



# backbone: An R package for extracting the backbone of bipartite projections



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PLoS ONE: <https://doi.org/10.1371/journal.pone.0244363>

All things backbone: [www.zacharyneal.com/backbone](http://www.zacharyneal.com/backbone)

# Backbone



The **backbone** of a bipartite projection is an unweighted or signed network that keeps only the **most important edges** of the projection.

The **backbone** R package provides different methods for extracting this backbone!

You can install it from CRAN and load it like this:

```
install.packages("backbone")  
library(backbone)
```




# Why use the backbone package?




Let's say we use a **co-sponsorship** network to measure political **collaboration** between senators.





When projected, **edge weights = number of co-sponsored bills**



SENATORS

BILLS 

1	0	1



	100	

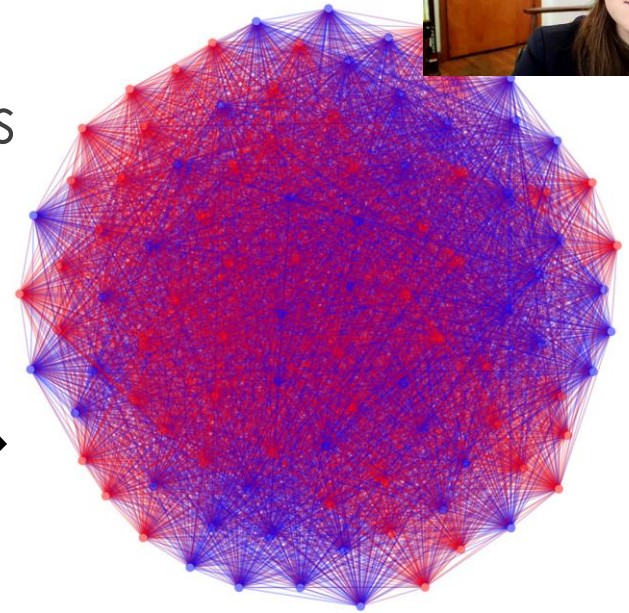
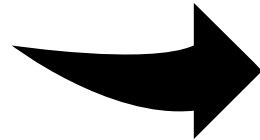
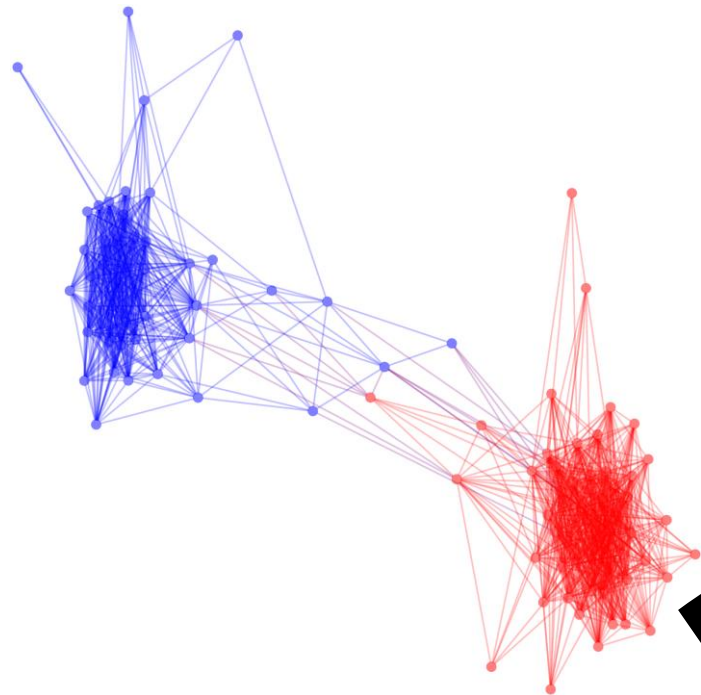
Edge weight depends on how many bills each person sponsors, and how many sponsors each bill tends to have.



# Why use the backbone package?



Here's a visual of the co-sponsorship network of US Senators in the 114<sup>th</sup> session. An edge was kept if two Senators co-sponsored at least one bill together.



Here's that same network using the backbone package's SDSM model! Most important edges are found to exist within party.

