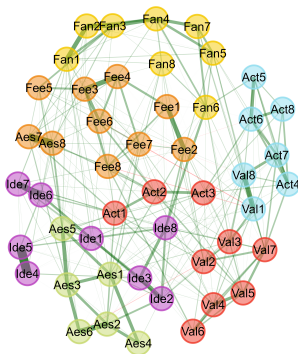


# EGA Framework

PSY-GS 8875 Behavioral Data Science



Overview: Week 13

## Readings (Optional)

- Christensen and Golino - 2021 - bootEGA
- Christensen et al. - 2023 - UVA
- Christensen and Golino - 2021 - loadings
- Jamison et al. - 2022
- Jimenez et al. - 2023
- Samo et al. - 2023

- Stability of communities and items
- Local dependence detection
- Network loadings
- Metric invariance
- Hierarchical dimensions



Stability of Communities and Items

# Stability of Communities and Items

**reliability:** are your measurements consistent (i.e., can they be repeated)?

- **internal consistency:** whether your items are interrelated – that is, moderate ( $r \geq 0.30$  to strongly correlated  $r \geq 0.50$ )
- **test-retest:** true “reliability” – whether your items can be repeated and are consistent each time you measure them

# Stability of Communities and Items

## Internal Consistency

$$\text{Cronbach's } \alpha = \frac{k}{k-1} \left( \frac{\sum_{i=1}^k \sigma_{x_i}^2}{\sigma_x} \right),$$

where

- $k$  = number of items
- $\sigma_{x_i}^2$  = variance of item  $i$
- $\sigma_x$  = variance associated with sum total of items  $x = \sum_{i=1}^k x_i$

💡 For more internal consistency measures, see [McNeish \(2018\)](#)

## Homogeneity

- Whether a set of items reflect a single underlying construct
- Often implicitly assumed and not usually tested (e.g., unidimensionality)

# Stability of Communities and Items

What seems stronger to be a **stronger** statement?

- Ⓐ internal consistency: items are interrelated
- Ⓑ homogeneity: items reflect a single underlying construct

# Stability of Communities and Items

What seems stronger to be a **stronger** statement?

- Ⓐ internal consistency: items are interrelated
- Ⓑ homogeneity: items reflect a single underlying construct

Both psychometric characteristics are important for measurement but are usually tested in a “silo”

# Stability of Communities and Items



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The question we usually want to answer is:

Do the items hang together in their representative dimensions  
*taking into account* the other items and dimensions?



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*and* homogeneous in a multivariate, multidimensional context

# Stability of Communities and Items

The question we usually want to answer is:

Do the items hang together in their representative dimensions  
*taking into account* the other items and dimensions?

That is, we want to know that our items are internally consistency  
*and* homogeneous in a multivariate, multidimensional context

 Traditional psychometric approaches do not consider multidimensionality

# Stability of Communities and Items

We also want to know whether our dimensions and the items placed in those dimensions are likely to **generalize**

# Stability of Communities and Items

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## Bootstrap Exploratory Graph Analysis

- Bootstrap using resampling with replacement (non-parametric) or multivariate normal data based on correlation matrix (parametric)
- Apply EGA to the replicate bootstrap sample

⚙ The community detection algorithm places items into dimensions *automatically*

# Stability of Communities and Items

From the bootstraps, we can...

- Determine how frequent the empirical number of dimensions appear across the bootstraps
- Determine how often items are placed into their empirical (or other) dimension
- Determine how often a dimension replicates *exactly* across bootstraps

# Stability of Communities and Items

## Implementation

```
# Load packages
library(EGAnet); library(psychTools)

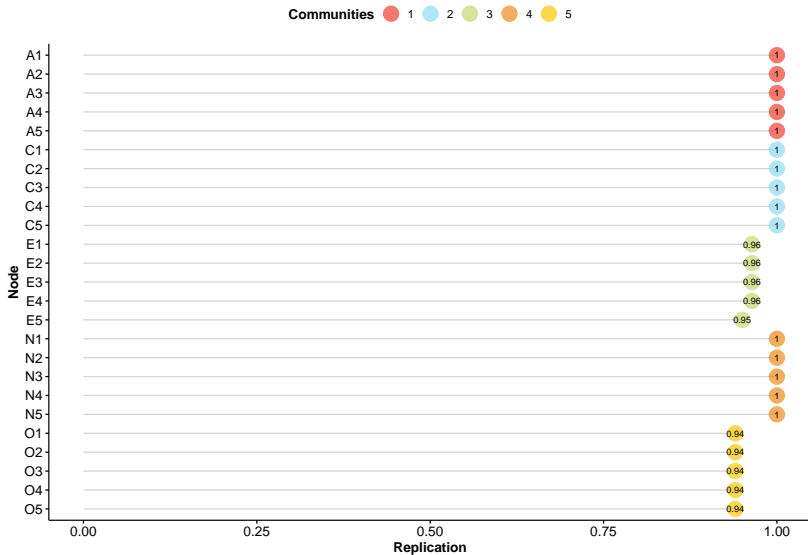
# Load data
data <- bfi[,1:25]

# Implement bootstrap EGA (empirical automatically computed)
bfi_boot <- bootEGA(data, seed = 42, ncores = 2)
# Seeds are set independent of R

# Print summary
summary(bfi_boot)

# Print dimension stability summary
summary(bfi_boot$stability)
```

# Stability of Communities and Items





# Stability of Communities and Items

Model: GLASSO (EBIC)  
Correlations: auto  
Algorithm: Walktrap  
Unidimensional Method: Louvain

