Motifs and bipartite networks

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Outline

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Bipartite network

Two types of actors.

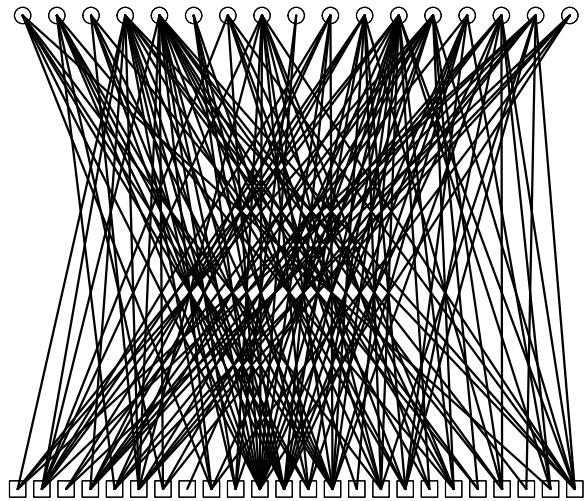
- Mutualistic: plant-pollinator
- Antagonistic: host-parasite

Topological analysis:
understanding the network organisation

Local: node or edge properties (degree, betweenness)

Global: density, connected components, nestedness

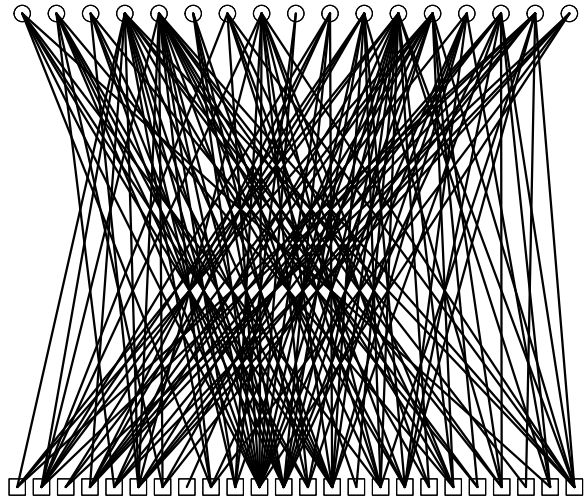
Zackenberg network: [SROB16]



Bipartite network: notations

Species.

- $i = 1, \dots m$ pollinators = rows = bottom nodes
- $j = 1, \dots n$ plants = columns = top nodes



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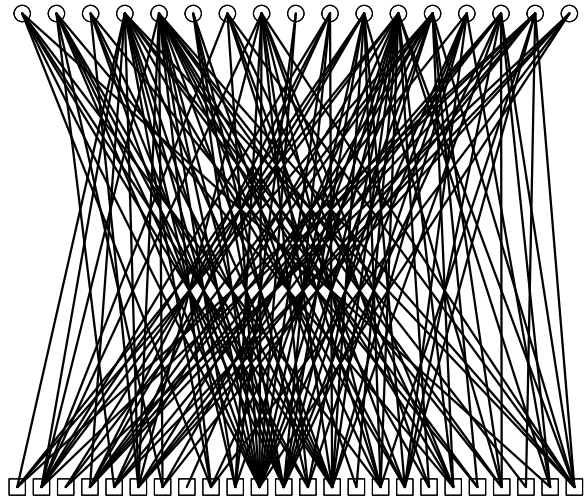

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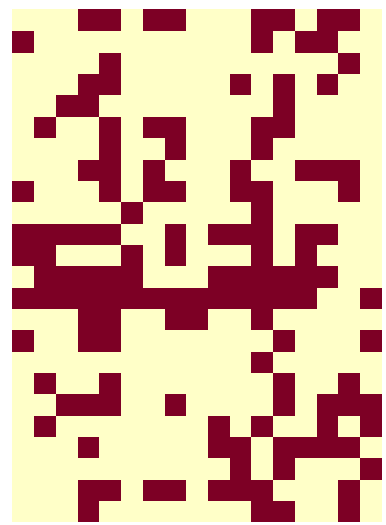
Interactions.

- $A_{ij} = 1$ if pollinator i interacts with plant j, 0 otherwise

$$A_{ij} = 1 \quad \Leftrightarrow \quad i \sim j$$

- adjacency matrix : $m \times n$

$$A = [A_{ij}]_{1 \leq i \leq m, 1 \leq j \leq n}$$



Bipartite motifs

'Meso-scale' analysis. [SCB+19]

- Motifs ='building-blocks'
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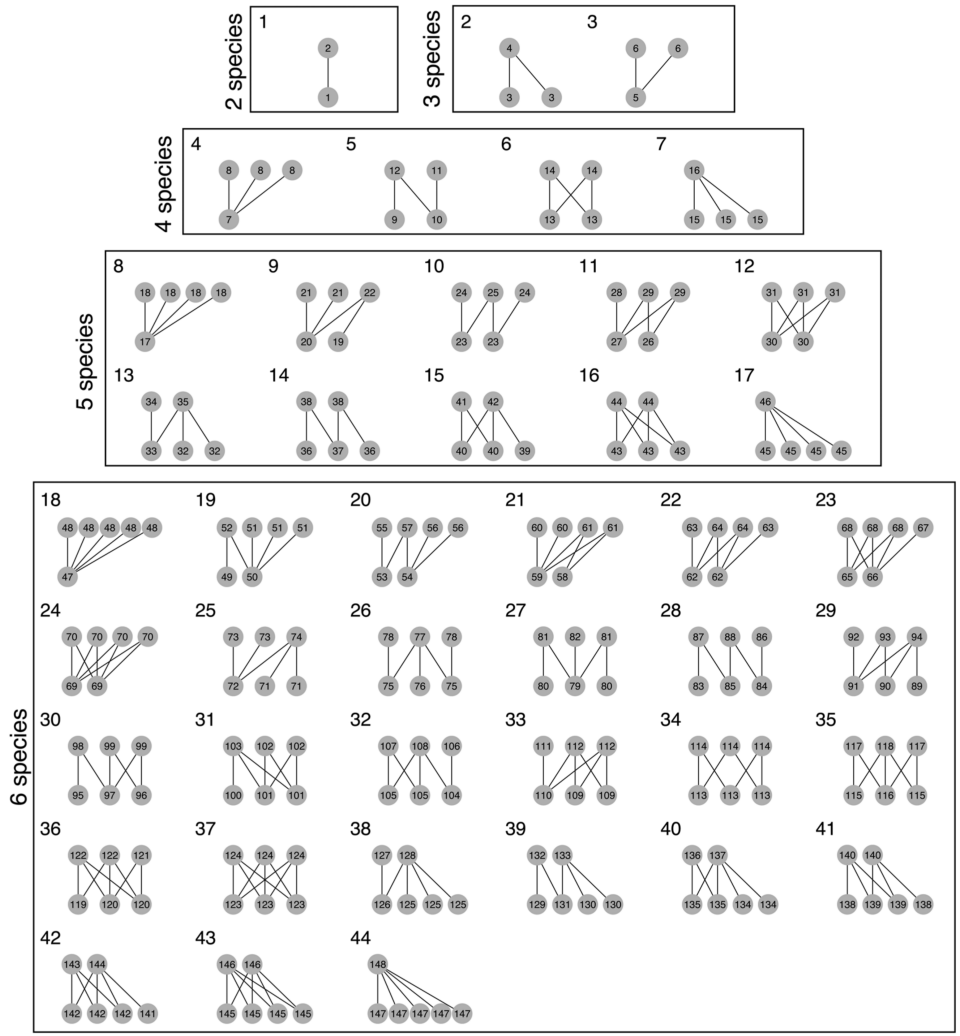

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- Enables network comparison
- Even when the nodes are different

