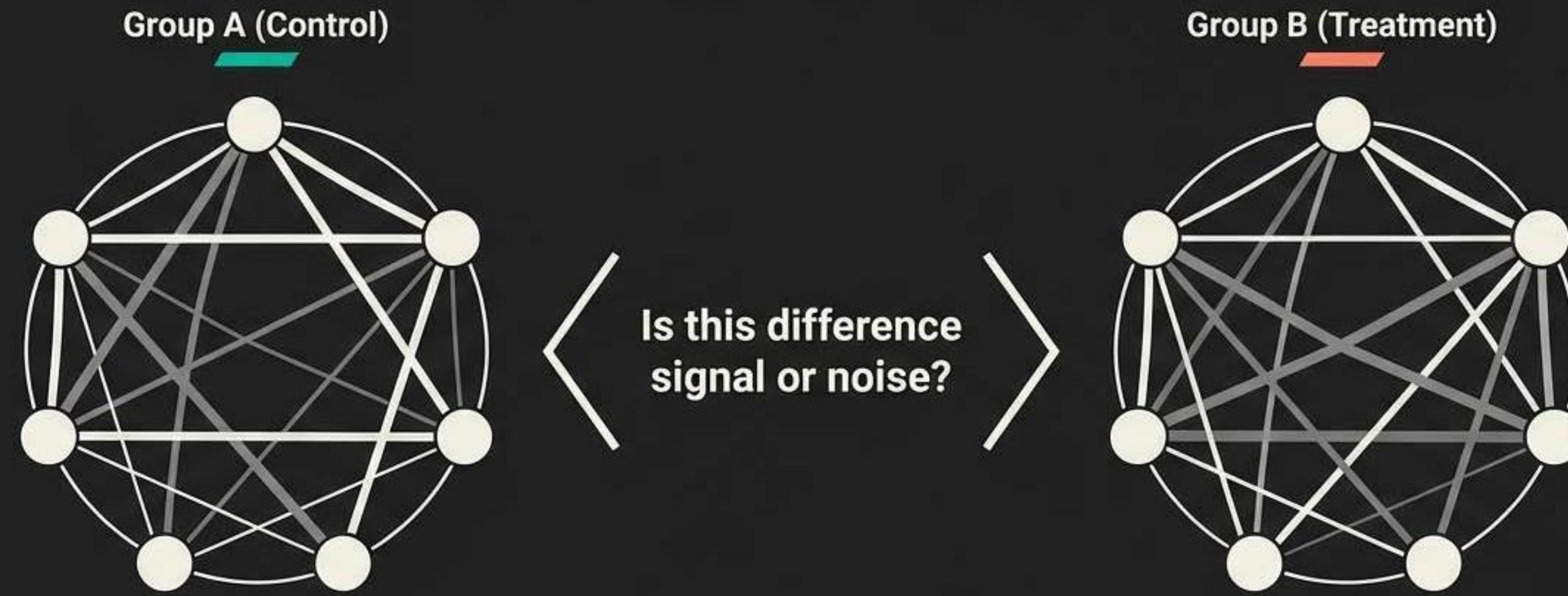


# Testing for Group Differences in Multilevel Vector Autoregressive Models

From 'Eyeballing' to Inference using  mIVAR

# The Problem with 'Eyeballing' Dynamics



Psychological research often asks: Is the dynamic network of Group A different from Group B?

Current State: Visual comparison is subjective.

The Gap: Without formal inference, we cannot distinguish population heterogeneity from sampling variation.