----

EGA Type: EGA  
Bootstrap Samples: 500 (Parametric)

	4	5
Frequency:	0.096	0.904

Median dimensions: 5 [4.42, 5.58] 95% CI

# Stability of Communities and Items

EGA Type: EGA

Bootstrap Samples: 500 (Parametric)

Proportion Replicated in Dimensions:

A1	A2	A3	A4	A5	C1	C2	C3	C4	C5	E1	E2	E3
1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	0.964	0.964	0.964
E4	E5	N1	N2	N3	N4	N5	O1	O2	O3	O4	O5	
0.964	0.950	1.000	1.000	1.000	1.000	1.000	0.940	0.940	0.940	0.940	0.940	

----

Structural Consistency:

1	2	3	4	5
1.00	1.00	0.95	1.00	0.94

# Stability of Communities and Items

## Guidelines

- Empirical solution frequency should be majority
  - Item stability (replication)  $> 0.75$
  - Dimension stability  $> 0.75$
- 💡 Resampling (non-parametric) will tend to produce equal or lower estimates to multivariate normal (parametric)

# Stability of Communities and Items

R Script

## Causes of Instability

- Smaller sample sizes
- Local dependence
  - Items will form “minor factors” where a major factor will split into two or more communities
- Multidimensional
  - Items will replicate relatively evenly across two or more communities

# Local (In)dependence

Local (In)dependence

## Latent Variable Definition

Variables are unrelated after conditioning on a latent variable

- Shared semantic references (e.g., similar item phrasing)
- Shared substantive causes *not* related to the latent variable (e.g., social desirability)
- Conventional psychometric practices such as maximizing Cronbach's  $\alpha$

## Network Psychometrics

- Components of the network are defined as “unique causal systems”
- Components are *unique* such that they are causally autonomous (i.e., distinct causal process)
- **Consequence:** variables in the network should be *unique* and **not** redundant



# Local (In)dependence

Take a network with many variables that are fairly unique but you have the two items

- ❶ I like to be the center of attention
- ❷ I don't like attention

These two variables will be **strongly** connected (i.e., large edge weight)

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When evaluating the *node strength* or the sum of the connections to each node in the network, these two variables will likely have *inflated* values

Node strength quantifies how well connected a node is in the network and many researchers take this meaning as “importance”

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When evaluating the *node strength* or the sum of the connections to each node in the network, these two variables will likely have *inflated* values

Node strength quantifies how well connected a node is in the network and many researchers take this meaning as “importance”

*A question arises: Is the strength of these two nodes because they are indeed important or because they are redundant*

## Unique Variable Analysis

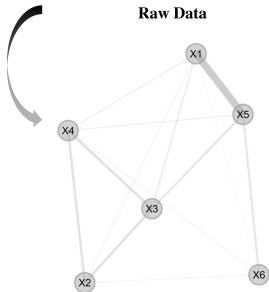
To assess whether there local dependence, Unique Variable Analysis (UVA) can be applied:

- ➊ Estimate a network (usually EBICglasso)
- ➋ Compute weighted topological overlap (wTO) on the network
- ➌ Apply a cut-off ( $\geq 0.25$ ) to determine redundant pairs
- ➍ Eliminate pairs based on some heuristics

# Local (In)dependence

	$X_1$	$X_2$	$X_3$	$X_4$	$X_5$	$X_6$
$P_1$	3	1	4	1	3	1
$P_2$	5	5	5	3	5	2
...	...	...	...	...	...	...
$P_i$	$x_{i,1}$	$x_{i,2}$	$x_{i,3}$	$x_{i,4}$	$x_{i,5}$	$x_{i,6}$

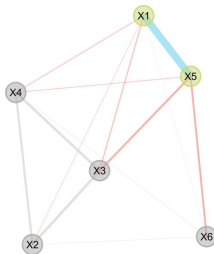
Raw Data



Network

$$\omega_{ij} = \frac{\sum_u a_{iu} a_{uj} + a_{ij}}{\min\{k_i, k_j\} + 1 - a_{ij}}$$

Weighted Topological Overlap (wTO)



Network with wTO

# Local (In)dependence

After cut-off, heuristics are used to eliminate redundant variable sets down to a single variable

**2 variables:** variable with the *lowest* maximum wTO to all *other* variables is retained

**3 or more variables:** variable with the *highest* mean wTO to all other variables in the *redundant set* is retained

# Local (In)dependence

## Implementation

```
# Apply UVA
bfi_uva <- UVA(
  data, key = as.character(bfi.dictionary$Item[1:25])
)

# Print summary
summary(bfi_uva)
```

# Local (In)dependence

Variable pairs with wTO > 0.30 (large-to-very large redundancy)

node_i	node_j	wto
Get angry easily.	Get irritated easily.	0.431

----

Variable pairs with wTO > 0.25 (moderate-to-large redundancy)

----

Variable pairs with wTO > 0.20 (small-to-moderate redundancy)

node_i	node_j	wto
Don't talk a lot.	Find it difficult to approach others.	0.226
Am exacting in my work.	Continue until everything is perfect.	0.225
Am indifferent to the feelings of others.	Inquire about others' well-being.	0.219
Do things in a half-way manner.	Waste my time.	0.209
Know how to comfort others.	Make people feel at ease.	0.207
Get angry easily.	Have frequent mood swings.	0.205
Have frequent mood swings.	Often feel blue.	0.204
Inquire about others' well-being.	Know how to comfort others.	0.203



# Local (In)dependence

R Script

## Effects of Reducing Redundancy

- ➊ More accurate dimension estimation: resolves issues associated with “minor factors” (i.e., smaller dimensions that form because of high shared variance between a smaller set of variables intend to form a dimension in a larger set)
- ➋ More accurate edge weights: associations between variables are due less to redundancy and more to their actual contribution to the network (assuming the network captures all variables of interest)

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Is reducing redundancy always necessary?

