

# Understanding Evolution:

## A Methodology for Evaluating the Extensibility of Boolean Networks' Structure and Function

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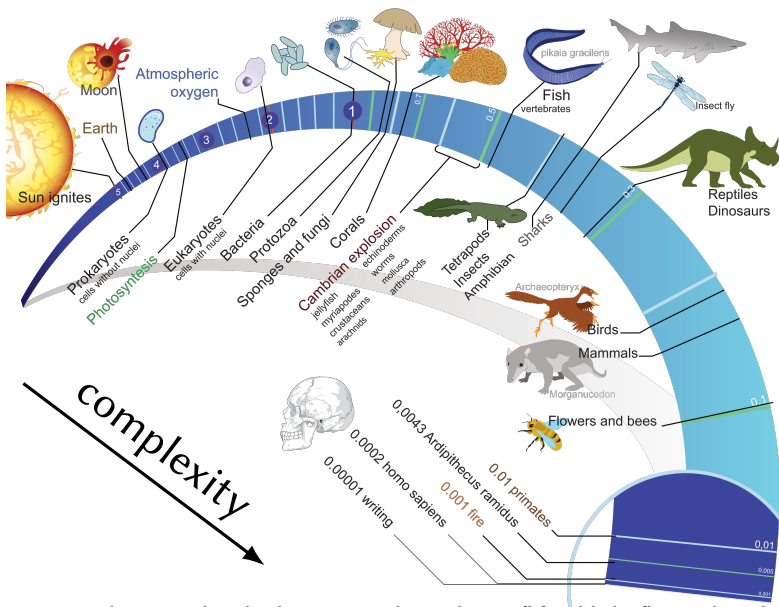
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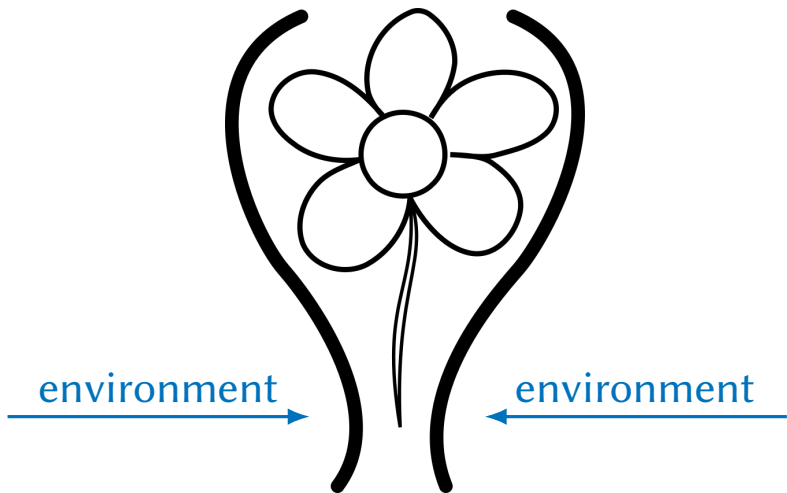
# Evolution



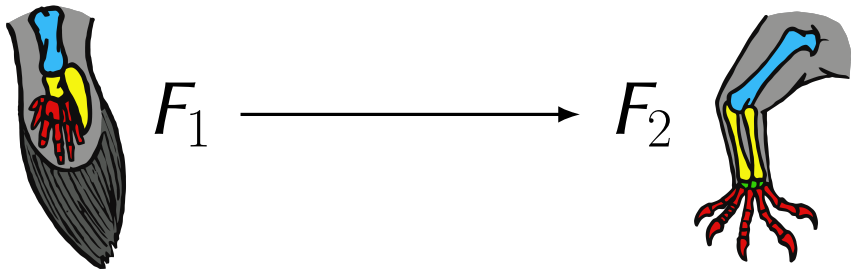
<https://openclipart.org/download/320330/timelineevolutionoflife-pd-ladyofhats-wikimedia.svg>

# Natural selection

reinforces the features best suited to the environment.