# Defining the Mathematical Object

$$y_{t,s} = \mu_s + B_s(y_{t-1,s} - \mu_s) + \zeta_{t,s}$$

State of subject s at time t

Subject-specific mean

The Temporal Network  
(Lagged coefficients)

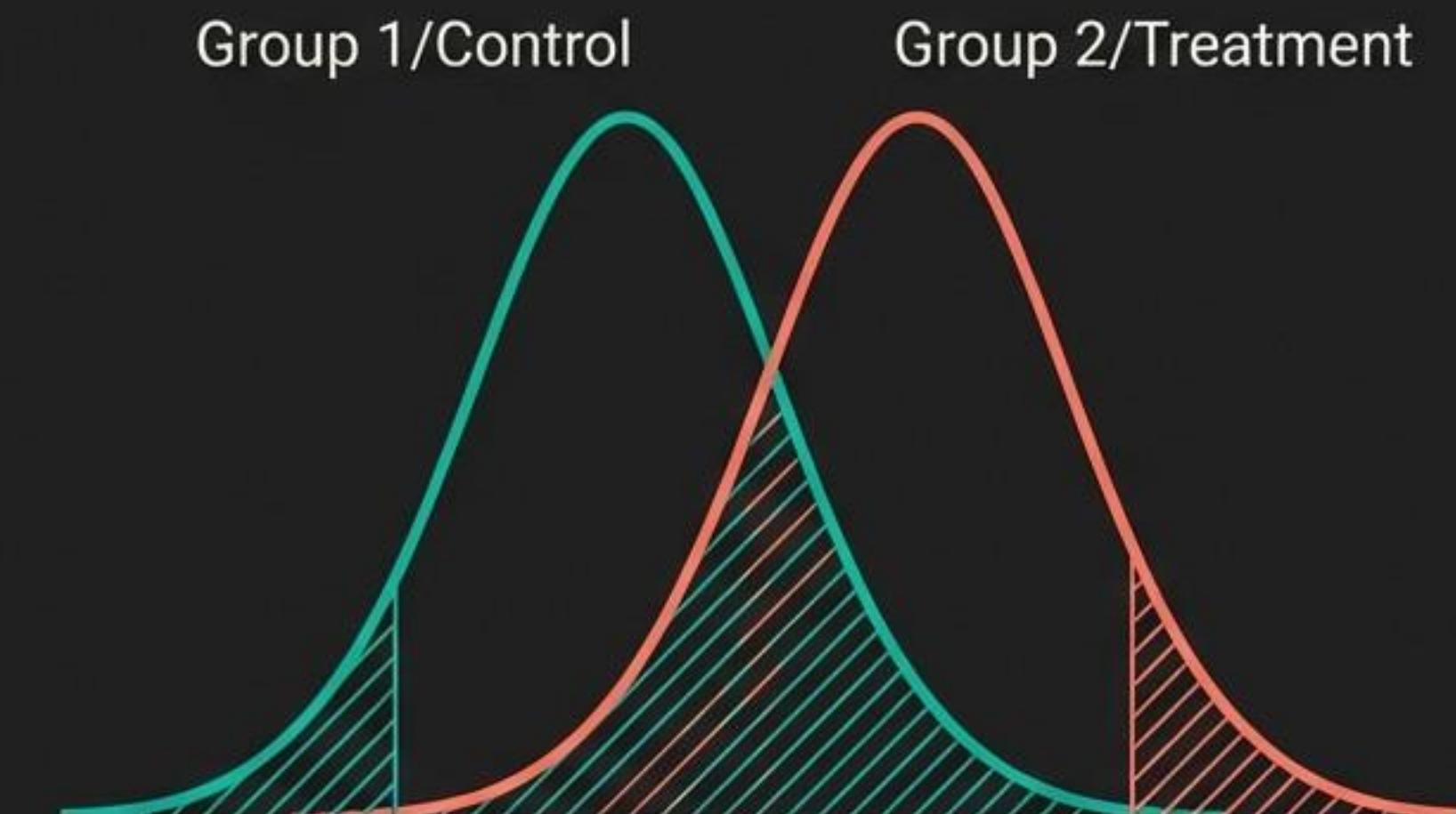
Primary target of testing

Innovation (residuals)

We test for differences in the distributions of these parameters across groups.