Network Loadings

# Network Loadings

Statistically consistent with factor/component loadings

## Loading Definitions

- **factor**: how much one item is related to the factor or how well an item represents and *measures* the latent factor
- **network**: each node's contribution to the *emergence* of a coherent dimension in the network

*In most applied circumstances, there is little difference*

In network science, network measures are more common:

- local = a node's position in the network (e.g., centrality)
- meso-scale = sub-structures such as communities
- global = overall structure of the network (e.g., average shortest path length)

Centrality (local) measures are still the most commonly applied measures in psychometric networks:

- node strength = absolute sum of a node's connections to other nodes in the network
- expected influence = signed sum of a node's connections to other nodes in the network

💡 There are hundreds of centrality measures but most are problematic with respect to psychometric interpretations (see [Bringmann et al., 2019](#))

Node Strength

$$S_i = \sum_{j=1}^n |w_{ij}|$$

Expected Influence

$$E_i = \sum_{j=1}^n w_{ij}$$



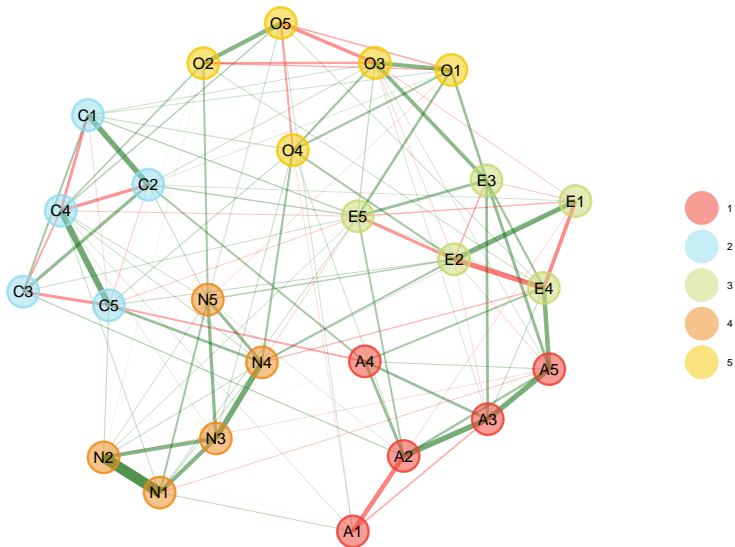
# Network Loadings

- Node strength is commonly used as a measure of “influence”
- In psychopathology, many have proposed symptoms highest in node strength as intervention targets
- These interpretations are misleading...
  - Assumes between-person model applies to all individuals in the sample
  - Assumes the network is unidimensional
  - Assumes all variables are unique (i.e., node strength is not due to redundancy)

# Network Loadings

```
# Apply EGA  
bfi_ega <- EGA(data)  
  
# Compute node strength  
sort(colSums(abs(bfi_ega$network)))
```

# Network Loadings



# Network Loadings

A1	C3	O2	A4	E1	C1	O4	O5	N5	O1	N3
0.43	0.63	0.64	0.64	0.66	0.73	0.74	0.75	0.76	0.80	0.92

A5	N2	C2	C5	A3	A2	E3	E5	N4	O3	C4
0.93	0.93	0.94	0.96	1.00	1.01	1.01	1.02	1.03	1.04	1.07

N1	E4	E2
1.07	1.11	1.16

Recall that... *Get angry easily* (N1) and *Get irritated easily* (N2) were determined to be locally dependent

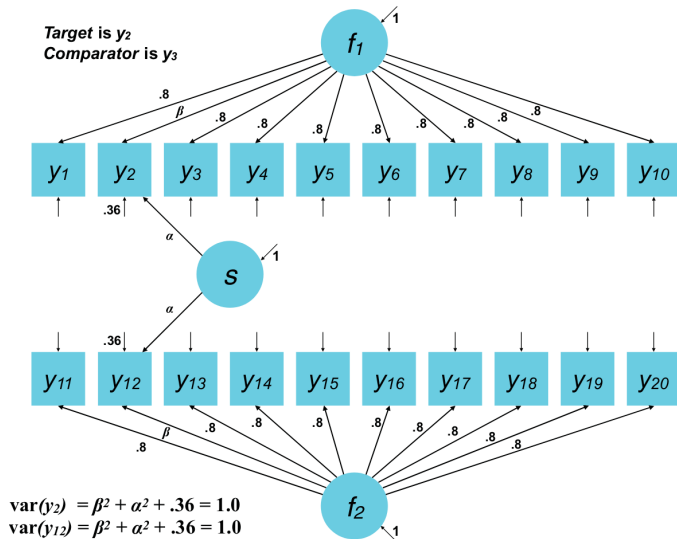
## Connections to Factor Loadings

Hallquist, Wright, and Molenaar (2019)

CFA Model	Strength	Closeness	Betweenness
One-factor	0.98	0.94	0.74
Orthogonal Two-factor	0.98	0.42	0.37
Correlated Two-factor	0.97	0.51	0.44
Orthogonal Three-factor	0.98	0.42	0.31
Correlated Three-factor	0.97	0.55	0.41

