

Ask not what algebra can do for biology –  
ask what biology can do for algebra

Workshop on  
*Model Theory of Differential Equations, Algebraic Geometry,  
and their Applications to Modeling*

BIRS

June 5, 2020

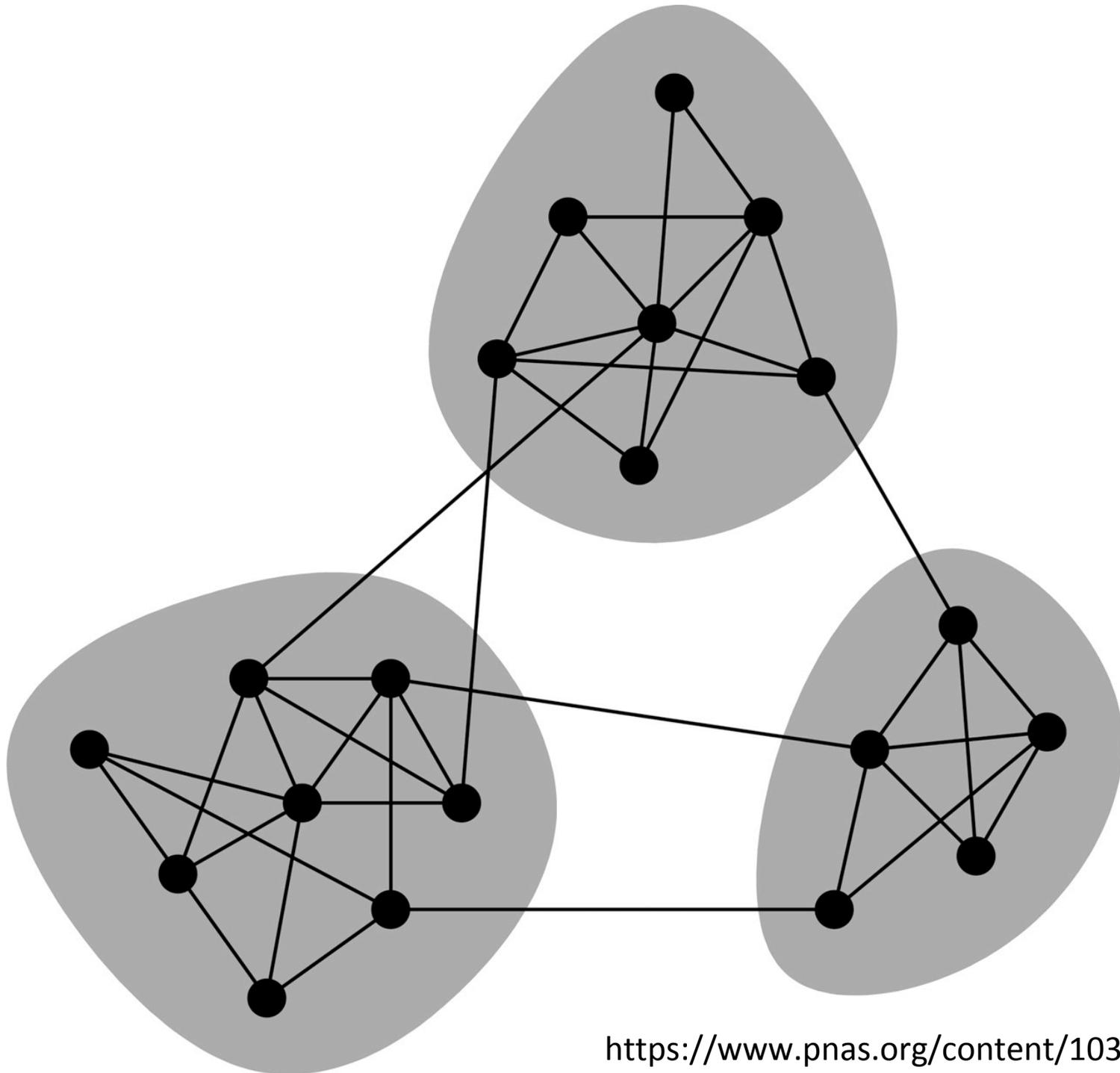
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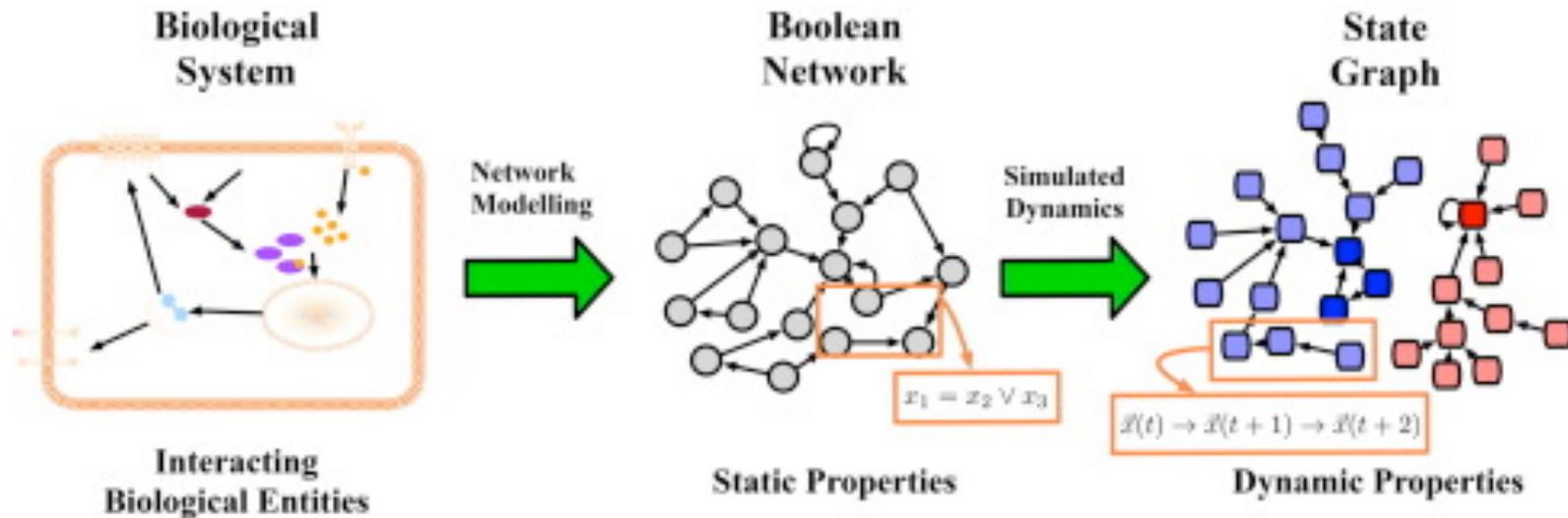


