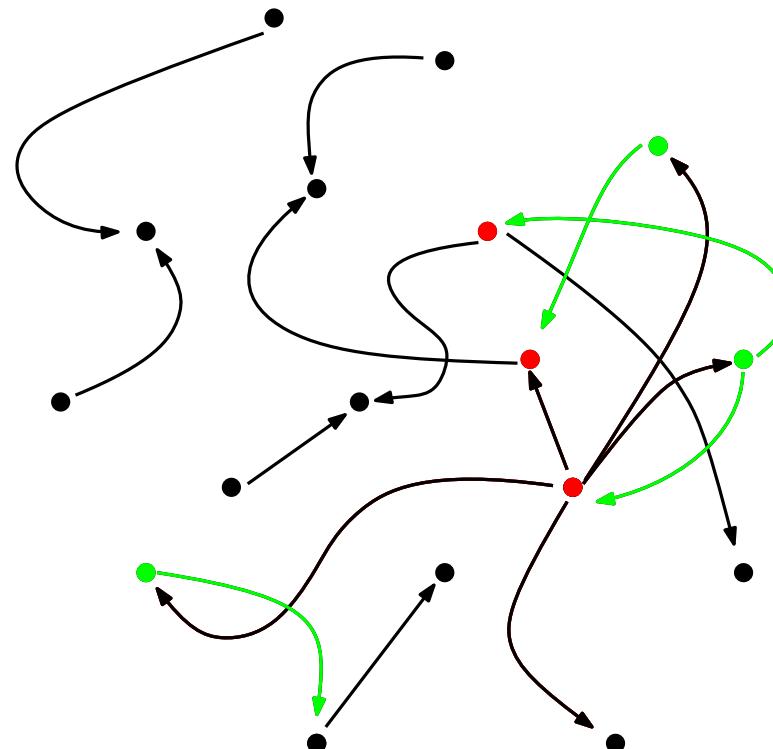
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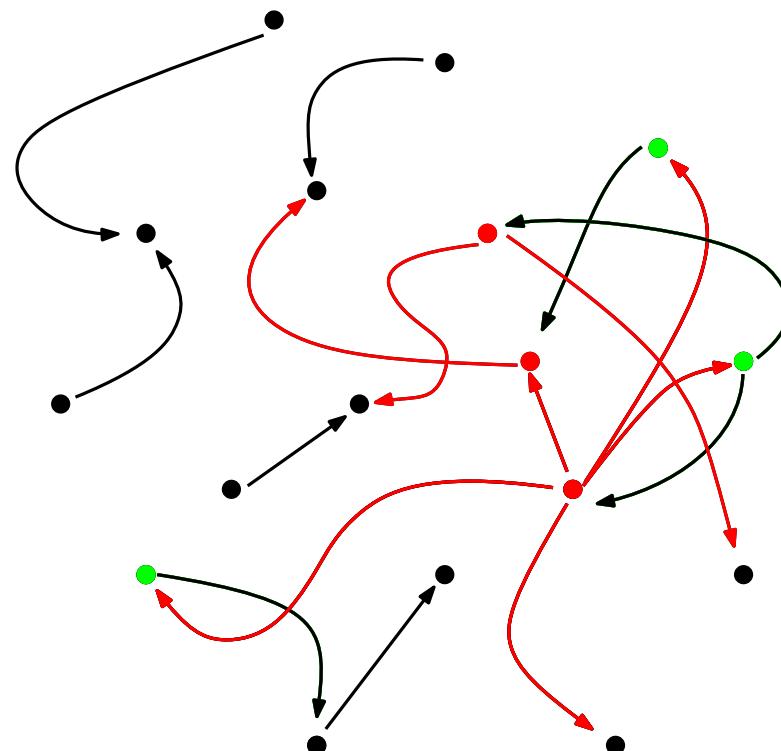
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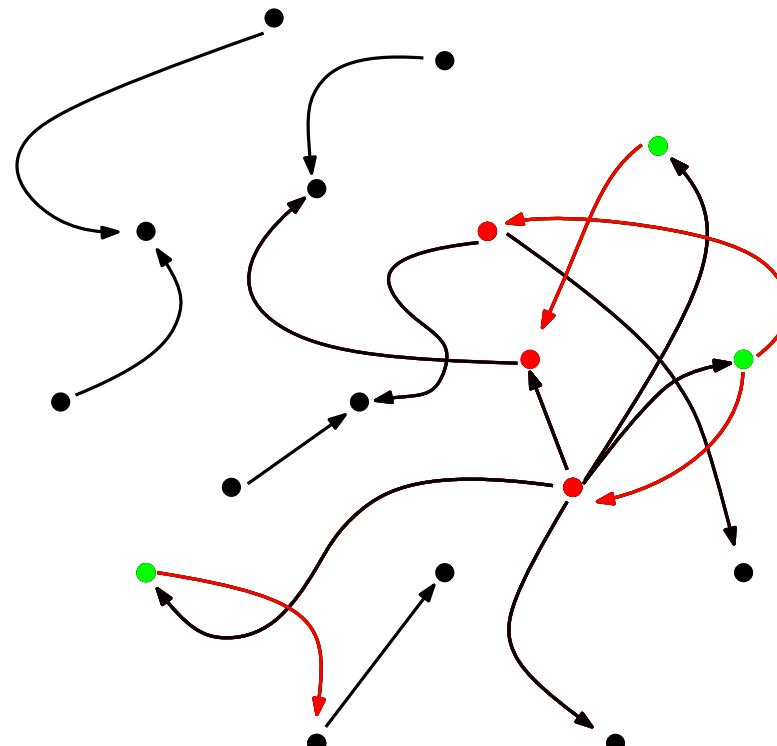
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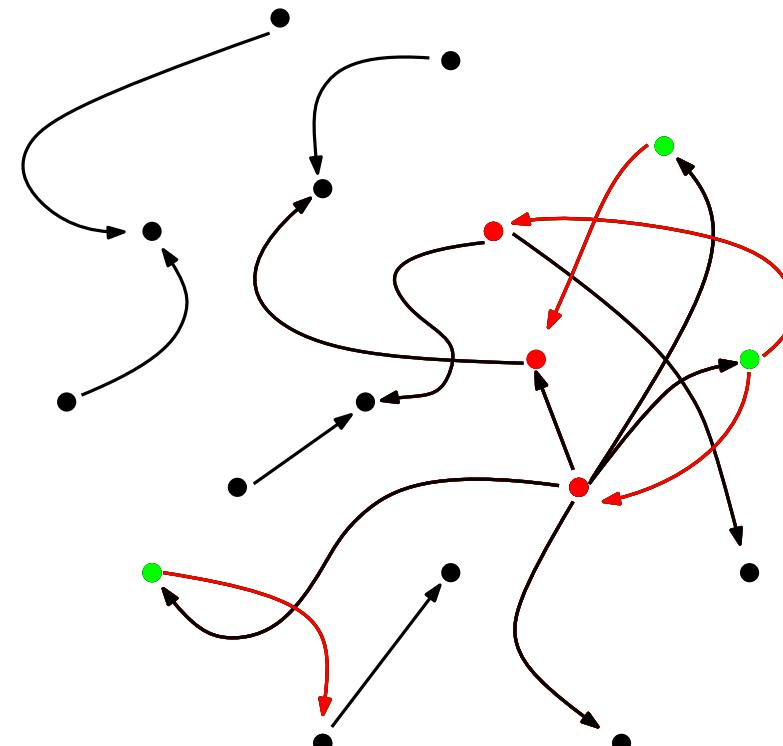
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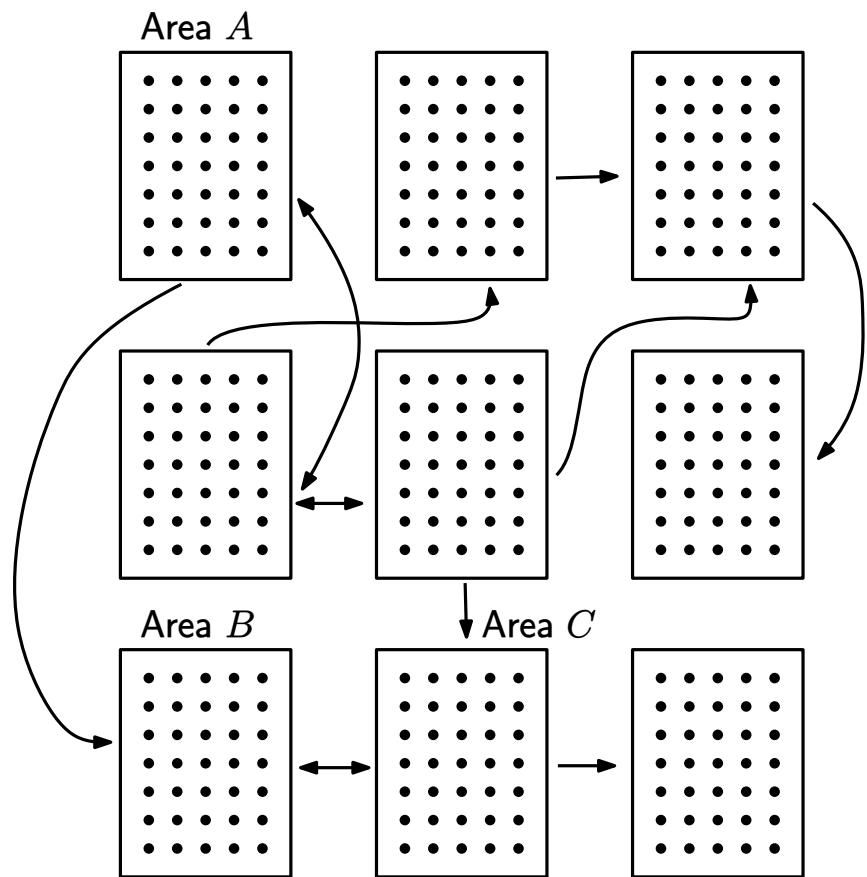
$$\text{weight} = \text{old\_weight} \cdot (1 + \beta)$$

**Hebbian plasticity**

**Assembly**: *stable* set of  $k$  neurons

# Operations

The brain and its areas

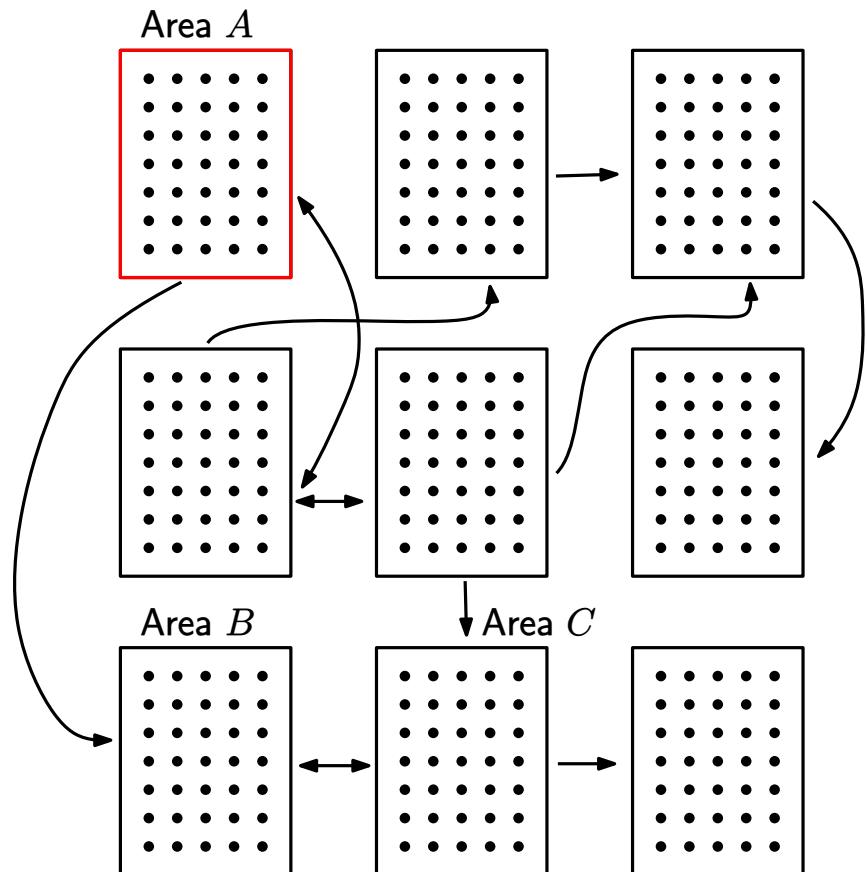


# Operations

- **Inhibition:** an area can be **inhibited**, that is, its neurons cannot fire

Inhibition is accomplished through populations of **inhibitory neurons**, whose firing prevents other neurons from firing [Mitropolsky et al., TACL 2021]

The brain and its areas



For areas  $A, B, C$ :

- `inhibitArea(A)`

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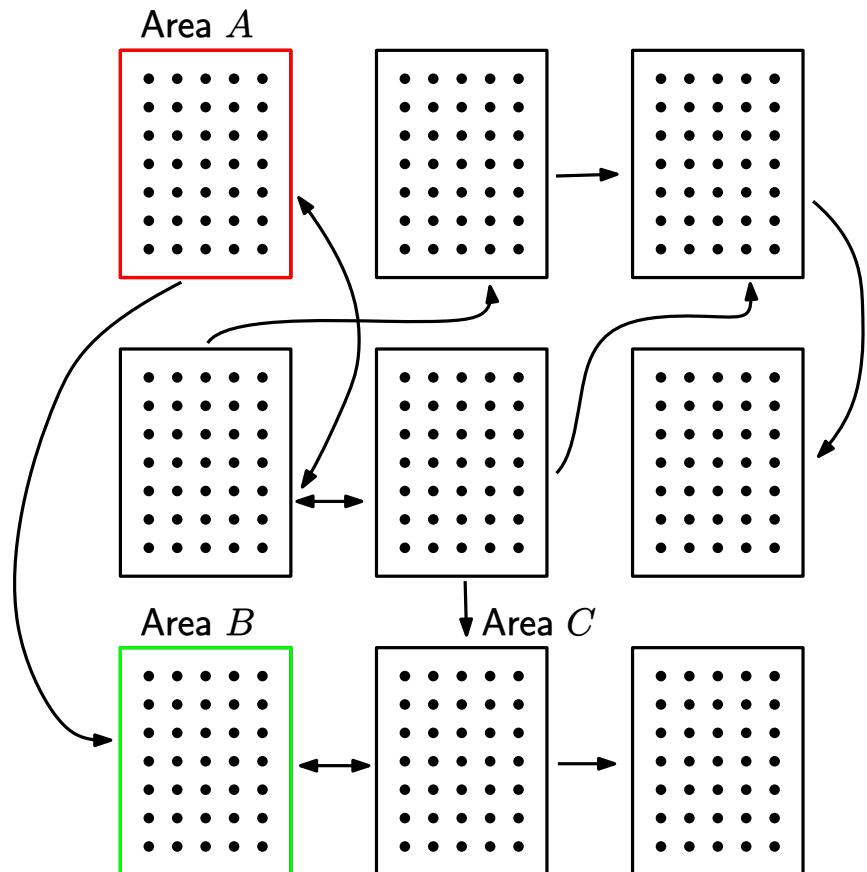
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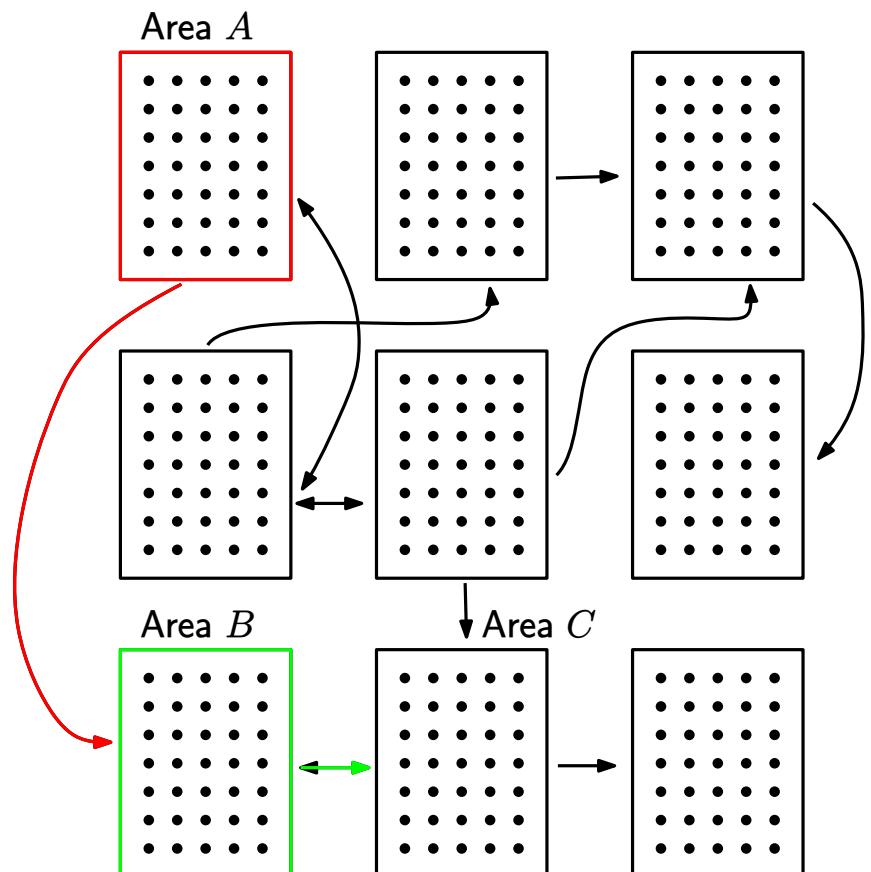
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We assume we can do the same with **fibers**

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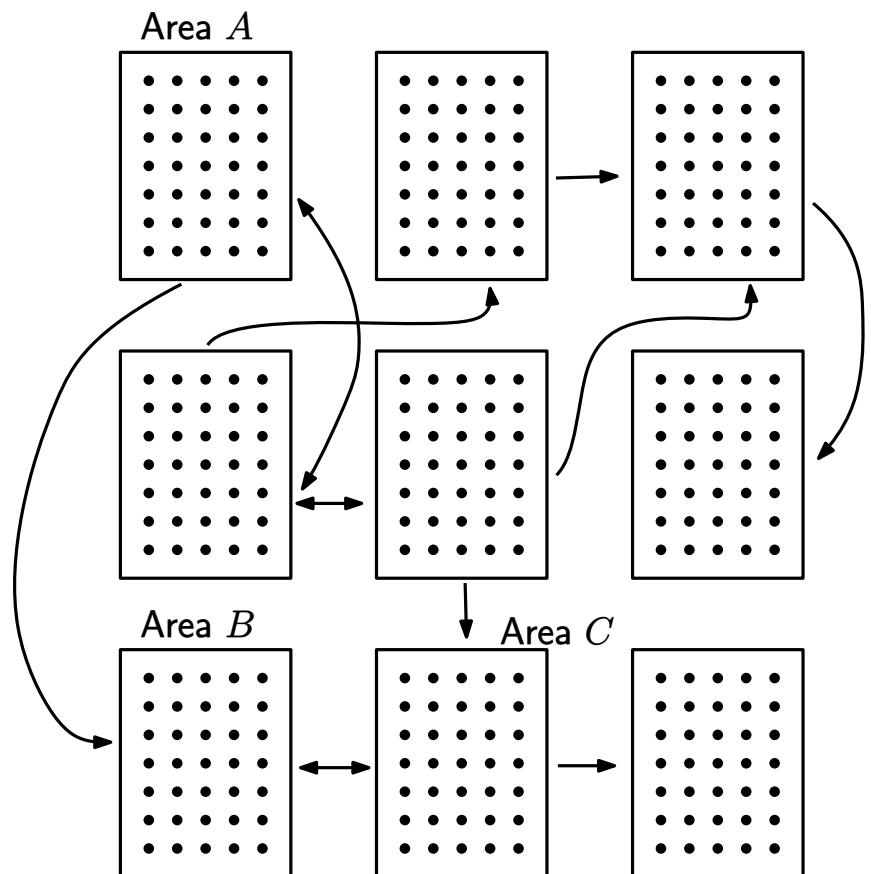
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The brain and its areas



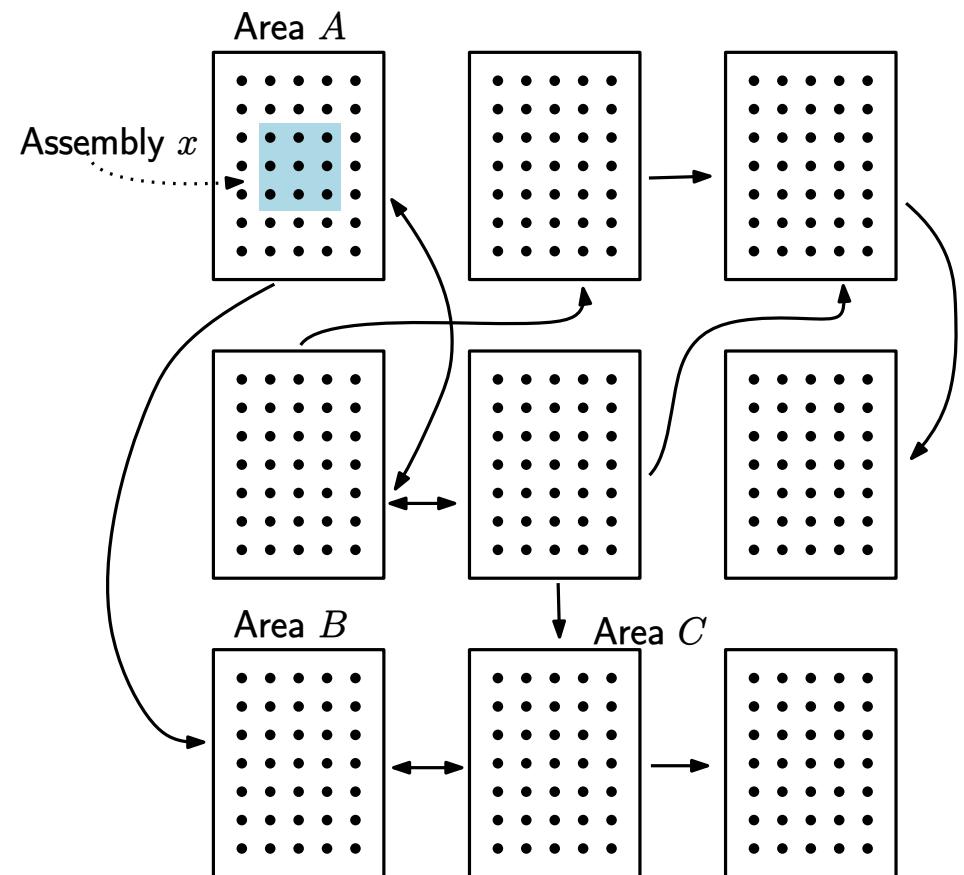
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The brain and its areas



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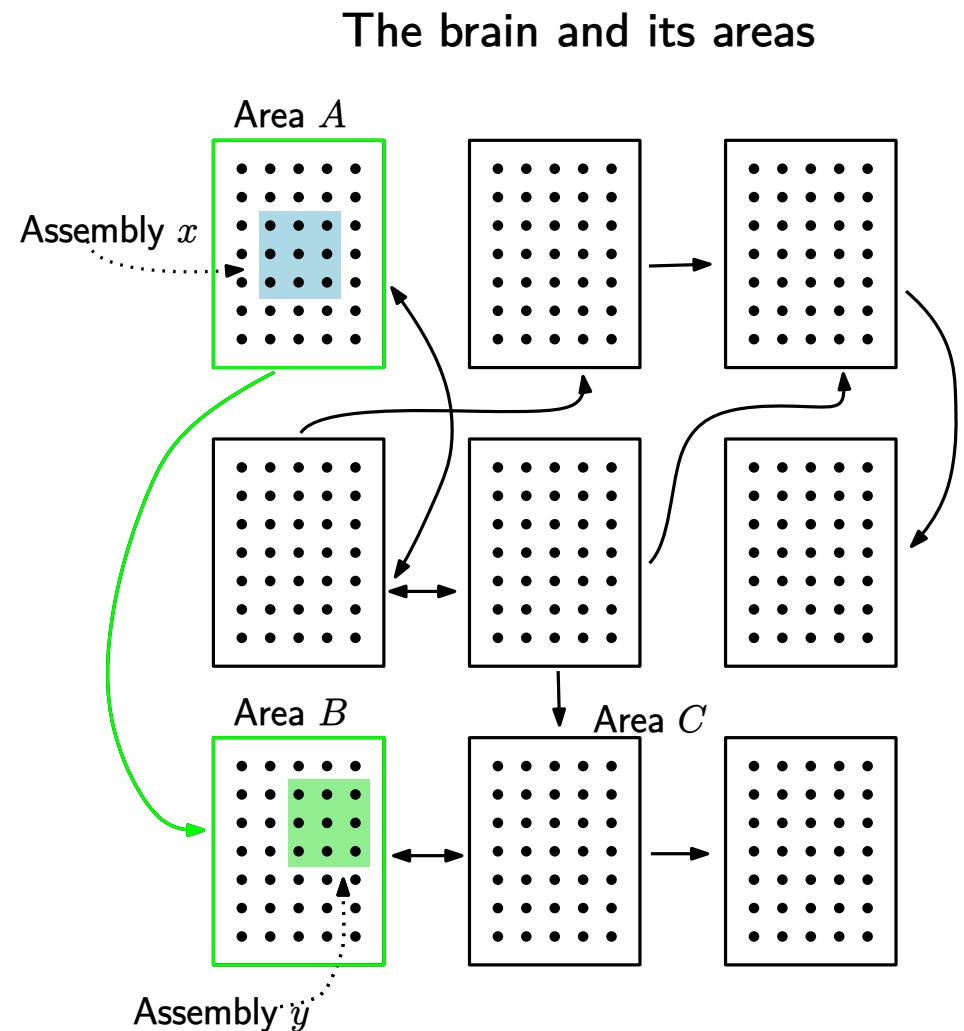
The brain and its areas