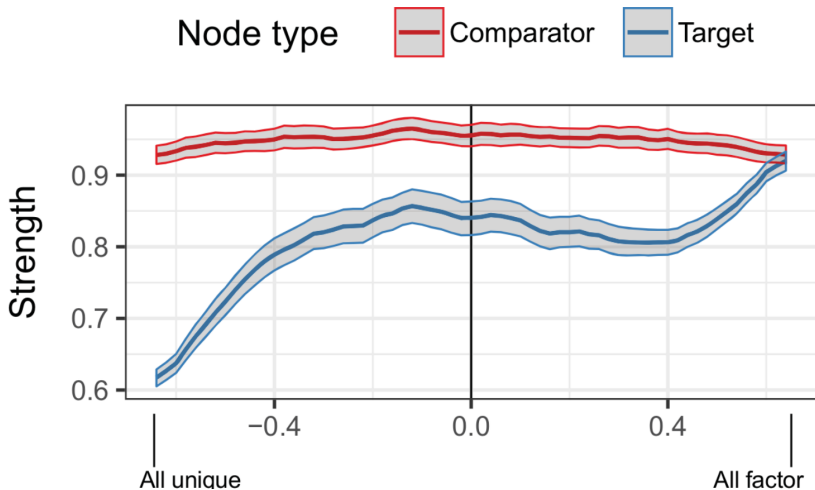
# Network Loadings

Hallquist, Wright, and Molenaar (2019)



# Network Loadings

Hallquist, Wright, and Molenaar (2019)