“Nature (likely) has Structure. And many models in natural sciences inherit a part of it. Therefore, understanding and exploiting the structure of a model at hand might be crucial to make the model useful. Algebra and logic offer a variety of tools to work with structures and greatly benefit from new types of structures and structural questions coming from other areas.”

‘Modularity is a widespread property in biological systems.’



**Concepts in Boolean network modeling :  
What do they all mean?**



## Definition

A *polynomial dynamical system (PDS)* over a finite field  $k$  is a function

$$f = (f_1, \dots, f_n) : k^n \longrightarrow k^n,$$

with the coordinate functions  $f_i : k^n \longrightarrow k$  in the polynomial ring  $k[x_1, \dots, x_n]$ .

Iteration of  $f$  results in a time discrete dynamical system over the space  $k^n$ .

**Note:** Any function  $k^n \longrightarrow k$  can be expressed as a polynomial.

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ORIGINAL ARTICLE

## The Dynamics of Conjunctive and Disjunctive Boolean Network Models

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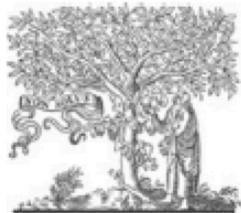
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### The number of periodic points

**Theorem 3** *Let  $f$  be a conjunctive Boolean network whose dependency graph is strongly connected and has loop number  $c$ . If  $c = 1$ , then  $f$  has the two fixed points  $(0, 0, \dots, 0)$  and  $(1, 1, \dots, 1)$  and no other limit cycles of any length. If  $c > 1$  and  $m$  is a divisor of  $c$ , then the number of periodic states of period  $m$  is*

$$|A(m)| = \sum_{i_1=0}^1 \cdots \sum_{i_r=0}^1 (-1)^{i_1+i_2+\cdots+i_r} 2^{p_1^{k_1-i_1} p_2^{k_2-i_2} \cdots p_r^{k_r-i_r}},$$

where  $m = \prod_{i=1}^r p_i^{k_i}$  is the prime factorization of  $m$ , that is  $p_1, \dots, p_r$  are distinct primes and  $k_i \geq 1$  for all  $i$ .