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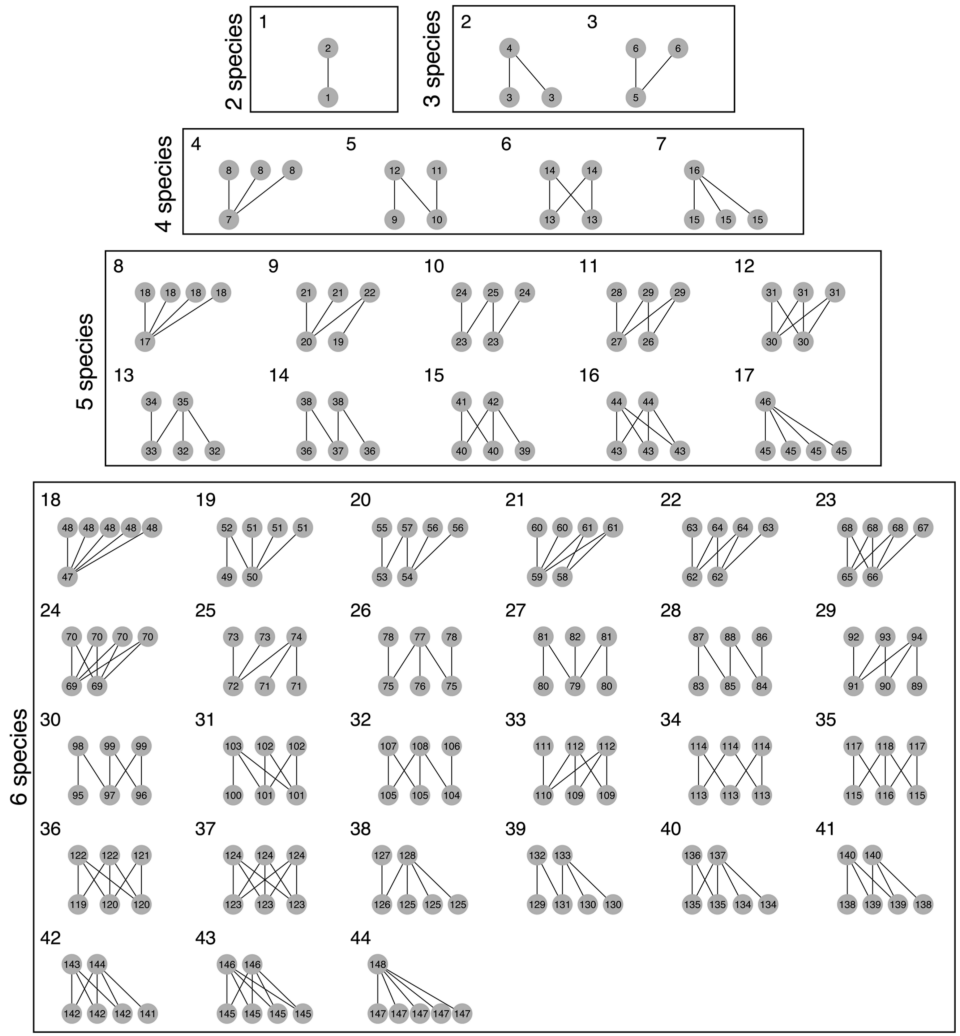

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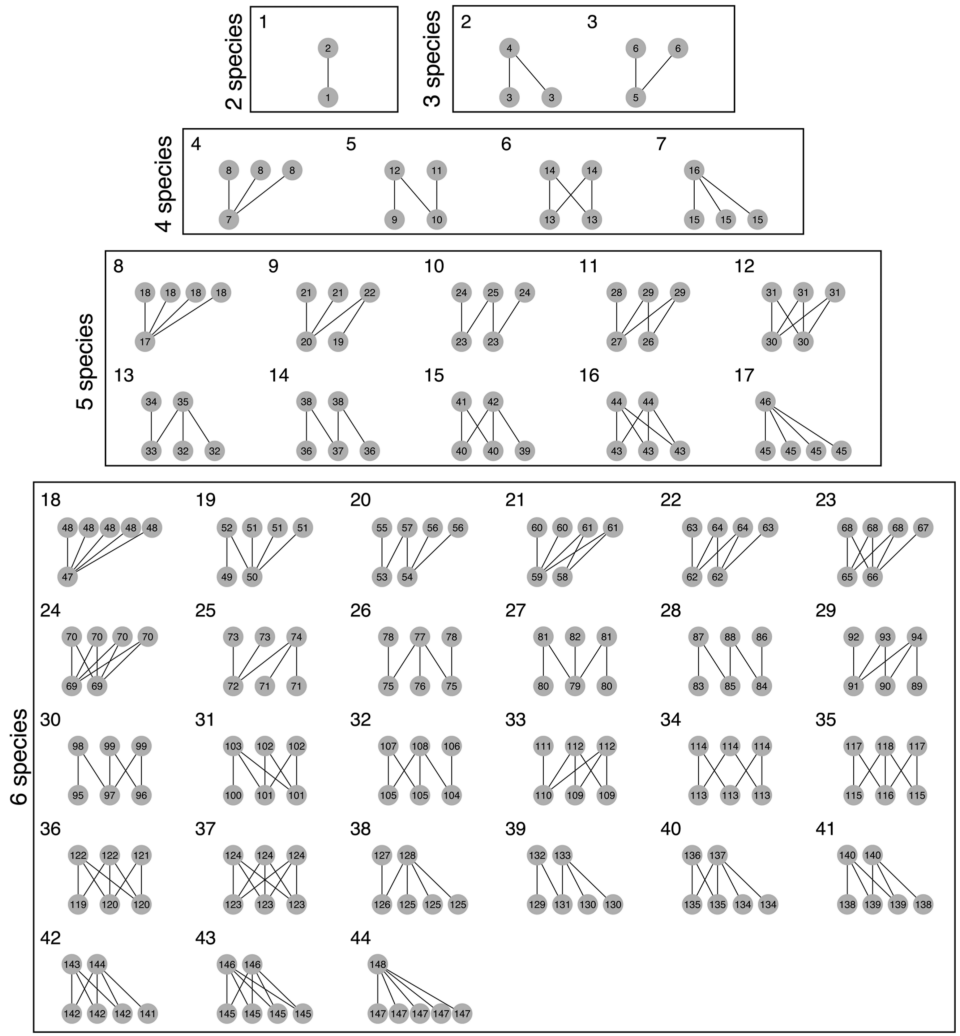
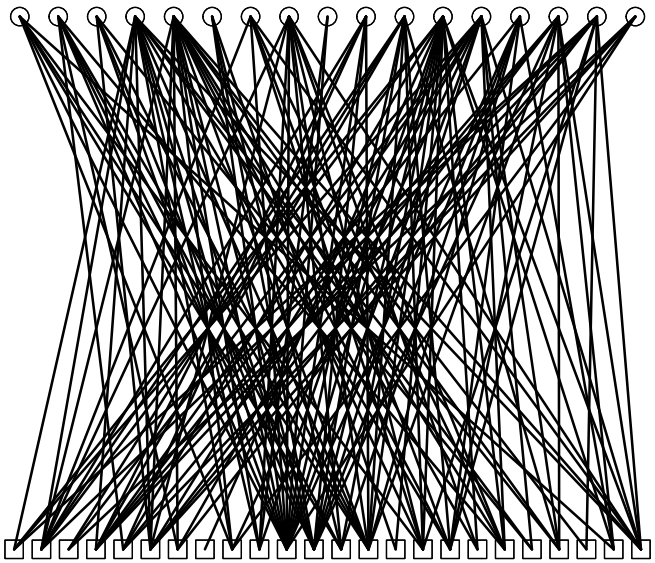
Existing tool. `bmotif` package [SSS+19]: counts motif occurrences



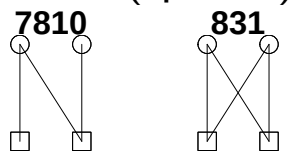
Example

Plant-pollinator network [SROB16]

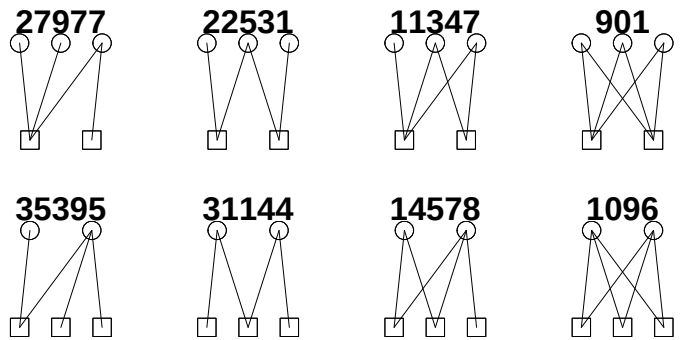


Motif counts.

4 nodes (species)



5 nodes



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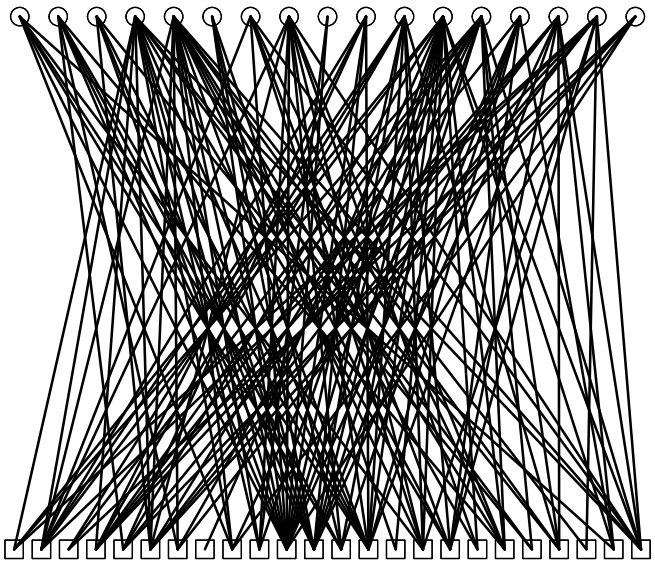

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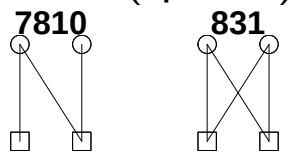

