

Practical: Phenotypic Factor Analysis

Big 5 dimensions Neuroticism & Extraversion in 361 female UvA students

- Exploratory Factor Analysis (**EFA**) using R (factanal) with Varimax and Promax rotation
- Confirmatory Factor Analysis (**CFA**) using OpenMx

Neuroticism: identifies individuals who are prone to psychological distress

n1 - Anxiety: level of free floating anxiety

n2 - Angry Hostility: tendency to experience anger and related states such as frustration and bitterness

n3 - Depression: tendency to experience feelings of guilt, sadness, despondency and loneliness

n4 - Self-Consciousness: shyness or social anxiety

n5 - Impulsiveness: tendency to act on cravings and urges rather than delaying gratification

n6 - Vulnerability: general susceptibility to stress

Extraversion: quantity and intensity of energy directed outwards into the social world

e1 - Warmth: interest in and friendliness towards others

e2 - Gregariousness: preference for the company of others

e3 - Assertiveness: social ascendancy and forcefulness of expression

e4 - Activity: pace of living

e5 - Excitement Seeking: need for environmental stimulation

e6 - Positive Emotions: tendency to experience positive emotions

Part 1: read the data**# clear the memory**

```
rm(list=ls(all=TRUE))
```

load OpenMx

```
library(OpenMx)
```

set workingdirectory

```
setwd("YOUR_WORKING_DIRECTORY")
```

read the data

```
datb5=read.table('rdataf') # read the female data
```

assign variable names

```
varlabs=c('sex',  
  'n1', 'n2', 'n3', 'n4', 'n5', 'n6',  
  'e1', 'e2', 'e3', 'e4', 'e5', 'e6',  
  'o1', 'o2', 'o3', 'o4', 'o5', 'o6',  
  'a1', 'a2', 'a3', 'a4', 'a5', 'a6',  
  'c1', 'c2', 'c3', 'c4', 'c5', 'c6')  
colnames(datb5)=varlabs
```

select the variables of interest

```
isel=c(2:13) # selection of variables n1-n6, e1-e6  
datb2=datb5[,isel] # the data frame that we'll use below.
```

Part 2: summary statistics

```
Ss1=cov(datb2[,1:12]) # calculate the covariance matrix in females
print(round(Ss1,1))
```

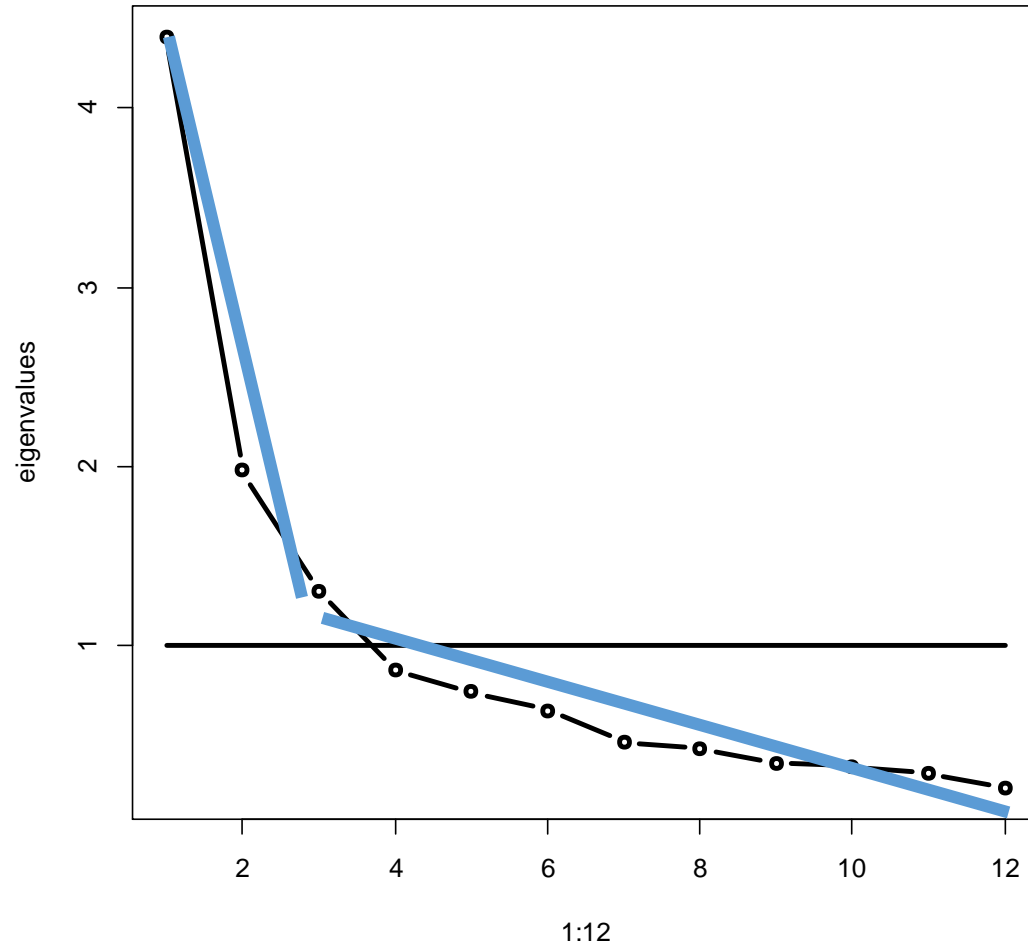
```
Rs1=cov2cor(Ss1) # convert to correlation matrix
print(round(Rs1,2))
```

```
Ms1=apply(datb2[,1:12],2,mean) # females means
print(round(Ms1,2))
```