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Brief paper

## Dynamics of semilattice networks with strongly connected dependency graph<sup>☆</sup>

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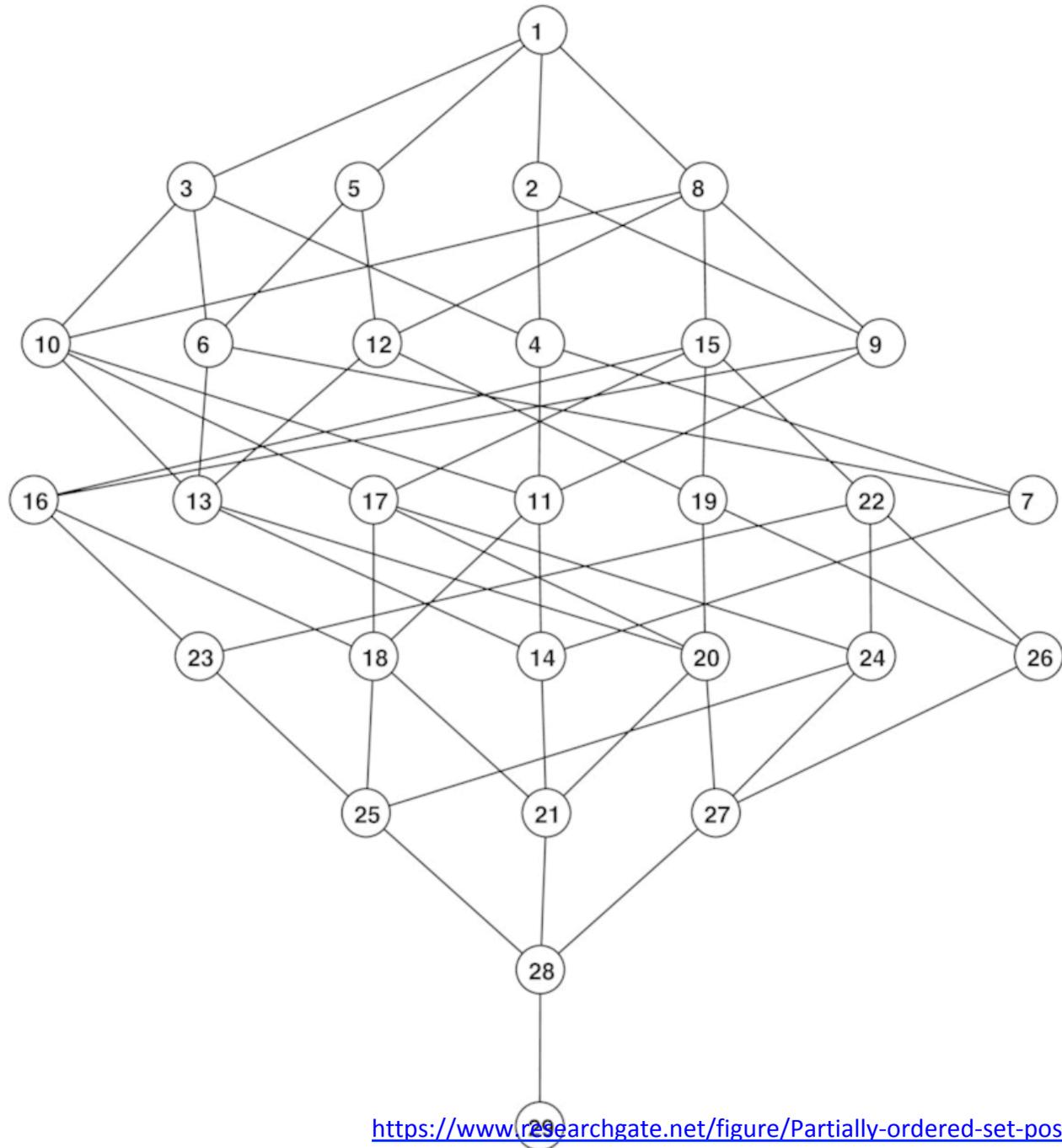
**Theorem 6.2.** *Consider the function*

$$\mathcal{L}(z_1, \dots, z_t) := \sum_{\mathcal{J} \subseteq \Omega} (-1)^{|\mathcal{J}|+1} \prod_{j \in \bigcap_{J \in \mathcal{J}} J} z_j.$$