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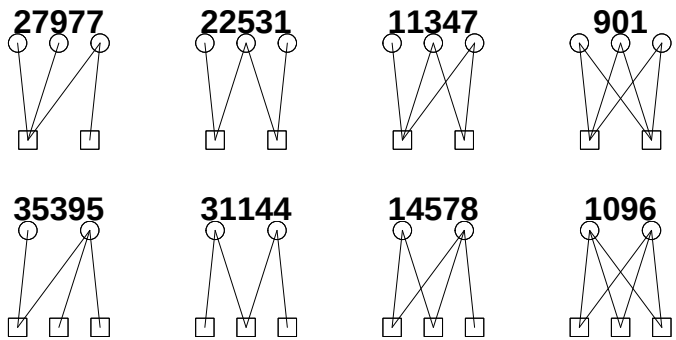
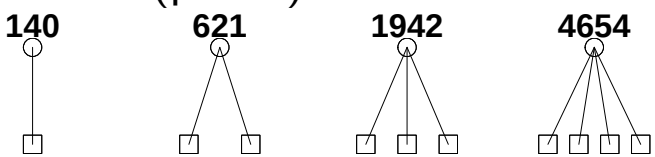
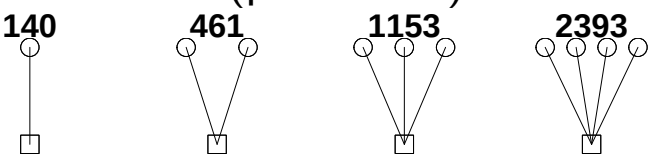


top 'stars' (plants)



bottom 'stars' (pollinators)



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Need for a null model

Motif counts obviously depend on

- the size of the network: $n \times m$
- the density of the network
- the imbalance between bottom-node degrees (specialist vs generalist pollinators)
- the imbalance between top-node degrees (specialist vs generalist plants)

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Bipartite expected degree distribution (BEDD) model: (in words)

- Consider m pollinators ($i = 1, \dots m$):
  each plant i has a specific propensity to interact (degree of generalism)
- Consider n plants ($j = 1, \dots n$):
  each plant j has a specific propensity to interact (idem)
- The probability for pollinator i and plant j to interact is proportional to the product of their respective propensities.

BEDD model

Bipartite expected degree distribution (BEDD) model: (formaly)

- $\rho =$ network density
- $g =$ top node degree imbalance ($\int g = 1$)
- $h =$ bottom node degree imbalance ($\int h = 1$)

$$\{U_i\}_{i=1,\dots m} \text{ iid } \sim \mathcal{U}[0,1] \qquad \{V_j\}_{j=1,\dots n} \text{ iid } \sim \mathcal{U}[0,1]$$

$$\mathbb{P}\{i \sim j \mid U_i, V_j\} = \rho\, g(U_i)\, h(V_j)$$

(Bipartite version of the EDD model [CL02])

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Model parameters:

$$\theta = (\rho, g, h).$$

BEDD model

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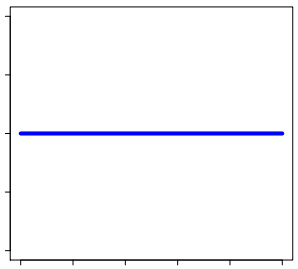
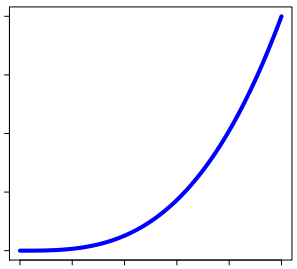
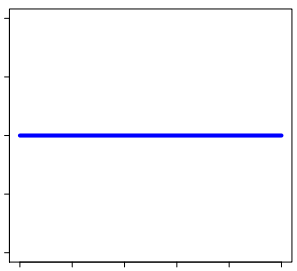
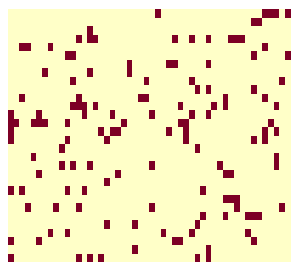
$$\mathbb{E}(D_i \mid U_i) = n\, \rho\, g(U_i)$$

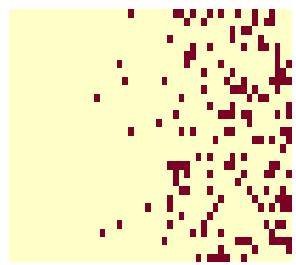
$$\mathbb{E}(D_j \mid V_j) = m\, \rho\, g(V_i)$$

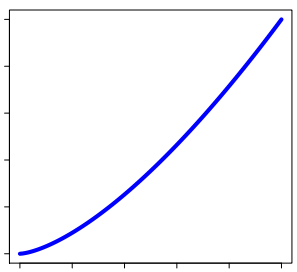
$h_0(v) =$ 

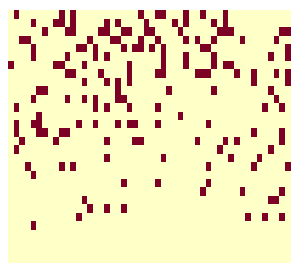
$h(v) =$ 

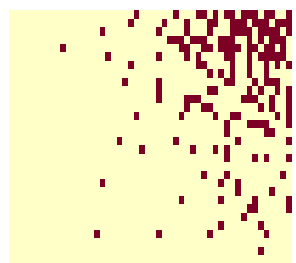
$g_0(u) =$ 





$g(u) =$ 





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Assumptions.

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Sufficient statistics to fit BEDD:

- Pollinator degrees + plant degrees
- or, equivalently, star (single edge, top, bottom) frequencies

Outline

Counting motifs

Number of 'positions'.

- Choose p nodes among m
- Choose q nodes among n
- Try all *automorphisms*

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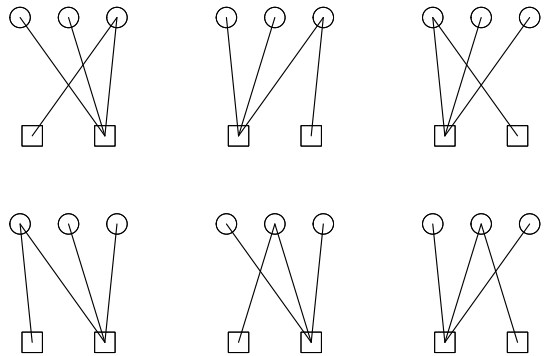

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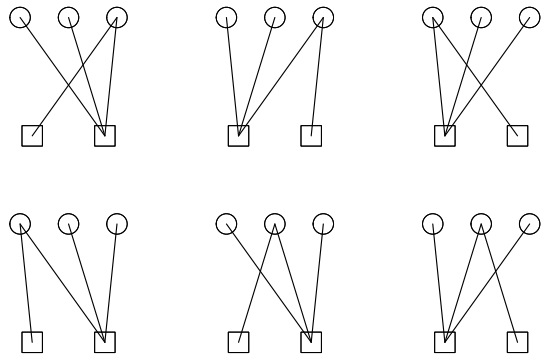

Motif count. Try all positions $\alpha = 1, \dots c_s$, define

$$Y_{s\alpha} = 1 \text{ if match}, \qquad 0 \text{ otherwise},$$

then count the number of matches:

$$N_s = \sum_{\alpha} Y_{s\alpha}$$

$\rightarrow$ Motif frequency: $F_s := N_s / c_s$

Motif probability

Occurrence probability $\overline{\phi}_s = \mathbb{P}\{Y_{s\alpha} = 1\}$. Under the B-EDD model [OLR22]:

$$\left(\quad \right) = \frac{\qquad\qquad\qquad\qquad\qquad\qquad\qquad\qquad\qquad\qquad\qquad\qquad}{\qquad\qquad\qquad\qquad\qquad\qquad\qquad\qquad\qquad\qquad\qquad\qquad}$$

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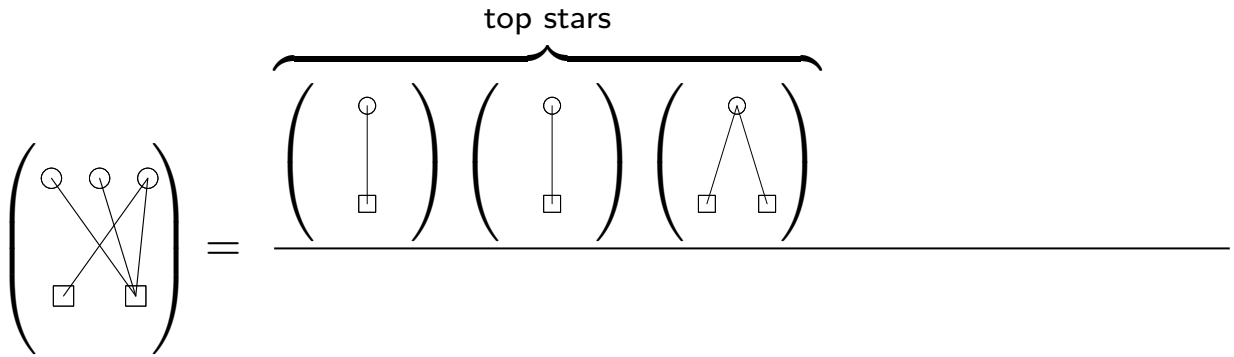