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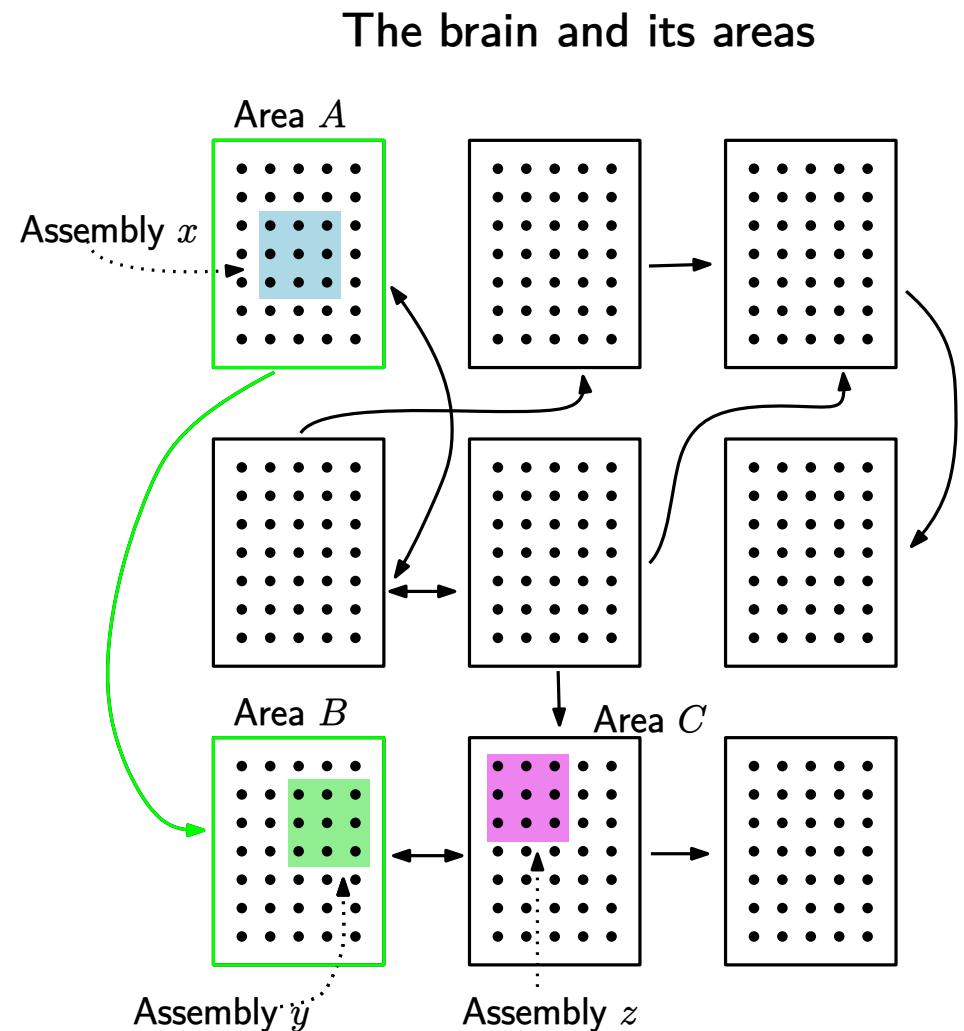
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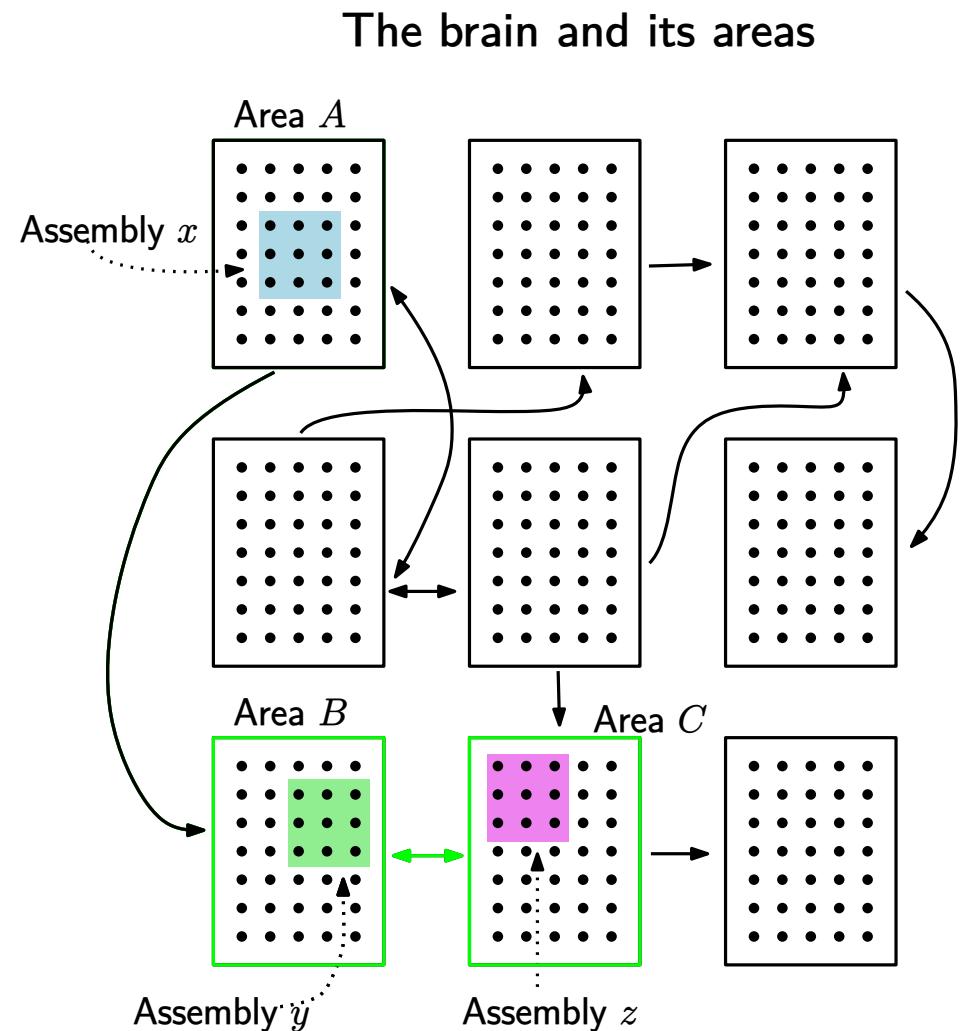
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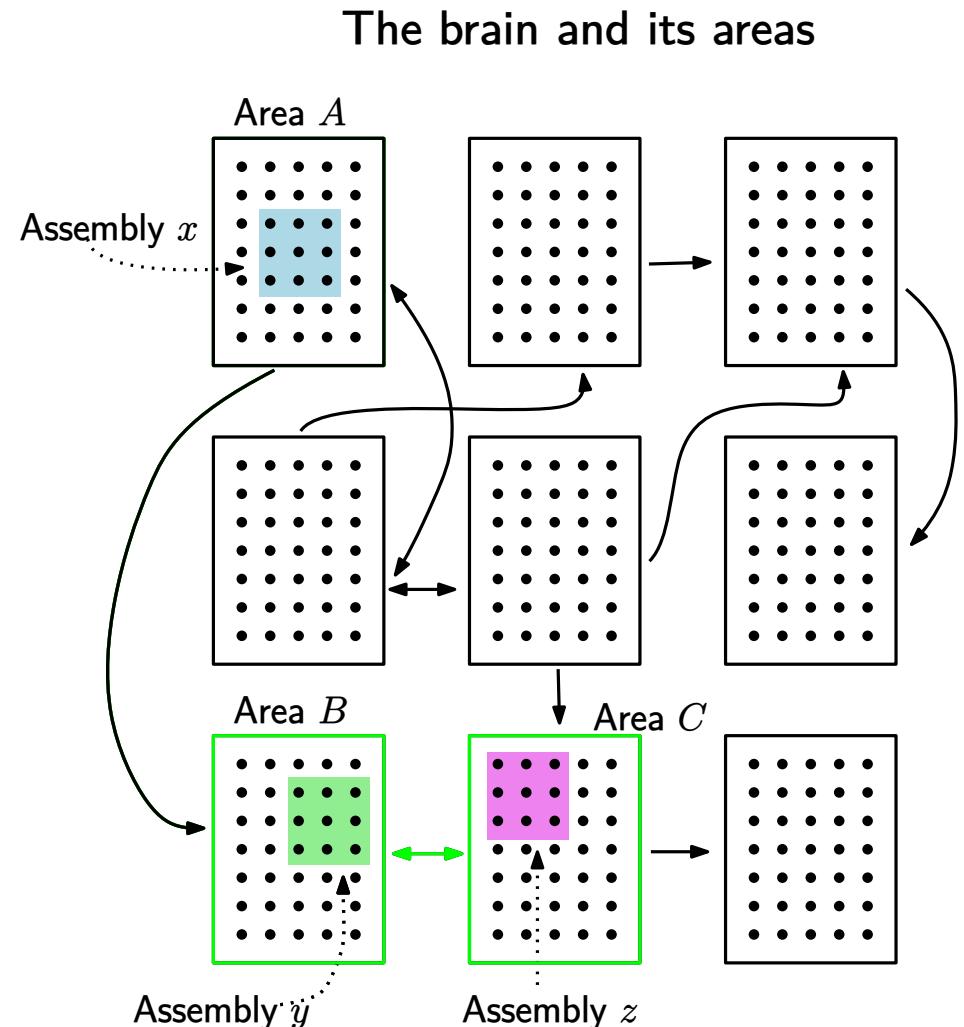
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For areas  $A, B, C$ :

- project  $(A, B)$
- associate  $(B, C)$



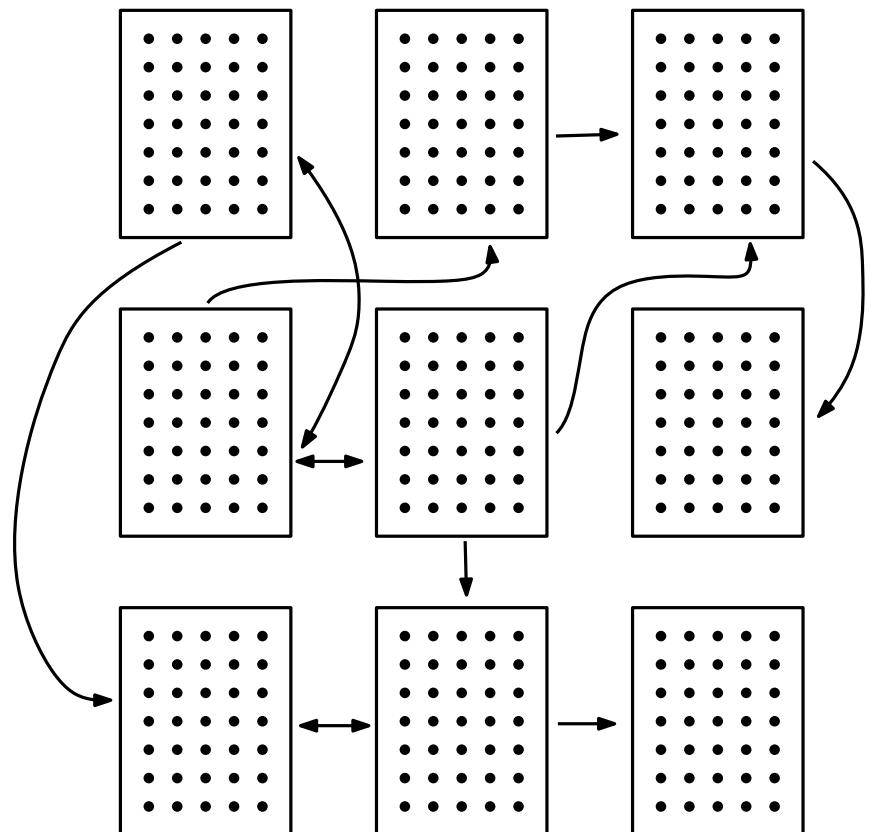
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- **Strong projection:** enhancement of the projection operation

Dynamics on the **subgraph induced** by all **disinhibited areas** and **fibers**:

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The brain and its areas



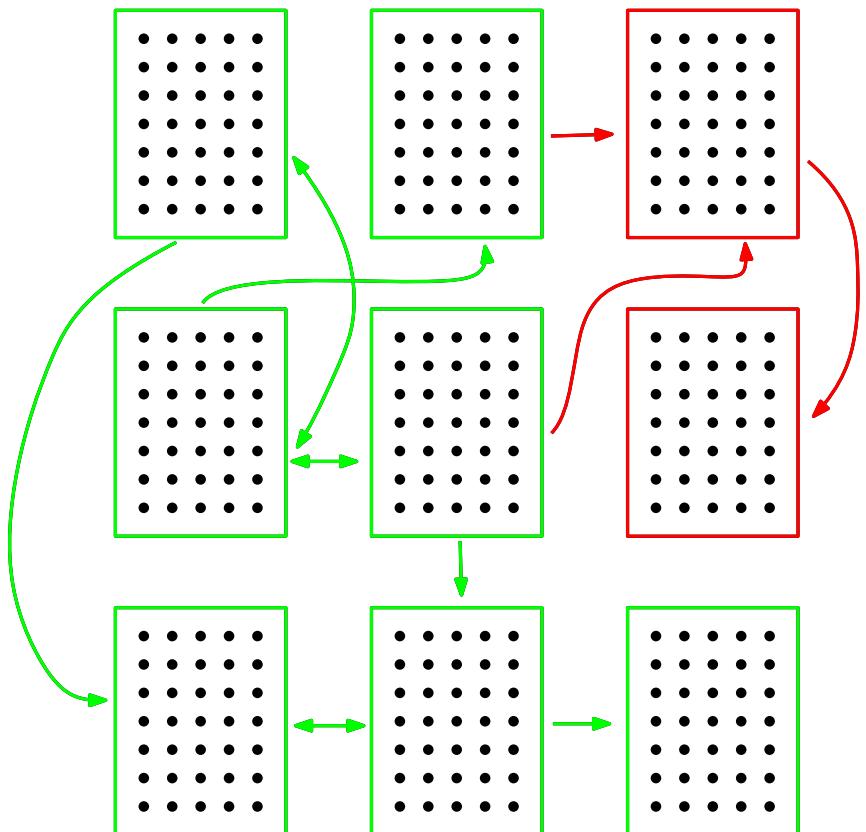
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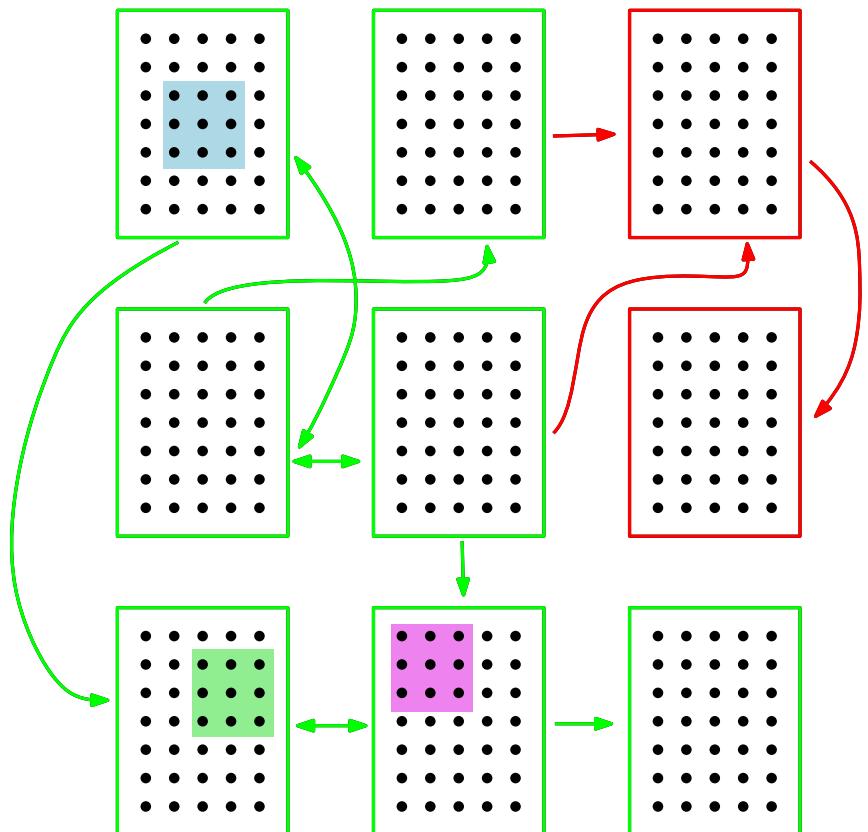
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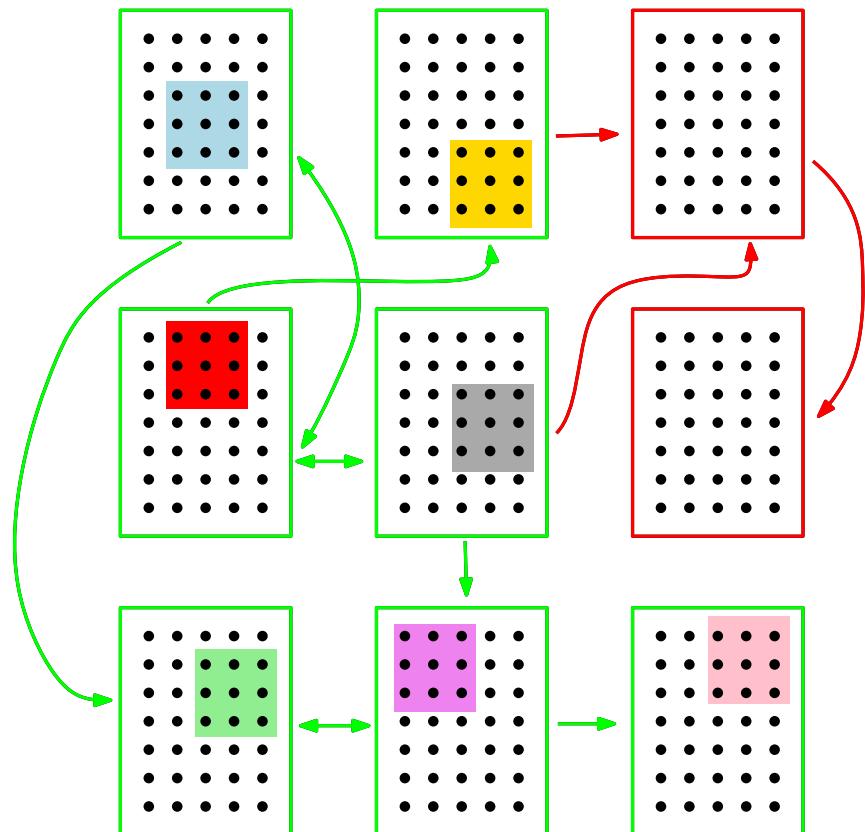
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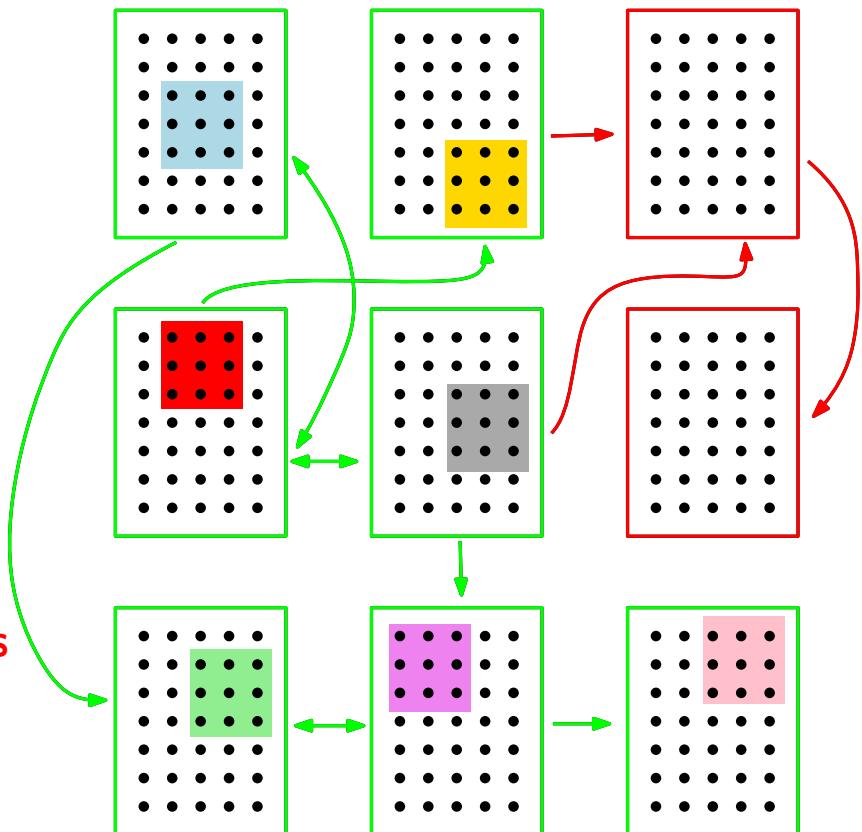
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The brain and its areas



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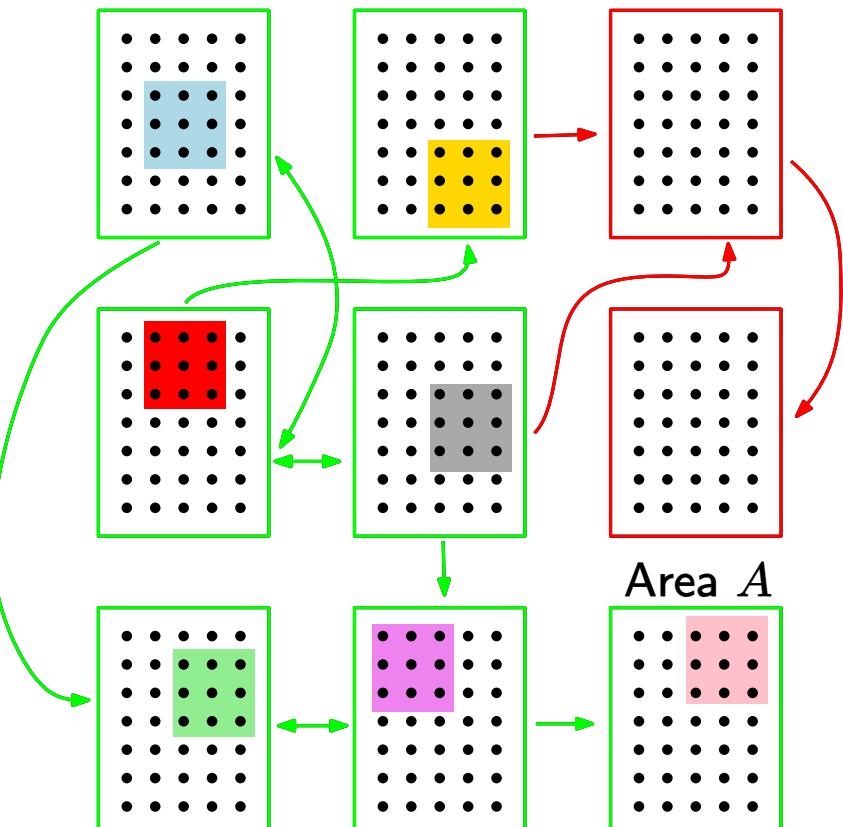
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We write **strongProject()** and **read( $A$ )** for a brain area  $A$