# Network Loadings

**Solution:** split node strength by community,  $c$

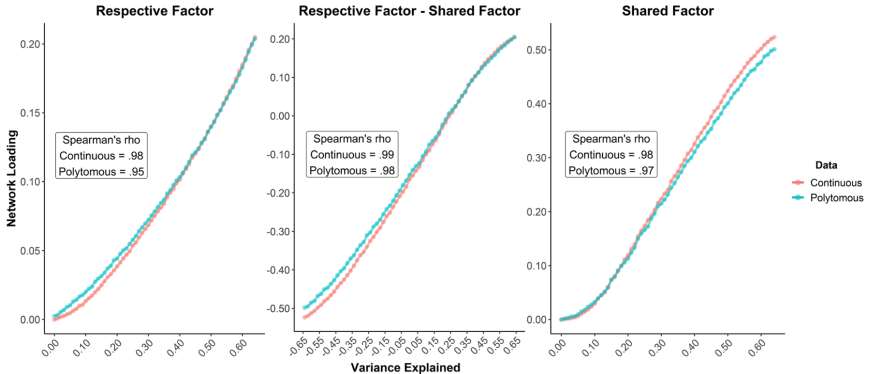
$$L_{ic} = \sum_{j \in c}^C |w_{ij}|$$

$$N_{ic} = \frac{L_{ic}}{\sqrt{\sum L_{.c}}}$$

with signs added afterward

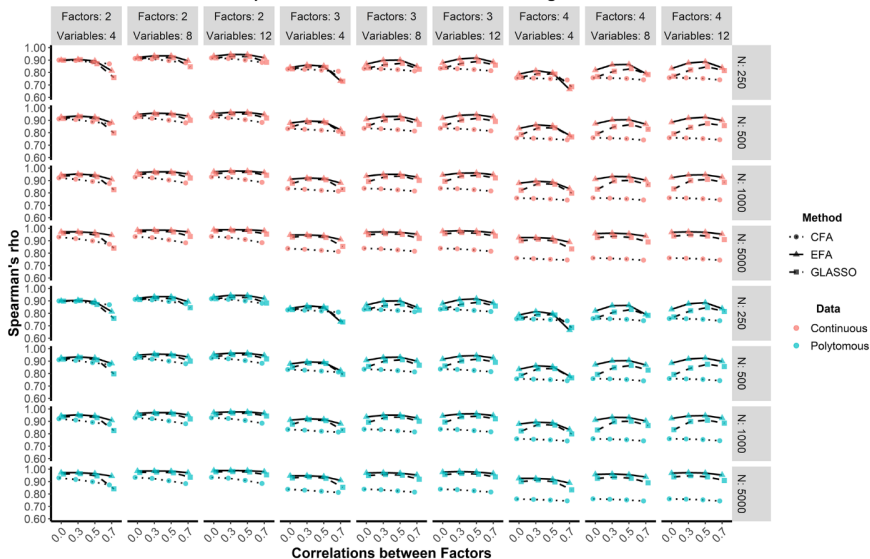


# Network Loadings

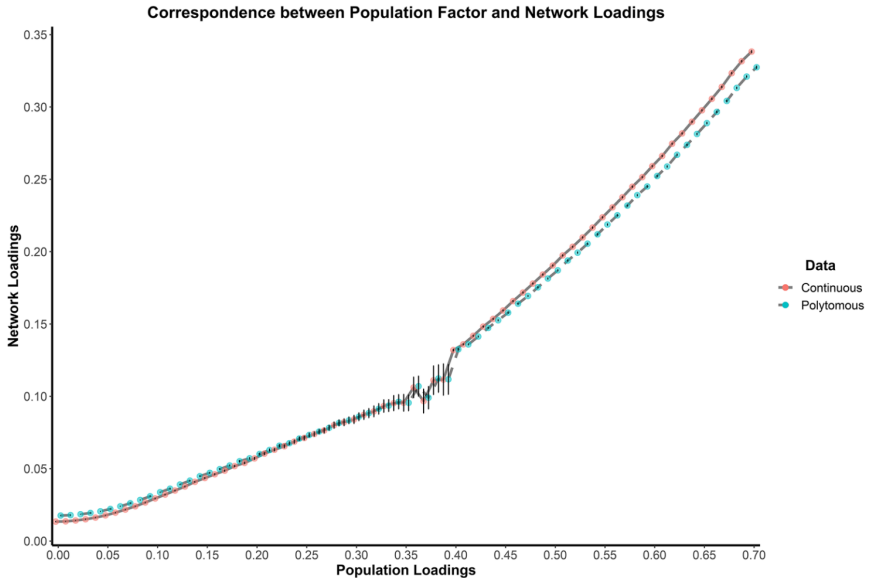


# Network Loadings

Comparison of Factor and Network Loadings



# Network Loadings



There were some lingering issues though...

- ❶ Negative signs were added post-hoc and in a way that didn't always align
- ❷ Community-assigned loadings were sometimes *smaller* than their cross-loadings (impossible with factor analysis)
- ❸ Magnitudes are significantly affected by number of variables per community

# Network Loadings

## Revised Loadings

$$\text{within } \ell_{i,c} = n_c \left( \frac{\sum_{j=1}^{n_c} t_{i,j}}{n_c - 1} \right),$$

and

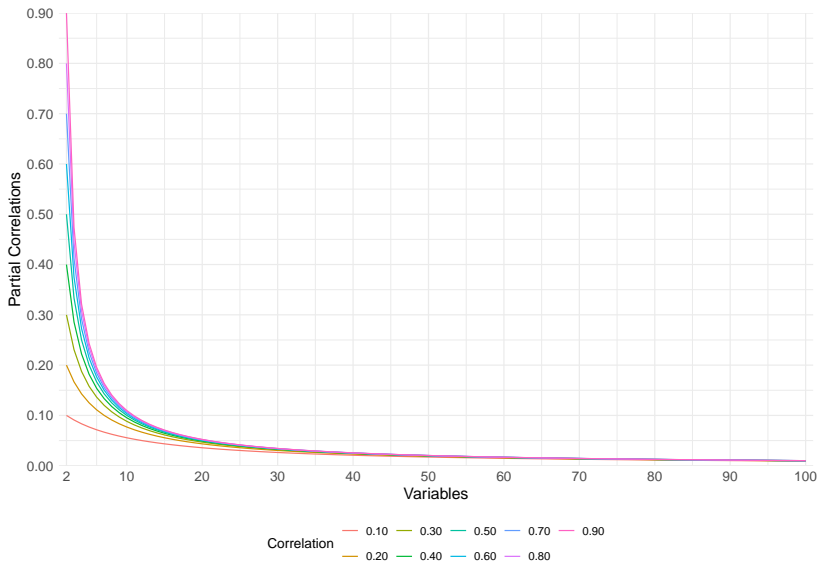
$$\text{between } \ell_{i,k} = \sum_{i=1}^{n_c} \sum_{j=1}^{n_k} w_{i \in c, j \in k},$$

where

- $t_{i,j}$  = target community sub-matrix with node  $i$  and  $j$  in community  $c$
- $n_c$  = number of nodes in the assigned community,  $c$
- $n_k$  = number of nodes in a community,  $k$ , that is not  $c$

# Network Loadings

💡 Guttman (1953): as  $n_c \rightarrow \infty$ , then  $r_{xy|z} \rightarrow 0$



## Revised Loadings

$$N_{i,c} = \frac{\ell_{i,c}}{\sqrt{\log(\zeta n_c) \sum_{j=1}^{n_c} |\text{within } \ell_{j,c}|}},$$

where

- $\log(n_c)$  = natural logarithm of the number of variables in community  $c$
- $\zeta$  = scaling factor for loading size (defaults to 2)

# Network Loadings

There were some lingering issues though...

- ❶ Negative signs were added post-hoc and in a way that didn't always align ✓
- ❷ Community-assigned loadings were sometimes *smaller* than their cross-loadings (impossible with factor analysis) ✓
- ❸ Magnitudes are significantly affected by number of variables per community ⚙️



# Network Loadings

Okay... but why network loadings at all?

❶ Need for community-aligned loadings

→ Number of communities in factor analysis does not guarantee alignment with variable assignments

❷ Network loadings are *unrotated*

❸ Networks have fewer assumptions than factor models

→ Psychometric reference for when factor models don't work

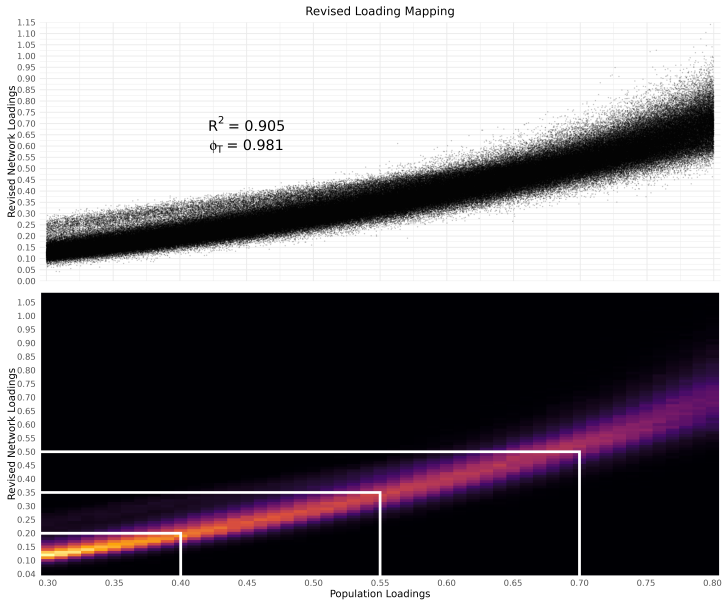
# Network Loadings

```
# Compute network loadings  
bfi_loadings <- net.loads(  
  bfi_ega, loading.method = "experimental"  
)$std[colnames(bfi_ega$network),] # standardized
```