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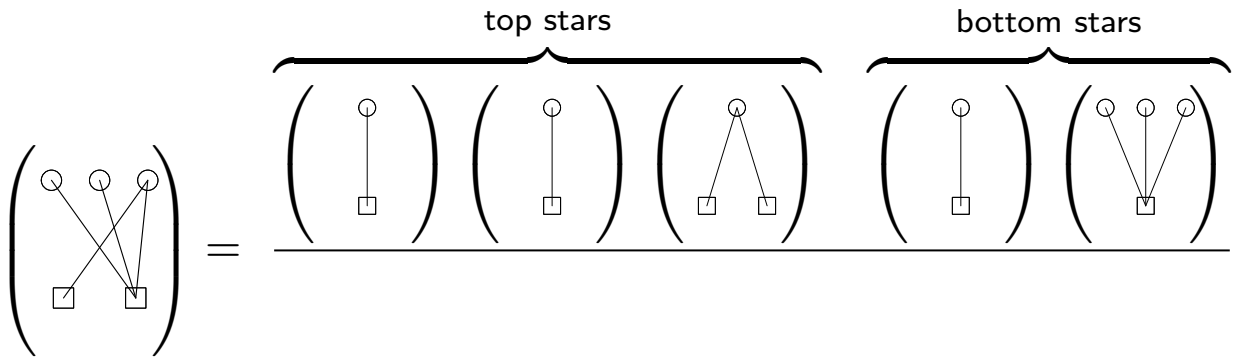

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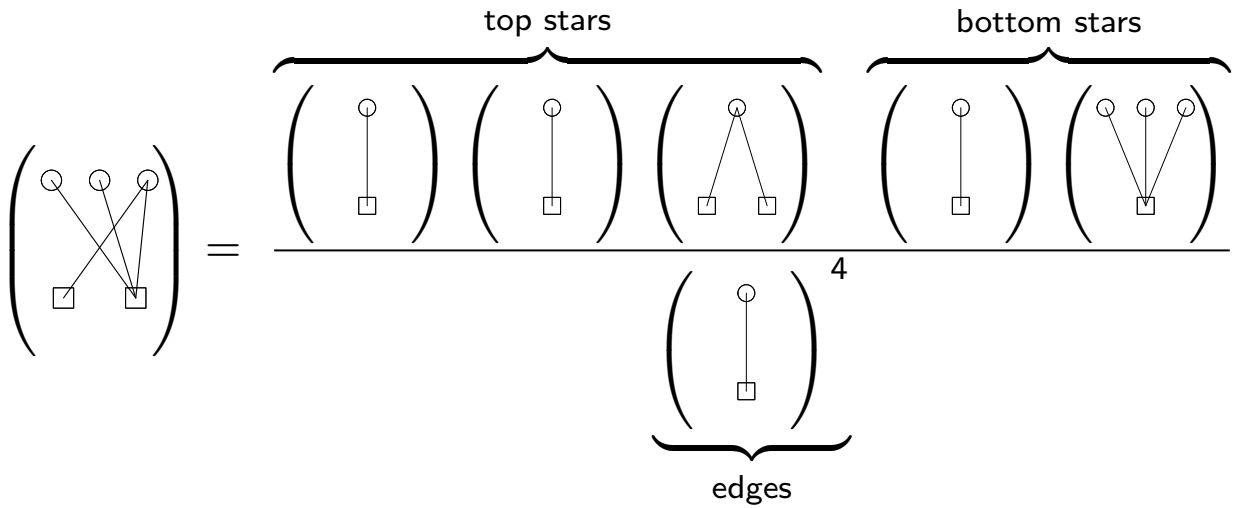

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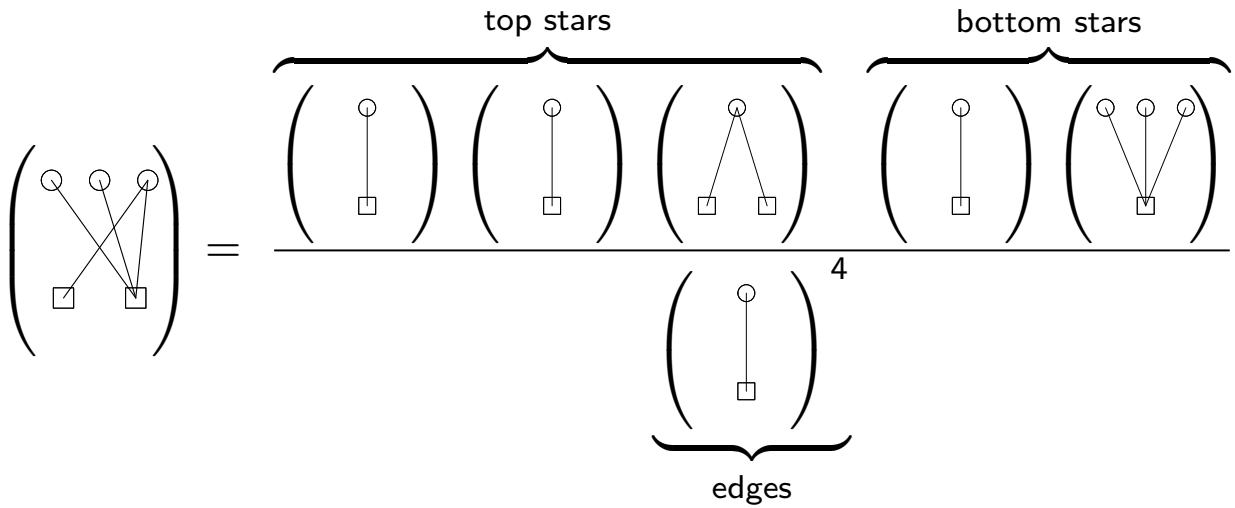

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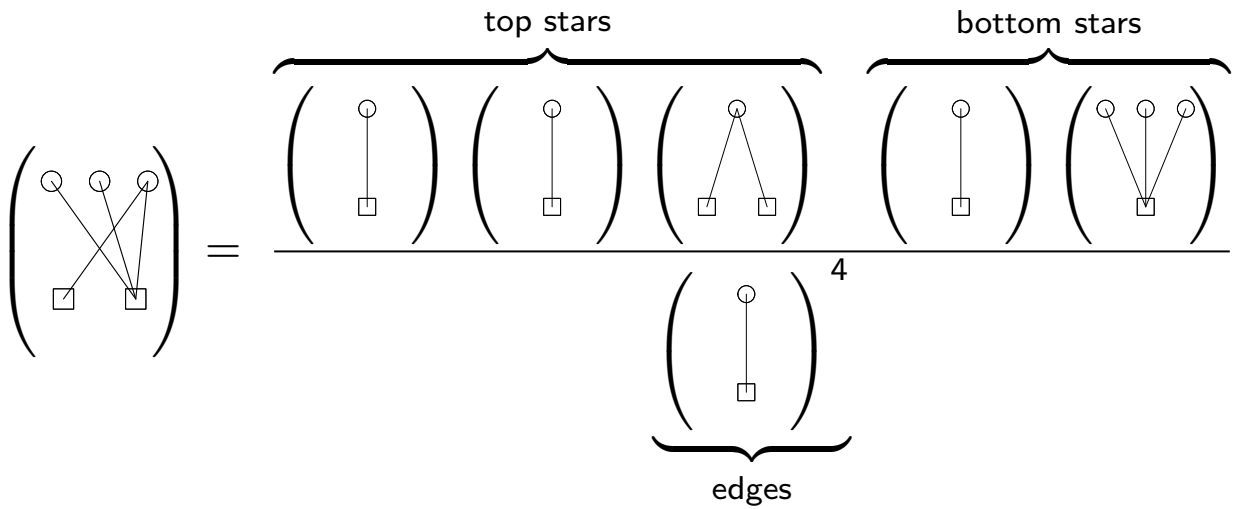

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Estimated probability $\overline{F}_s$.

$$\overline{\phi}_s := \frac{\phi_2\phi_4}{\phi_1} \quad \rightarrow \quad \overline{F}_s := \frac{F_2F_4}{F_1}$$

where F_1, F_2, F_4 = observed frequencies of edges, top stars and bottom stars.