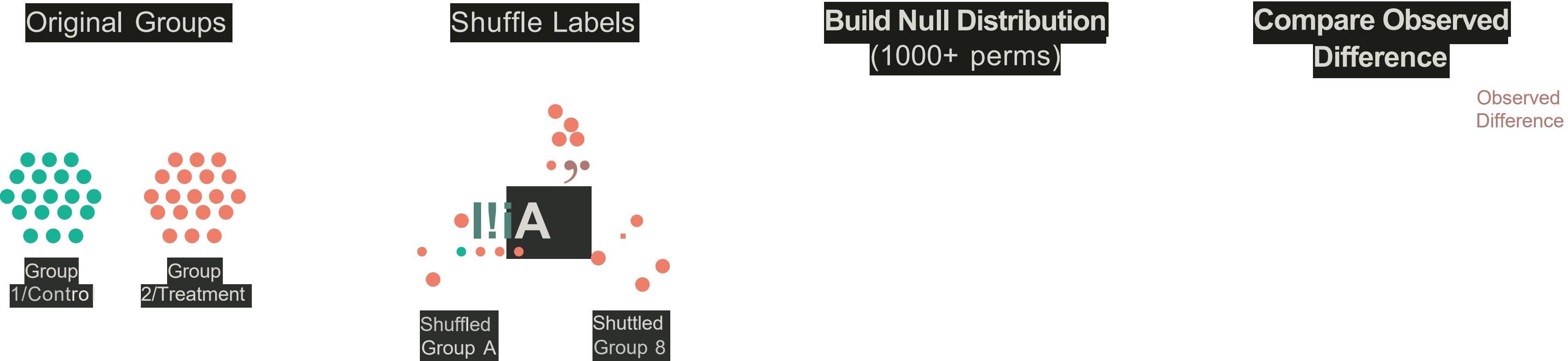
# Path 1: The Parametric Test

- **Logic:** Uses standard errors to compute t-tests on parameters.
- **Assumption:** Sampling distribution of group differences is Gaussian (Central Limit Theorem).
- **Pros:** Fast computation.
- **Cons:** Assumes normality (often violated in psych data). Cannot test Random Effects variances in mlVAR .



Comparing Means:  $\beta_{group1}$  vs  $\beta_{group2}$

# Path 2: The Nonparametric Permutation Test (Recommended)



Computationally heavy, but robust to non-normal data and applicable to random effects.

# mnet

**Modeling Group Differences and Moderation  
Effects in Statistical Network Models**

```
> install.packages('mnet')
> library(mnet)
```

The bridge between mlVAR estimation and hypothesis testing.

# Step 1: Data Preparation

Subject Identifier

ID	Time	Happy	Group
1	1	5	0
1	2	6	0
2	1	2	1
2	2	3	1

Variables of Interest

Must be a binary indicator (e.g., 0 vs 1).  
Continuous moderators must be binarized.

# Step 2: The Function Call

```
output <- mlVAR_GC( _____  
  data = data,  
  vars = c('Happy', 'Sad', 'Anxious'),  
  idvar = 'ID',  
  dayvar = 'Day',  
  beepvar = 'Beep',  
  groups = 'GroupVar',  
  test = 'permutation', - - - - -  
  nP = 1000  
)
```

Group Comparison  
Function

Critical Choice:  
Parametric vs  
Permutation

# Step 3: Critical Arguments

-  ``test``: Choose 'permutation' (robust) or 'parametric' (fast).
-  ``paired``: FALSE for independent samples (Patients vs Controls).  
TRUE for within-subject (Pre vs Post).
-  ``nCores``: Crucial for performance. Use parallel processing (e.g.,  
`nCores = 12`) to reduce runtime from days to minutes.
-  ``nP``: Number of Permutations. Recommend  $\geq 1000$  for stable  
p-values.