# Network Loadings

	1	2	3	4	5
A1	<b>-0.24</b>	-0.01	0.00	0.03	-0.03
A2	<b>0.58</b>	0.04	0.07	0.01	0.02
A3	<b>0.57</b>	0.00	0.12	0.00	0.01
A4	<b>0.25</b>	0.11	0.06	0.00	0.00
A5	<b>0.33</b>	0.00	0.24	-0.03	0.01
C1	0.00	<b>0.39</b>	0.04	0.00	0.07
C2	0.06	<b>0.46</b>	0.03	0.02	0.03
C3	0.04	<b>0.37</b>	0.02	0.00	0.00
C4	-0.01	<b>-0.52</b>	-0.02	0.06	-0.10
C5	-0.06	<b>-0.37</b>	-0.05	0.10	0.03
E1	-0.01	0.01	<b>-0.41</b>	0.01	-0.02
E2	-0.02	-0.03	<b>-0.56</b>	0.09	0.05
E3	0.17	0.00	<b>0.29</b>	0.00	0.20
E4	0.23	0.00	<b>0.44</b>	-0.04	-0.04
E5	0.06	0.14	<b>0.29</b>	0.02	0.12
N1	-0.05	-0.02	0.02	<b>0.59</b>	0.00
N2	-0.01	-0.03	0.02	<b>0.54</b>	0.00
N3	0.00	-0.01	0.00	<b>0.56</b>	0.03
N4	0.00	-0.10	-0.13	<b>0.35</b>	0.07
N5	0.01	0.01	-0.05	<b>0.30</b>	-0.09
O1	0.00	0.03	0.16	-0.01	<b>0.37</b>
O2	0.00	-0.05	0.02	0.06	<b>-0.31</b>
O3	0.02	0.04	0.18	0.00	<b>0.48</b>
O4	0.03	0.00	-0.06	0.08	<b>0.26</b>
O5	-0.02	-0.04	0.02	0.01	<b>-0.45</b>

# Network Loadings



# Network Loadings

Network loadings open the door for many different traditional psychometric procedures

- **group comparison (with dimensionality)**
- **network scores (and hierarchical dimensionality)**
- conversion of loadings to IRT parameters ([Muraki & Carlson, 1995](#))

Metric Invariance

Group comparison is often a goal in the social sciences

Many methods have been developed to make group comparisons in network psychometrics

- Fused GLASSO
- Network Comparison Test
- Group-as-Moderator
- Bayesian Posteriors

*All of these methods implicitly treat the network as unidimensional*

## Motivating Example

Using the Big Five data as an example, let's say we want to examine whether there are any personality differences between those with a college degree and those without

```
# Obtain groups
groups <- ifelse(bfi[, "education"] < 4, "Non-grad", "Grad")

# Filter for missing groups
group_data <- data[!is.na(groups),]
groups <- na.omit(groups)

# Frequencies
table(groups)
```

```
groups
  Grad Non-grad
   812   1765
```



## Procedure

A permutation-based procedure can be employed to test for differences between groups (assuming some structure holds for both groups):

- 1 Estimate networks and network loadings for both groups
- 2 Compute the difference between the *assigned* loadings ( $\tau$ )
- 3 Permutation: shuffle group label and repeat steps 1. and 2. for  $P$  times (e.g., 500;  $\tau_{R_p}$ )
- 4 Compute  $\sum_{p=1}^P |\tau| \geq \tau_{R_p}$  to obtain  $p$ -values

**Significant** differences ( $p < 0.05$ ) suggest non-invariance (group differences exist) whereas  $p > 0.05$  suggest invariance (group differences **do not** exist)

R Script

# Metric Invariance

```
# Perform metric invariance
bfi_invariance <- invariance(
  data = group_data, groups = groups,
  structure = rep(1:5, each = 5), # theoretical structure
  loading.method = "experimental", # use latest loadings
  ncores = 8, seed = 42
)

# Summary
summary(bfi_invariance)

# Plot
plot(bfi_invariance, p_type = "p_BH")
```

# Metric Invariance

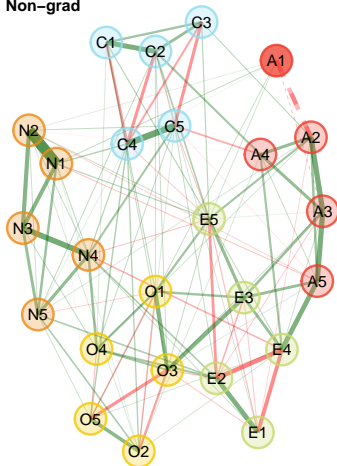
	Membership	Difference	p	p_BH	sig	Direction
A1	1	0.181	0.002	0.050	**	Non-grad > Grad
A2	1	0.094	0.070	0.438	.	
A3	1	-0.032	0.570	0.679		
A4	1	0.042	0.286	0.596		
A5	1	-0.036	0.376	0.609		
C1	2	-0.076	0.136	0.486		
C2	2	-0.084	0.096	0.480	.	
C3	2	0.003	0.940	0.940		
C4	2	-0.036	0.504	0.630		
C5	2	0.004	0.936	0.940		
E1	3	0.033	0.414	0.609		
E2	3	0.122	0.018	0.150	*	Non-grad > Grad
E3	3	0.038	0.410	0.609		
E4	3	0.044	0.318	0.609		
E5	3	0.048	0.264	0.596		
N1	4	0.042	0.278	0.596		
N2	4	-0.004	0.914	0.940		
N3	4	-0.052	0.128	0.486		
N4	4	0.010	0.796	0.905		
N5	4	0.039	0.234	0.596		
O1	5	0.038	0.470	0.630		
O2	5	0.042	0.360	0.609		
O3	5	0.035	0.484	0.630		
O4	5	-0.064	0.158	0.494		
O5	5	0.222	0.008	0.100	**	Non-grad > Grad