The brain and its areas



# Related Works

- [Legenstein et al., ITCS 2018] shows analytically how assemblies emerge from stimuli
- [Papadimitriou et Vempala, ITCS 2019] analyzes the convergence of processes defined by operations with assemblies
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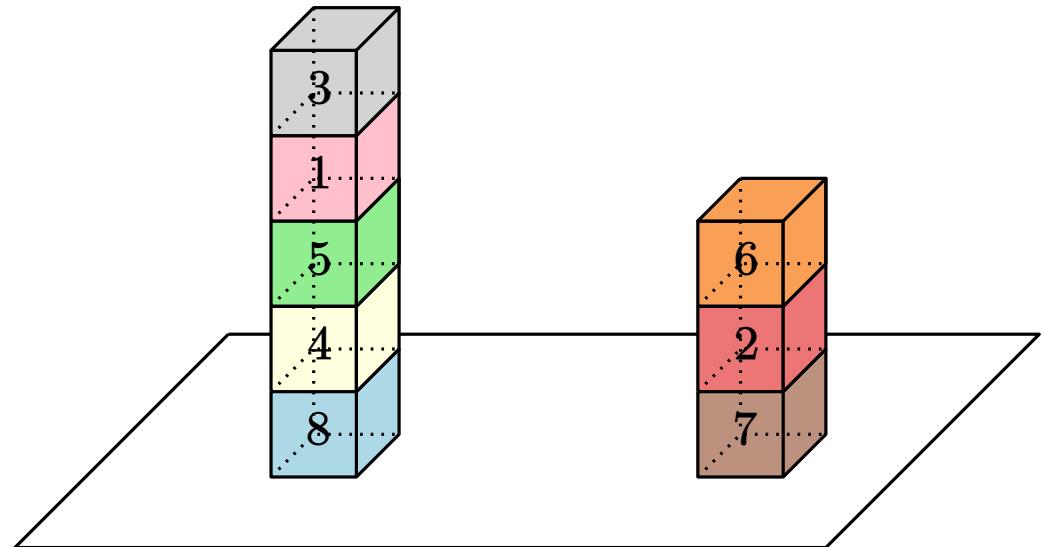
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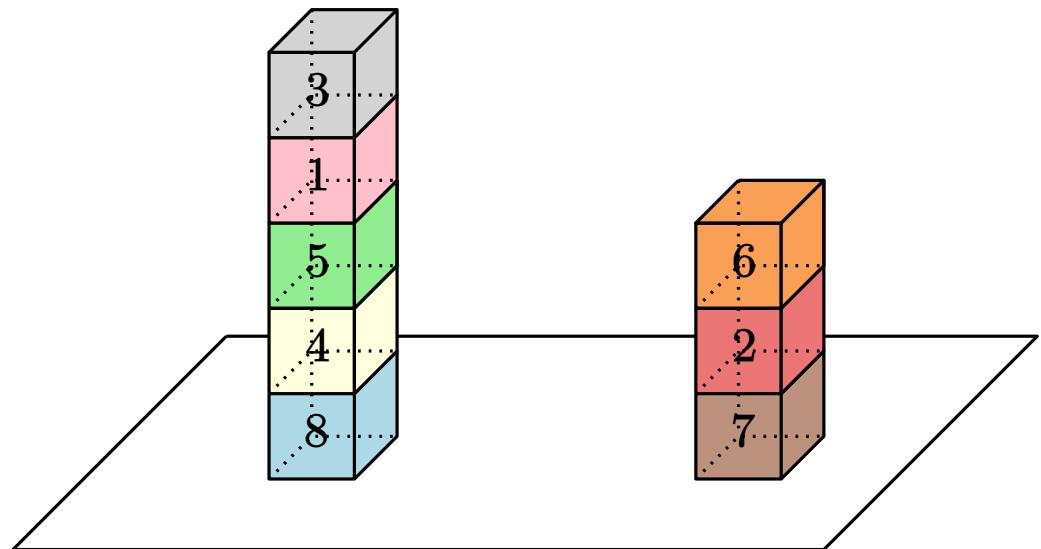


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**Blocks world:**  $n$  unique blocks **labelled** in  $\{1, \dots, n\}$ , organized as follows

- set of stacks  $\{S_i\}_i$
- each stack is a sequence of blocks  
 $S_i = (b_1^{(i)}, \dots, b_{k_i}^{(i)})$
- $\cup_i \{b_1^{(i)}, \dots, b_{k_i}^{(i)}\} = \{1, \dots, n\}$



# The Planning Task

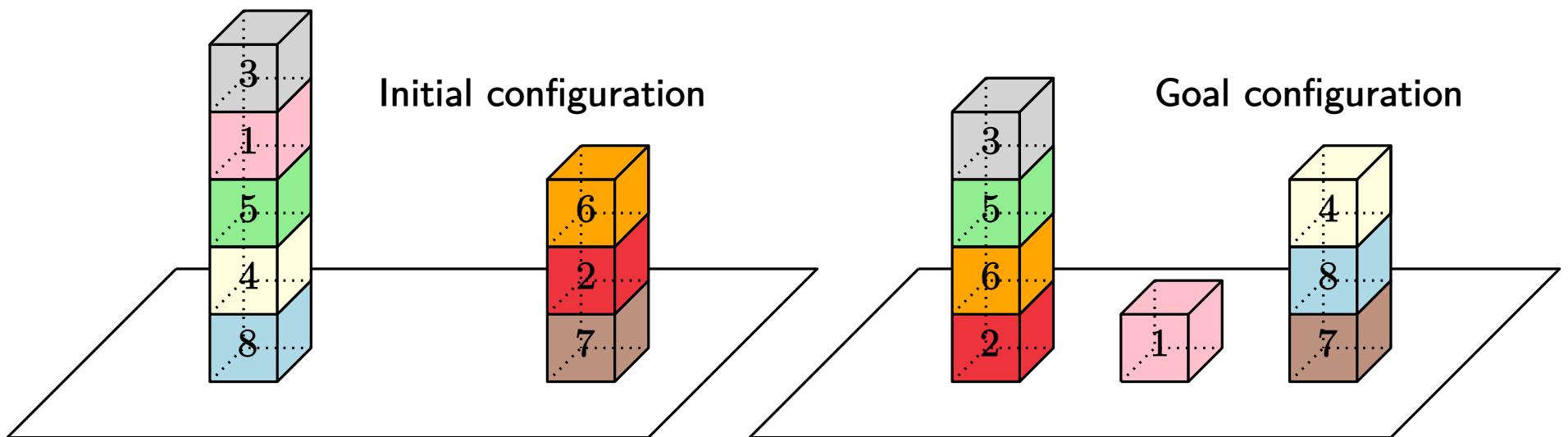
- Possible moves:**
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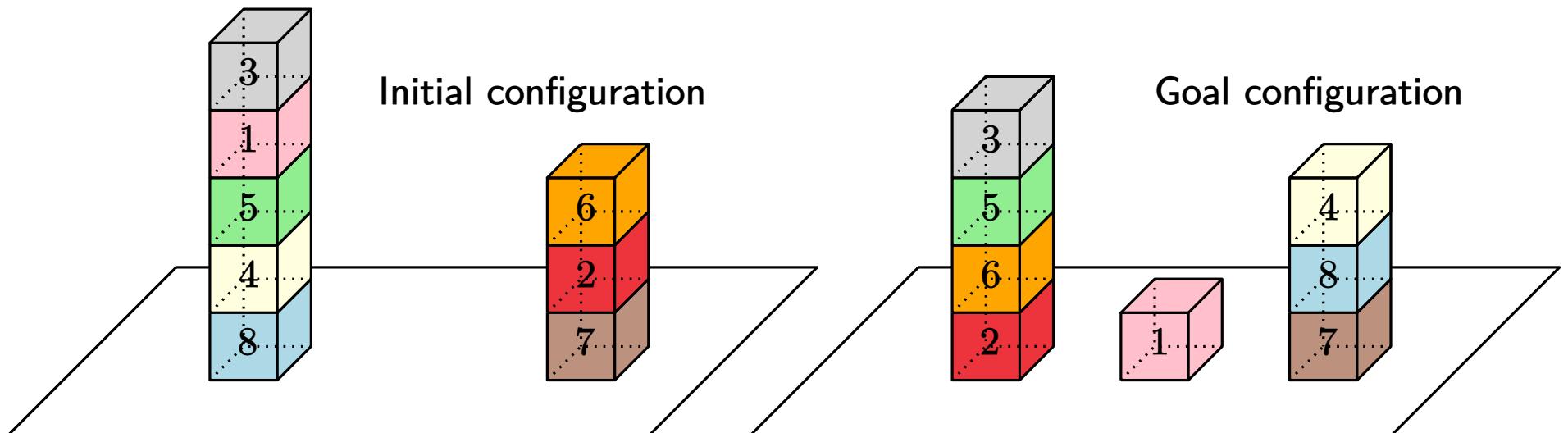
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**Output:**

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We implement both of them...

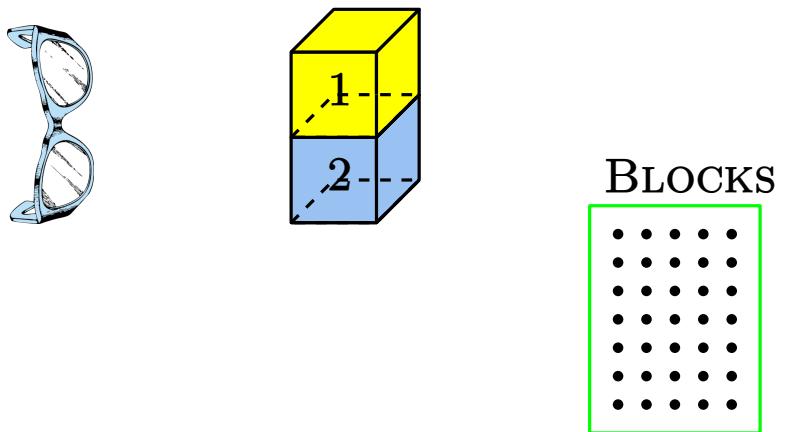
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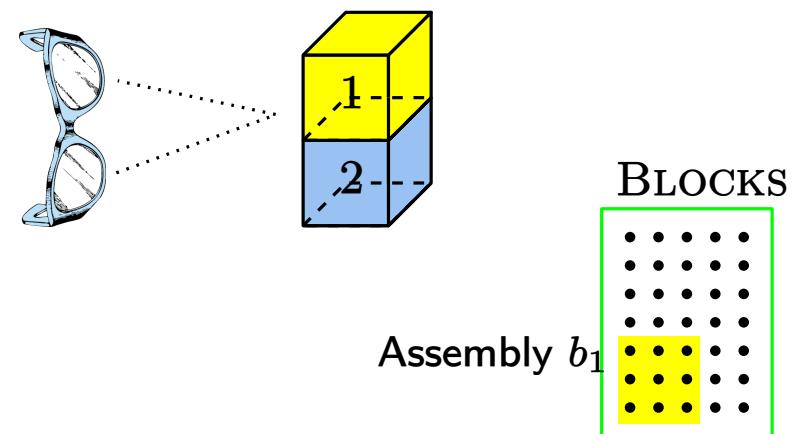
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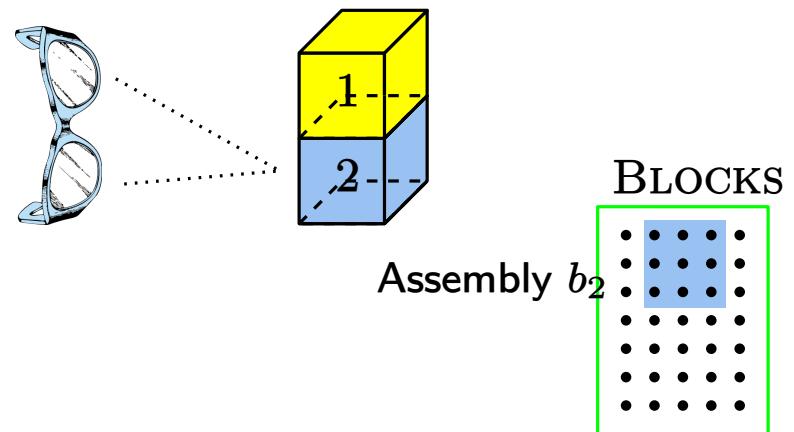
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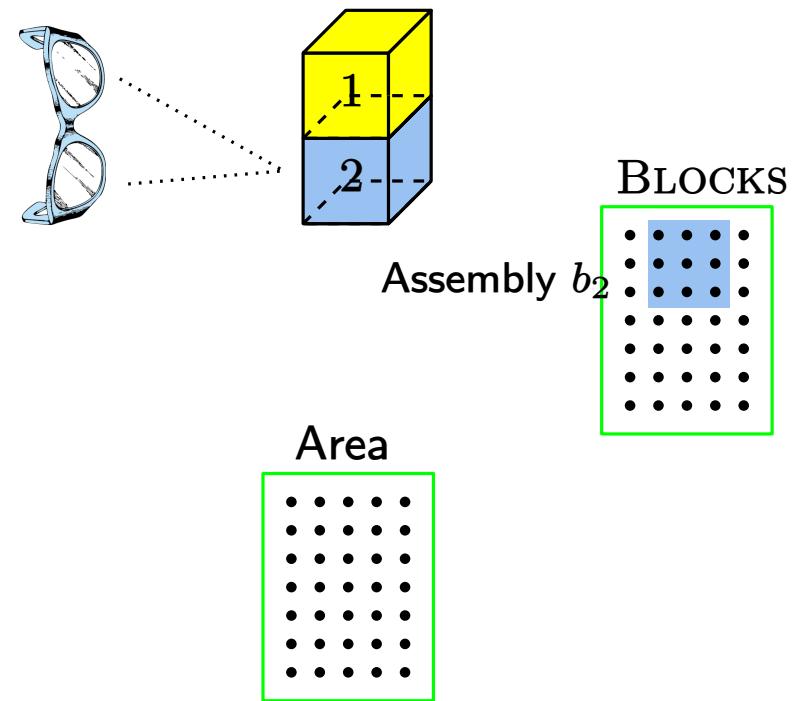


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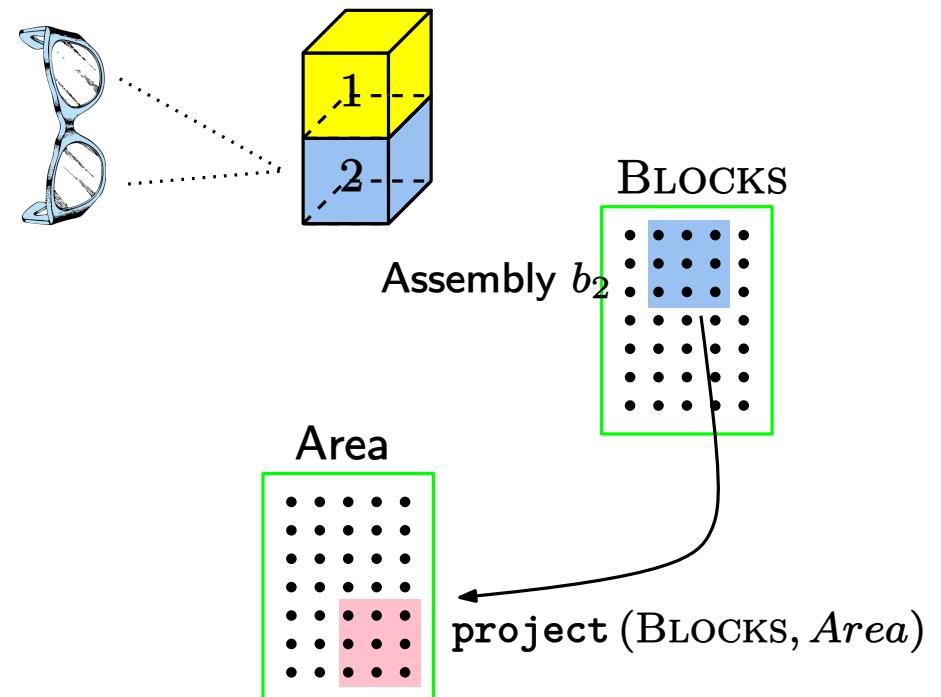


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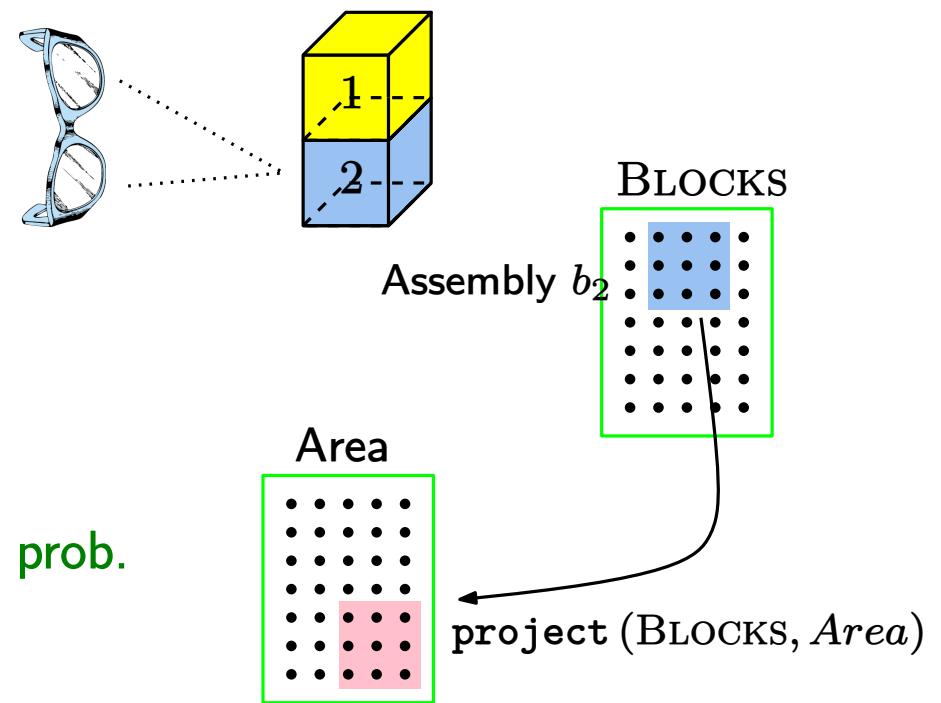
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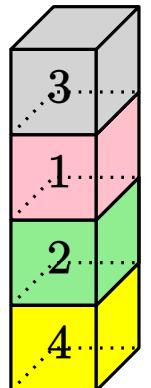
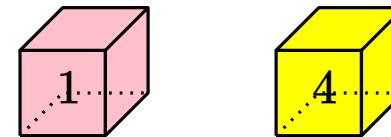
Each **area** has  $n$  **neurons**, the same **Erdős-Renyi prob.**  $p$ , the same  $k$  for the **k-CAP**



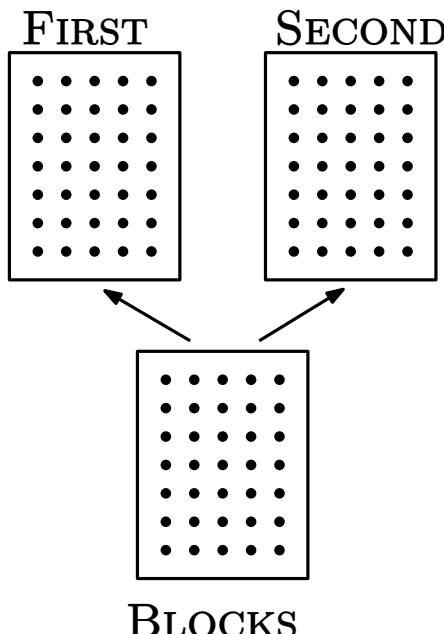
# A Toy Example

Given a **stack** of blocks and two of its **blocks**  $a, b$ , is  $a$  **above**  $b$ ?

E.g., is block 1 above block 4?