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- Covariance: Same game to compute $\mathbb{C}\text{ov}(N_s, N_{s'})$

Distribution of the count

Asymptotic normality for non-star motifs. Under BEDD (and sparsity conditions):

$$(F_s - \overline{F}_s) \Big/ \sqrt{\widehat{\mathbb{V}}(F_s)} \quad \overset{m,n\to\infty}{\longrightarrow} \quad \mathcal{N}(0,1)$$

Proof [OLR22]:

$$\text{decompose} \qquad F_s - \overline{F}_s = \underbrace{(F_s - \phi_s)}_{\text{random fluctuations}} + \underbrace{(\phi_s - \overline{\phi}_s)}_{\text{null under BEDD}} + \underbrace{(\overline{\phi}_s - \overline{F}_s)}_{\text{estimation error} \to 0},$$

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Consequence. We know

- the expected behavior (mean, variance, distribution) of any motif count
- under the BEDD model (= 'null model').

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Example.

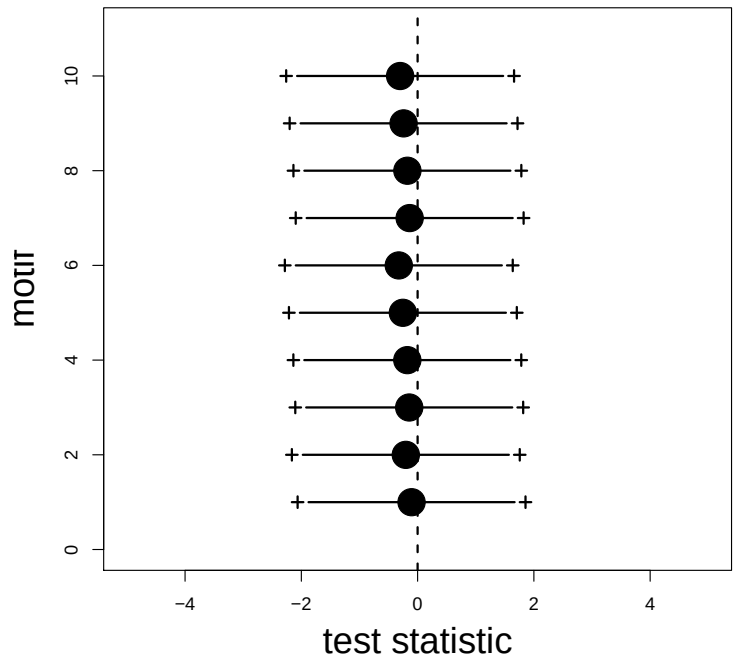
1. Data = observed plant-pollinator network
2. Statistic T = motif count N_s
3. Model = BEDD

Goodness-of-fit (GOF) of the BEDD model

Zackenberg network.

Raw statistic:

$$T_s = \frac{N_s - \widehat{\mathbb{E}} N_s}{\sqrt{\widehat{\mathbb{V}} N_s}}$$



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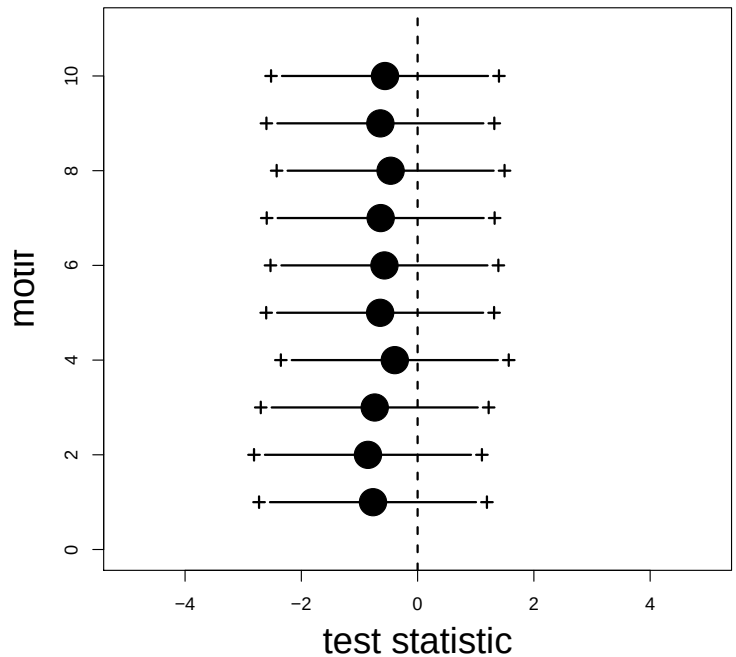
Corrected stat.: accounts for the estimation error in $\widehat{\mathbb{E}} N$

$$T'_s = \frac{N_s - (\widehat{\mathbb{E}} N_s - \widehat{\mathbb{B}}(\widehat{\mathbb{E}} N_s))}{\sqrt{\widehat{\mathbb{V}}(N_s - \widehat{\mathbb{E}} N_s)}}$$



Testing degree imbalance

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Statistical test.

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$$H_0 = \{h = 1\}$$
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Example. (only one significant difference)

plant-pollinator

s	5	6	10	15	16
W_s	-6.45 10^{-2}	9.96 10^{-1}	-6.63 10^{-2}	7.52 10^{-1}	2.43

seed dispersal

s	5	6	10	15	16
W_s	-2.14 10^{-1}	-2.14 10^{-1}	-2.93 10^{-1}	-2.95 10^{-1}	-3.56 10^{-1}

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with $\mathcal{N}(0,1)$

Example. (no significant difference)

s	5	6	10	15	16
F_s^A	$9.21\ 10^{-5}$	$1.00\ 10^{-5}$	$8.12\ 10^{-6}$	$3.32\ 10^{-7}$	$4.47\ 10^{-8}$
$\widehat{\mathbb{E}}_0 F_s^A$	$1.96\ 10^{-4}$	$3.75\ 10^{-5}$	$1.74\ 10^{-5}$	$4.25\ 10^{-6}$	$1.33\ 10^{-6}$
F_s^B	$5.13\ 10^{-4}$	$1.15\ 10^{-4}$	$5.07\ 10^{-5}$	$1.79\ 10^{-5}$	$5.96\ 10^{-6}$
$\widehat{\mathbb{E}}_0 F_s^B$	$2.66\ 10^{-4}$	$2.92\ 10^{-5}$	$2.85\ 10^{-5}$	$1.50\ 10^{-6}$	$1.69\ 10^{-7}$
W_s	-1.56	-1.56	-0.97	-1.28	-0.96

Outline

Network embedding: Multivariate analysis

Analysing multiple networks. Principle

- 'Embed' each network into a convenient space (e.g. $\mathbb{R}^d$)
- Use standard multivariate analysis (clustering, PCA, MDS, ...)

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$$(\text{Network})_k \quad \to \quad (N_1^k, \ldots, N_S^k) \in \mathbb{R}^S$$

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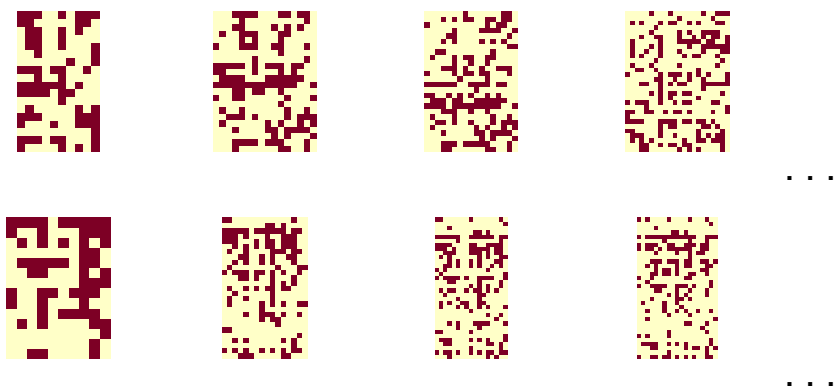
Zackenberg dataset. $K = 46$ networks

- 2 years
- 1 network observed every few days

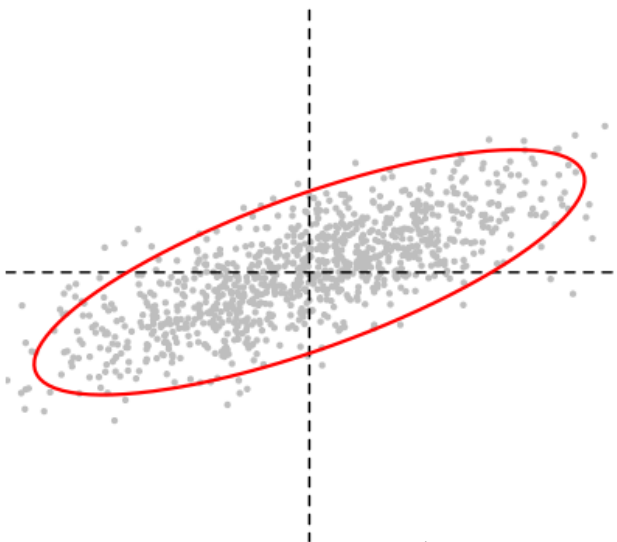


Choleski transform

Aim: 'Remove' correlation and variance heterogeneity

Covariance matrix of (X_1, X_2):

$$\Sigma_{X_1,X_2} = \begin{bmatrix} \sigma_1^2 & \sigma_{12} \\ \sigma_{12} & \sigma_2^2 \end{bmatrix}$$