# Evolution is gradual

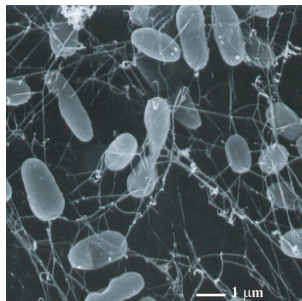
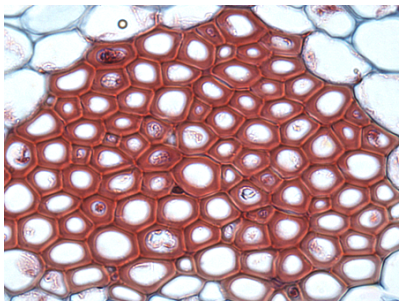


The cards are rarely entirely reshuffled: maintaining the functioning of the whole system is essential for survival.

[https://en.wikipedia.org/wiki/File:Crossopterygii\\_fins\\_tetrapod\\_legs.svg](https://en.wikipedia.org/wiki/File:Crossopterygii_fins_tetrapod_legs.svg)

# Evolution and cancer

Atavistic theory of cancer: cancerous cells devolve to primitive unicellular mode.



<sup>1</sup> [https://en.wikipedia.org/wiki/File:Plant\\_cell\\_type\\_sclerenchyma\\_fibers.png](https://en.wikipedia.org/wiki/File:Plant_cell_type_sclerenchyma_fibers.png)

<sup>2</sup> [https://en.wikipedia.org/wiki/File:Thermophile\\_bacteria2.jpg](https://en.wikipedia.org/wiki/File:Thermophile_bacteria2.jpg)

How does an organism grow more complex, while maintaining original structures and functions?

Which structures are the most extensible?

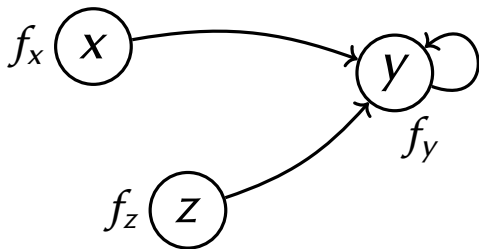
What language to answer this question?

- 1 Threshold/Sign Boolean networks (BN)
- 2 Extensibility in Sign BN
- 3 Complexity of Sign BN
- 4 Preliminary results and analysis



# Boolean networks

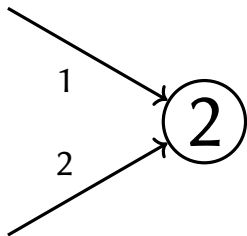
Boolean nodes with Boolean update functions.



$x, y, z \in \{0, 1\}$ , a state:  $\{x : 0, y : 1, z : 0\} \equiv 010$

$$f_x, f_y, f_z : \{0, 1\}^3 \rightarrow \{0, 1\}$$

# TBF: Threshold Boolean Functions



$$f(\mathbf{x}_1, \mathbf{x}_2) \equiv \mathbf{w}_1 \mathbf{x}_1 + \mathbf{w}_2 \mathbf{x}_2 > \theta$$

$$f = (\mathbf{w}_1, \mathbf{w}_2, \theta)$$

$$\mathbf{w}_1 = 1$$

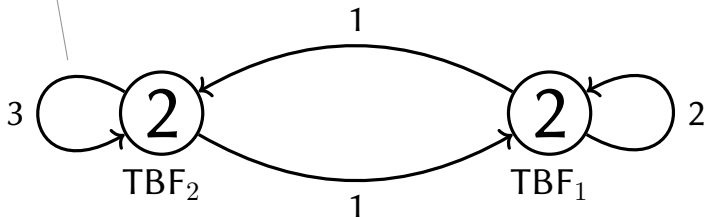
$$\mathbf{w}_2 = 2$$

$$\theta = 2$$

$w_1$	$w_2$	
$x_1$	$x_2$	$f$
0	0	0
0	1	0
1	0	0
1	1	1

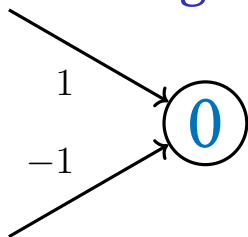
# TBN: Threshold Boolean networks

autoactivation



interaction graph

# SBF/N: Sign Boolean functions/networks



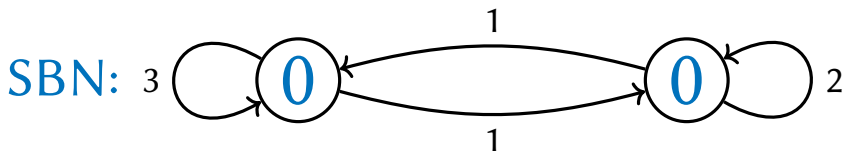
$$f(\mathbf{x}_1, \mathbf{x}_2) \equiv w_1 x_1 + w_2 x_2 > 0$$