**Assumptions:** 3 brain areas, BLOCKS, FIRST, SECOND

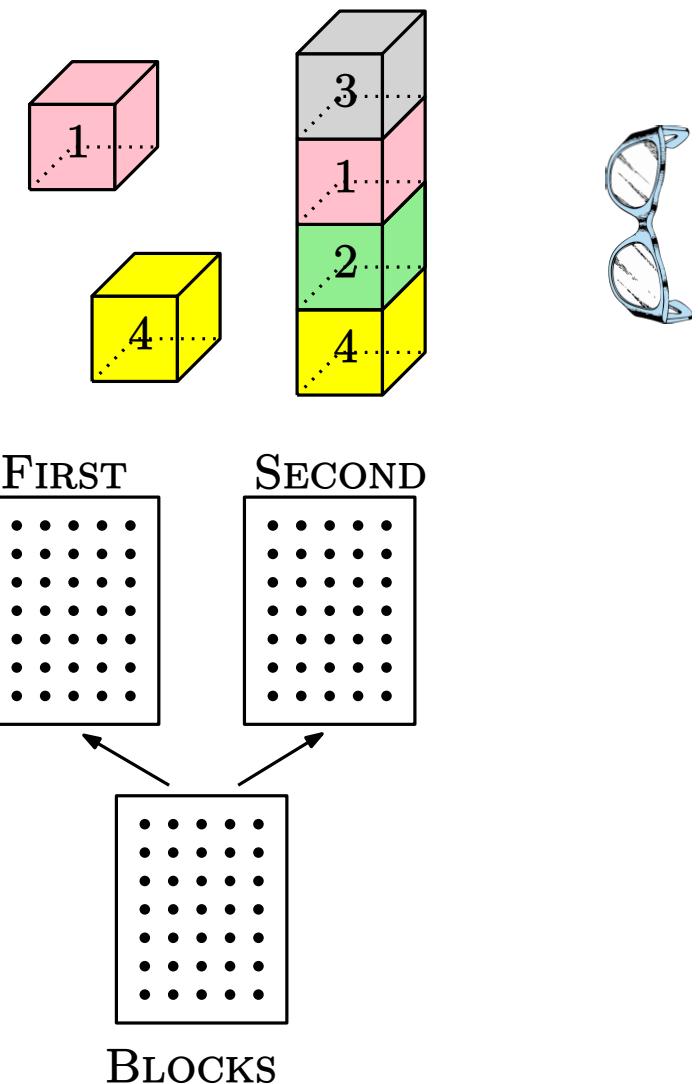


# IS ABOVE Algorithm

## Algorithm 1: IS ABOVE ( $S, x, y$ )

```
input: a stack  $S$  of blocks  $b_1, b_2, \dots, b_s$ ;  
      two blocks  $x, y$  with  $x, y \in S$ .  
  
1 disinhibitArea ({BLOCKS, FIRST});  
2 disinhibitFiber ({(BLOCKS, FIRST)});  
3 fire ( $x$ );  
4 strongProject();  
5 inhibitArea ({FIRST});  
6 disinhibitArea ({SECOND});  
7 disinhibitFiber ({(BLOCKS, SECOND)});  
8 fire ( $y$ );  
9 strongProject();  
10 foreach  $i$  with  $2 \leq i \leq s$  do  
11     inhibitArea ({SECOND});  
12     disinhibitArea ({FIRST});  
13     fire ( $b_i$ );  
14     if read (FIRST) then return true ;  
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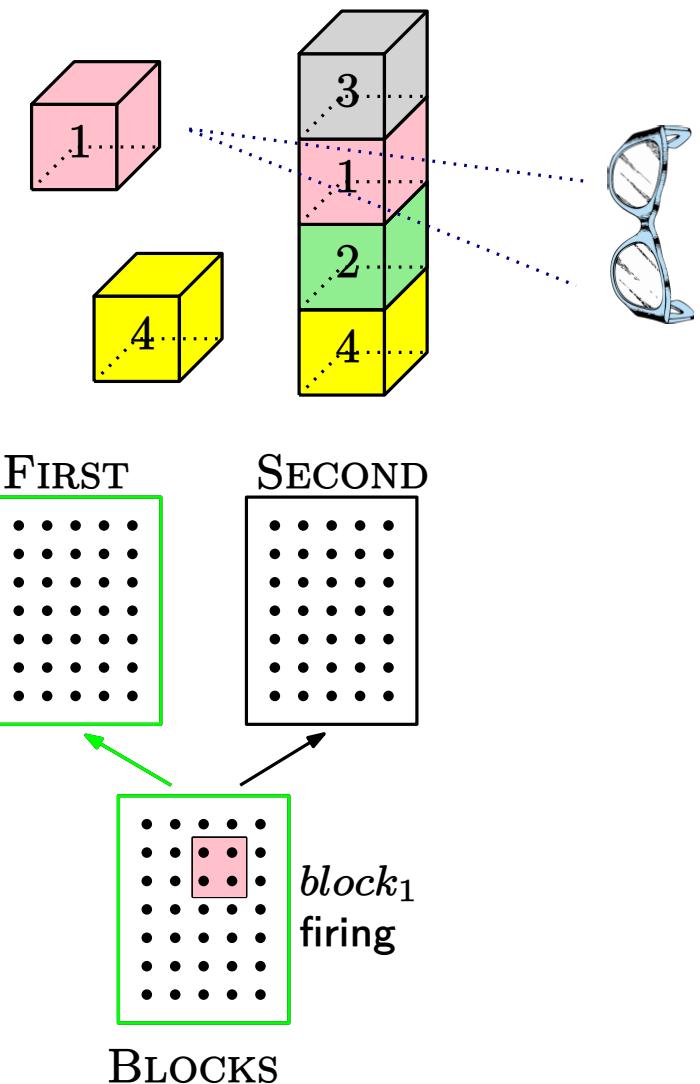


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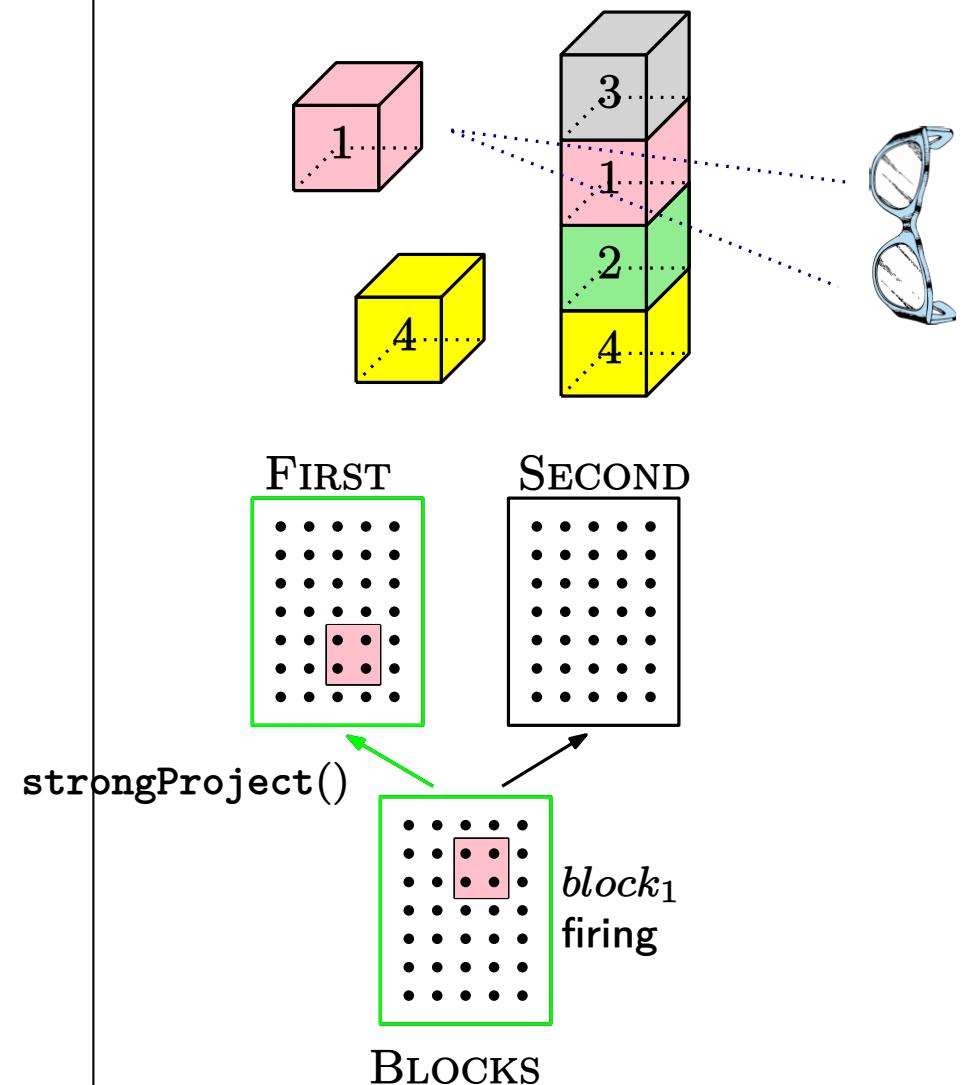
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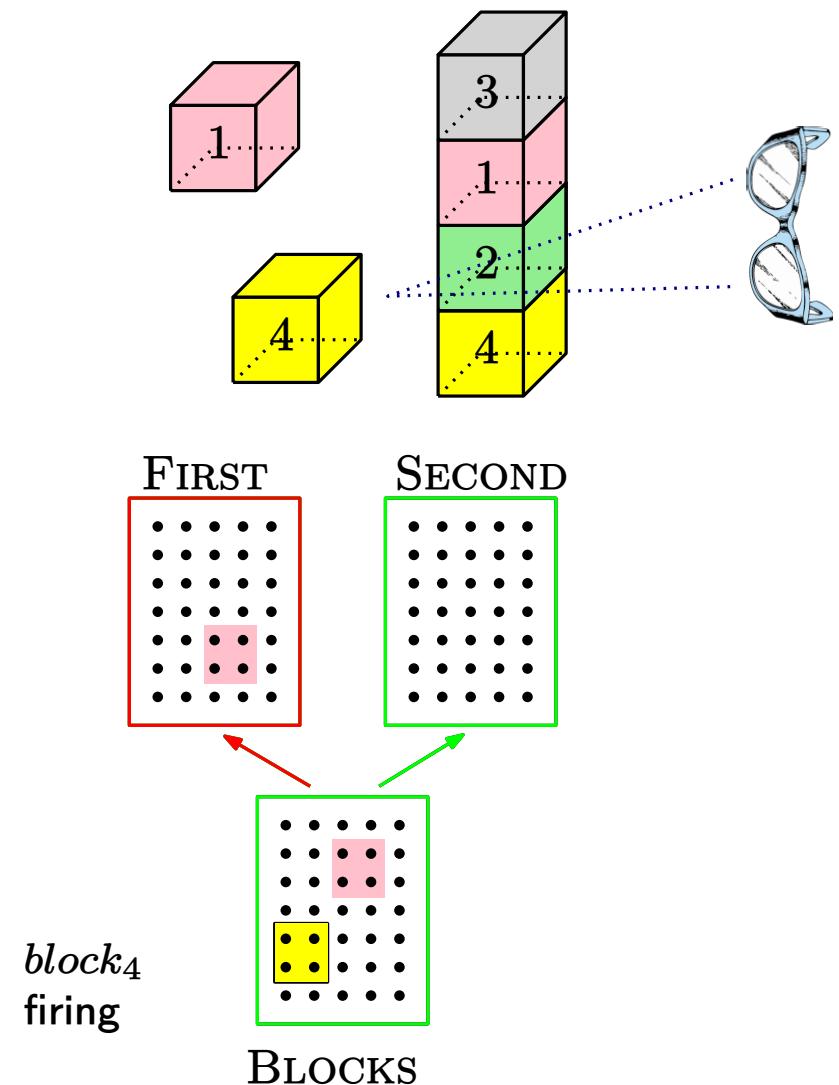
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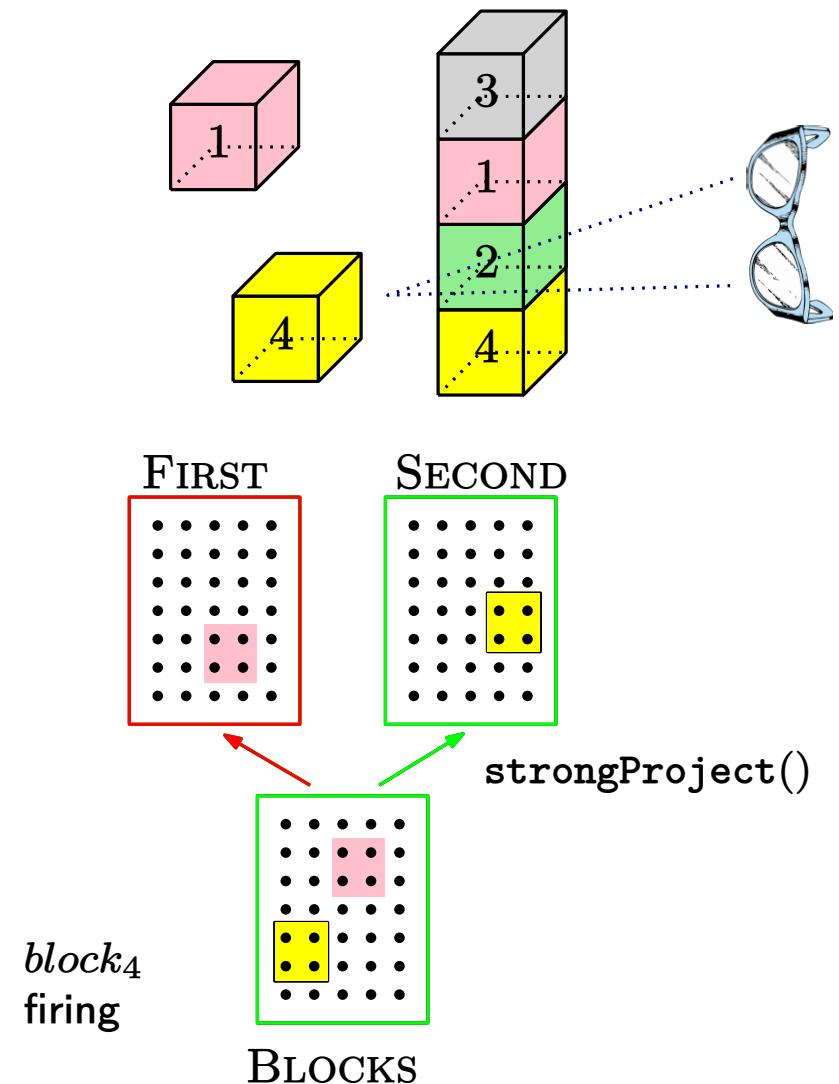
## Algorithm 1: IS ABOVE ( $S, x, y$ )

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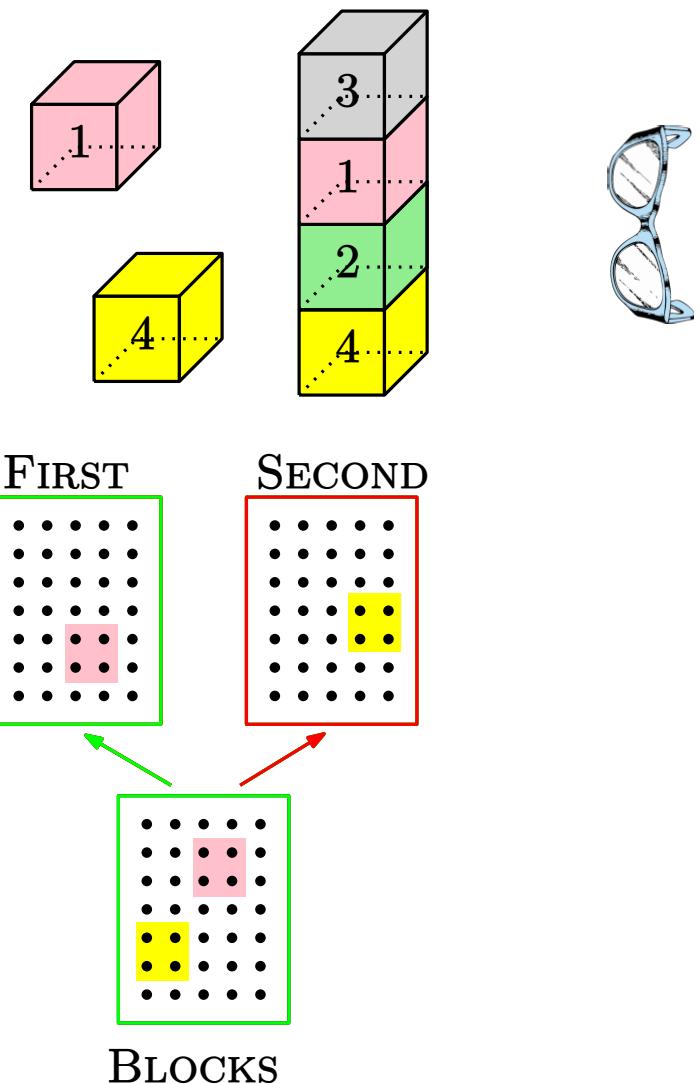
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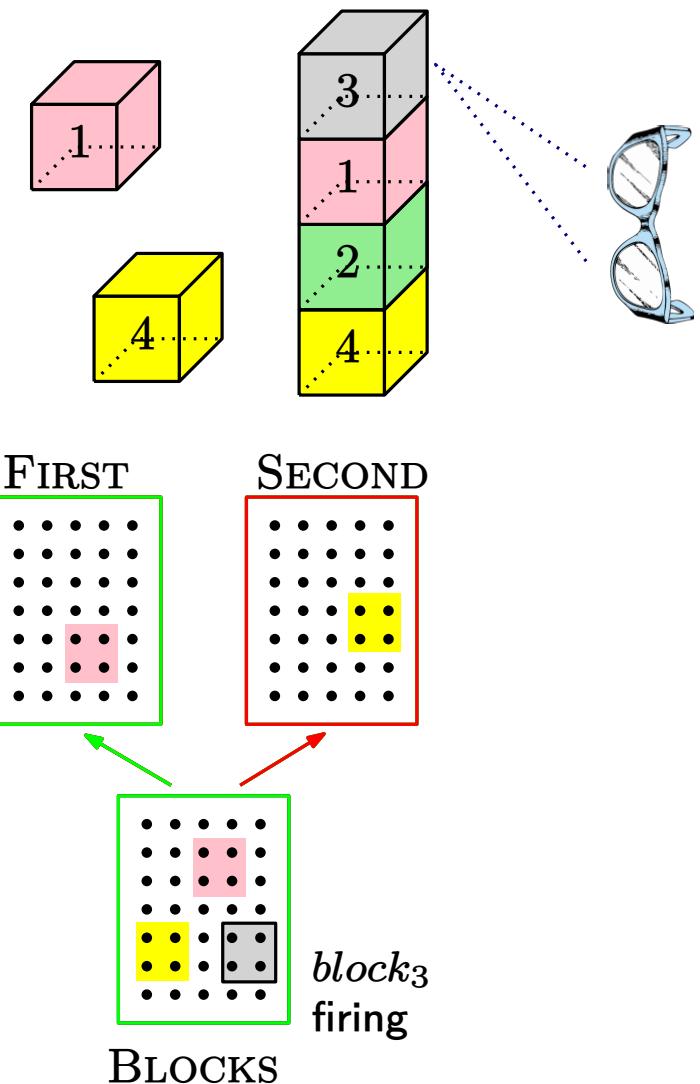
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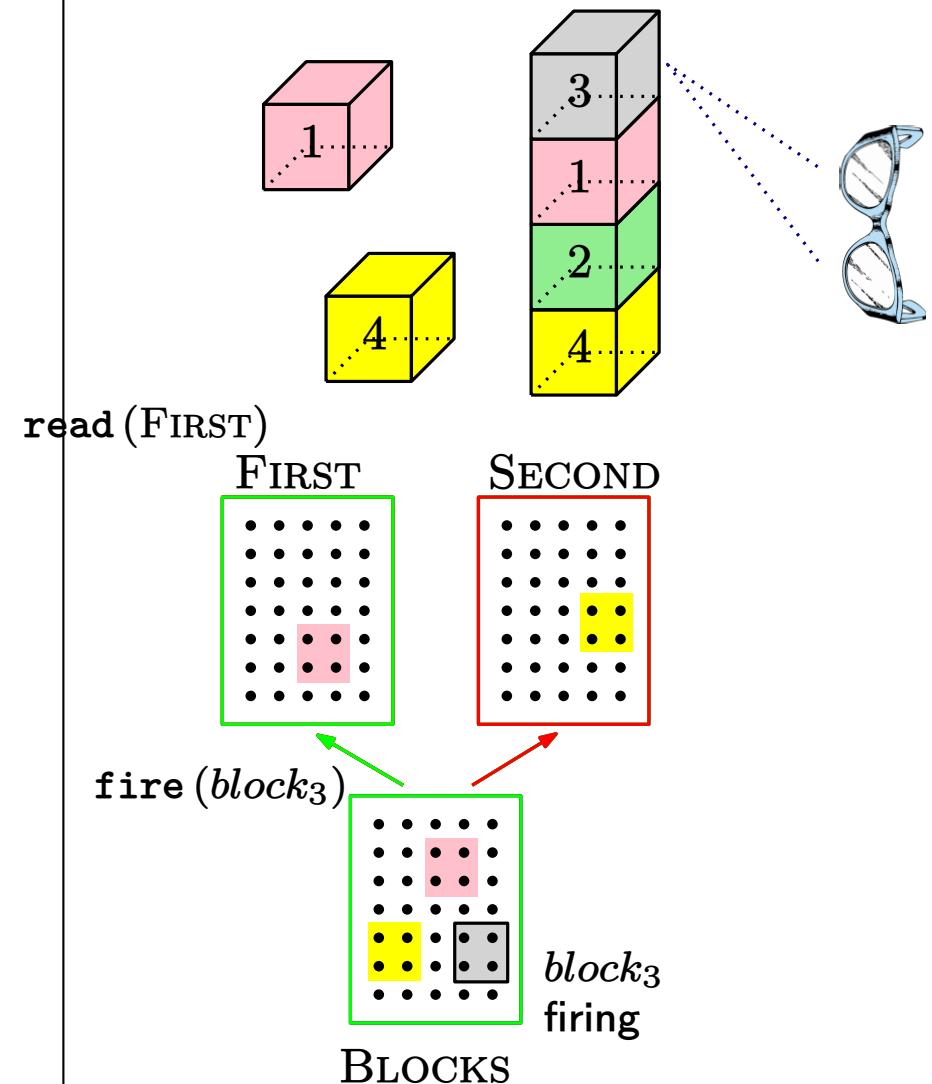