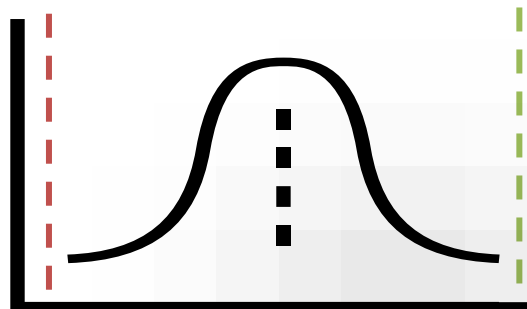


# How Does It Work?



Backbone uses **statistical tests** to compare an edge's observed weight in the bipartite projection to the **distribution of its weights** expected under a **null model**.

An edge's observed weight is **statistically significant** if it is in the upper or lower tail of the distribution.



Upper tail = **positive** edge in the backbone

Lower tail = **negative** edge in the backbone



# Three Different Types of Null Models

Compares the observed edge weight to a distribution of random bipartite graphs where...



## Hypergeometric: `hyperg`

...the row sums exactly match the observed bipartite, *but* column sums are allowed to vary.

## Stochastic Degree Sequence Model: `sds`

...the row sums *and* column sums are allowed to vary, but match on average.

## Fixed Degree Sequence Model: `fdsm`

...the row sums and column sums *exactly* match the observed bipartite.



# Using the backbone package



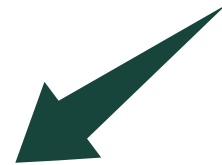
## Your Data

- Adjacency matrix
- A sparse Matrix object
- Bipartite igraph object
- Bipartite network object
- An edgelist



## Apply Null Model

**hyperg**  
**sds**  
**fdsm**



## Get backbone Object

**positive** - matrix with entries equal to the *probability* of an edge under the null distribution having weight equal to or above the weight observed in the projection.

**negative** - probability of equal to or below...



## Extract Backbone

Plug in backbone object to **backbone.extract()** to find a positive or signed backbone at a given significance level.





# Extracting the Backbone



```
backbone.extract(backbone, ←  
                  signed = TRUE, ←  
                  alpha = .01, ←  
                  fwer = "none", ←  
                  class = "original", ←  
                  narrative = TRUE) ←
```

**fwer**: Family-wise error rate test correction. Choose from “none”, “Bonferroni”, and “Holm”

**backbone**: a backbone class object as returned by `hyperg`, `sds`, `fdsm`

**signed**: TRUE for a signed backbone, FALSE for a positive backbone

**alpha**: significance test value (always a two-tailed test)

**class**: type of graph object you’d like back: `matrix`, `igraph`, `network`, `edgelist`

**narrative**: suggested manuscript text and citations





# Hypergeometric Distribution

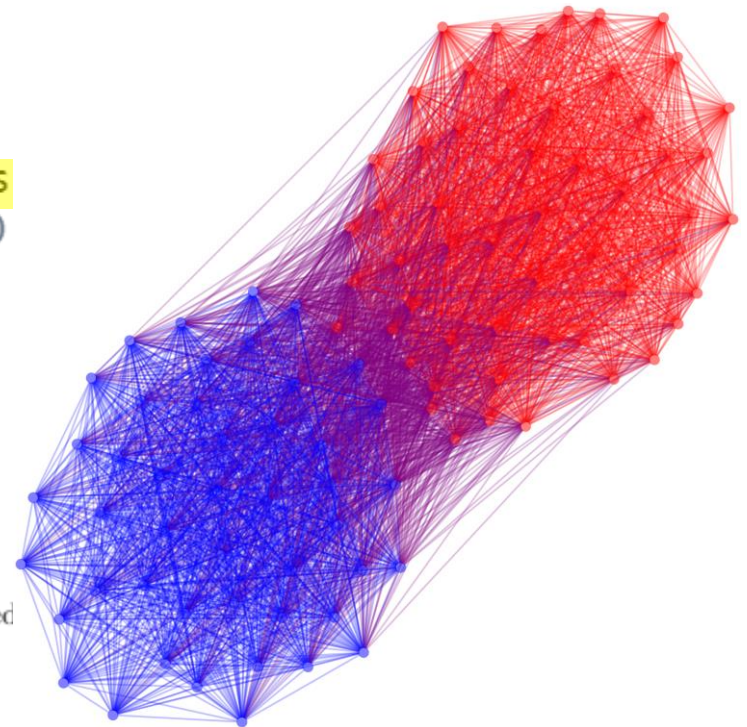


Senate co-sponsorship: fixes the number of bills that each senator sponsored, while allowing each bill to be sponsored by a varying number of senators.