$x_1$	$x_2$	$f$
0	0	0
0	1	0
1	0	1
1	1	0

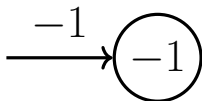
$$f = (w_1, w_2)$$

$$w_1 = 1$$

$$w_2 = -1$$



# SBF < TBF



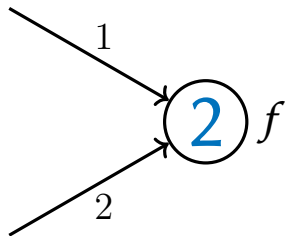
$$\begin{aligned} f &= (w_1 = -1, \theta = -1) \\ &\equiv \neg x_1 \end{aligned}$$

$x_1$	$f$
0	1
1	0

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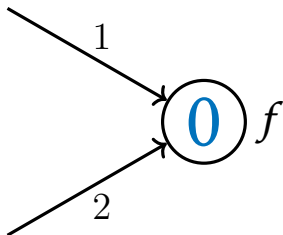
$$\forall f \in \text{SBF} : f(\mathbf{0}) = 0$$

# SBN vs. TBF



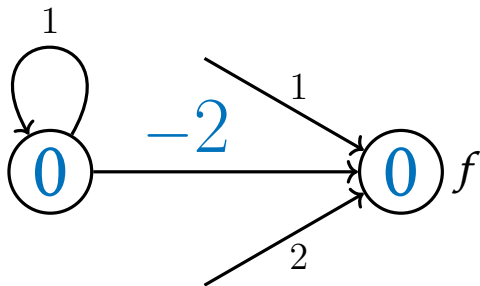
$x_1$	$x_2$	$f$
0	0	0
0	1	0
1	0	0
1	1	1

# SBN vs. TBF



$x_1$	$x_2$	$f$
0	0	0
0	1	1
1	0	1
1	1	1

# SBN vs. TBF



$x_1$	$x_2$	$f$
0	0	0
0	1	0
1	0	0
1	1	1

TBF = SBF + a hidden SBF



# TBF vs. propositional BF

$w_1$	$\theta$	1	1	0		1	1	1
$-1$	$-1$	$x_1$	$x_2$	$x_1 \vee x_2$		$x_1$	$x_2$	$x_1 \wedge x_2$
$x_1$	$\neg x_1$	0	0	0		0	0	0
0	1	1	0	1		1	0	0
1	0	1	1	1		1	1	1

$\{\neg, \vee, \wedge\}$  is a basis for BF

$\implies$  any BF = a combination of TBF (a TBN)

# SBN vs. TBN vs. propositional BN

**Propositional BN:** much power hidden in the formula

**TBN:** the power is explicit in the connections

↪ the interaction graph completely specifies the SBN

**SBN:** a more symmetric and explicit model

↪ all numerical parameters have the same role

↓ syntactic simplicity

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$$\text{power(SBN)} = \text{power(TBN)} = \text{power(propositional BN)}$$

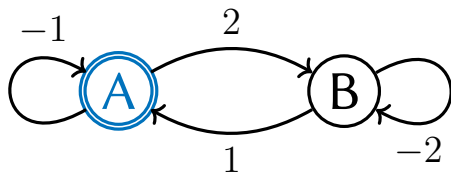
# Update modes

We typically consider the **synchronous** mode

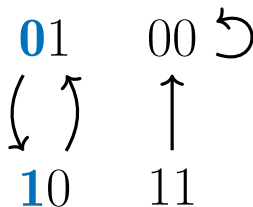
↔ all variables are updated at every step.

Other modes can be considered, too.

