

# Setting up a CFA

FACTOR ANALYSIS IN R

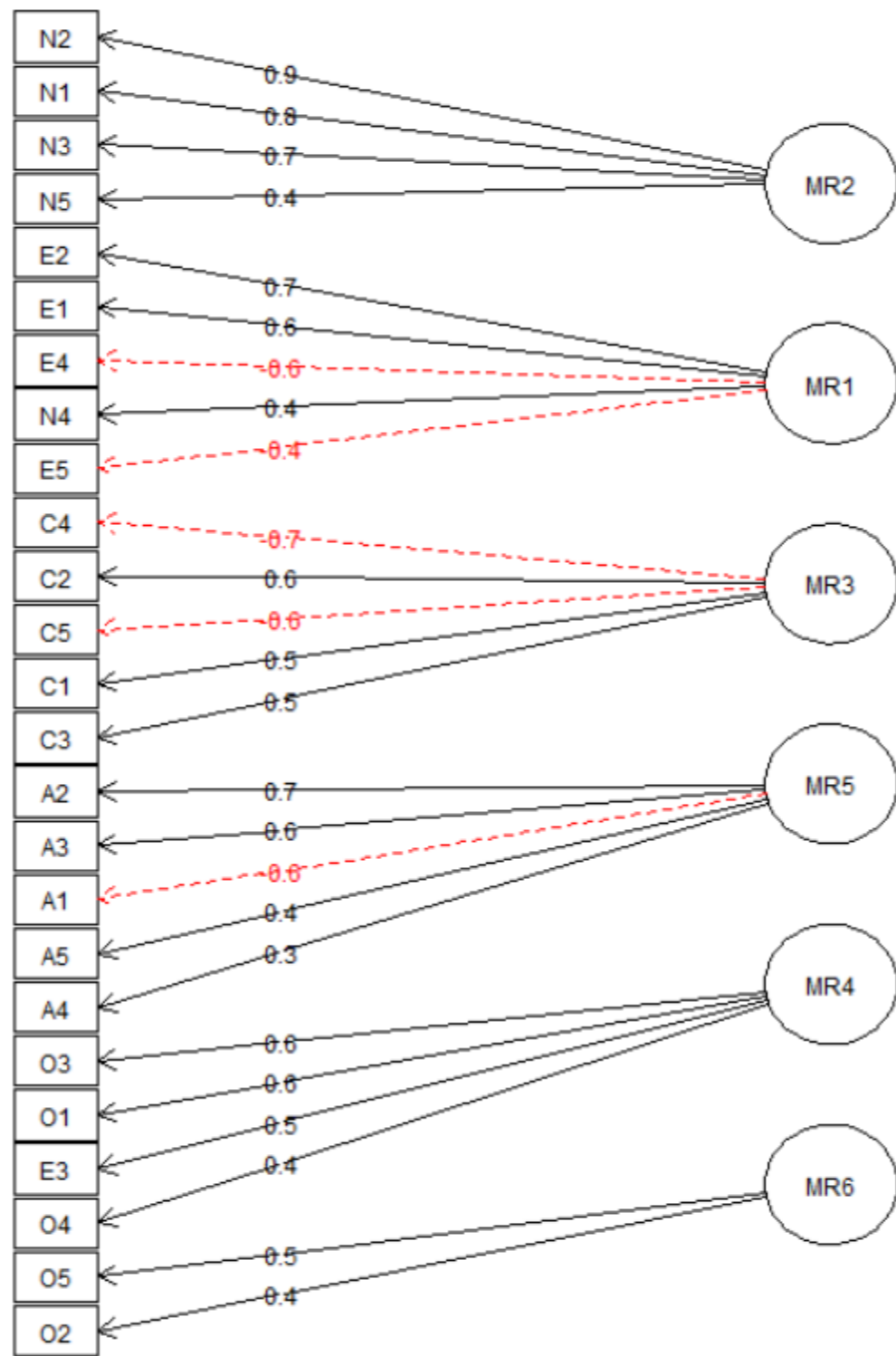


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# Why a confirmatory analysis?

Benefits of a confirmatory analysis:

- Explicitly specified variable/factor relationships
- Testing a theory that you know in advance
- This is the right thing to publish when you are developing a new measure!



# Using the wrapper function to set up a CFA

```
EFA_syn <- structure.sem(EFA_model)
EFA_syn
```

```
      Path      Parameter Value
[1,] "MR5->A1"  "F4A1"    NA
[2,] "MR5->A2"  "F4A2"    NA
[3,] "MR5->A3"  "F4A3"    NA
[4,] "MR5->A4"  "F4A4"    NA
[5,] "MR5->A5"  "F4A5"    NA
[6,] "MR3->C1"  "F3C1"    NA
[7,] "MR3->C2"  "F3C2"    NA
[8,] "MR3->C3"  "F3C3"    NA
[9,] "MR3->C4"  "F3C4"    NA
[10,] "MR3->C5" "F3C5"    NA
[11,] "MR1->E1" "F2E1"    NA
...
```

# Syntax created from the wrapper function

```
EFA_syn
```

```
      Path      Parameter Value  
[1, ] "MR5->A1"  "F4A1"    NA
```

- Factor 4 (F4) = Factor MR5 from the EFA
- Examinees' level of a factor predicts item responses
- Wrapper function automatically names parameters
- NA Value = starting value chosen at random

# Creating CFA syntax from your theory

```
# Set up syntax specifying which items load onto each factor
theory_syn_eq <- "
AGE: A1, A2, A3, A4, A5      #Agreeableness
CON: C1, C2, C3, C4, C5      #Conscientiousness
EXT: E1, E2, E3, E4, E5      #Extraversion
NEU: N1, N2, N3, N4, N5      #Neuroticism
OPE: O1, O2, O3, O4, O5      #Openness
"
```