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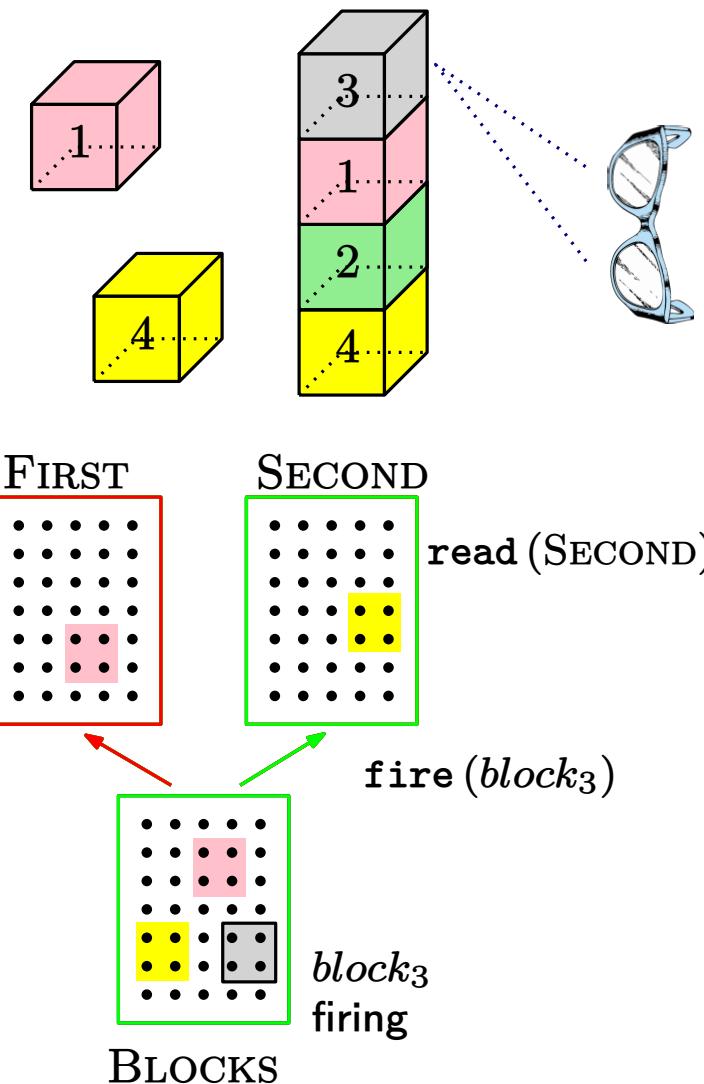
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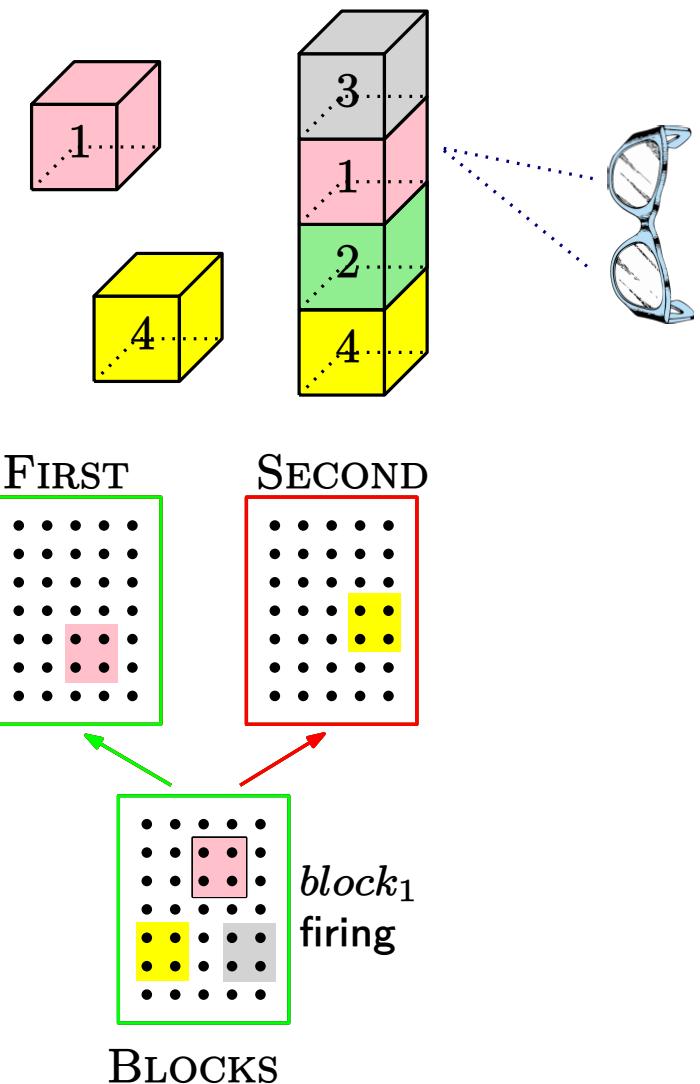
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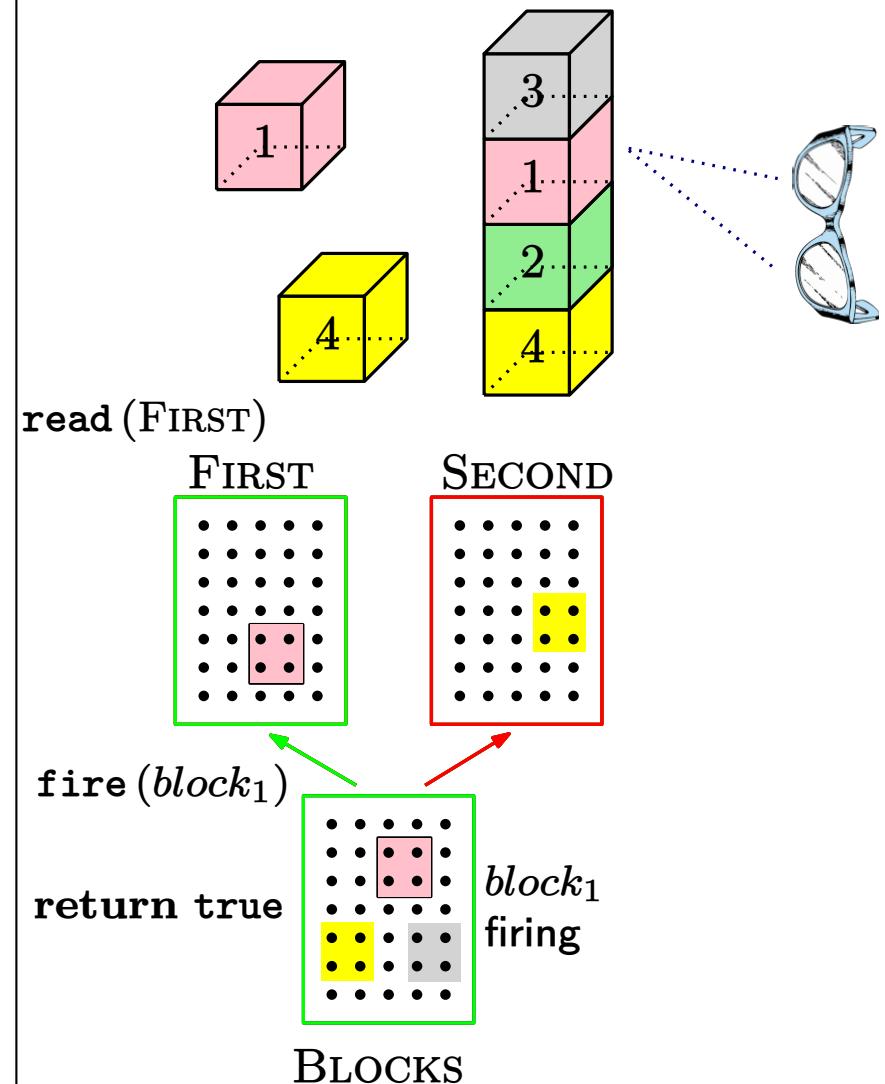
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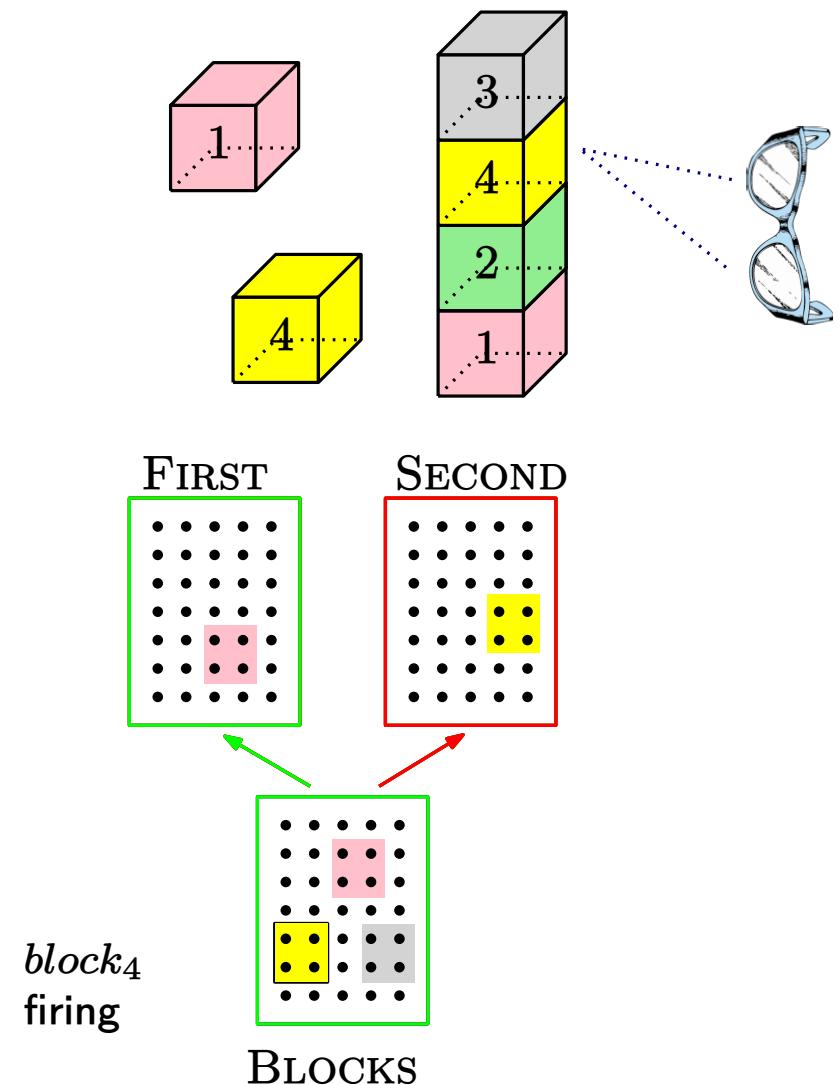
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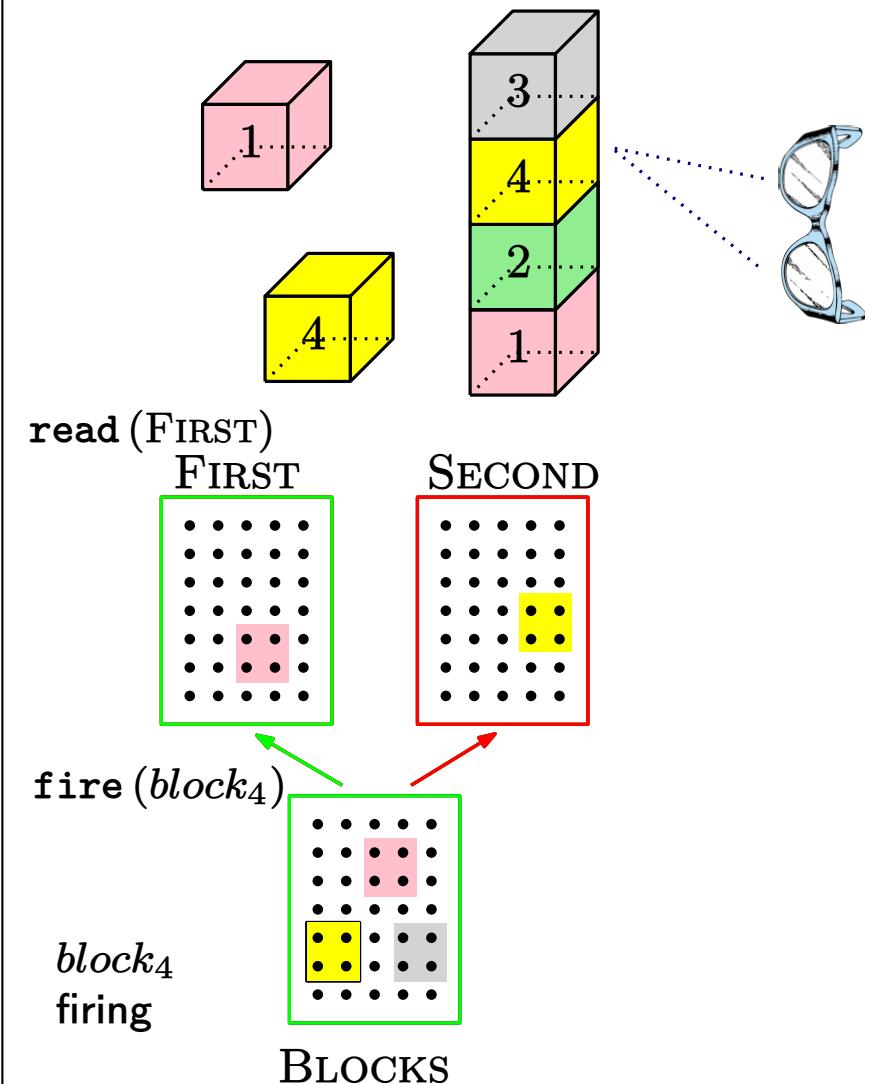
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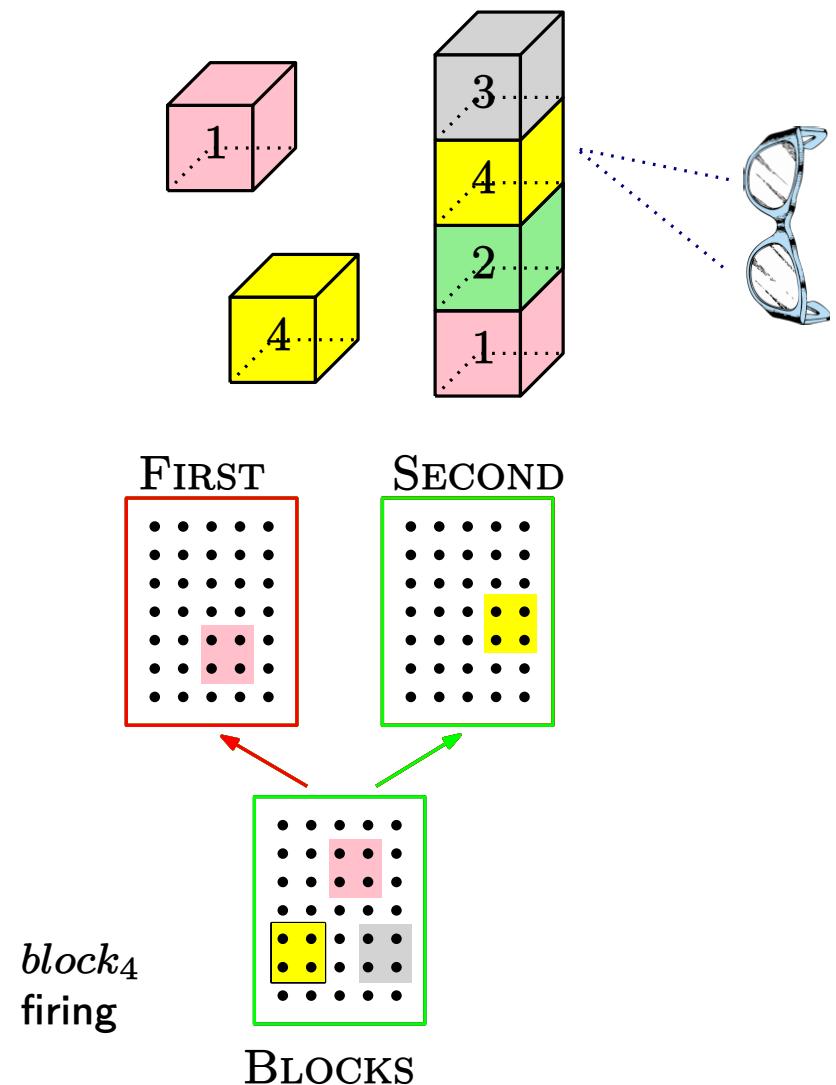
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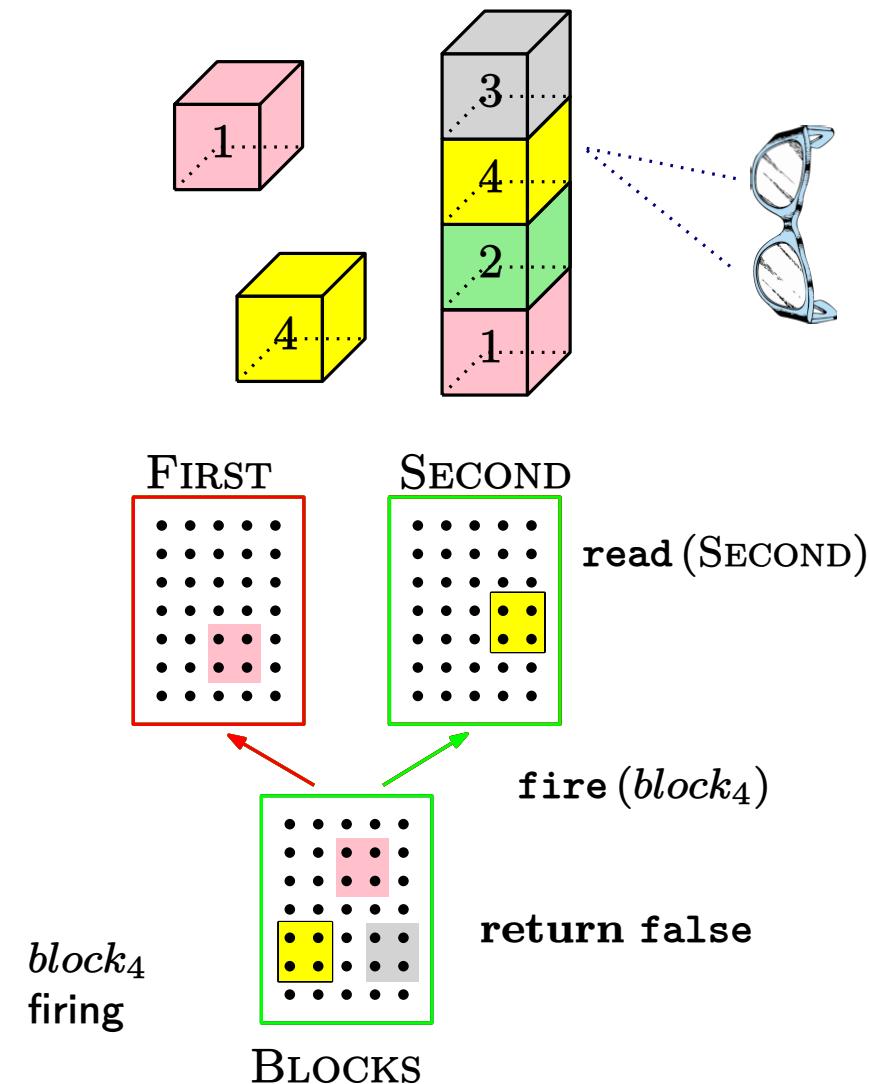
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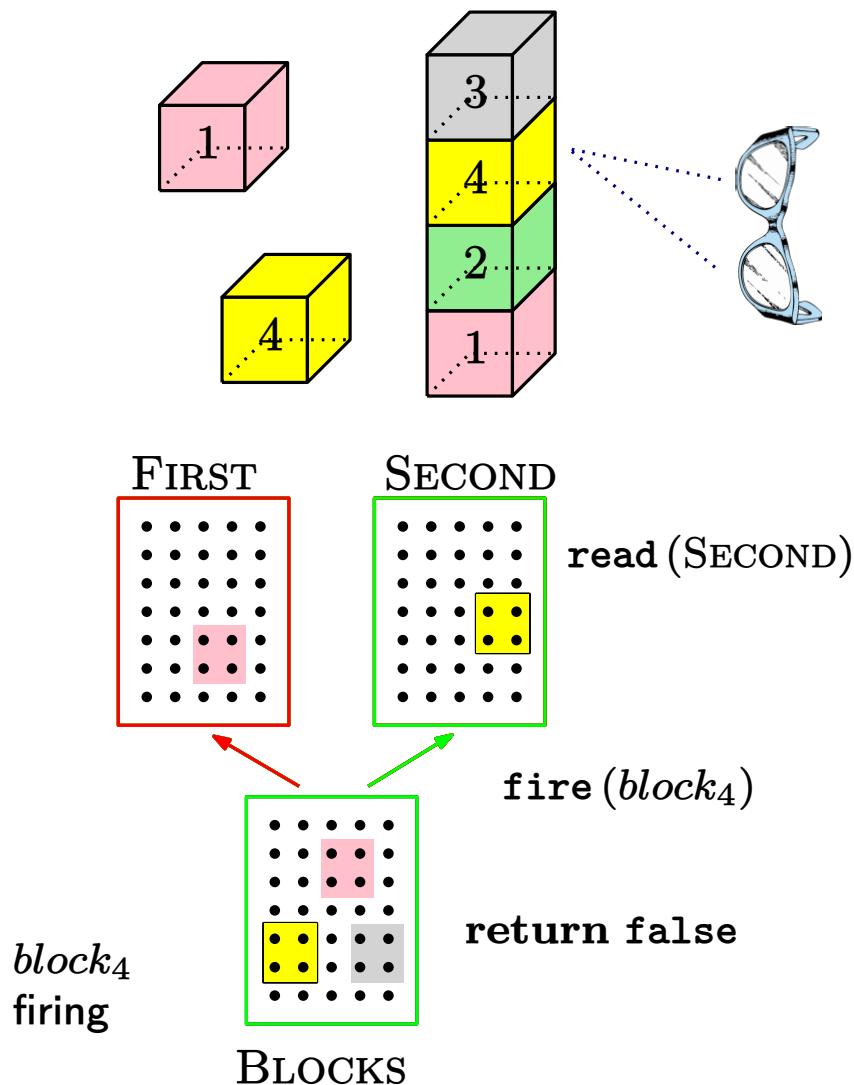
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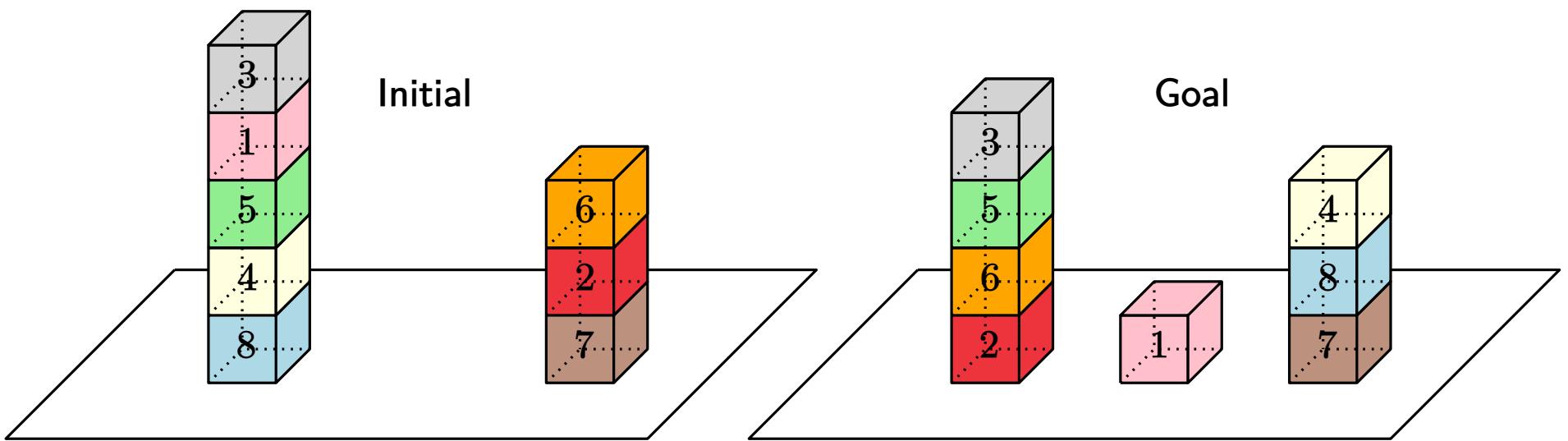
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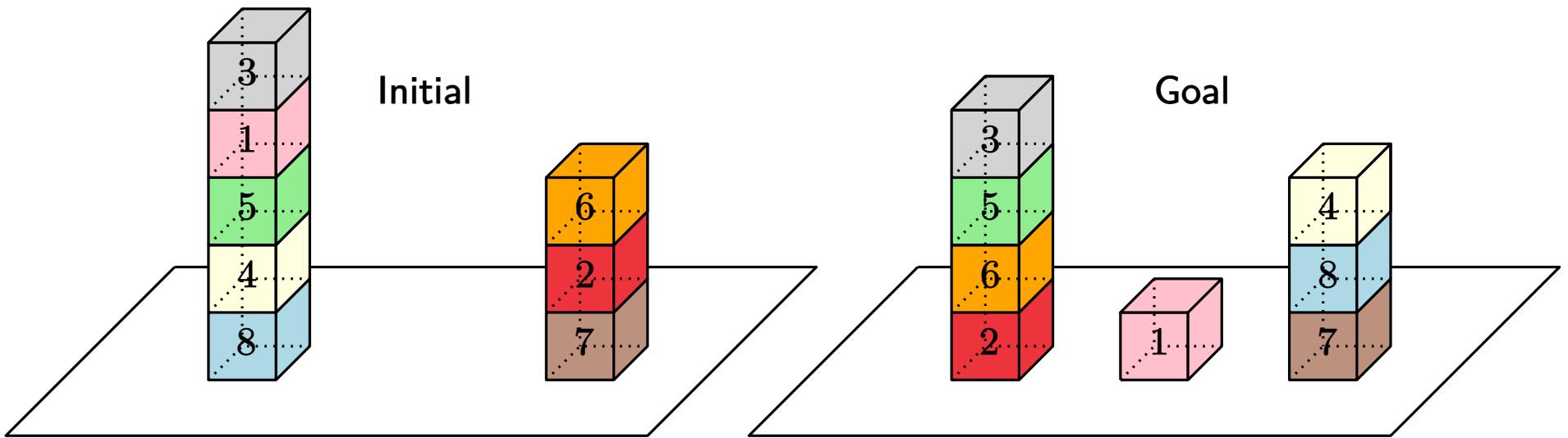
How does the brain  
understand a positive  
(negative) answer?

The answer corresponds to an  
assembly activating in a particular  
brain area

# Back to the Planning Problem

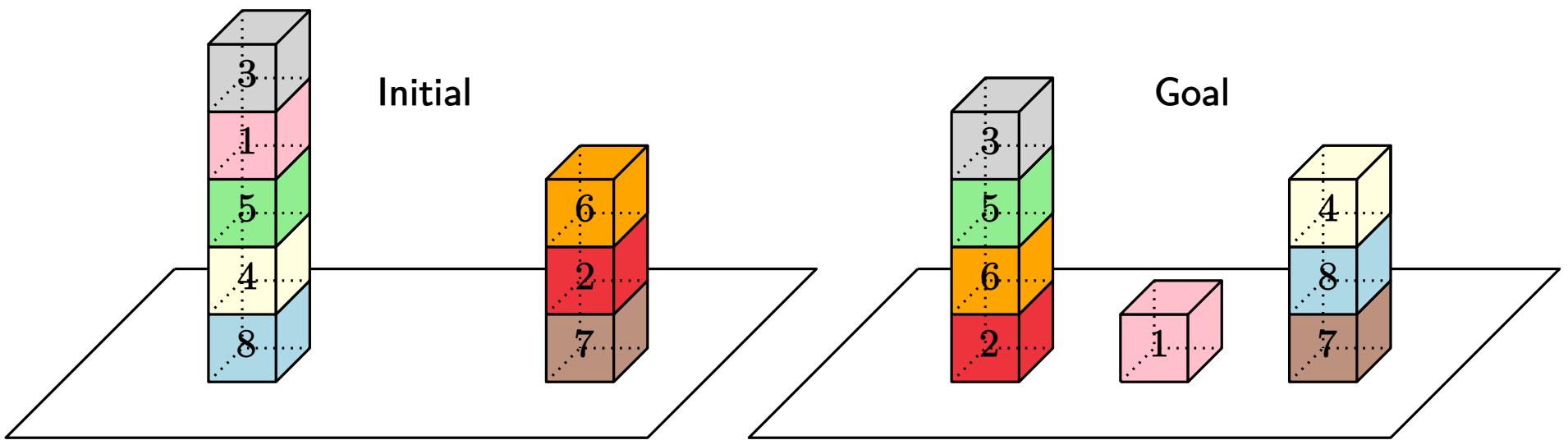


# Back to the Planning Problem



- **First:** way to represent the **initial** and the **goal** stacks in the brain

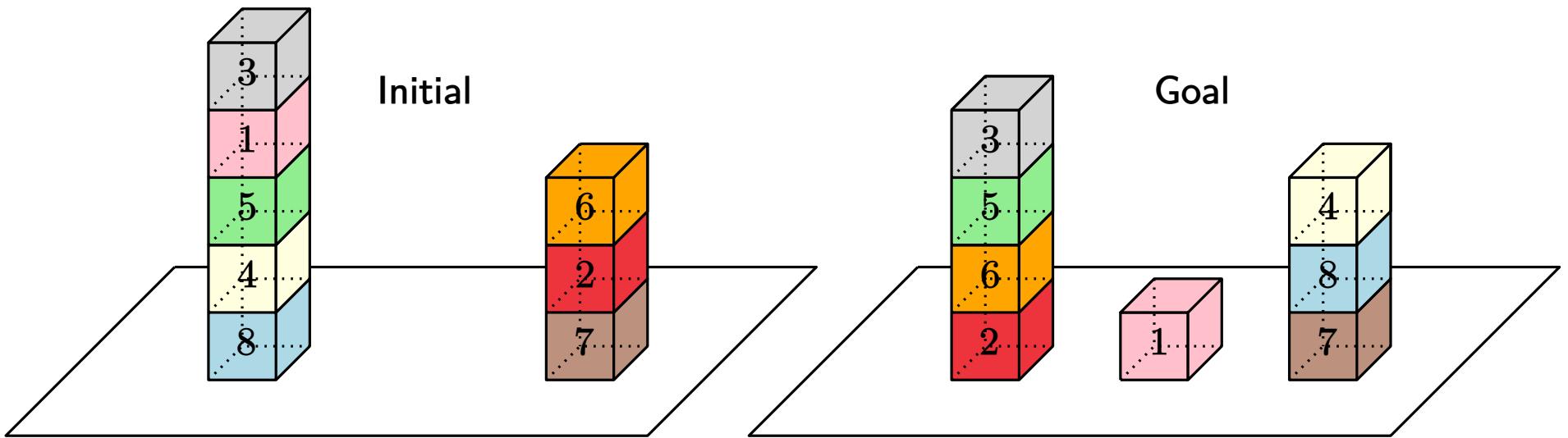