# SBN: Transition graphs and outputs



interaction graph



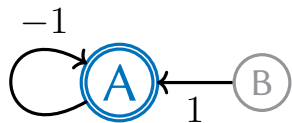
transition graph

Output = the binary sequence observed on **A**

Synchronous mode  $\implies$  outputs  $\subseteq uv^*$

$u \in \{0, 1\}^k$  – the pre-period       $v \in \{0, 1\}^m$  – the period

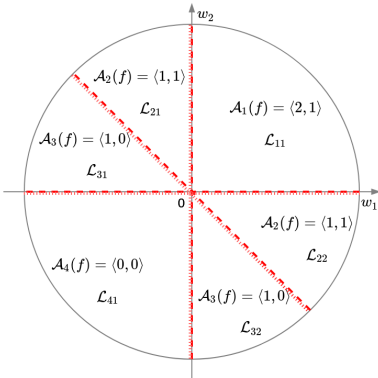
# Equivalence by truth tables



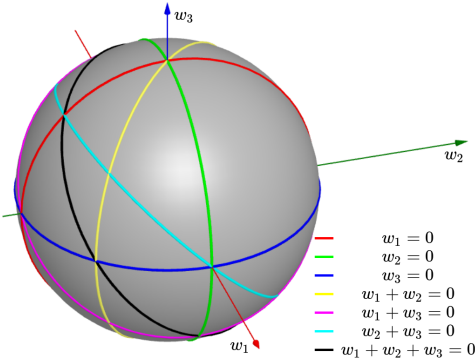
A	B	A	
0	0	0	$0 \leq 0$
0	1	1	$w_B > 0$
1	0	0	$w_A \leq 0$
1	1	0	$w_A + w_B \leq 0$

The inequalities determine subspaces of the parameter space = equivalence classes

# Equivalence classes



$d = 2$



$d = 3$

1 Threshold/Sign Boolean networks (BN)

2 Extensibility in Sign BN

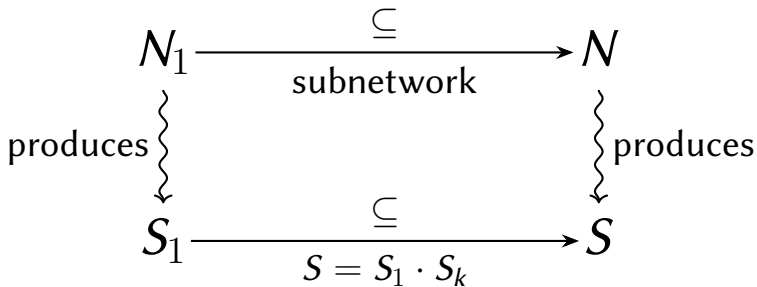
3 Complexity of Sign BN

4 Preliminary results and analysis

# Definition of extensibility in SBN

How can an SBN gain a new function, maintaining

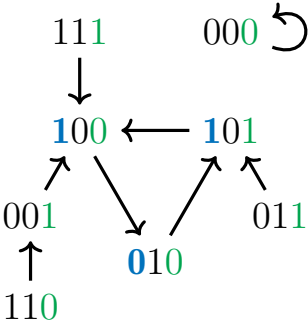
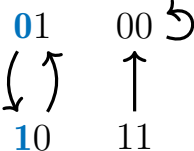
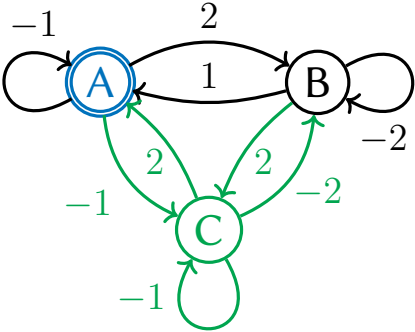
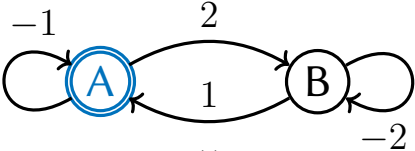
- the prior structures? (interaction graph)
- the prior functions?



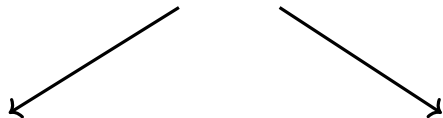
commutative diagram



# Extensibility in SBN: an example



## Growth of a network



more robustness?

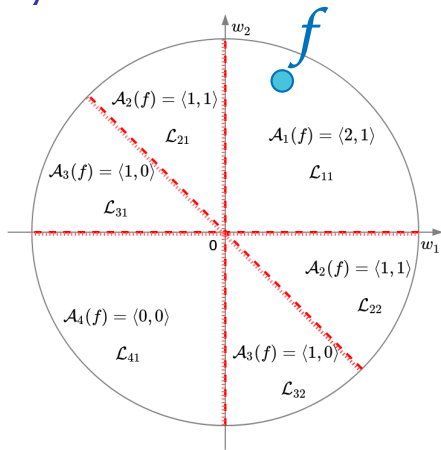
more functions?

