

# Overview of Network Models

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Network Science and Social Networks at the U (NETSNAU)

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# Contents

## Modeling Networks

## Taxonomy of Methods

- Modeling Structure

- Modeling Behavior

- Modeling Structure x Behavior

## More on Modeling Complexity

You can download the slides from [ggv.cl/slides/netsnau0](http://ggv.cl/slides/netsnau0)

# The question

“I have data ABC and hypotheses XYZ...

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What **method** should I use to  
**address these questions?”**

# Modeling Networks

## Taxonomy of Methods

Modeling Structure

Modeling Behavior

Modeling Structure x Behavior

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# Modeling Networks

Today (in a non-comprehensive survey)

Network Models

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Modeling Networks

Taxonomy of Methods

Modeling Structure

Modeling Behavior

Modeling Structure x  
Behavior

More on Modeling  
Complexity

Referencias

# Modeling Networks

Today (in a non-comprehensive survey)

- ▶ Classical statistical analysis assumes observations distribute *independently* and *identically* to each other

Network Models

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Modeling Networks

Taxonomy of Methods

Modeling Structure

Modeling Behavior

Modeling Structure x  
Behavior

More on Modeling  
Complexity

Referencias

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Modeling Networks

Taxonomy of Methods

Modeling Structure

Modeling Behavior

Modeling Structure x  
Behavior

More on Modeling  
Complexity

Referencias



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Nonetheless...

# Modeling Networks (cont. 1)

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- ▶ And many communities, e.g., and , accelerating the science.

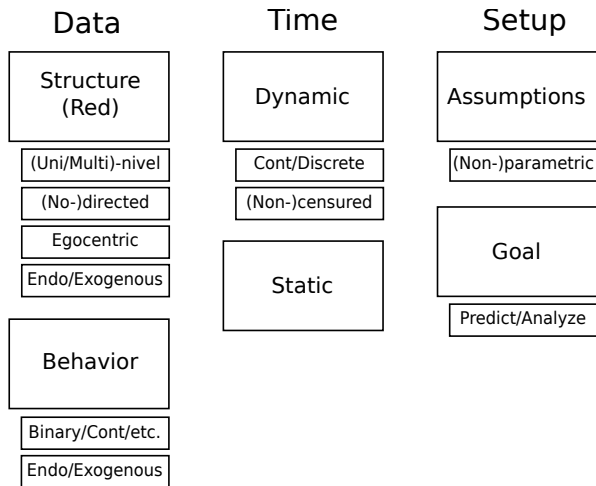
# Modeling Networks (cont. 1)

- ▶ We now have more data.
- ▶ More computational power (Hofman y col., 2021; Lazer y col., 2020).
- ▶ And many communities, e.g., and , accelerating the science.

We can (and should) have a more systematic view of modeling networks and complex systems.



# Modeling Networks Complexity (cont. 2)



**Figura:** Different dimensions of SNA (a view from the data.) All components interact with each other.

## Modeling Networks

### Taxonomy of Methods

Modeling Structure

Modeling Behavior

Modeling Structure x Behavior

### More on Modeling Complexity

# Taxonomy of Methods: A proposal

Two key dimensions, **structure** vs **behavior**, in particular

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Modeling Networks

**Taxonomy of Methods**

Modeling Structure

Modeling Behavior

Modeling Structure x  
Behavior

More on Modeling  
Complexity

Referencias

# Taxonomy of Methods: A proposal

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# Taxonomy of Methods: A proposal

Two key dimensions, **structure** vs **behavior**, in particular

- ▶ Structure: We just care about the network.
- ▶ Behavior: Behavior given the existence of a network.
- ▶ Structure x Behavior: We want to understand the co-evolution/distribution of network and behavior.