# Back to the Planning Problem



- First: way to represent the initial and the goal stacks in the brain

A Parser

# Back to the Planning Problem



- **First:** way to **represent** the **initial** and the **goal** stacks in the brain

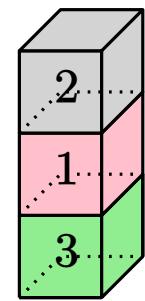
A Parser

- **Second:** way to **implement actions** and **represent** them

# The Parser

## Assumptions:

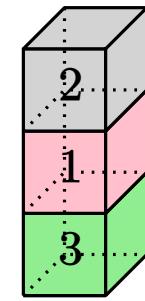
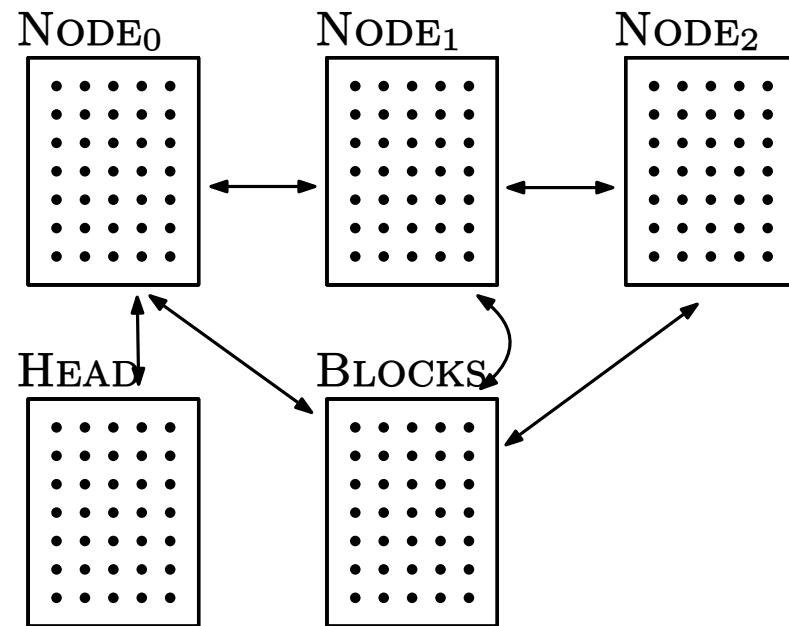
- single stack
- brain as **undirected graph**
- 5 brain areas, BLOCKS, NODE<sub>0</sub>, NODE<sub>1</sub>, NODE<sub>2</sub>, HEAD



# The Parser

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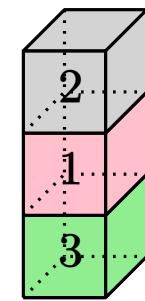
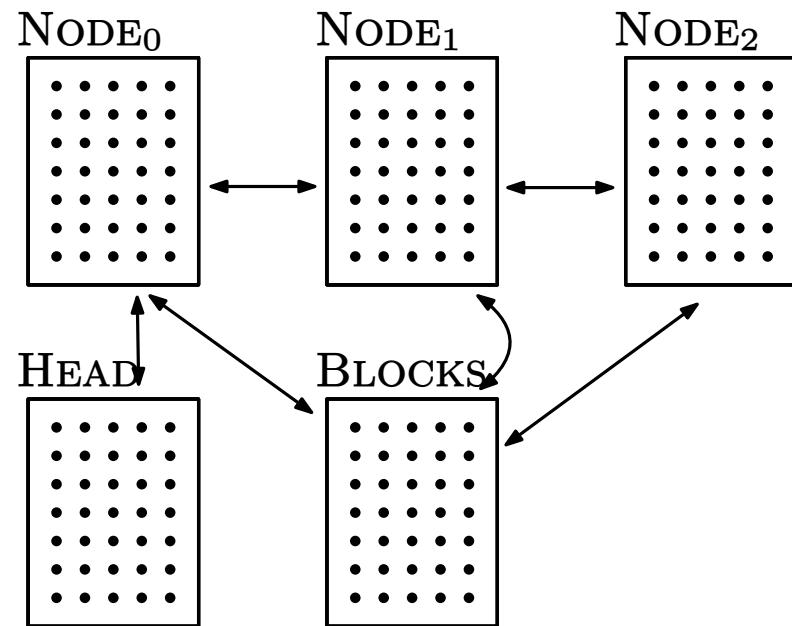
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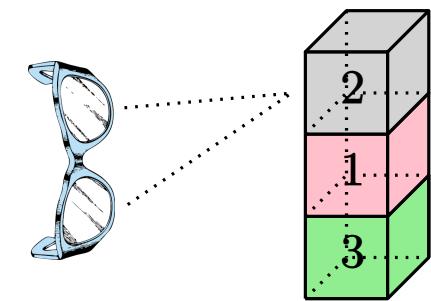
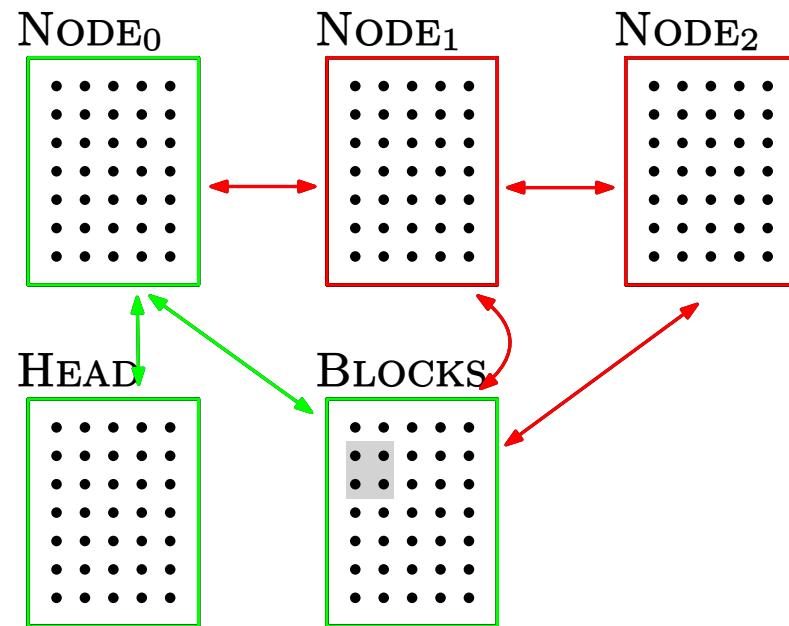
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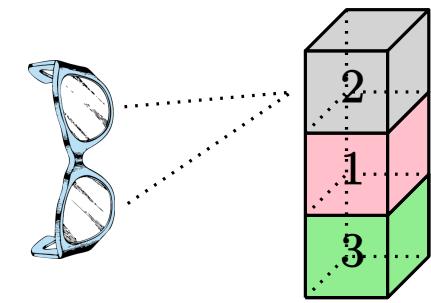
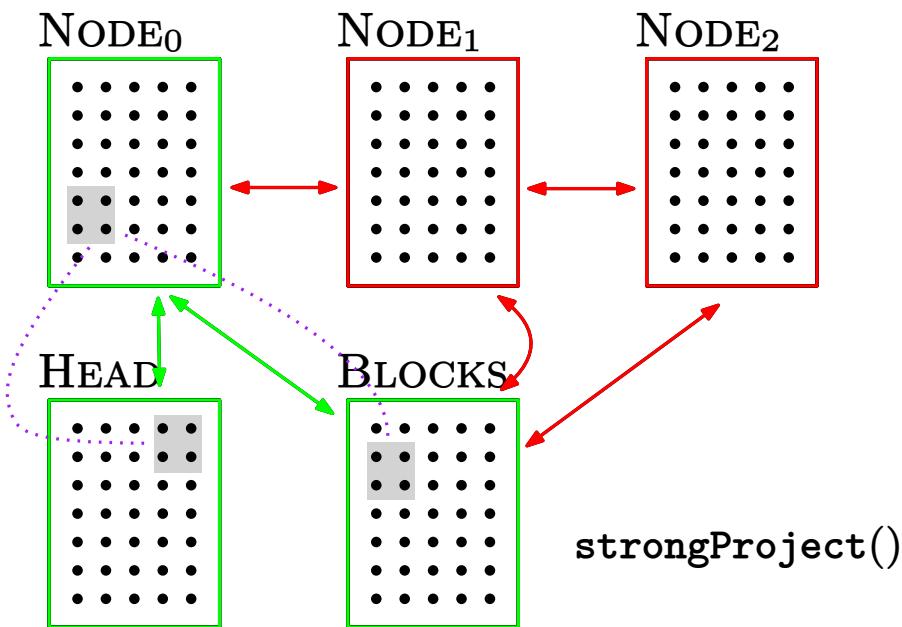
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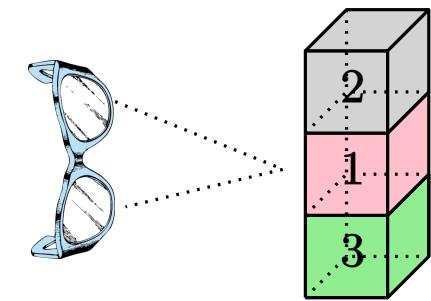
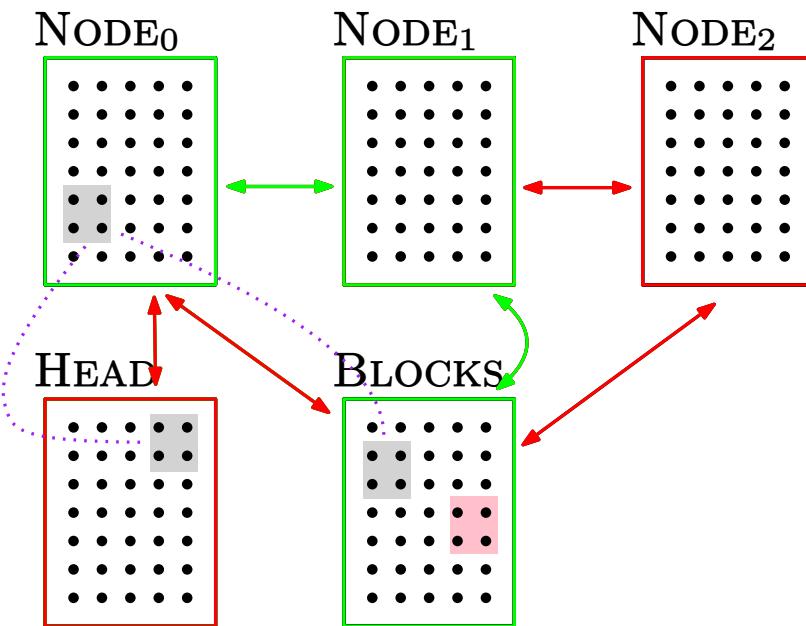
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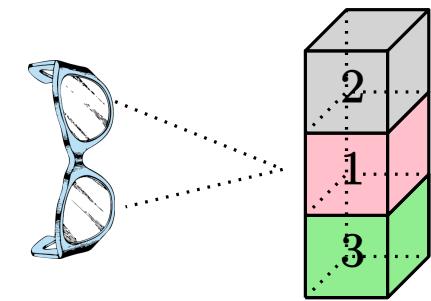
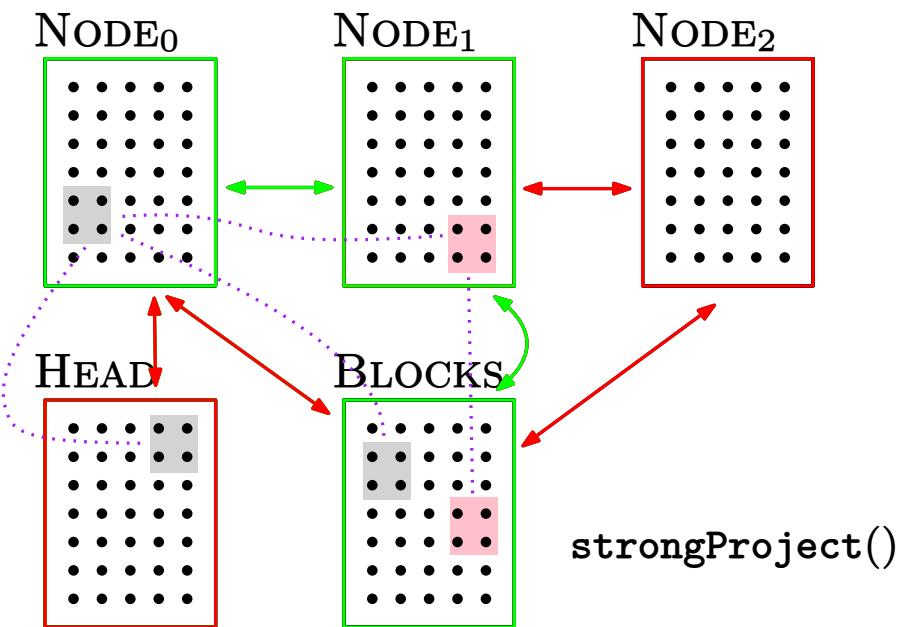
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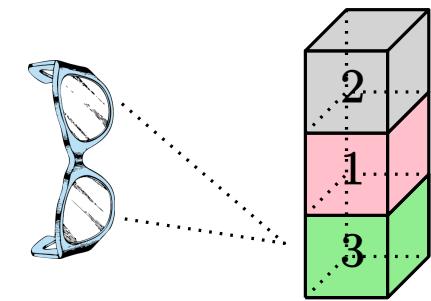