# Step 4: Interpreting the Output Object

output

\$EmpDiffs

Matrix of observed  
differences  
(Group 1 - Group 2)

\$ModelsEmp

\$Pvals

Matrix of  
significance values

\$Runtime\_min

Execution time in  
minutes

Original mIVAR  
objects for both  
groups

# Case Study: Emotion Dynamics in Depression

Source: Koval et al. (2013)

**Subjects: 95**

Undergraduate students

**Data: ESM**

10 times/day for 7 days

**Variables: 7**

(Happy, Relaxed, Sad, Angry,  
Anxious, Depressed, Stressed)

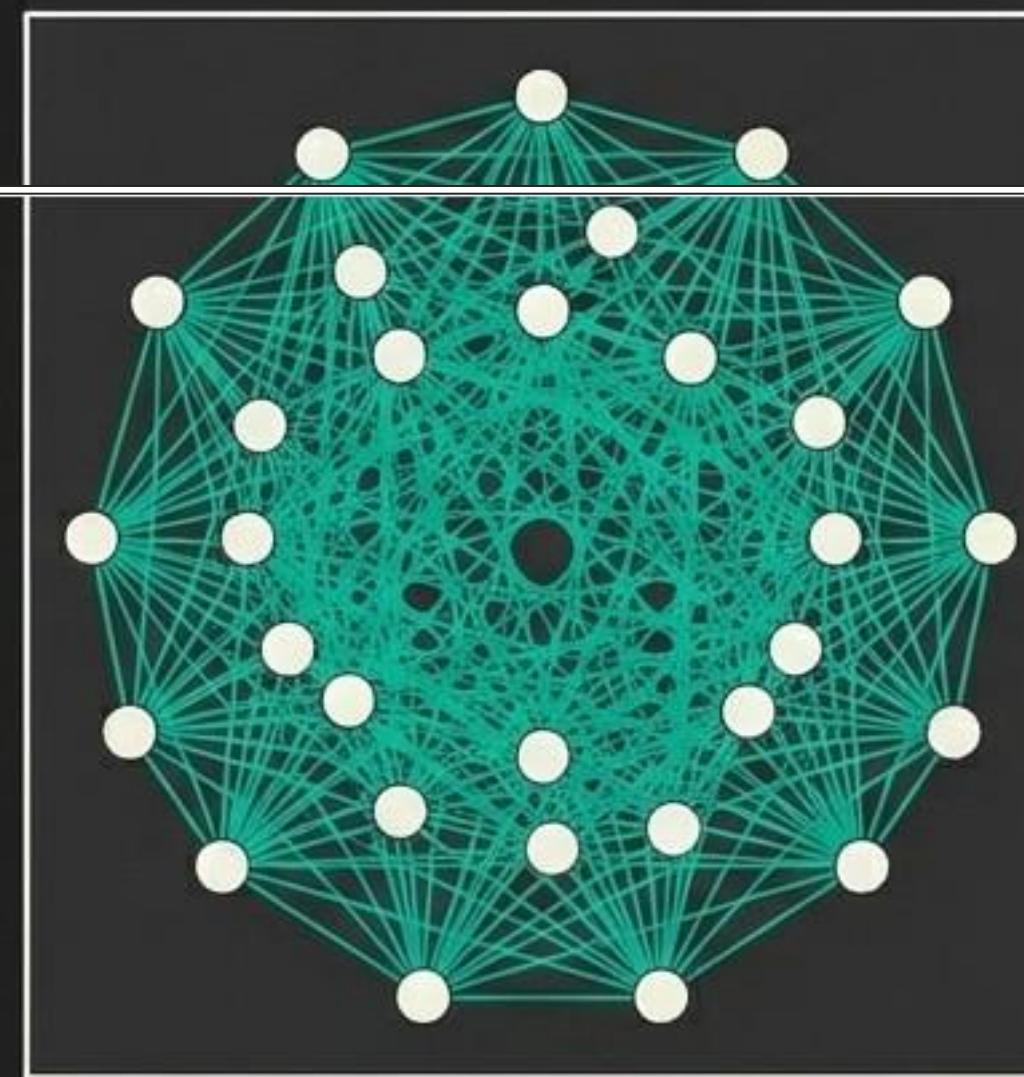
 Group 1  
(Low Depressive)

Defined by CES-D > 16.