# Taxon: Structure

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Modeling Networks

Taxonomy of Methods

Modeling Structure

Modeling Behavior

Modeling Structure x  
Behavior

More on Modeling  
Complexity

Referencias

# Taxon: Structure

## Non-parametric

- ▶ **Network Bootstrap** Standard errors and graph-level contrasts.
- ▶ **Network rewiring algorithms (e.g. CUG test)** *Motifs* detection conditioning on observables (e.g., sequence or degree distribution)
- ▶ **Quadratic Assignment Procedure (QAP)** Label permutation for hypothesis testing.



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## Parametric

- ▶ **Exponential Random Graph Models (ERGMs)** Including all its flavors, like, TERGMs, BERGMs, ERGMitos, etc.
- ▶ **Relational Event Models (REMs) y Dynamic Actor Network Models (DyNAMs)** Sequence of interactions throughout time.

Ref: Butts (2008a, 2008b), Caimo y Friel (2014), Krackhardt (1988), Robins y col. (2007), Snijders y Borgatti (1999), Stadtfeld y Block (2017) y Vega Yon y col. (2021)

# Taxon: Behavior

Network Models

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Modeling Networks

Taxonomy of Methods

Modeling Structure

**Modeling Behavior**

Modeling Structure x  
Behavior

More on Modeling  
Complexity

Referencias

# Taxon: Behavior

## Non-parametric

- ▶ **Permutation tests:** Simple or conditioned (ej. en *in-degree*)

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## Parametric

- ▶ **Spatial Autocorrelation:** Like Moran's I.
- ▶ **Spatial Autoregressive Models:** Like GLMs, but assuming that errors are jointly distributed in a graph-like fashion.
- ▶ **Lagged regressions** Entities' behavior is a function of previous exposure to peers.

Ref: Butts (2008a), LeSage (2008), Moran (1950) y Valente y col. (2019)

# Taxon: Structure x Behavior

Network Models

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Modeling Networks

Taxonomy of Methods

Modeling Structure

Modeling Behavior

Modeling Structure x  
Behavior

More on Modeling  
Complexity

Referencias

# Taxon: Structure x Behavior

## Non-parametric

- ▶ **Agent-Based Models (ABM)** Simulation of Complex systems.
- ▶ **(idem)** used for parameter estimation through, e.g., likelihood-free MCMC and Approximate Bayesian Computation (ABC).

# Taxon: Structure x Behavior

## Non-parametric

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## Parametric

- ▶ **Stochastic Actor Oriented Model (SAOM)** Continuous Markov process in which individuals behavior and embeddedness interact throughout time.
- ▶ **Discrete Exponential-Family Models (DEFMs)** Individuals make multiple decisions on behavior and ties simultaneously (on development by yours truly...).

Ref: Marjoram y col. (2003), Ry y col. (2010), Snijders y col. (2010) y Tisue y Wilensky (2004)

# More on Modeling Complexity

Too many other methods/themes out there:

- ▶ Modeling evolution using dynamic programming (likelihoods on directed graphs) (biology and computer science.) (Felsenstein, 1981)
- ▶ Identifying key entities in networks (centrality measures, structural holes, etc.) (Bringmann y col., 2019)
- ▶ Micro to macro behavior in Cellular Automata. (Wolfram, 1983)
- ▶ Bayesian Networks (treating variables as nodes.) (Heckerman, 2008)
- ▶ Networks of Gene products. (Hecker y col., 2009)
- ▶ Stochastic block models (multi-group networks.) (Holland y col., 1983)
- ▶ Signed graphs (e.g., balance theory, i.e., the friend of my friend is my friend.) (Cisneros-Velarde y col., 2021)
- ▶ Survey and sampling methods for networks (e.g., snowball sampling.) (Goodman, 1961)
- ▶ Statistical mechanics of complex systems (Albert & Barabasi, 2002; Barabási & Pálfai, 2016)
- ▶ ...



# Thank you!

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## Network Models

## Modeling Networks

## Taxonomy of Methods

## Modeling Structure

## Modeling Behavior

## More on Modeling Complexity

## Referencias

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