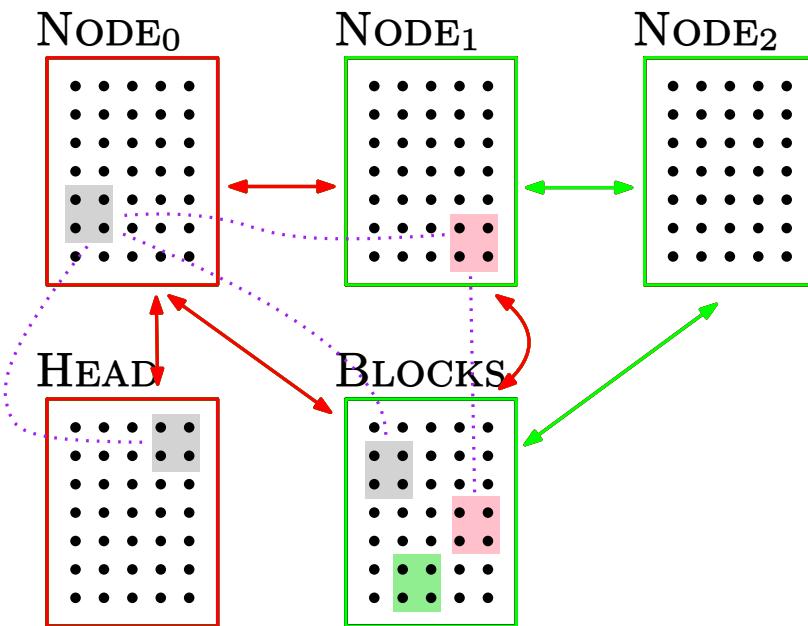
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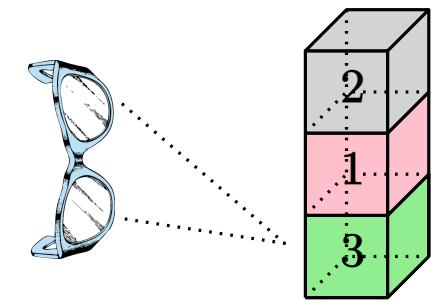
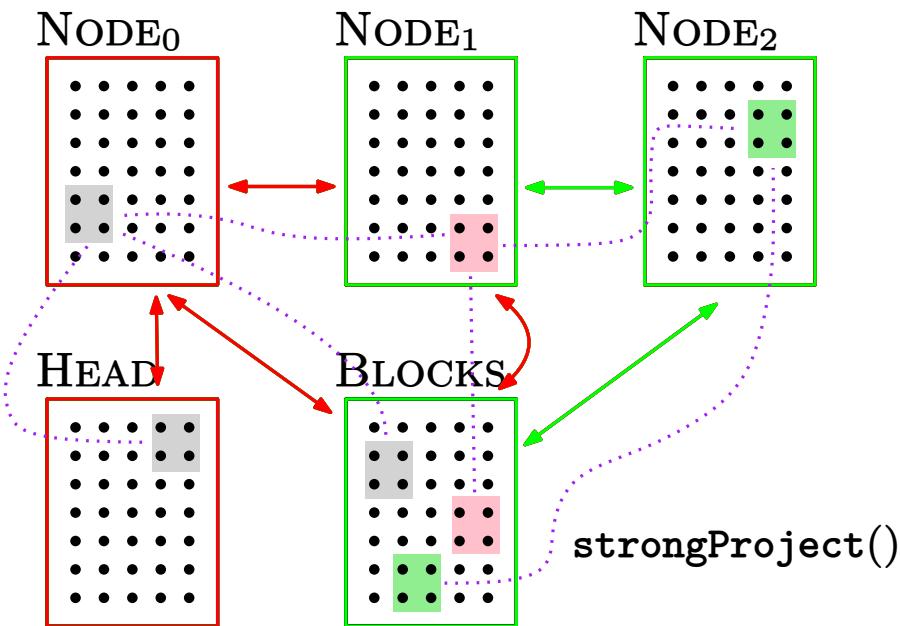
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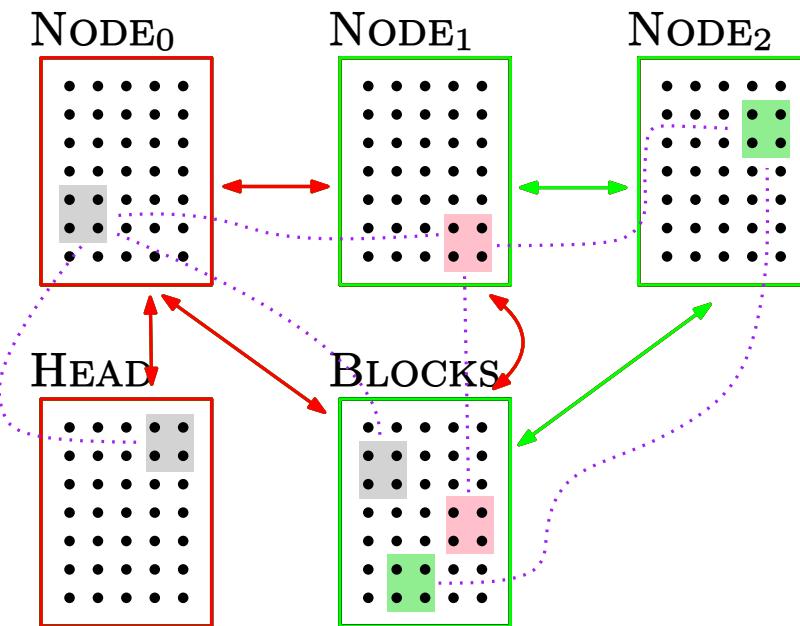
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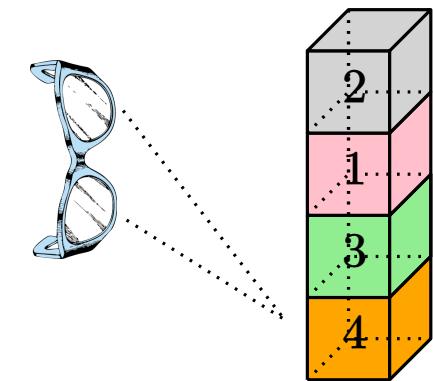
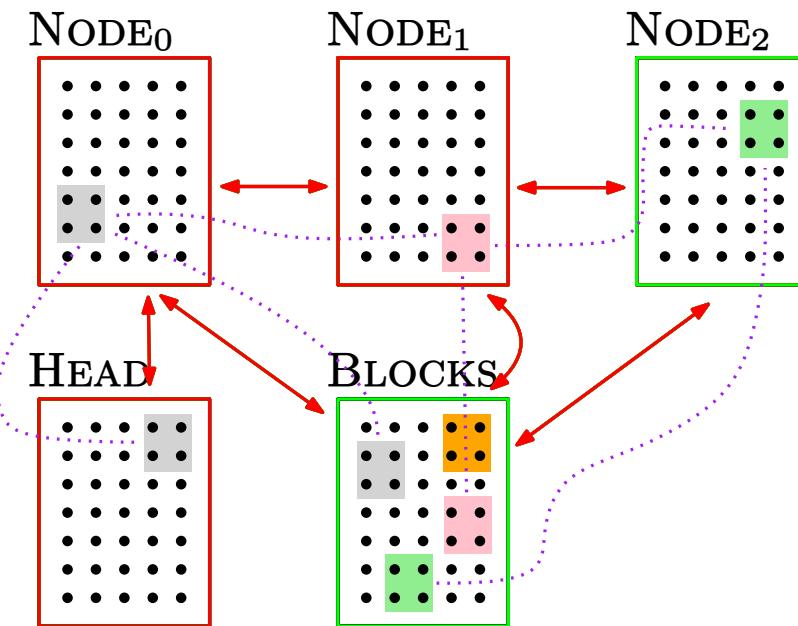
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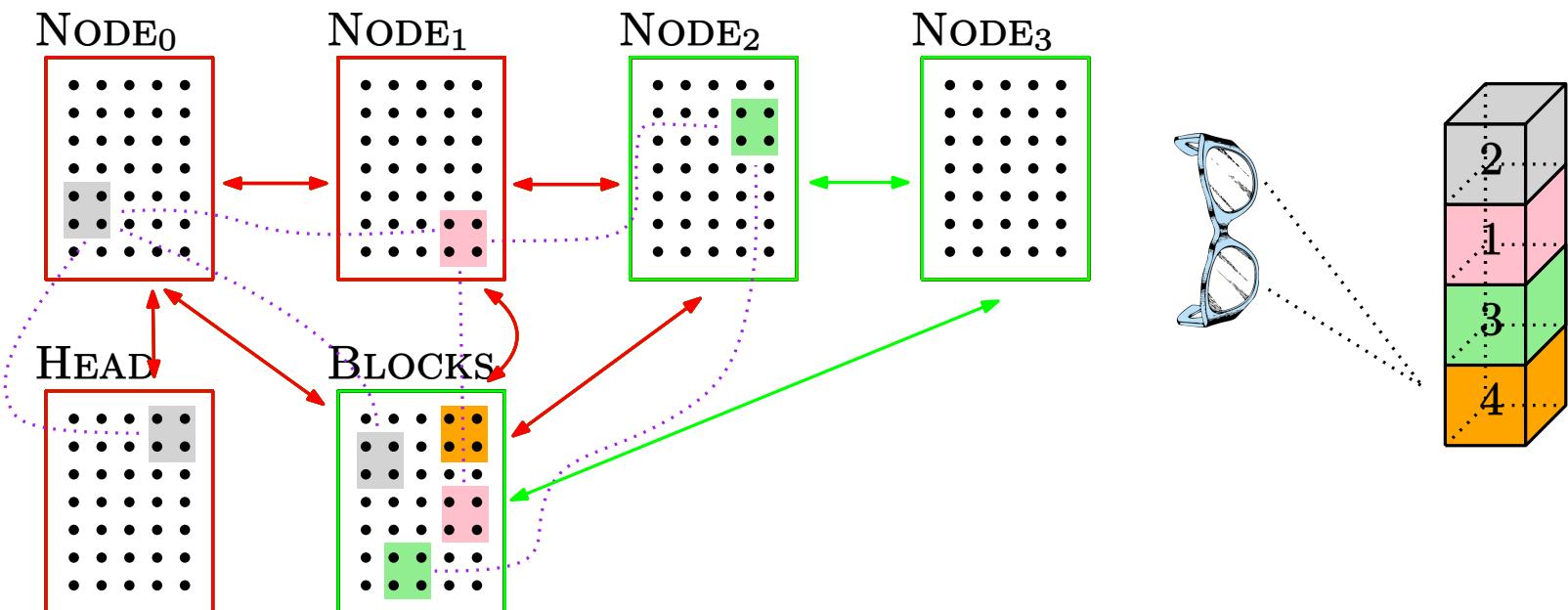
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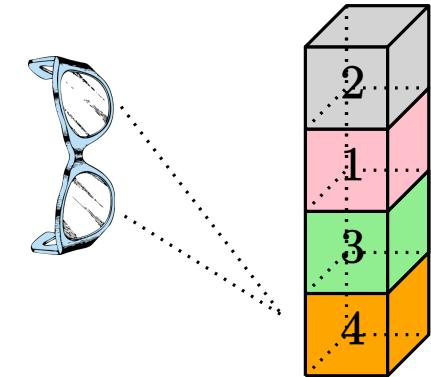
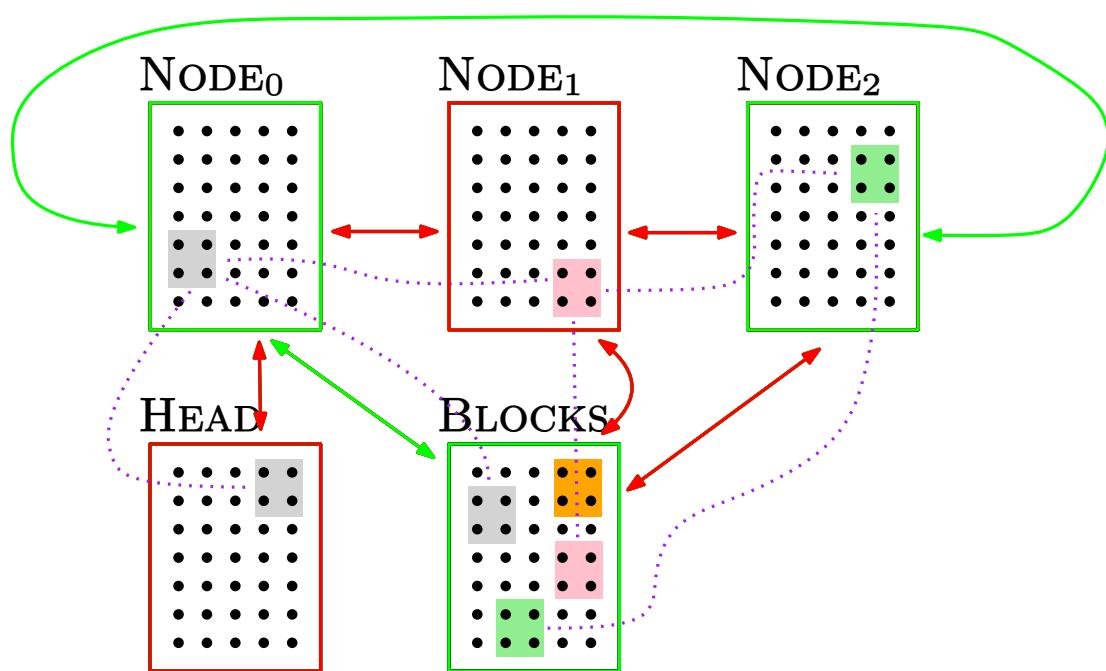


As many areas as the number of  
blocks... :(

# The Parser

## Assumptions:

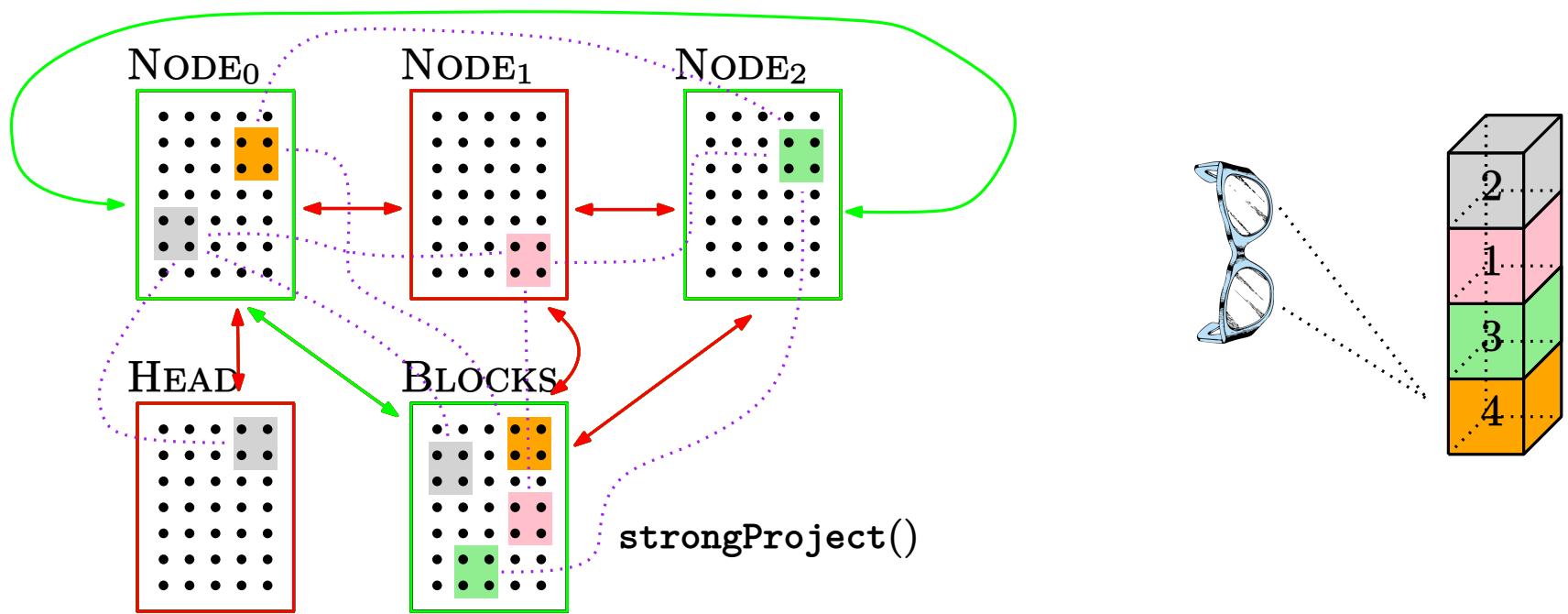
- single stack
- brain as **undirected graph**
- 5 brain areas, BLOCKS, NODE<sub>0</sub>, NODE<sub>1</sub>, NODE<sub>2</sub>, HEAD



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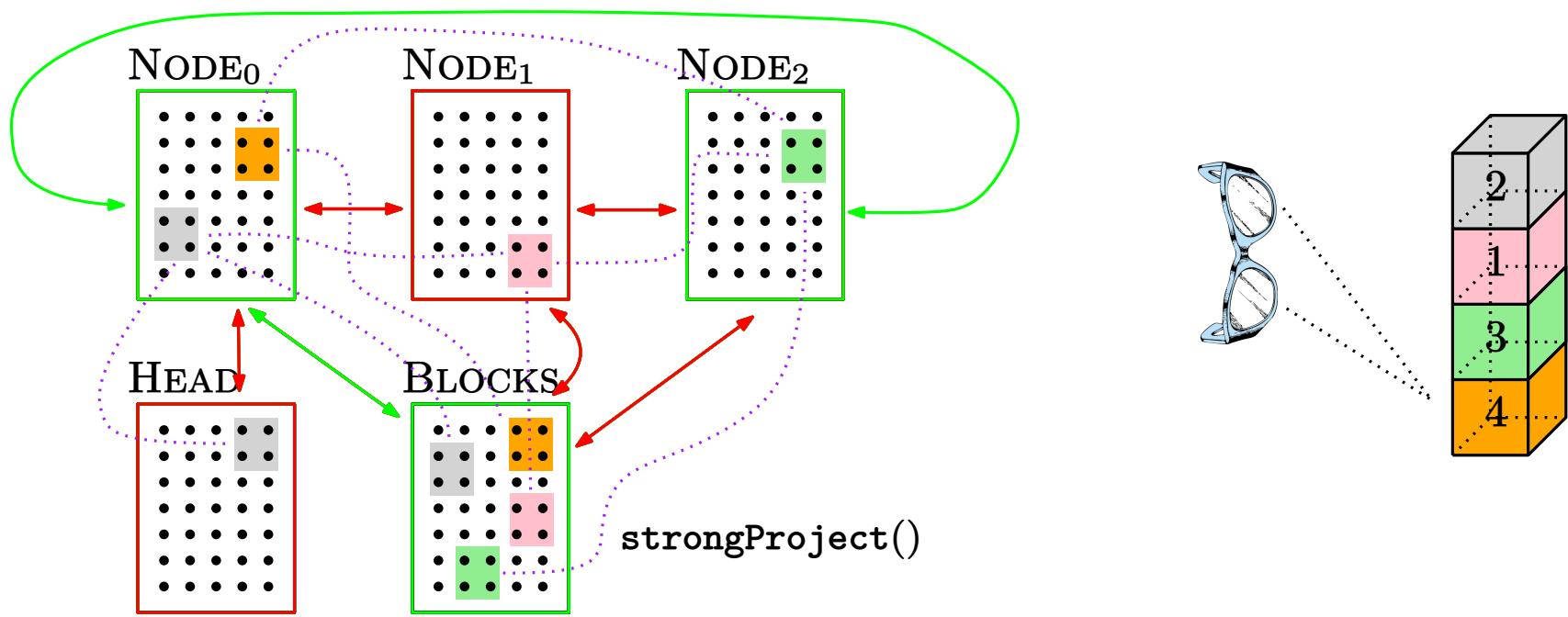
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- single stack
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**Chaining:** novelty. **Number of blocks** represented depends only on  $n$  (**number of neurons per area**) and  $k$  (**the CAP**)

# Parser Code

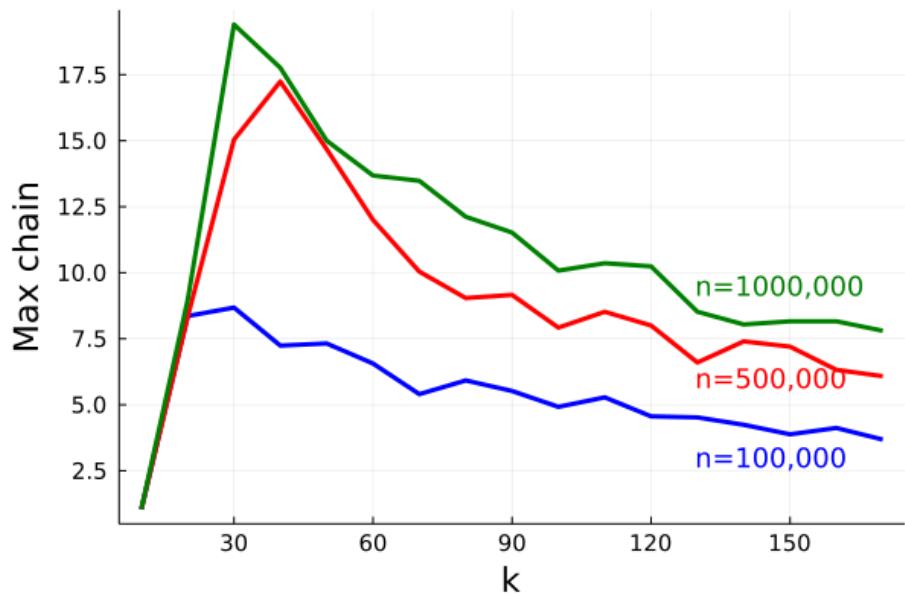
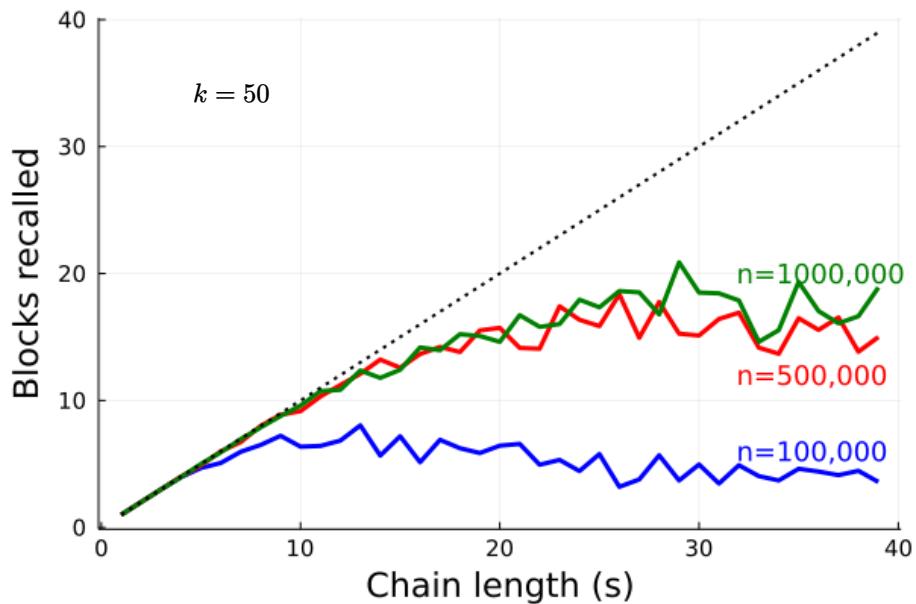
## Algorithm 2: PARSER ( $S$ )