On the Senate co-sponsorship data from the 114<sup>th</sup> session, we get:

```
hyperg_probs <- hyperg(data)
hyperg_bb <- backbone.extract(hyperg_probs,
                              alpha = .01)
```

“backbone” class object that contains the probabilities.  
Plug entire thing into the backbone.extract function



Tumminello M, Miccichè S, Lillo F, Piilo J, Mantegna RN. Statistically Validated Networks in Bipartite Complex Systems. PLOS ONE. 2011;6(3):e17994. doi:10.1371/journal.pone.0017994.



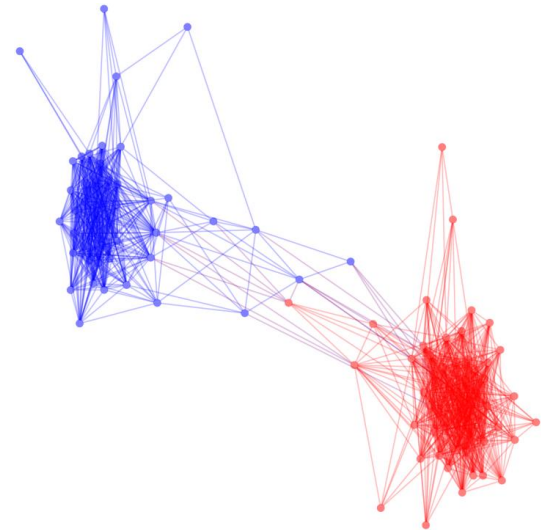
# Stochastic Degree Sequence Model



Senate co-sponsorship: compares the observed values to the distribution where each senator sponsors roughly the same number of bills, and each bill is sponsored by roughly the same number of people.

```
sdsd <- sdsd(data)
sdsd_bb <- backbone.extract(sdsd, alpha = .01, narrative = TRUE)
```

- 1) computes probabilities for each cell in the bipartite matrix to be equal to 1
- 2) applies the **Poisson Binomial distribution** to the weighted projection.



# Stochastic Degree Sequence Model

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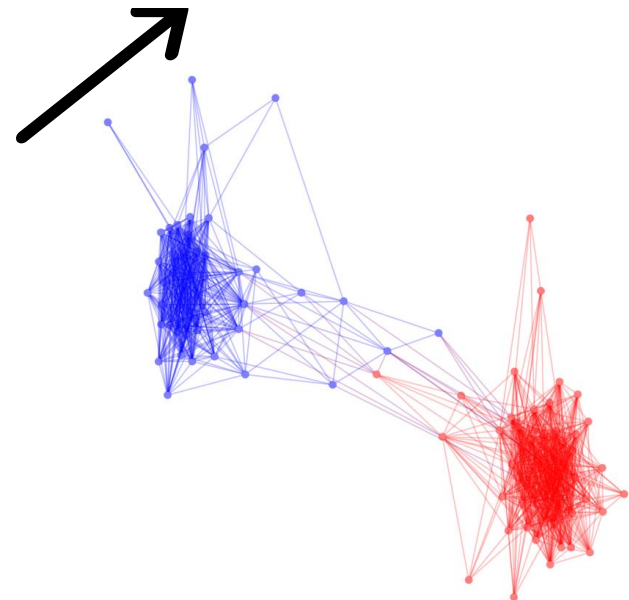
```
sdsbm <- sdsbm(data)
sdsbm_bb <- backbone.extract(sdsbm, alpha = .01, narrative = TRUE)
```

Suggested manuscript text and citations:

From a bipartite graph containing 100 agents and 3589 artifacts, we obtained the weighted bipartite projection, then extracted its signed backbone using the backbone package (Domagalski, Neal, & Sagan, 2021). Edges were retained in the backbone if their weights were statistically significant ( $\alpha = 0.01$ ) by comparison to a null Stochastic Degree Sequence Model (Neal, 2014).

Domagalski, R., Neal, Z. P., and Sagan, B. (2021). backbone: An R Package for Backbone Extraction of Weighted Graphs. PLoS ONE. <https://doi.org/10.1371/journal.pone.0244363>

Neal, Z. P. (2014). The backbone of bipartite projections: Inferring relationships from co-authorship, co-sponsorship, co-attendance and other co-behaviors. Social Networks, 39, 84-97. <https://doi.org/10.1016/j.socnet.2014.06.001>



# Fixed Degree Sequence Model



Senate co-sponsorship: compares the observed values to a distribution where each senator sponsors the exact same number of bills and each bill is sponsored by the exact same number of people.