-----

Signif. code: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 'n.s.' 1

# Metric Invariance

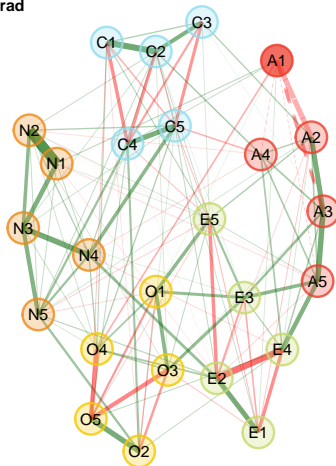
Non-grad



Invariant ( $p_{\text{adj.}} > 0.05$ )



Grad



Noninvariant ( $p_{\text{adj.}} < 0.05$ )



## Non-invariant Items

Item	Description	Significance	Direction
A1	Am indifferent to the feelings of others.	p_BH	Non-grad > Grad
E2	Find it difficult to approach others.	p	Non-grad > Grad
O5	Will not probe deeply into a subject.	p	Non-grad > Grad

# Metric Invariance

Group differences can be examined with the network psychometric framework *accounting for* the community structure

Tends to show comparable accuracy to traditional methods (e.g., SEM) with some advantage for disparate sample sizes (see Jamison et al., 2022)

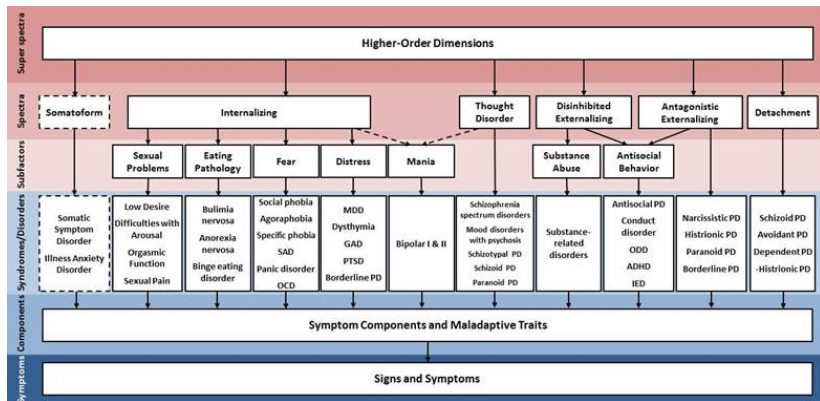
Scores can be computed based on the network loadings using **X** (available using `net.scores()`)

Hierarchical Dimensions



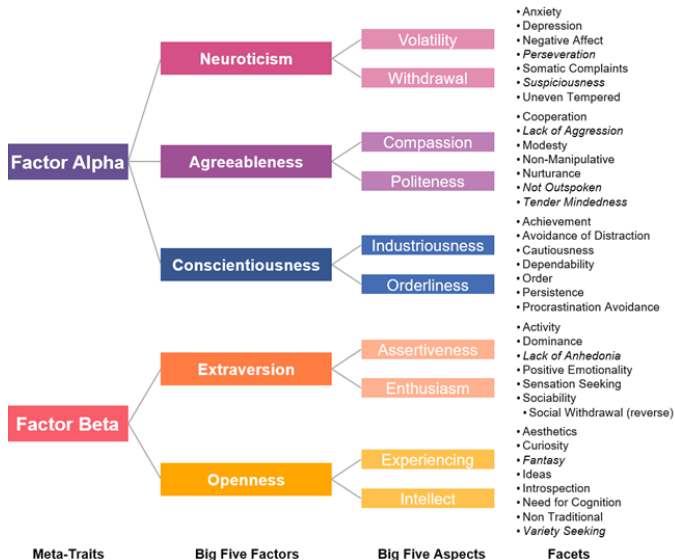
# Hierarchical Dimensions

Many psychological phenotypes are theorized to be hierarchically structured



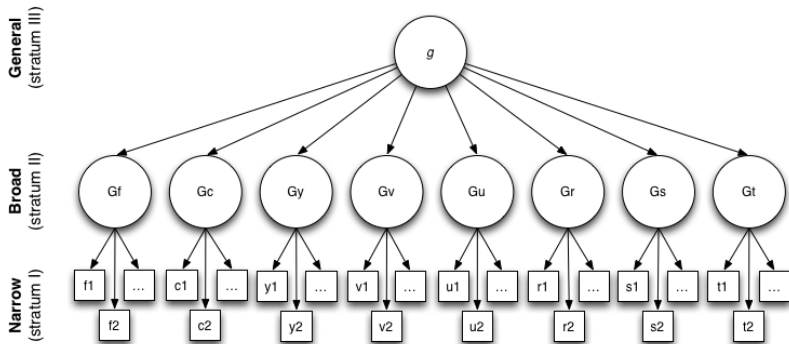
# Hierarchical Dimensions

Many psychological phenotypes are theorized to be hierarchically structured



# Hierarchical Dimensions

Many psychological phenotypes are theorized to be hierarchically structured



## **Motivating Example:** Synthetic Aperture Personality Assessment

The SPI ([SAPA Personality Inventory](#)) is a set of 135 items primarily selected from the [International Personality Item Pool](#)