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Which structures are the most extensible?

**Complexity** is a good way to discriminate between SBN.

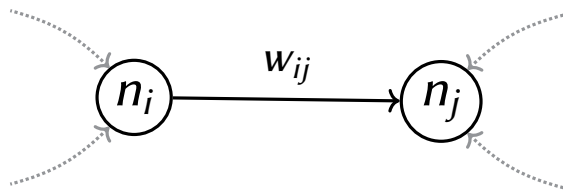
# Complexity of an SBF



$\mathcal{C}^f = \text{complexity}(f) = \text{probability of not picking } f$   
 $\hookrightarrow$  uniform distribution over the unit ball

Complex function  $\implies$  low probability

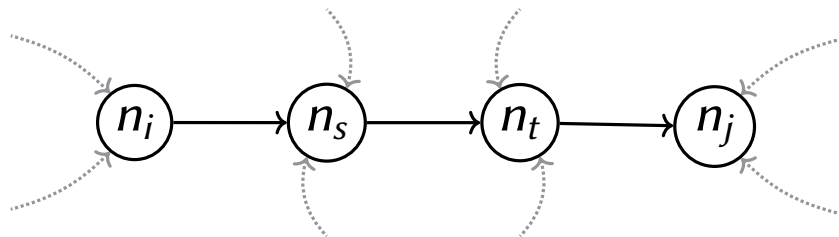
# Complexity of an edge $E_{ij}$ in SBN



The probability that  $(n_i)$  influences  $(n_j)$  via  $E_{ij}$ :

$$P^I(E_{ij}) = \frac{|w_{ij}|}{d \sum_{k=1} |w_{kj}|}$$

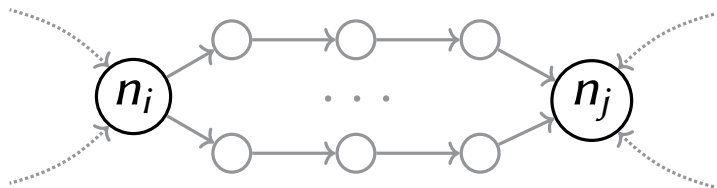
# Complexity of a path in SBN



normalize by the length of the path

$$P^l(Path_{ab}) = \sqrt[3]{P^l(E_{is}) \cdot P^l(E_{st}) \cdot P^l(E_{tj})}$$

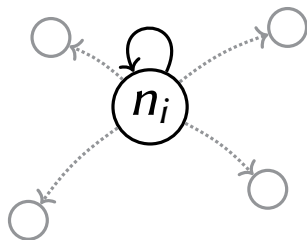
# Probability that $n_i$ influences $n_j$



$$P_{ab}^I = P \left( \bigcup_{Path \in Paths_{ij}} Path \right)$$



# A centrality of a node



centrality( $n_i$ ) = probability that  $n_i$  influences at least one other node, including itself

$$C_i^s = P \left( \bigcup_{k=1}^d \left( \bigcup_{Path \in Paths_{ik}} Path \right) \right)$$

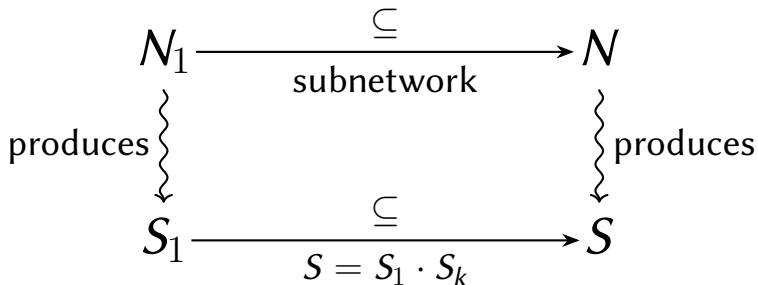
# Complexity of an SBN

Product of complexities of connected nodes, modulated by their centralities:

$$\mathcal{C}(SBN) = \prod_{i=1}^d \begin{cases} \mathcal{C}_i^f \times \mathcal{C}_i^s & \text{if } \mathcal{C}_i^s > 0 \\ 1 & \text{otherwise} \end{cases}$$