*Then for any conjunctive Boolean network  $f$  with subnetworks  $h_1, \dots, h_t$  and  $\Omega$  its set of maximal antichains in the poset of  $f$ , we have*

$$\mathcal{L}(\mathcal{C}(h_1), \dots, \mathcal{C}(h_t)) \leq \mathcal{C}(f). \tag{9}$$

*Here, the function  $\mathcal{L}$  is evaluated using the “multiplication” described in Corollary 3.5. This inequality provides a sharp lower bound on the number of limit cycles of  $f$  of a given length.]*



[https://www.researchgate.net/figure/Partially-ordered-set-poset-model-for-cognitive-functioning-of-mild-cognitive\\_fig4\\_236052149](https://www.researchgate.net/figure/Partially-ordered-set-poset-model-for-cognitive-functioning-of-mild-cognitive_fig4_236052149)

## **Modularity for dynamic biological systems/models**

- Given a model, compute its modules, their attractor structure, and information about the attractor structure of the model itself.
- Characterize the “degrees of freedom” to combine simple models.

# A “Hölder Program” for BNs

- Identify a class of “decomposable” BNs.
- Identify a class of decomposable BNs that are “simple” and sufficiently “rich.”
- Define a notion of “quotient” of a BN by a subnetwork.
- Show that each decomposable BN has a filtration by subnetworks so that each successive quotient is a simple network.
- Classify the different ways in which decomposable BNs can be built as extensions of two BNs that are simpler.
- Rigorous definition of “dynamic equivalence” of BNs.
- Develop a category-theoretic foundation for this program.



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## Nested analyzing, unate cascade, and polynomial functions<sup>☆</sup>

Abdul Salam Jarrah<sup>a,\*</sup>, Blessilda Raposa<sup>b</sup>, Reinhard Laubenbacher<sup>a</sup>

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<sup>b</sup> *Mathematics Department, De La Salle University, 2401 Taft Avenue, Manila, Philippines*

$$\begin{aligned}
f(x_1, x_2, \dots, x_n) &= (x_1 - a_1)[(x_2 - a_2)[\dots [(x_{n-1} - a_{n-1})(x_n - a_n) \\
&\quad + (b_n - b_{n-1})] + (b_{n-1} - b_{n-2})] \dots] \\
&\quad + (b_2 - b_1)] + b_1
\end{aligned}$$

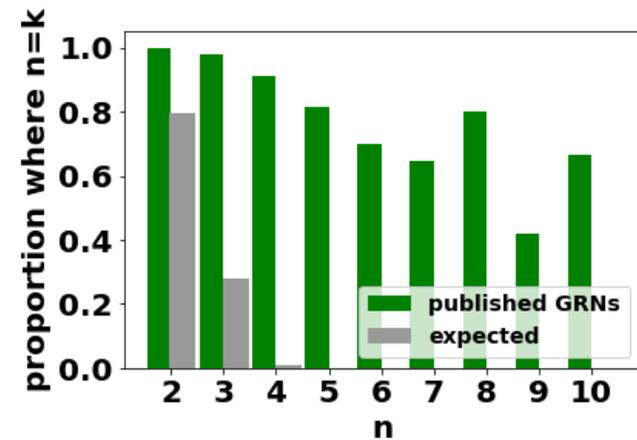
*or, equivalently,*

$$\begin{aligned}
f(x_1, x_2, \dots, x_n) &= \prod_{i=1}^n (x_i - a_i) \\
&\quad + \sum_{j=1}^{n-1} \left[ (b_{n-j+1} - b_{n-j}) \prod_{i=1}^{n-j} (x_i - a_i) \right] + b_1.
\end{aligned}$$

# Prevalence of canalization

		k										
		0	1	2	3	4	5	6	7	8	9	10
0	2											
1	0	1688										
2	0	0	744									
3	6	1	0	411								
4	23	2	0	0	251							
5	19	8	2	0	0	133						
6	17	5	6	0	1	0	66					
7	7	4	3	1	1	0	0	27				
8	4	1	0	2	0	0	0	0	26			
9	2	3	3	0	0	3	0	0	0	7		
10	0	0	1	0	1	0	0	0	0	0	4	

observed



- Nested canalizing functions (and therefore? canalizing functions) are overrepresented in GRNs.



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Advances in Applied Mathematics 30 (2003) 655–678

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# Decomposition and simulation of sequential dynamical systems

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## Future work

- Find a version of the classification of monomial networks in the language of computational algebra.
- Study the properties of nested canalizing polynomials.
- Carry out the Hölder Program for larger classes of networks, for instance, AND-NOT networks.

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