Extensive factor analytic and psychometric analyses ([Condon, 2017](#)) have arrived at the “Little” 27 lower-order and can be narrowed to a 70-item Big Five (e.g., last week’s AHA)

**Motivating Question:** Do we find the Little 27 and Big Five using hierarchical EGA?

## Hierarchical EGA

- ➊ Apply EGA using the *first pass* of the Louvain algorithm to obtain the lower order dimensions
- ➋ Estimate network loadings and compute network scores based on lower order dimensions
- ➌ Apply EGA to the network scores to obtain the higher order dimensions

## Caveat

Remember: the Louvain algorithm results can change with node ordering

This stochastic nature of the algorithm is more acute at the lowest level (i.e., first pass)

To mitigate this issue, an approach known as *consensus clustering* can be used

## Consensus Clustering ([Lancichinetti & Fortunato, 2012](#))

- ➊ Randomly shuffle node order
- ➋ Apply Louvain algorithm
- ➌ Repeat 1. and 2. for  $N$  times (e.g., 1000)
- ➍ Obtain *most common* community structure across  $N$  applications

**Result:** more consistent (and accurate) results

# Hierarchical Dimensions

## Application

```
# Obtain SAPA data
sapa <- psychTools::spi[,11:145]

# Apply hierarchical EGA
sapa_hier <- hierEGA(
  data = sapa,
  loading.method = "experimental",
  scores = "network"
)

# Summary
summary(sapa_hier)

# Plot
plot(sapa_hier)
```



R Script

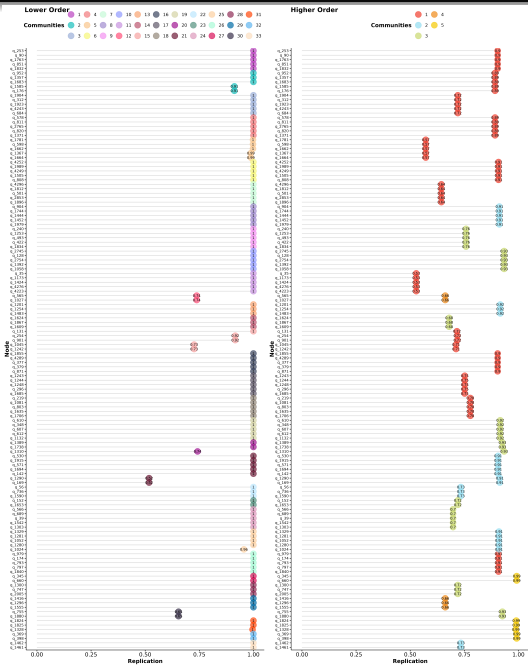
# Hierarchical Dimensions

## How Stable are these Dimensions?

```
# Apply bootstrap hierarchical EGA
sapa_hier_boot <- bootEGA(
  data = sapa, EGA.type = "hierEGA",
  loading.method = "experimental",
  scores = "network",
  ncores = 8, seed = 42
)

# Summary
summary(sapa_hier_boot)
```

# Hierarchical Dimensions



Summary

# Summary

`bootEGA` = determine the dimension and item stability as well as potential for problematic items

`UVA` = determine redundancies in the network (or local dependence in latent variable modeling)

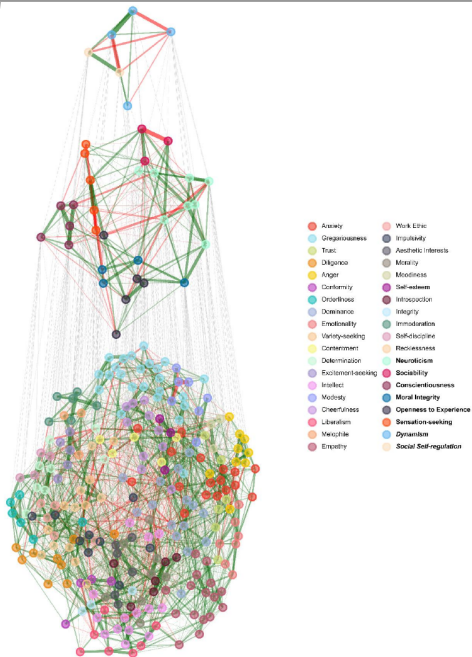
`net.loads` = estimate network loadings

`invariance` = compute metric invariance based on network psychometrics

`net.scores` = compute network scores based on network loadings

`hierEGA` = estimate hierarchical dimensionality

# Summary



Dynamic Readings

Derivatives: [Deboeck et al. \(2009\)](#)

Dynamic EGA: [Golino et al. \(2022\)](#)

Vector autoregression networks: [Epskamp et al. \(2018\)](#)

GIMME: [Beltz and Gates \(2017\)](#)

Heterogeneity in Dynamic Structures

- [Golino et al. \(2023\)](#)
- [Santoro and Nicosia \(2020\)](#)
- [De Domenico et al. \(2015\)](#)