End of part 2

```
> print(round(Ms1,2))
  n1    n2    n3    n4    n5    n6    e1    e2    e3    e4    e5    e6
23.22 20.38 23.35 23.99 27.62 20.00 31.23 28.81 23.36 25.43 27.78 31.13
> print(round(Rs1,2))
      n1    n2    n3    n4    n5    n6    e1    e2    e3    e4    e5    e6
n1  1.00  0.44  0.77  0.59  0.18  0.72 -0.25 -0.20 -0.29 -0.24 -0.14 -0.40
n2  0.44  1.00  0.45  0.30  0.27  0.44 -0.36 -0.22  0.07 -0.03  0.02 -0.33
n3  0.77  0.45  1.00  0.62  0.20  0.69 -0.28 -0.24 -0.36 -0.29 -0.11 -0.47
n4  0.59  0.30  0.62  1.00  0.15  0.59 -0.34 -0.24 -0.42 -0.19 -0.11 -0.39
n5  0.18  0.27  0.20  0.15  1.00  0.22  0.06  0.15  0.06  0.13  0.37  0.19
n6  0.72  0.44  0.69  0.59  0.22  1.00 -0.28 -0.14 -0.35 -0.27 -0.08 -0.43
e1 -0.25 -0.36 -0.28 -0.34  0.06 -0.28  1.00  0.46  0.13  0.28  0.15  0.59
e2 -0.20 -0.22 -0.24 -0.24  0.15 -0.14  0.46  1.00  0.14  0.19  0.37  0.38
e3 -0.29  0.07 -0.36 -0.42  0.06 -0.35  0.13  0.14  1.00  0.38  0.13  0.26
e4 -0.24 -0.03 -0.29 -0.19  0.13 -0.27  0.28  0.19  0.38  1.00  0.27  0.45
e5 -0.14  0.02 -0.11 -0.11  0.37 -0.08  0.15  0.37  0.13  0.27  1.00  0.30
e6 -0.40 -0.33 -0.47 -0.39  0.19 -0.43  0.59  0.38  0.26  0.45  0.30  1.00
```