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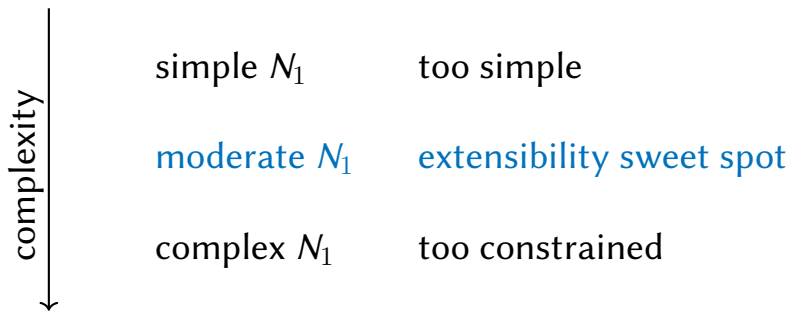
# Extensibility = function(complexity)



Which  $N_1$  yield the most extensions?

# Hypothesis

Which  $N_1$  yield the most extensions?



# Testing the hypothesis for $d = 2$

- 1 Enumerate all  $N_1$ .
- 2 For each  $N_1$ , fix an output node,  $S_1$ , and  $S$ .
- 3 Enumerate all  $N$  for which the diagram commutes:

$$\begin{array}{ccc} N_1 & \xrightarrow{\subseteq} & N \\ \downarrow \wr & & \downarrow \wr \\ S_1 & \xrightarrow{\subseteq} & S \end{array}$$


- 4 Compute the complexities of  $N_1$ ,  $N$ ,  $S_1$ , and  $S$ .

a version of Kolmogorov complexity  
adapted to short strings

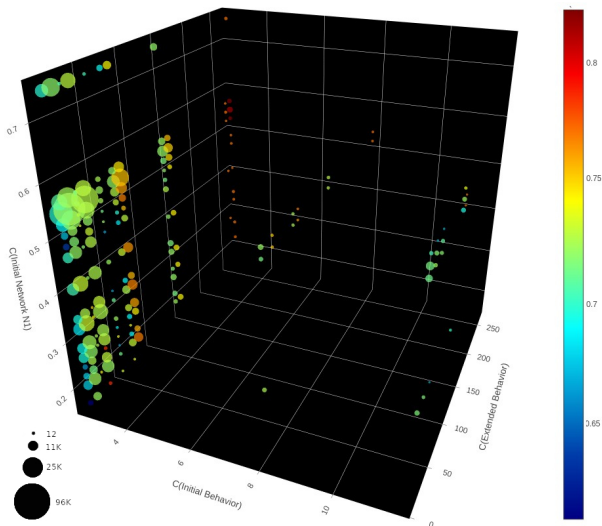
Soler-Toscano F., Zenil H., Delahaye J.-P., Gauvrit N.: Calculating Kolmogorov complexity from the output frequency distributions of small Turing machines; PLoS ONE 9(5): e96223 (2014)

# Combinatorics of the method

$d$	$SBF$	$SBN$	$\{N_1, S_1, S\}$	$N$
2	7	101	$96 \cdot 10^3$	777216
3	17	$206 \cdot 10^3$	$[180 \cdot 10^6 - 32 \cdot 10^9]$	$[1.5 \cdot 10^9 - 280 \cdot 10^9]$
4	47	$76 \cdot 10^9$		

  
estimations

# First results



Color = average complexity of SBN in the class

Point size = number of networks in the class (log-scale)



# Discussion

- Behaviours of most  $d = 2$  networks are not complex.
- Most extensions have moderate structural and behavioural complexity.
- Complex extended behaviors and networks are mostly obtained from networks of moderate complexity.

# Improvement directions

- Refine the analysis.
- Look into higher dimensions.
- Consider other modes of evolution:
  - ▶ increase in the number of attractors,
  - ▶ increase in the size of basins of attraction.

simple  $\frac{\triangleleft \triangleright}{??}$  complex

Rémi  
Segretain



Sergiu  
Ivanov



Laurent  
Trilling



Nicolas  
Glade



Special thanks to

Ibrahim  
Cheddadi



Communauté  
UNIVERSITÉ Grenoble Alpes

université  
PARIS-SACLAY

★ île de France

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