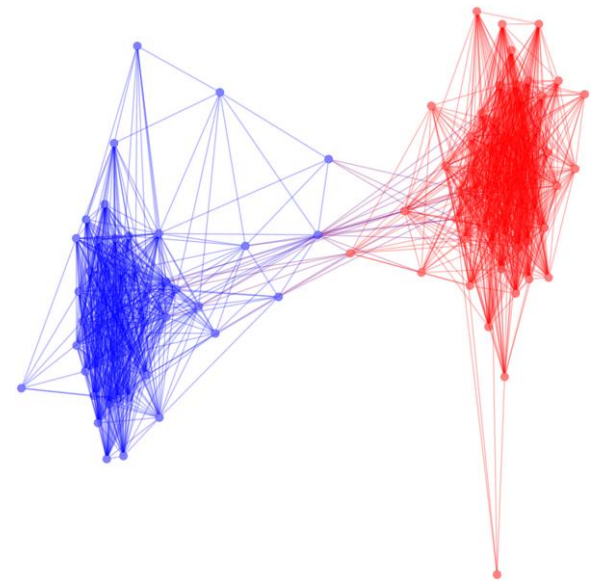
Samples this distribution by applying the **curveball** algorithm to the bipartite graph numerous times.

Choose the number of times it's applied with the “**trials**” parameter (more trials, more accurate)



```
fdsm <- fsm(data, trials = 1000)
```

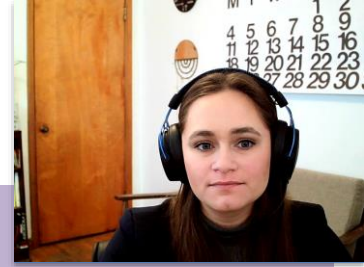
```
fdsm_bb <- backbone.extract(fdsm, signed = TRUE, alpha = 0.01)
```



Zweig KA, Kaufmann M. A systematic approach to the one-mode projection of bipartite graphs. *Social Network Analysis and Mining*. 2011;1(3):187–218. doi:10.1007/s13278-011-0021-0.

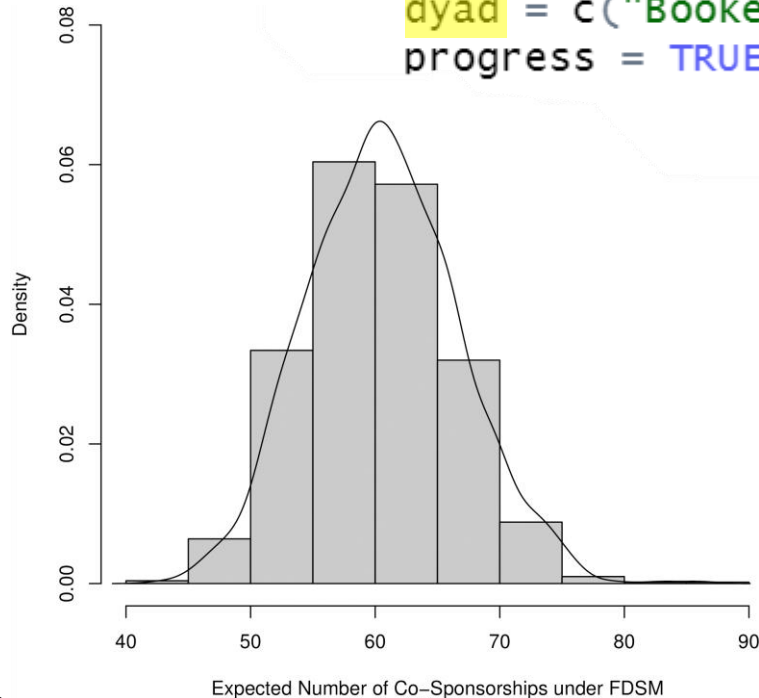


# Fixed Degree Sequence Model



Senate co-sponsorship: compares the observed values to a distribution where each senator sponsors the exact same number of bills and each bill is sponsored by the exact same number of people.

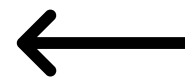
```
fdsm <- fsm(data, trials = 1000,  
            dyad = c("Booker, C. (NJ-D)", "Warren, E. (MA-D)"),  
            progress = TRUE)
```



Can also get the distribution for a chosen **dyad**.

```
fdsm$dyad_values
```

We can visualize this data in a histogram.