Diagonalization: $\Sigma = P\Lambda P^{-1}$

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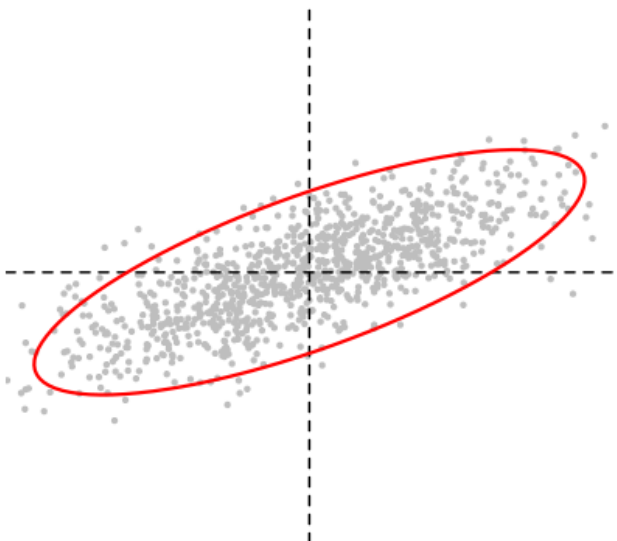
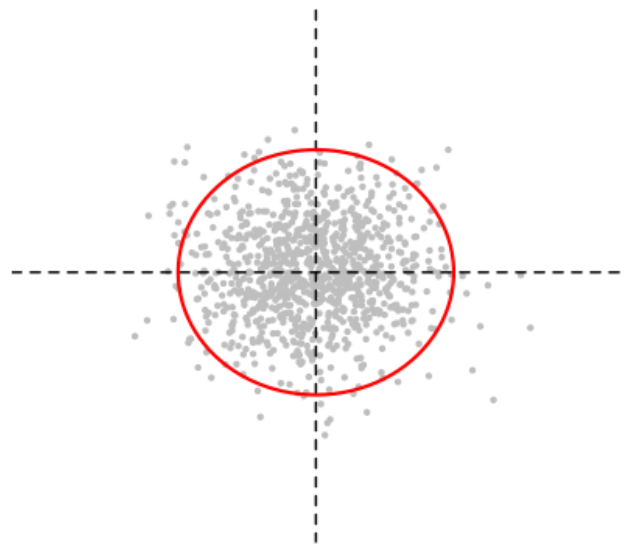
$$\begin{bmatrix} X_1' \\ X_2' \end{bmatrix} = \Sigma^{-1/2} \begin{bmatrix} X_1 \\ X_2 \end{bmatrix}$$





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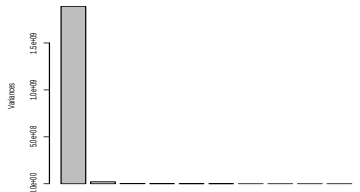
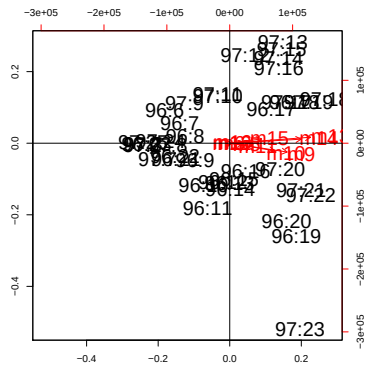
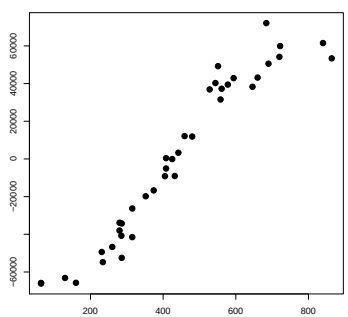
$$\Sigma_{X_1',X_2'} = \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix}$$

Network embedding: Zackenberg's data [SROB16]

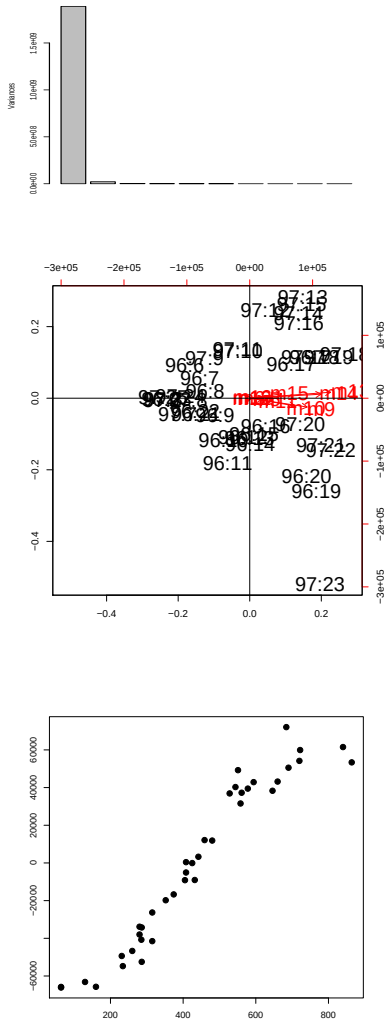
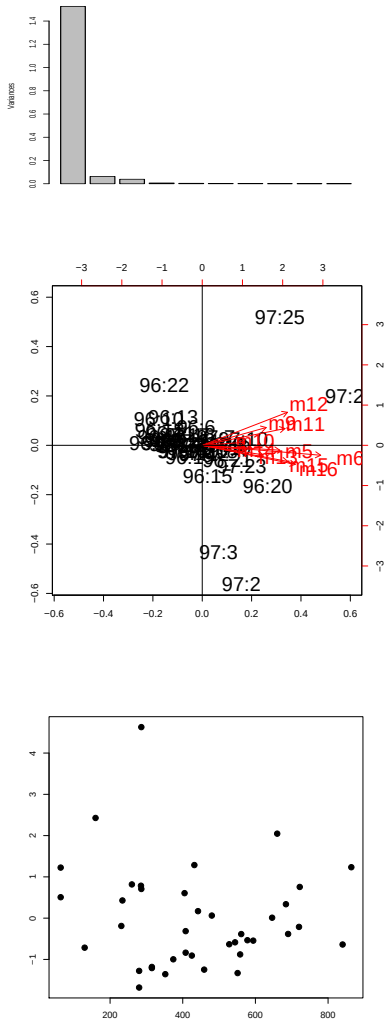
Raw counts



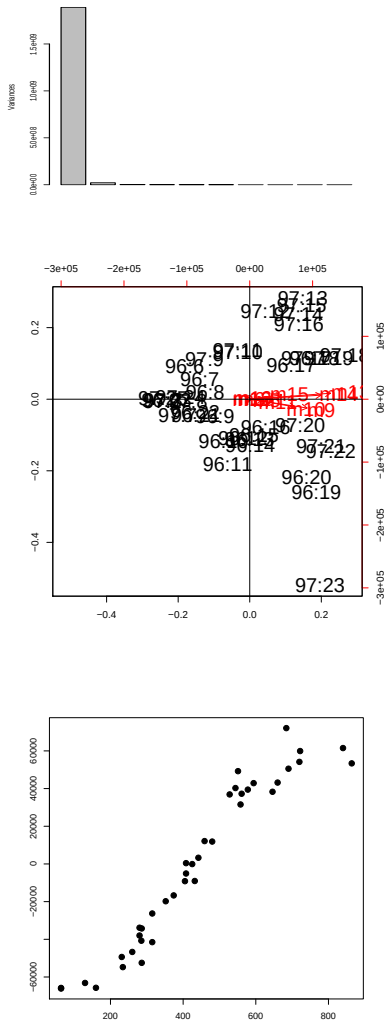
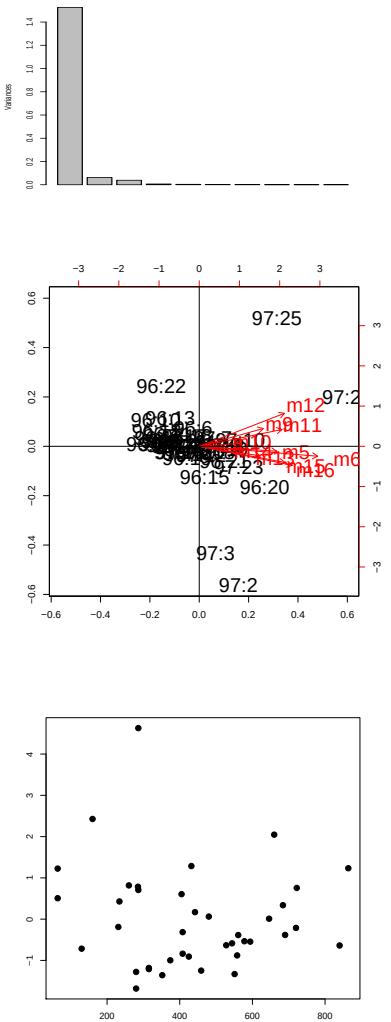
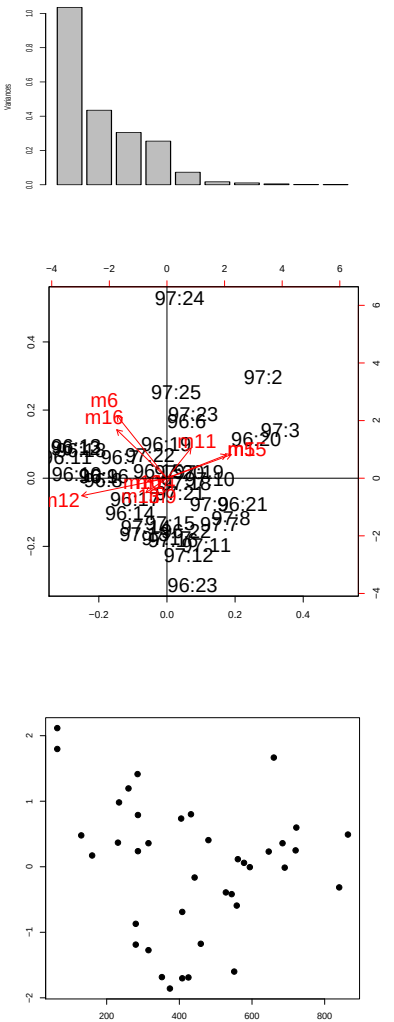