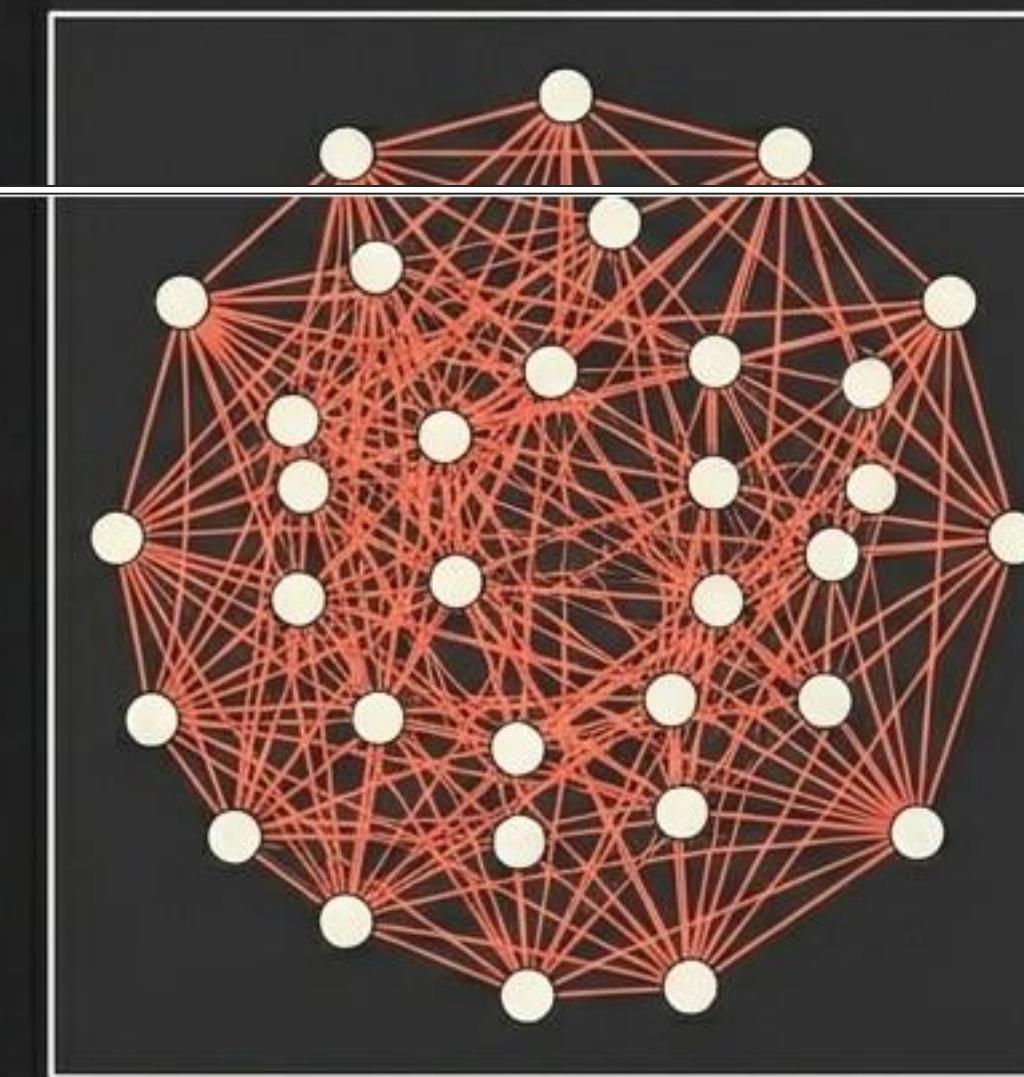
 Group 2  
(High Depressive)