Their actual number of cosponsorships is 98.





# Coming soon!



## Early February 2021: backbone 1.3.0

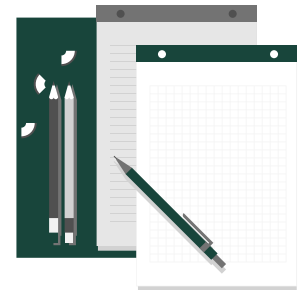
- Easier model selection / guidance
- Faster SDSM computations
- Bells & Whistles

## Summer 2021: backbone 2.0.0

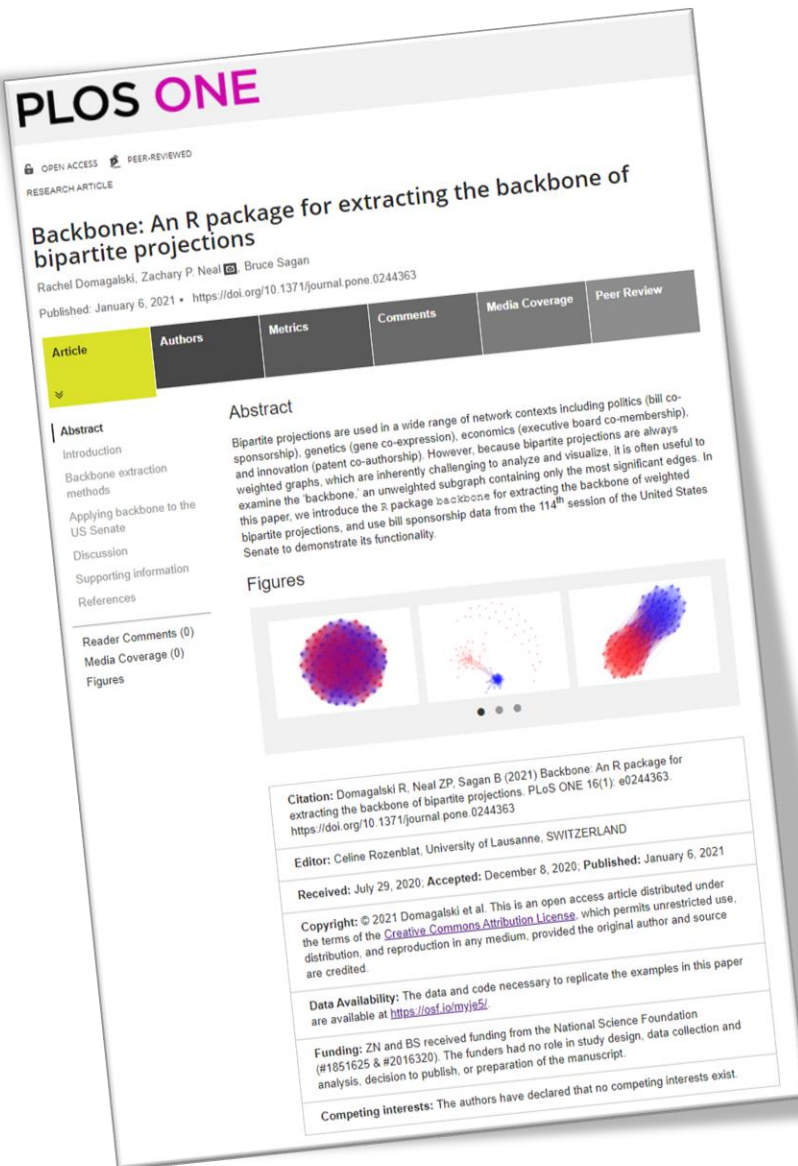
- Models for general weighted graphs (not necessarily bipartite projections)

## Manuscript to aid in model selection

- Accuracy of probabilities generated in SDSM
- Effects of degree distributions and significance levels
- Ability to recover group structure



# Backbone in PLoS ONE



Learn in detail about each null model and the mathematics behind them

Data and code available to replicate all the results and examples

<https://doi.org/10.1371/journal.pone.0244363>

[www.zacharyneal.com/backbone](http://www.zacharyneal.com/backbone)  
[github.com/domagal9/backbone](https://github.com/domagal9/backbone)

Tweet us any questions or comments!



@zpneal  
@rdomagalski



Backbone @ 2021 NASN

## THANK YOU!