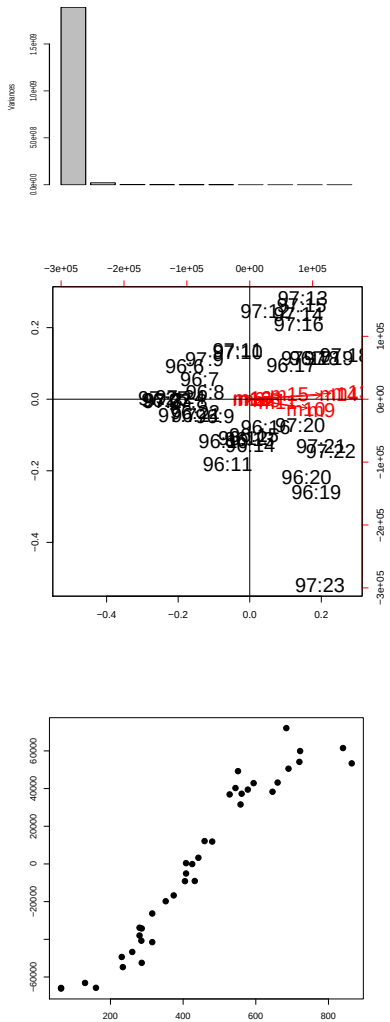
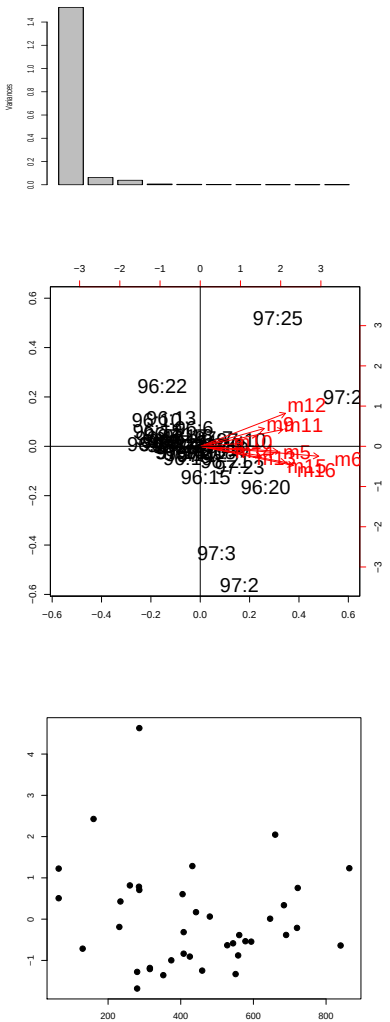
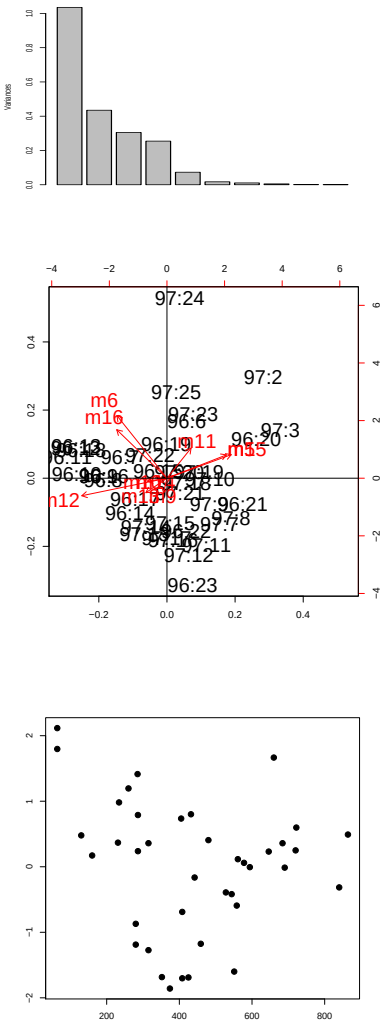
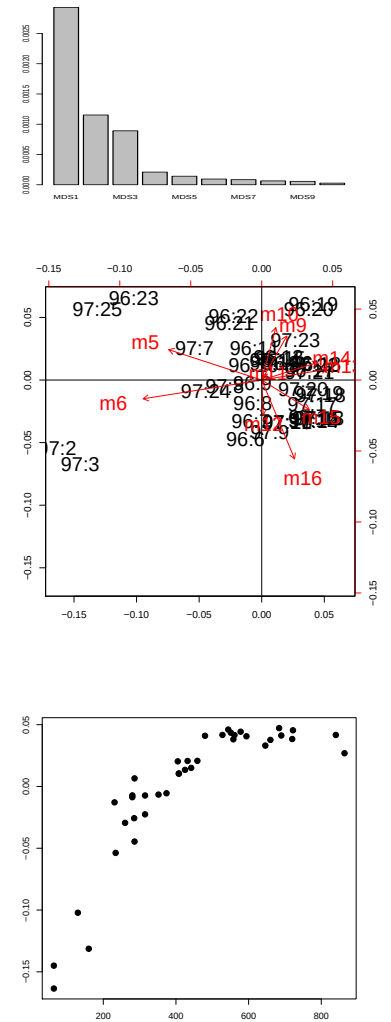
Network embedding: Zackenberg's data [SROB16]

Raw counts	Corrected stat.		
			

Network embedding: Zackenberg's data [SROB16]

Raw counts	Corrected stat.	Choleski
		

Network embedding: Zackenberg's data [SROB16]

Raw counts	Corrected stat.	Choleski	Bray-Curtis MDS
			

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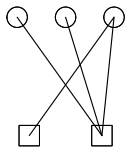
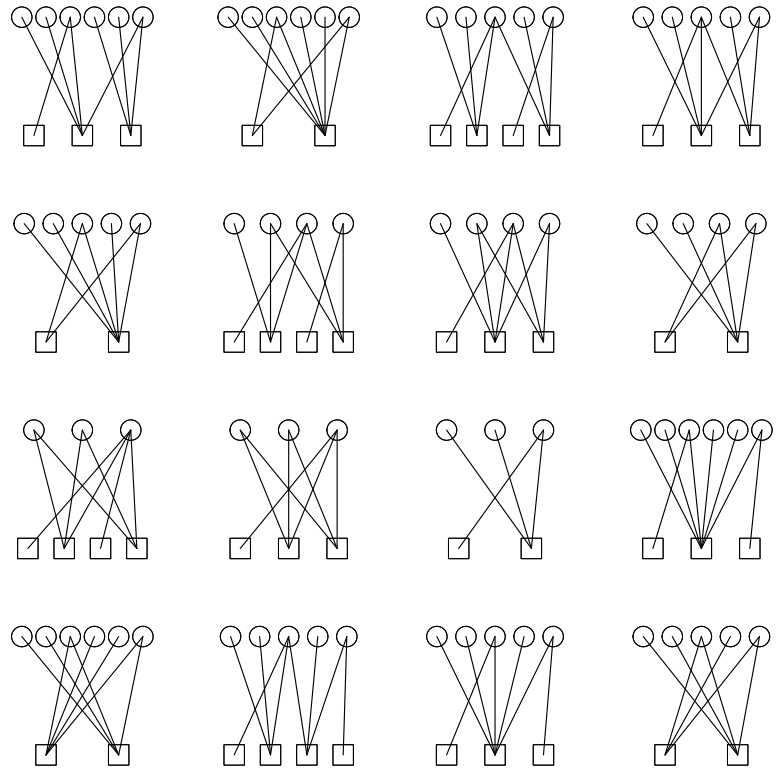
Super-motifs

Motif:



Some super-motifs:



. . . 396 super-motifs

Variance:

$$N_s^2 = \left(\sum_\alpha Y_{s\alpha}\right)^2$$

$$= \sum_{\alpha,\beta:\alpha\cap\beta=\emptyset} Y_{s\alpha} Y_{s\beta}$$

$$+ \sum_{\alpha,\beta:\alpha\cap\beta\neq\emptyset} \underbrace{Y_{s\alpha} Y_{s\beta}}_{\substack{\text{occurrence of} \\ \text{a super-motif}}}$$

Covariance: same game, for $Y_{s\alpha} Y_{s'\beta}$ with $s \neq s'$

Asymptotic distribution of the count

Estimated probability.

$$\overline{\phi}_s := \phi_2 \phi_4 \, / \phi_1 \qquad \rightarrow \qquad \overline{F}_s := F_2 F_4 \, / F_1$$

where F_1, F_2, F_4 = observed frequencies of top stars, bottom stars and edges.