# The Visual Inspection Trap

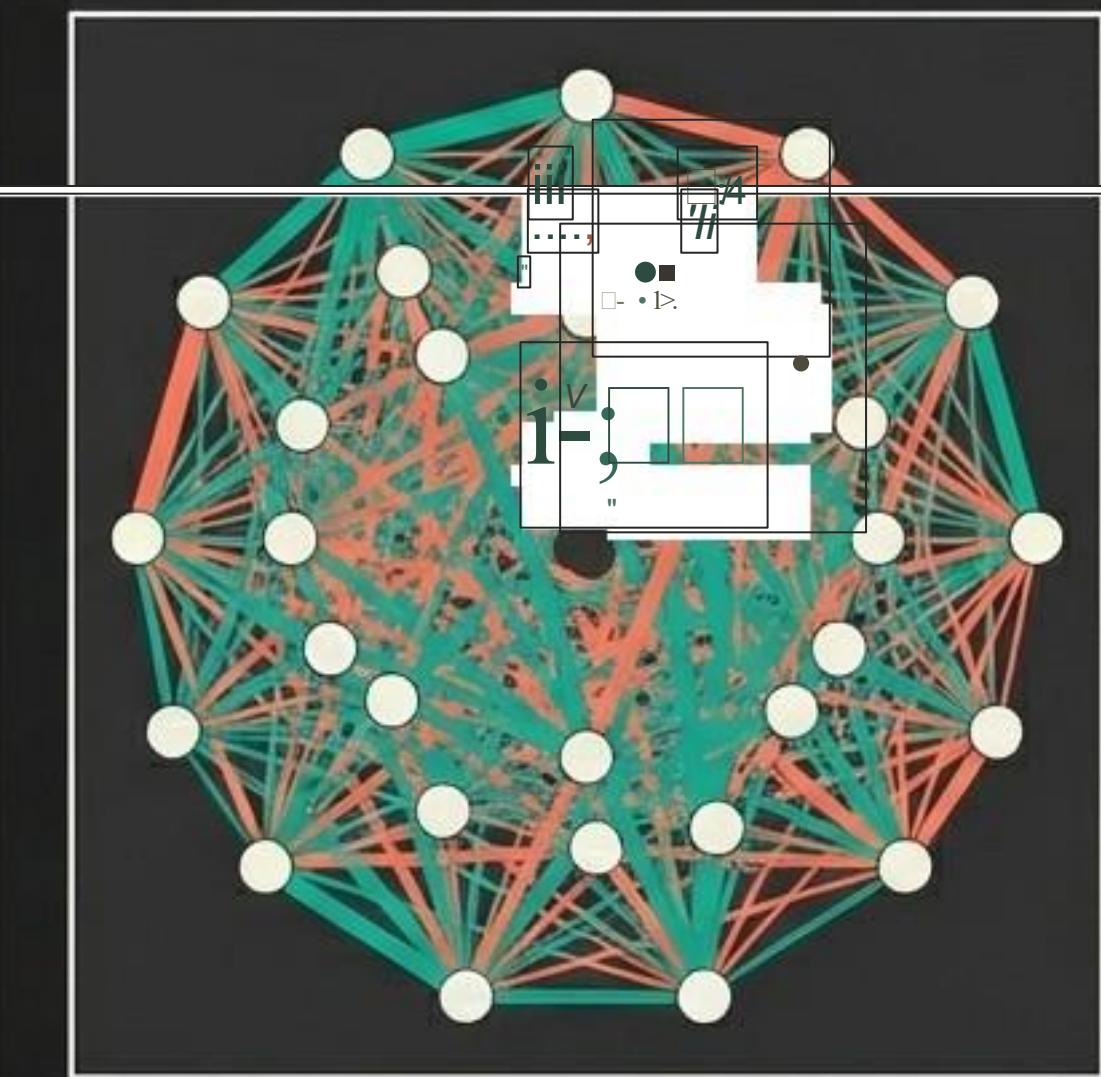
Group Low



Group High



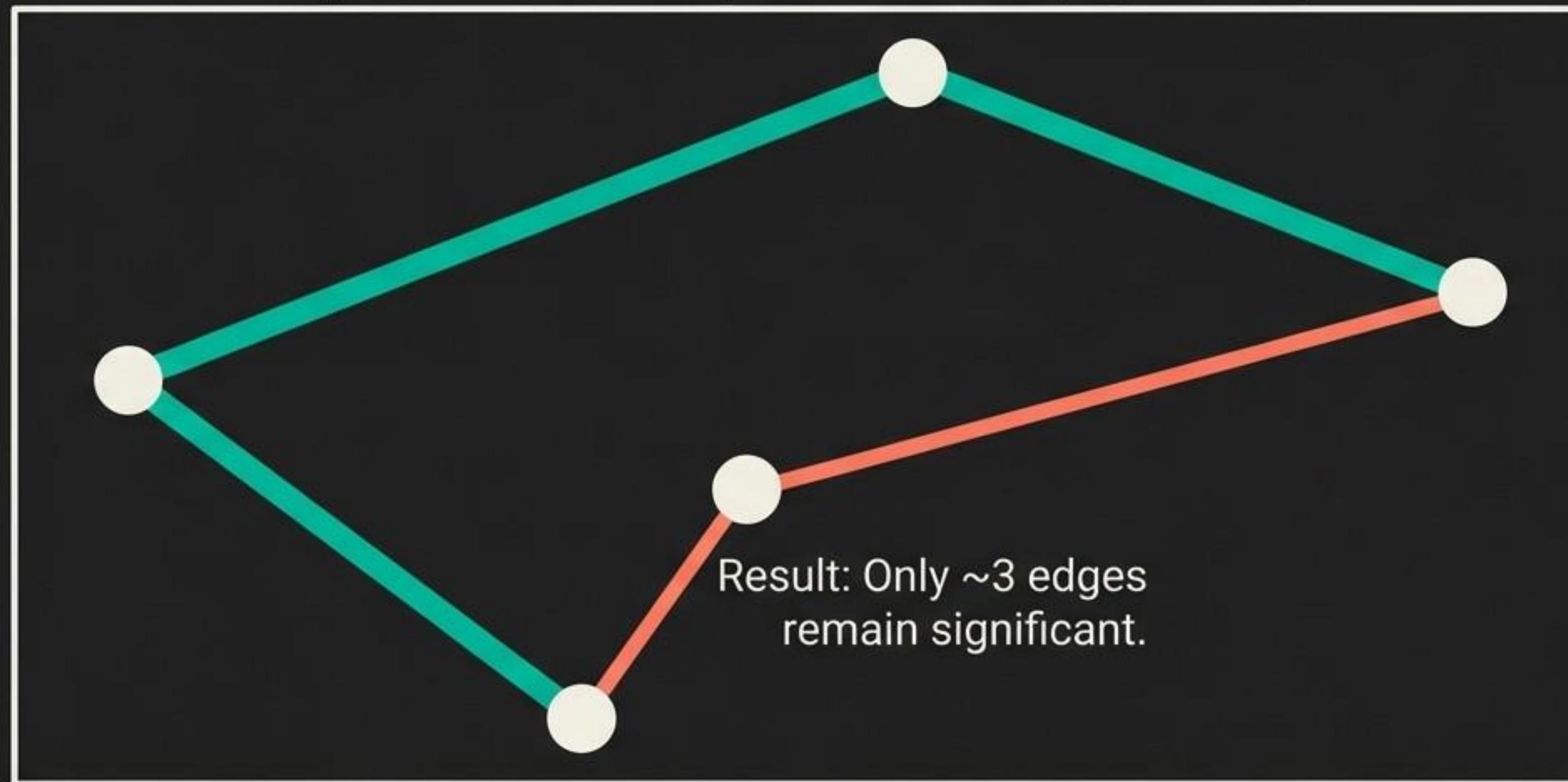
Raw Difference



Visually, the networks appear substantially different. Is it real?

# Filtering Signal from Noise

Significant Differences (Permutation Test, alpha = 0.05)



Most visual differences were due to sampling variation. Formal inference prevented false positives.

# Recommendations & Limitations

## ✓ Best Practices

- ✓ Use Permutation tests for robustness.
- ✓ Use  $\geq 1000$  permutations.
- ✓ Avoid visual 'eye-balling'.

## ⚠ Limitations

- ⚠ Computational cost is higher.
- ⚠ Caution with Between-Subject Networks (mlVAR estimation bias).

# Summary

mnet provides a methodological bridge from descriptive analysis to rigorous hypothesis testing.



GitHub: [jmbh/mlVARGD](https://github.com/jmbh/mlVARGD)

- Paper: Haslbeck, Epskamp, & Waldorp (2023)
- Package: `install.packages('mnet')`