```
1 input: a stack  $S$  of blocks  $b_1, b_2, \dots, b_s$ .
2 disinhibitArea ({BLOCKS, HEAD, NODE0});
3 disinhibitFiber ({(HEAD, NODE0), (NODE0, BLOCKS)} );
4 fire ( $b_1$ );
5 strongProject();
6 inhibitArea ({HEAD});
7 inhibitFiber ({(HEAD, NODE0), (NODE0, BLOCKS)} );
8 foreach  $i$  with  $2 \leq i \leq s$  do
9    $p = (i - 2) \bmod 3$ ;  $c = (i - 1) \bmod 3$ ;
10  disinhibitArea ({NODEc});
11  disinhibitFiber ({(NODEp, NODEc), (NODEc, BLOCKS)} );
12  fire ( $b_i$ );
13  strongProject();
14  inhibitArea ({NODEp});
15  inhibitFiber ({(NODEp, NODEc), (NODEc, BLOCKS)} );
16 end
17 inhibitArea ({BLOCKS, NODE(s-1) \bmod 3} );
```

# Chaining Experiments

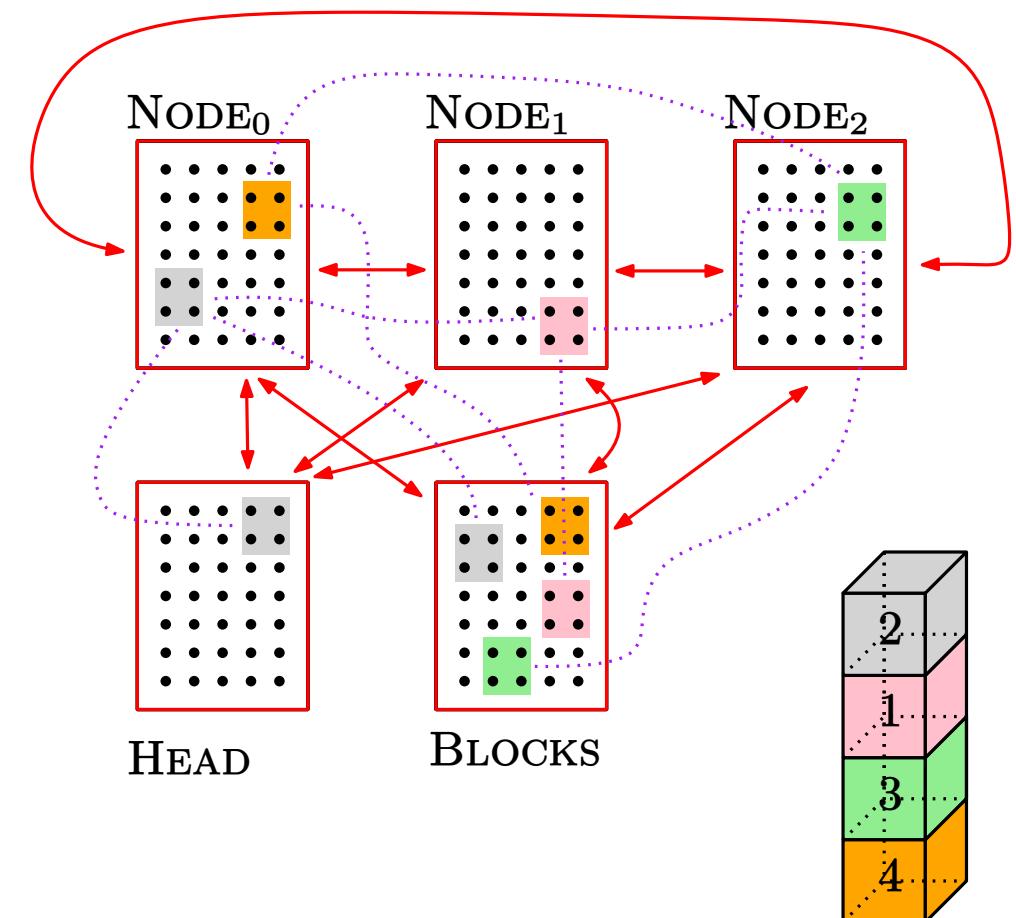
Due to **overlap** of assemblies, chaining may **fail**

Experiments conducted averaging over 50 trials



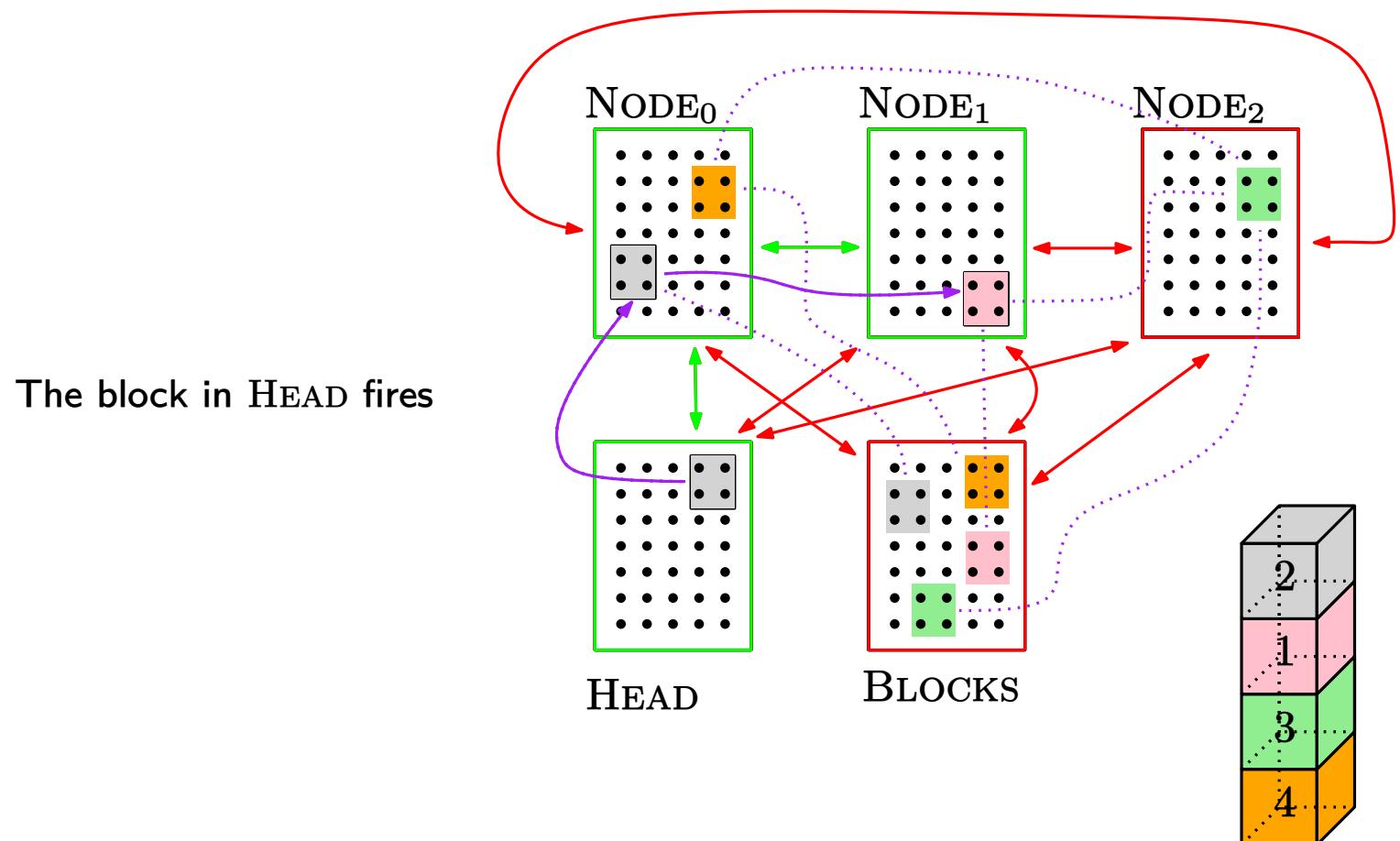
# Actions

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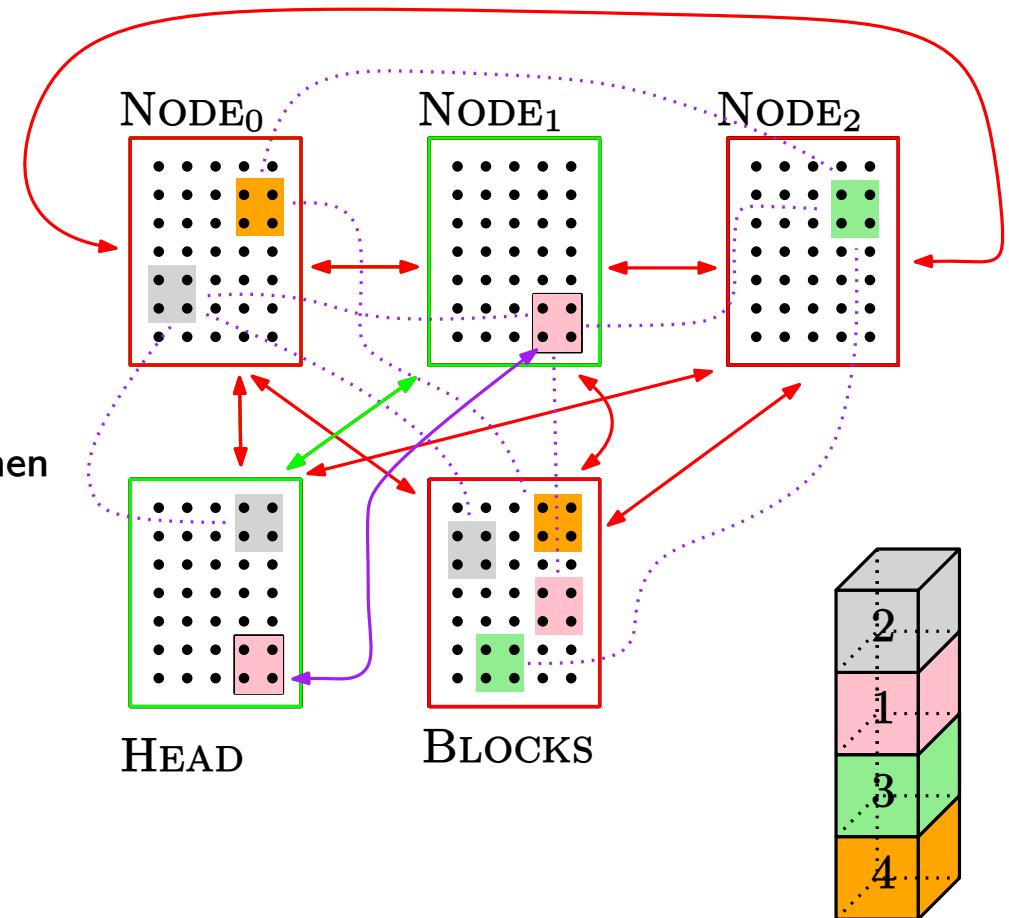
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- *Removing the top block*

The **last active assembly** in HEAD will point to the **new top**

The model **doesn't forget** old connections, it simply **doesn't use** them again

Now, project (NODE<sub>1</sub>, HEAD) and then project (HEAD, NODE<sub>1</sub>) to reinforce



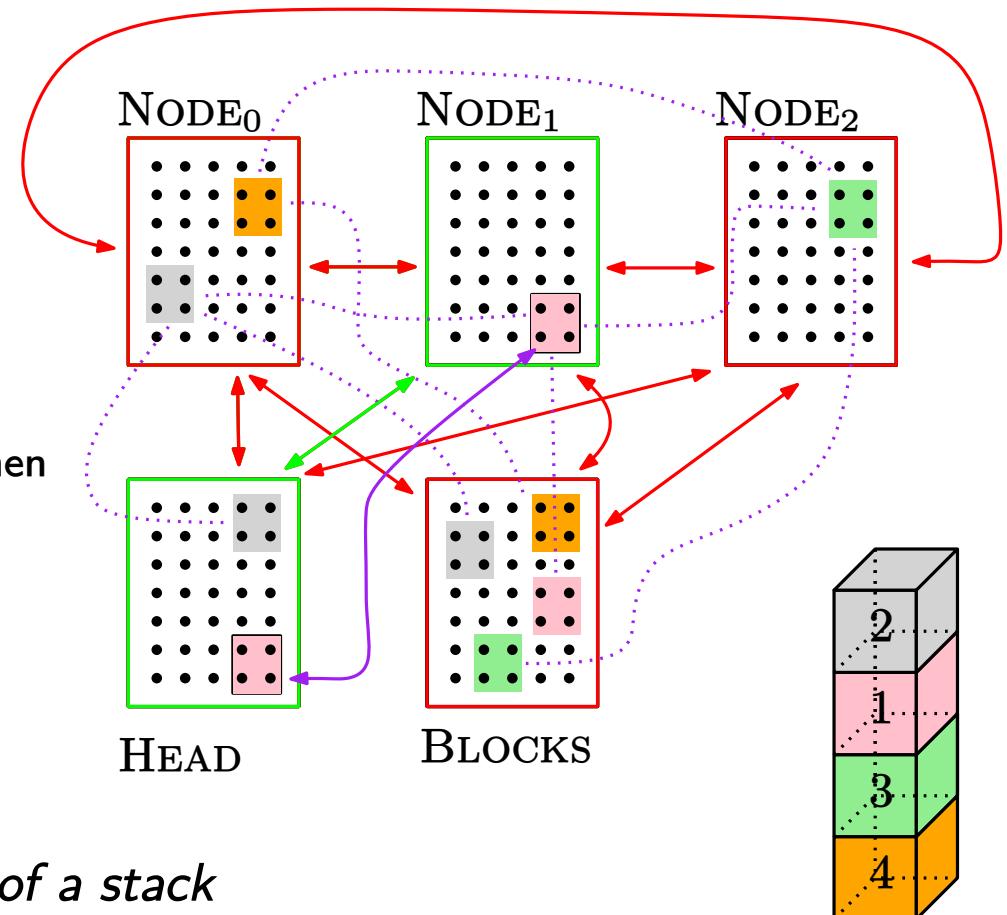
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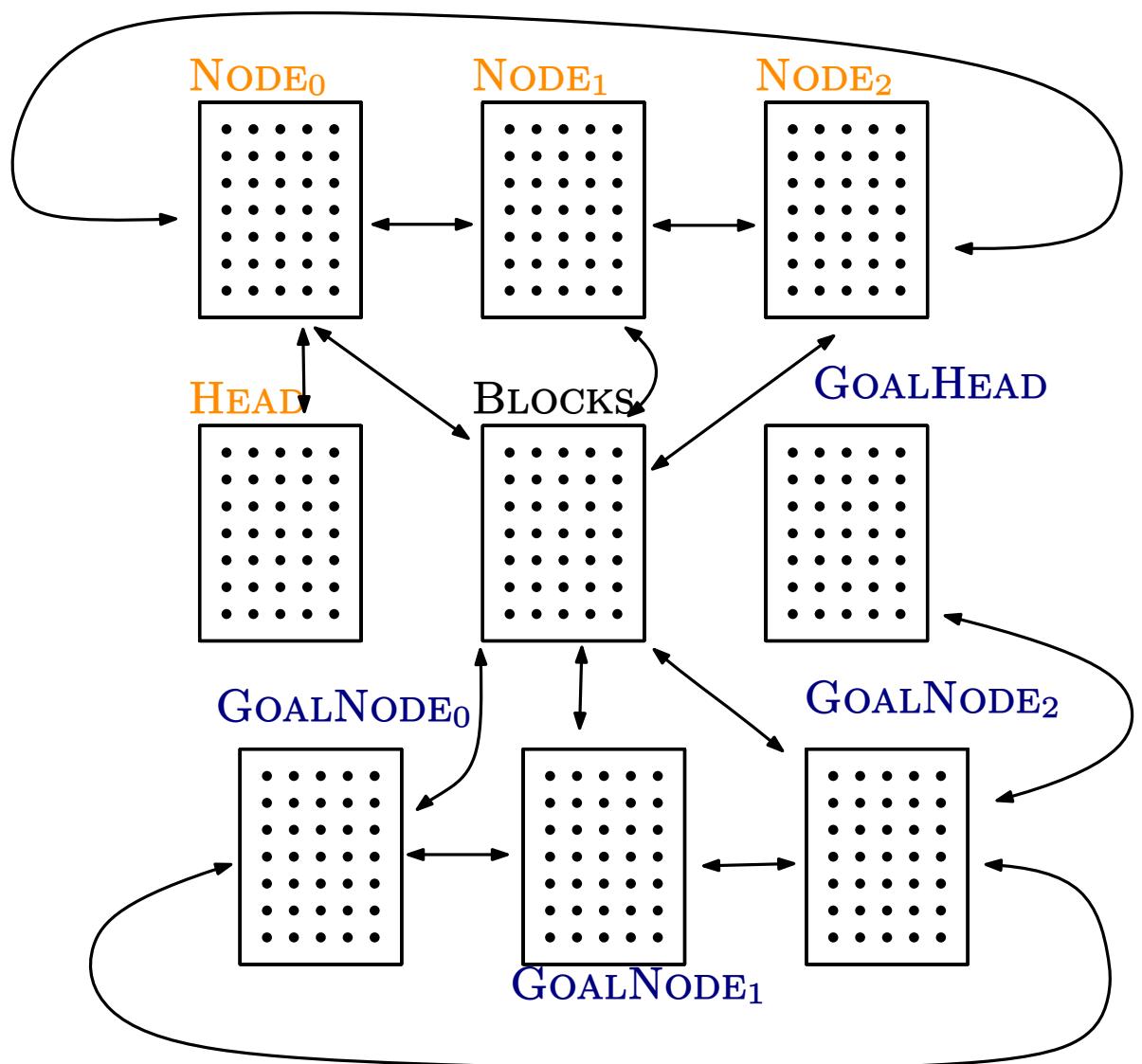
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- *Put a block from the table to the top of a stack*

# Comparisons

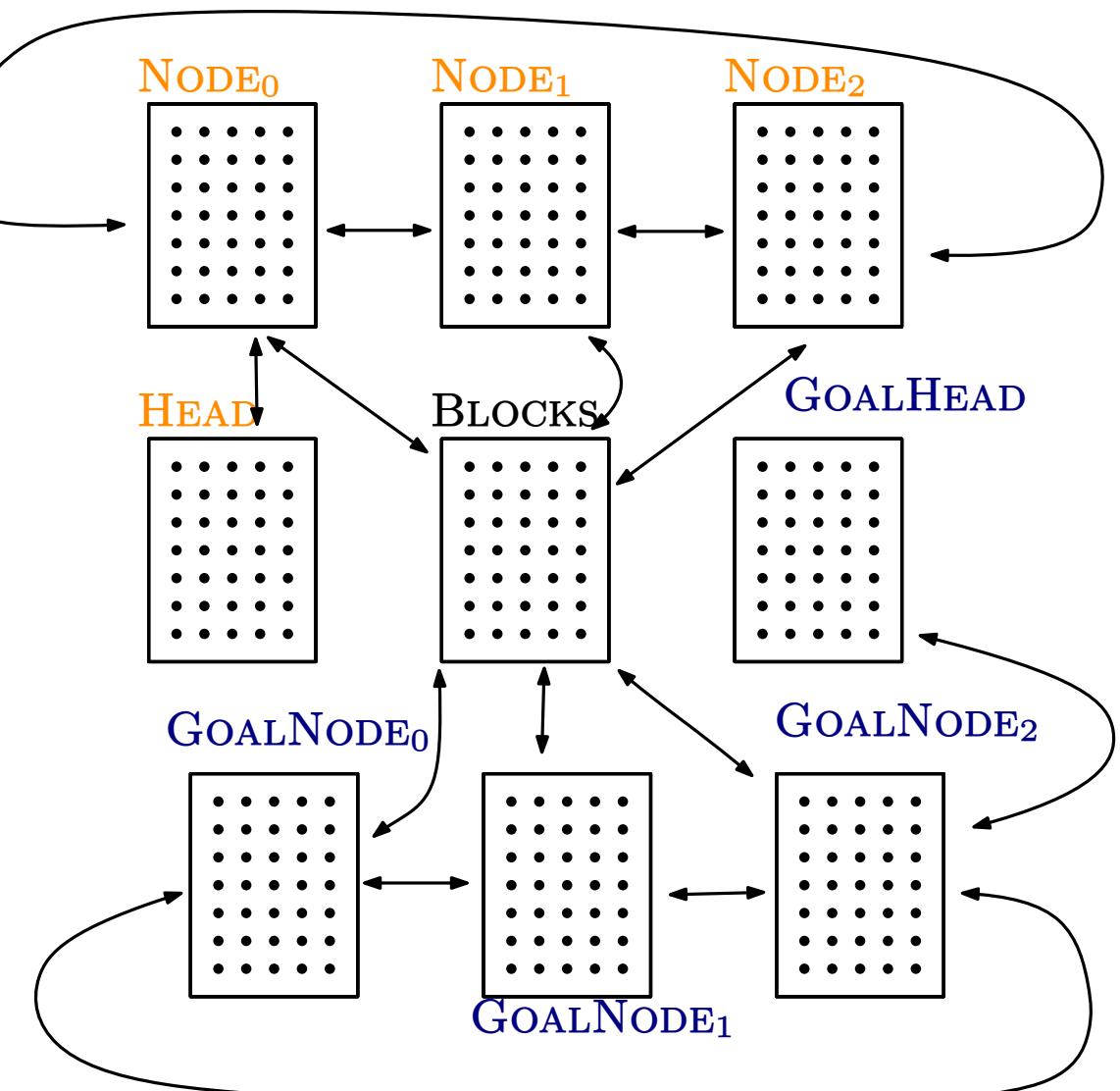
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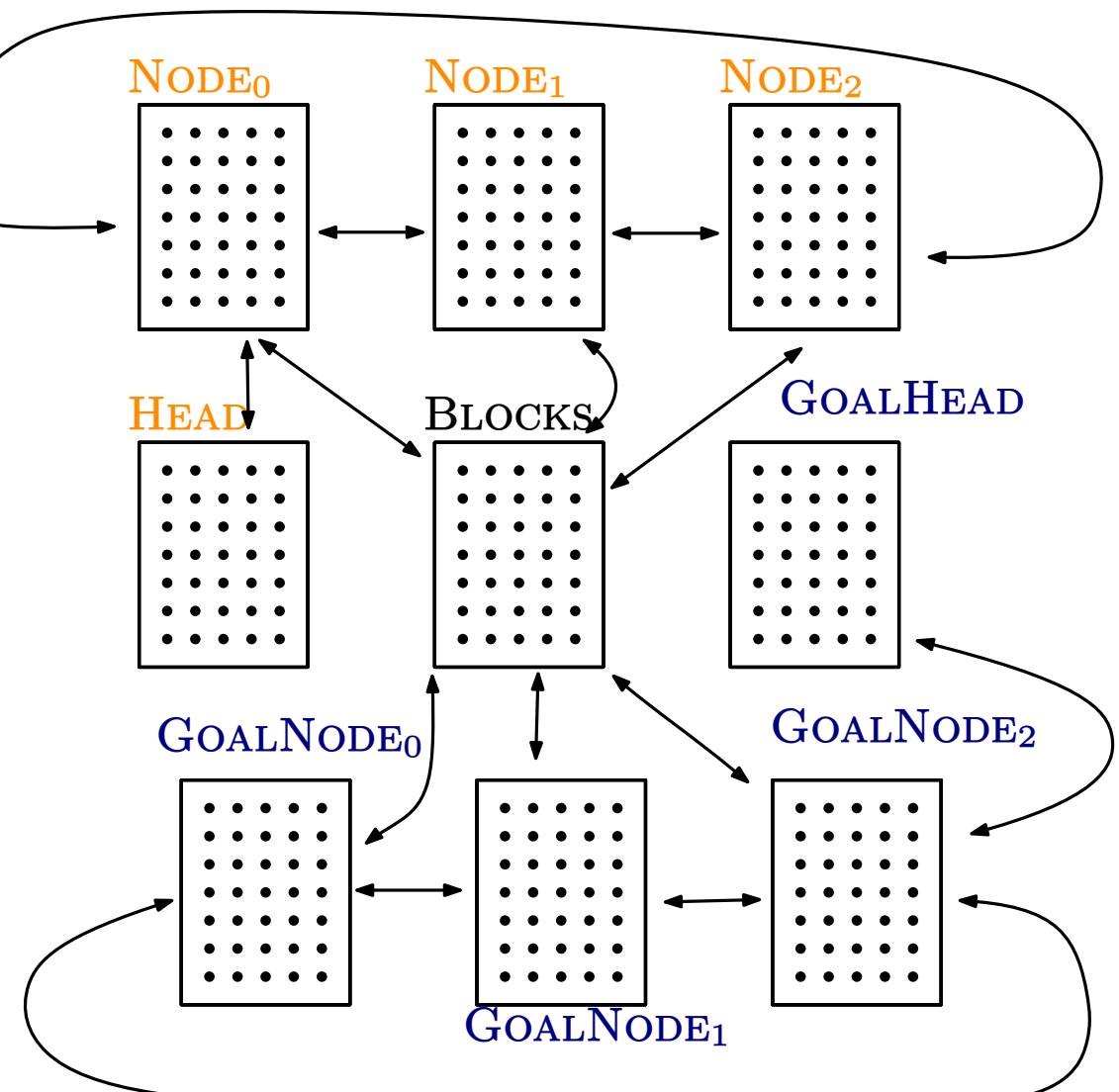
- *Intersect*: scrolls down the two **stacks** until reaching the **last block**, then starts **comparing** each pair of blocks by scrolling up



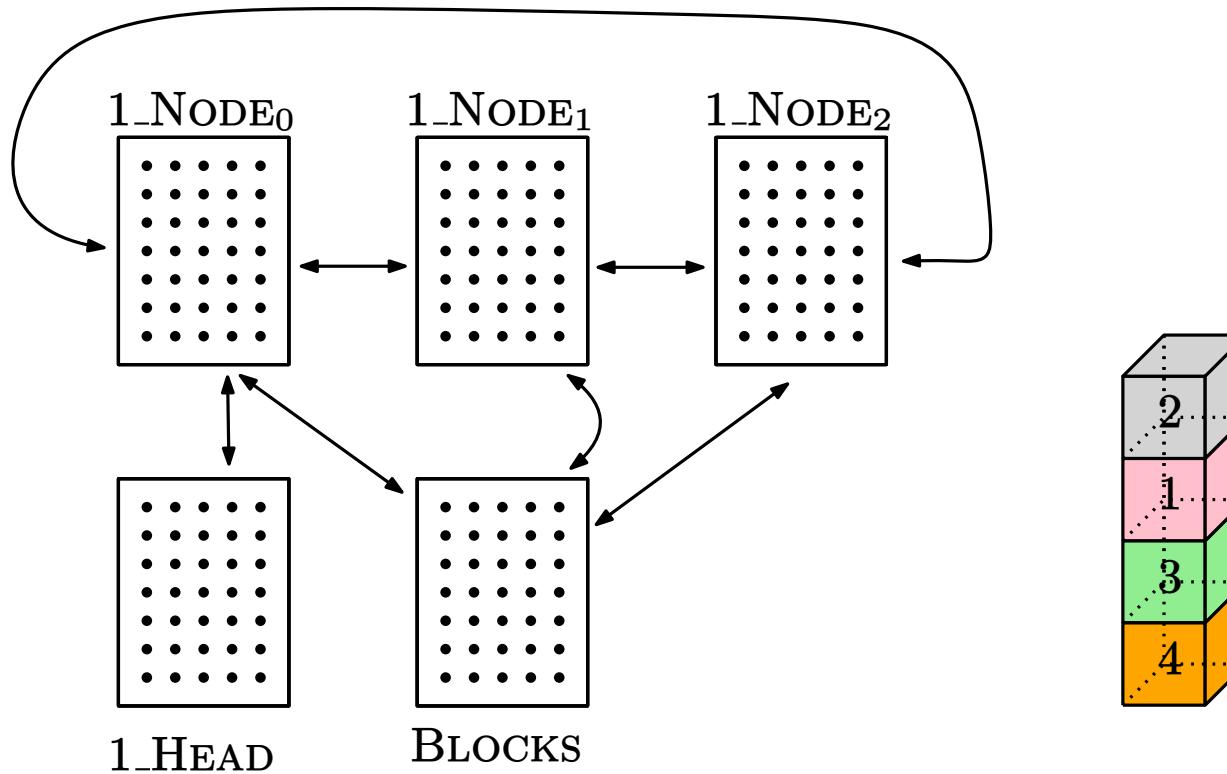
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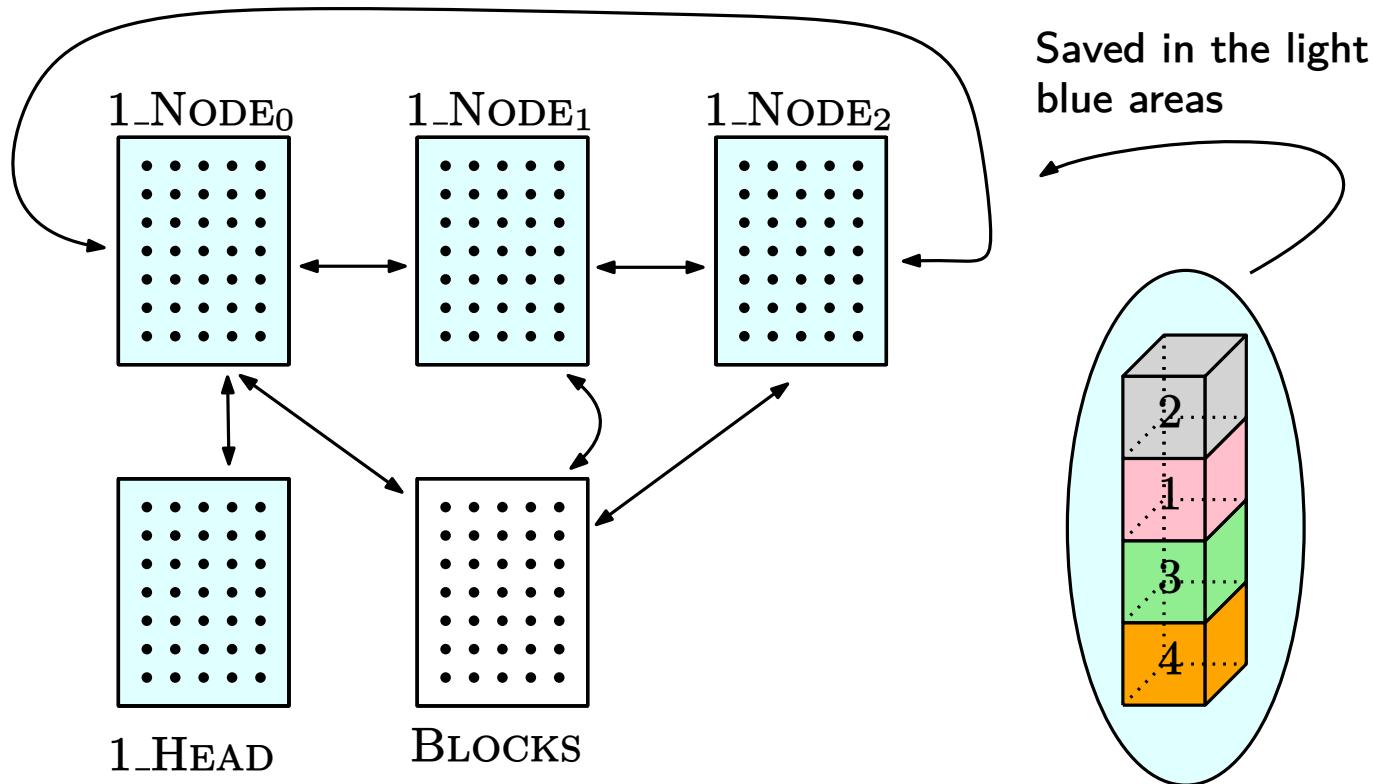
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- By combining these **operations** we implement the **planning strategies**



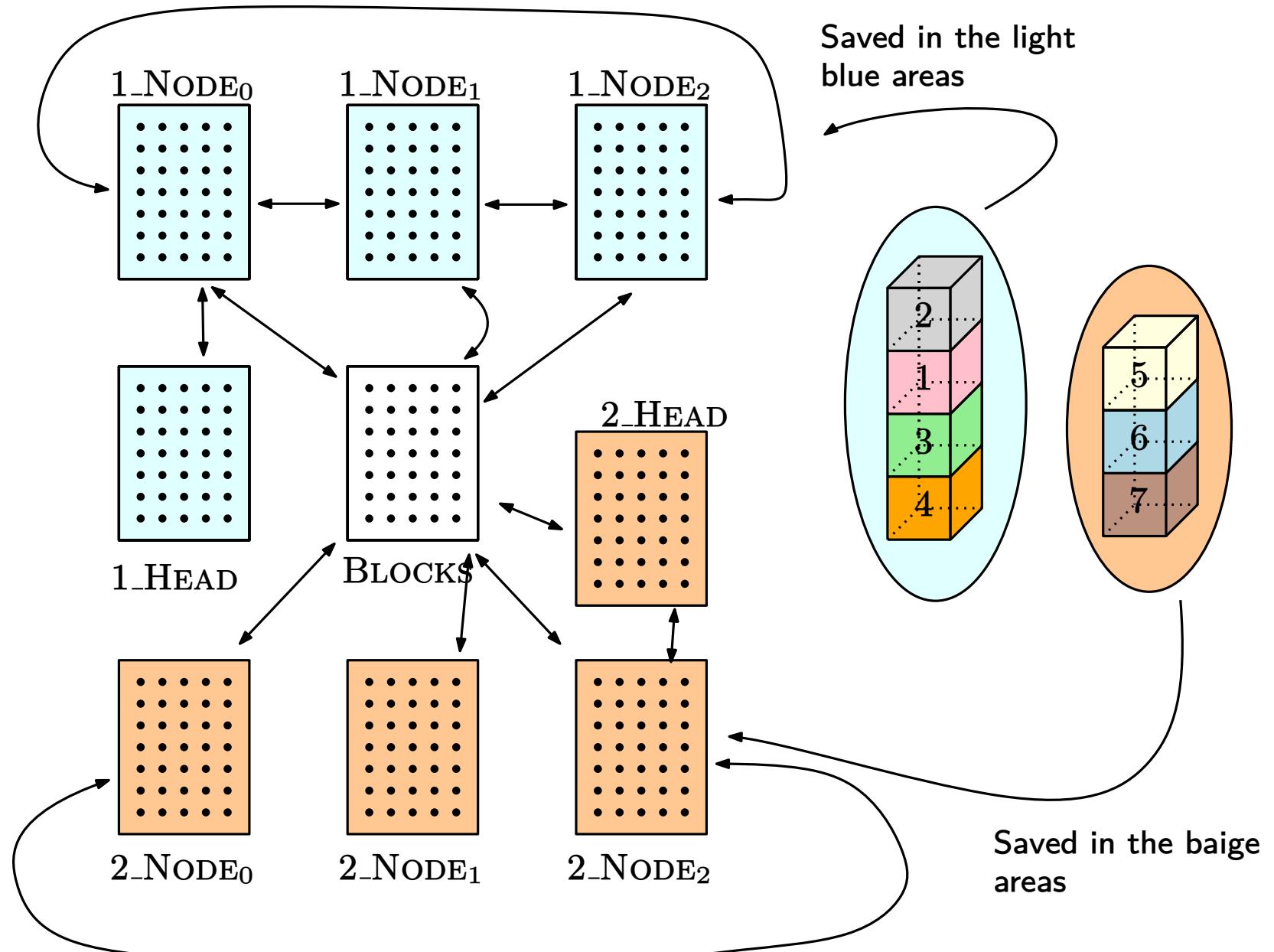
# Multiple Stacks Case



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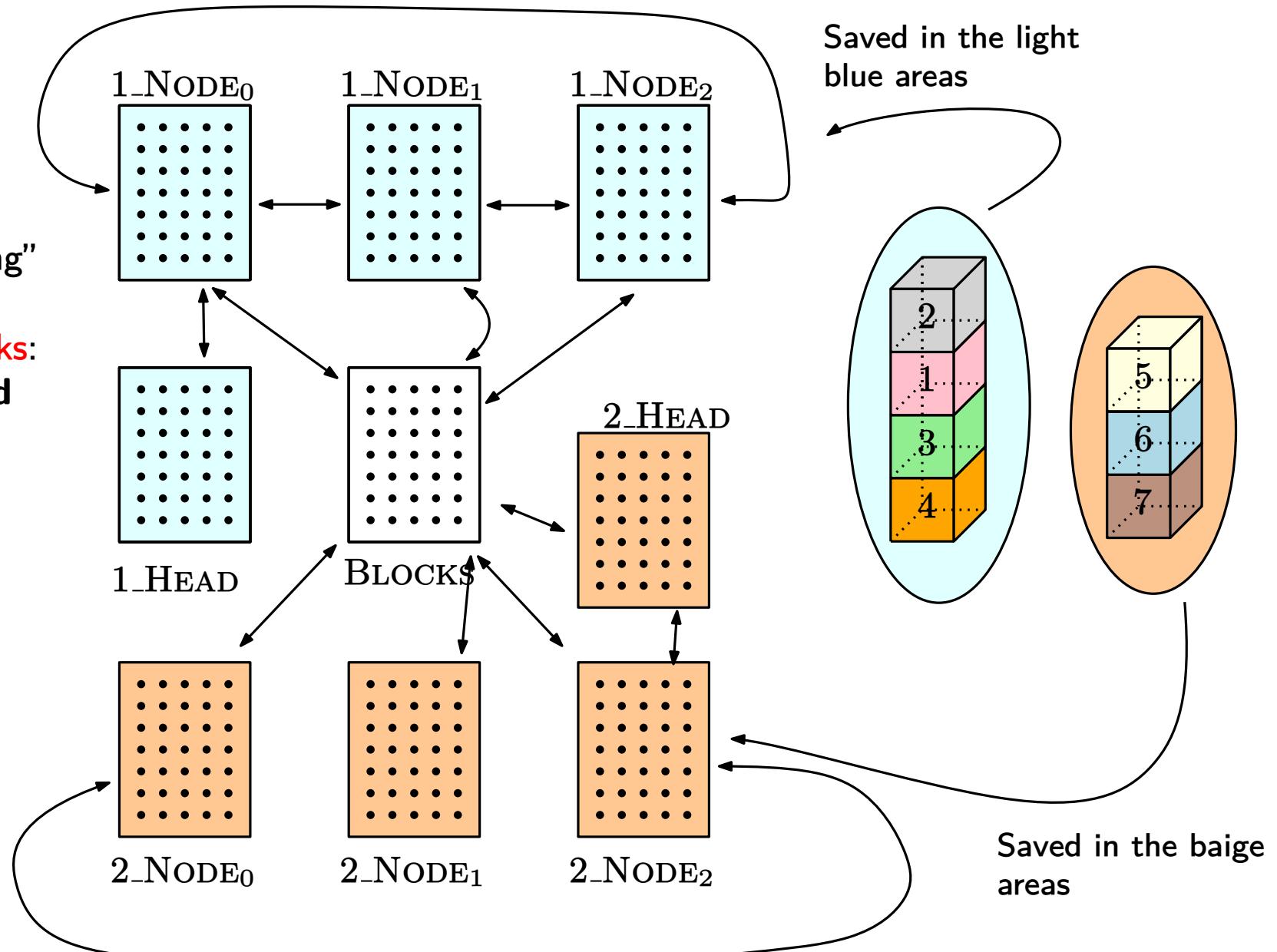


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As many “saving” regions as the  
**number of stacks:**  
**to be improved**  
(like chaining)



# Discussion

- Demonstrated experimentally that reasonably large and complex programs in the assembly calculus can execute correctly and reliably
- Shown the realization of a *list-like data structure* which makes use of a constant number of brain regions
- Shown how simple manipulations of the data structure (removing or putting a block) can be realized by making use of a constant number of brain regions
- Bottleneck: the parsing. Its reliability depends on the ratio between the number of neurons and the size of the assemblies in each region
  - Must be the object of *further investigation*