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Asymptotic normality for non-star motifs. Under BEDD (and sparsity conditions):

$$(F_s - \overline{F}_s) \Big/ \sqrt{\widehat{\mathbb{V}}(F_s)} \quad \overset{m,n \to \infty}{\longrightarrow} \quad \mathcal{N}(0,1)$$

Proof:

- decompose

$$F_s - \overline{F}_s = \underbrace{(F_s - \phi_s)}_{\text{random fluctuations}} + \underbrace{(\phi_s - \overline{\phi}_s)}_{\text{null under BEDD}} + \underbrace{(\overline{\phi}_s - \overline{F}_s)}_{\text{estimation error} \to 0},$$

- construct a counting martingale [GL17] for $F_s - \phi_s$

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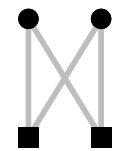
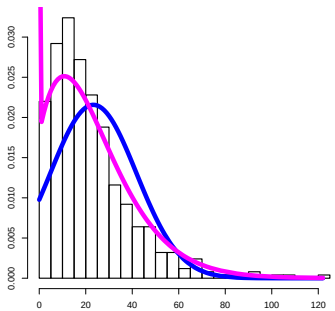
Test statistic. Under BEDD:

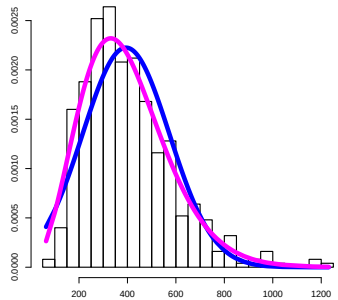
$$N_s \approx \mathcal{N}\left(\widehat{\mathbb{E}}(N_s), \widehat{\mathbb{V}}(N_s)\right) \qquad \Leftrightarrow \qquad \left(N_s - \widehat{\mathbb{E}}(N_s)\right) \Big/ \sqrt{\widehat{\mathbb{V}}(N_s)} \approx \mathcal{N}(0,1)$$

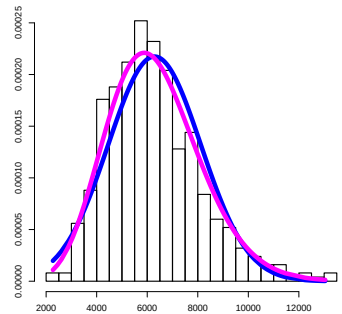
In practice: Asymptotic normality

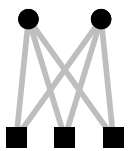
$(n = 2m/3)$ $m = 50$ $m = 100$ $m = 200$

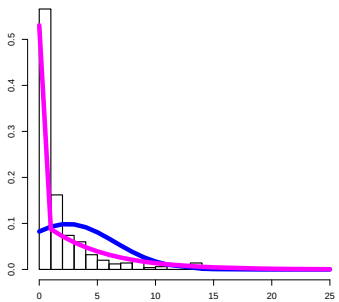


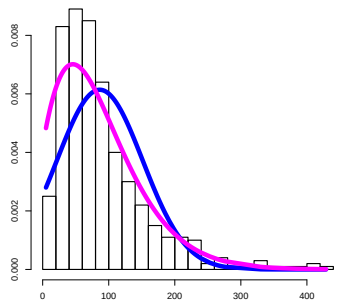


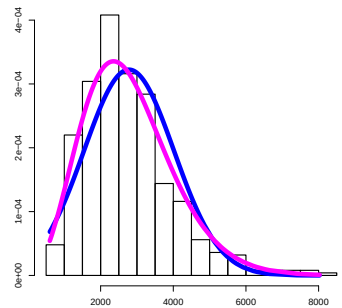












Normal distribution, Poisson-geometric distribution with same mean and variance [Sta01,PDK+08]

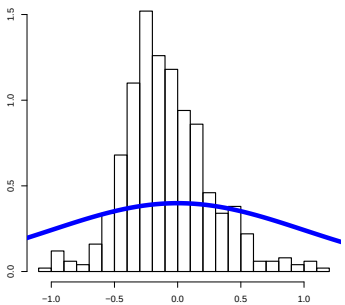
In practice: Test statistic

Need to account for the estimation error of $\widehat{\mathbb{E}}N$

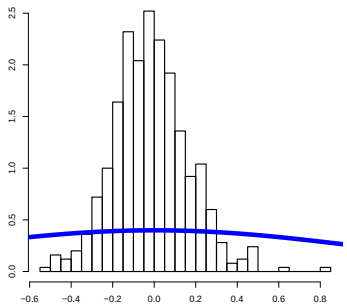
Regular stat.:

$$\frac{N - \widehat{\mathbb{E}}N}{\sqrt{\widehat{\mathbb{V}}N}}$$

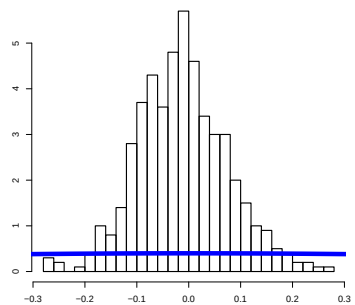
$m = 50$



$m = 100$



$m = 200$



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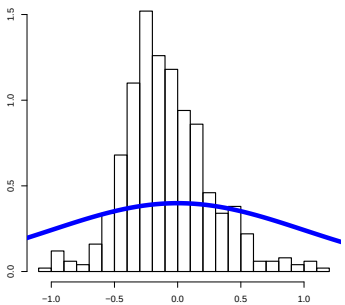
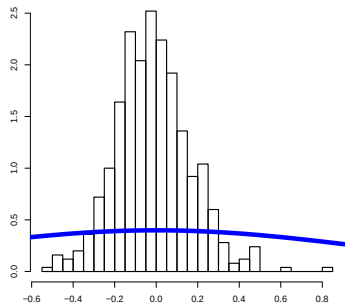
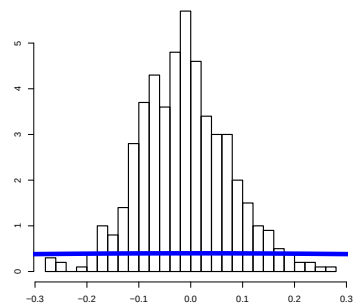
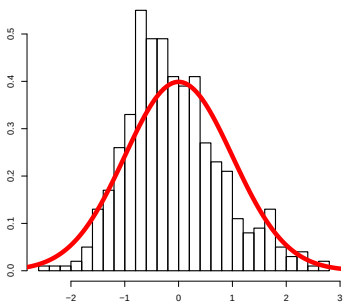
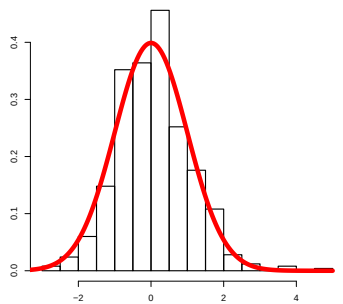
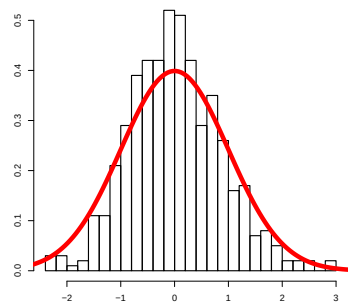
Need to account for the estimation error of $\widehat{\mathbb{E}}N$

	$m = 50$	$m = 100$	$m = 200$
Regular stat.: $\dfrac{N - \widehat{\mathbb{E}}N}{\sqrt{\widehat{\mathbb{V}}N}}$			
Correction: $\dfrac{N - (\widehat{\mathbb{E}}N - \widehat{\mathbb{B}}(\widehat{\mathbb{E}}N))}{\sqrt{\widehat{\mathbb{V}}(N - \widehat{\mathbb{E}}N)}}$			

- Need to evaluate $\mathbb{V}(N - \widehat{\mathbb{E}}(N))$ and $\mathbb{B}(\widehat{\mathbb{E}}N)$: resort to Taylor expansion ($\Delta-$method)