- Short, memorable factor names
- Factor name followed by colon
- Items in a comma-separated list

```
theory_syn <- cfa(text = theory_syn_eq, reference.indicators = FALSE)
```

# Let's create some syntax!

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# Understanding the `sem()` syntax

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# Relationships between variables and factors

theory\_syn

```
Path      Parameter  StartValue
1 AGE-> A1  lam[A1:AGE]
2 AGE-> A2  lam[A2:AGE]
3 AGE-> A3  lam[A3:AGE]
4 AGE-> A4  lam[A4:AGE]
5 AGE-> A5  lam[A5:AGE]
6 CON-> C1  lam[C1:CON]
7 CON-> C2  lam[C2:CON]
8 CON-> C3  lam[C3:CON]
9 CON-> C4  lam[C4:CON]
10 CON-> C5  lam[C5:CON]
11 EXT-> E1  lam[E1:EXT]
...
```

1. Path: Relationships between factors and items
2. Parameter: Automatically assigned names for each parameter
3. Starting value: Blank means they will be randomly generated

# Factor variances

```
theory_syn
```

```
Path          Parameter      StartValue
26 AGE <-> AGE <fixed>      1
27 CON <-> CON <fixed>      1
28 EXT <-> EXT <fixed>      1
29 NEU <-> NEU <fixed>      1
30 OPE <-> OPE <fixed>      1
```

# Factor covariances

theory\_syn

	Path	Parameter	StartValue
31	AGE <-> CON	C[AGE,CON]	
32	AGE <-> EXT	C[AGE,EXT]	
33	AGE <-> NEU	C[AGE,NEU]	
34	AGE <-> OPE	C[AGE,OPE]	
35	CON <-> EXT	C[CON,EXT]	
36	CON <-> NEU	C[CON,NEU]	
37	CON <-> OPE	C[CON,OPE]	
38	EXT <-> NEU	C[EXT,NEU]	
39	EXT <-> OPE	C[EXT,OPE]	
40	NEU <-> OPE	C[NEU,OPE]	

# Item variances

theory\_syn

```
Path      Parameter  StartValue
41 A1 <-> A1    V[A1]
42 A2 <-> A2    V[A2]
43 A3 <-> A3    V[A3]
44 A4 <-> A4    V[A4]
45 A5 <-> A5    V[A5]
46 C1 <-> C1    V[C1]
47 C2 <-> C2    V[C2]
48 C3 <-> C3    V[C3]
49 C4 <-> C4    V[C4]
50 C5 <-> C5    V[C5]
51 E1 <-> E1    V[E1]
52 E2 <-> E2    V[E2]
```

...

# Running the CFA

Actually running the CFA is *much* easier than setting up the syntax!

```
#Use the sem() function to run a CFA  
theory_CFA <- sem(theory_syn, data = bfi_CFA)
```

```
summary(theory_CFA)
```

```
Model Chi-square = 2212.032   Df = 265 Pr(>ChiSq) = 9.662018e-304
AIC = 2332.032
BIC = 326.618
Normalized Residuals
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
-5.5800 -0.3732  1.0350  1.1220  2.4710  8.9000
R-square for Endogenous Variables
  A1    A2    A3    A4    A5    C1    C2    C3    C4
0.1178 0.4475 0.5731 0.2994 0.4713 0.3006 0.3667 0.2947 0.4886
...
Parameter Estimates
      Estimate  Std Error  z value  Pr(>|z|)
lam[A1:AGE] -0.5011716 0.04487184 -11.168956 5.785714e-29 A1 <--- AGE
lam[A2:AGE]  0.8230960 0.03447831  23.872862 5.863008e-126 A2 <--- AGE
...
```

# Let's practice!

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# Investigating model fit

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# Default fit statistics

Chi-square test (aka the log likelihood test) is only default

```
summary(theory_CFA)
```

```
Model Chi-square = 2231.647    Df = 265 Pr(>ChiSq) = 1.695873e-307
```

- Often significant due to sample size
- Desired outcome is lack of significance

# Changing the options

```
options(fit.indices = c("CFI", "GFI", "RMSEA", "BIC"))
```

- $RMSEA < 0.05$
- $GFI$  (Goodness of Fit Index)  $> 0.90$
- $CFI$  (Comparative Fit Index)  $> 0.90$

# Absolute model fit

```
summary(theory_CFA)
```

```
Model Chi-square = 2305.159    Df = 271 Pr(>ChiSq) = 0  
Goodness-of-fit index = 0.8527977  
RMSEA index = 0.07815051    90% CI: (NA, NA)  
Bentler CFI = 0.7754574
```

# Relative fit

```
summary(theory_CFA)
```

```
Model Chi-square = 2305.159    Df = 271 Pr(>ChiSq) = 8.422189e-319  
Goodness-of-fit index = 0.8527977  
RMSEA index = 0.07815051    90% CI: (NA, NA)  
Bentler CFI = 0.7754574  
BIC = 377.0563
```

```
summary(theory_CFA)$BIC
```

```
326.618
```

# Relative fit: comparing models

```
summary(theory_CFA)$BIC
```

```
326.618
```

```
# Run a CFA using the EFA syntax you created earlier  
EFA_CFA <- sem(EFA_syn, data = bfi_CFA)  
summary(EFA_CFA)$BIC
```

```
377.0563
```

- Useful for nested models that are fit to the same dataset
- Don't use if these conditions are not met!

# Let's practice!

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