SCREEPLOT

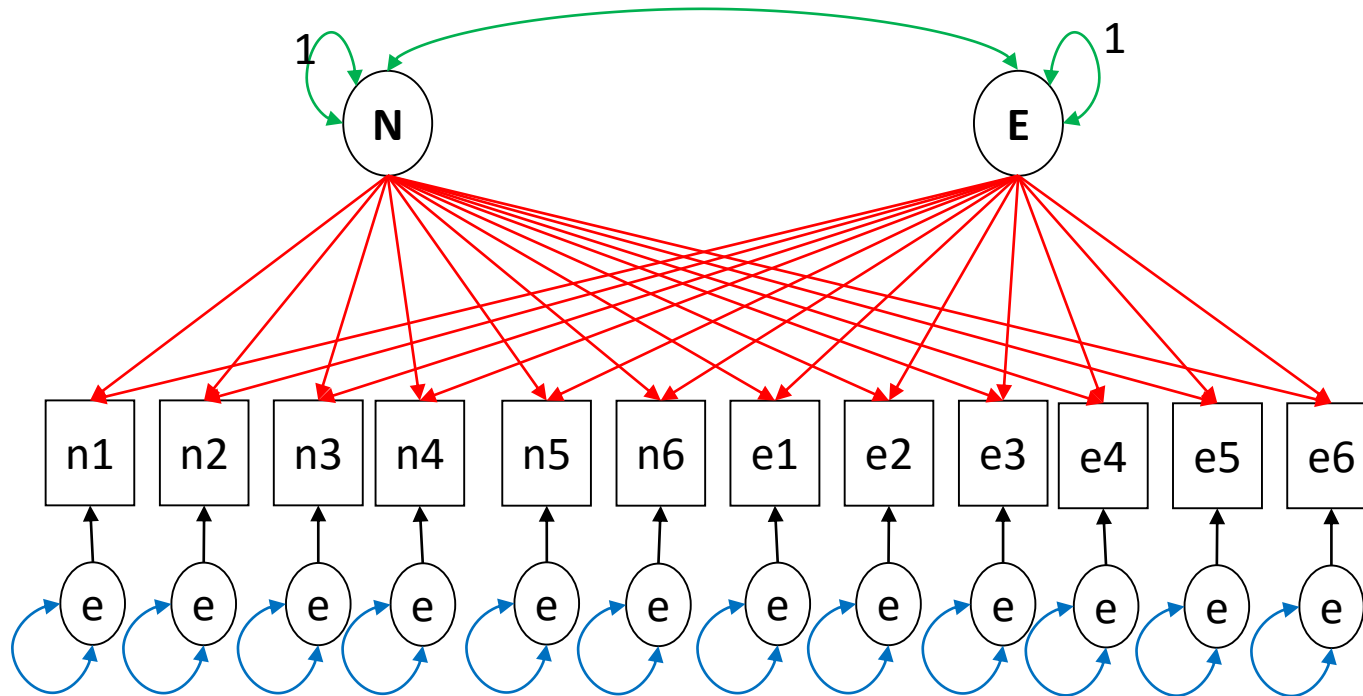


How many factors?

Ambiguous....

- Eigenvalues > 1 suggests 3 factors
- Elbow criterion suggests 2 factors

$$\Sigma_y = \Lambda \Psi \Lambda^t + \Theta$$



$$\Sigma_y = \Lambda \Psi \Lambda^t + \Theta$$

Matrix (12x2) of factor loadings Λ (note: |loading| < .1 not shown):

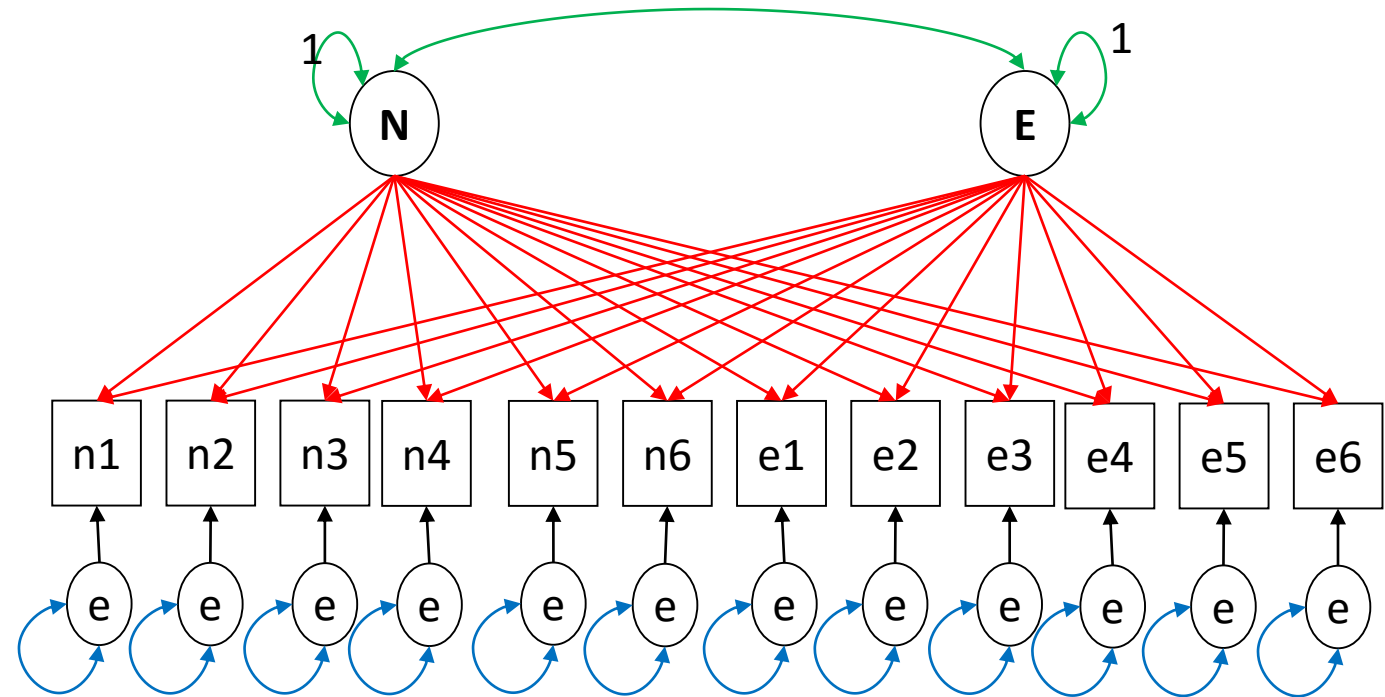
| | Factor1 | Factor2 |
|----|---------|---------|
| n1 | 0.851 | |
| n2 | 0.486 | |
| n3 | 0.838 | |
| n4 | 0.647 | -0.140 |
| n5 | 0.464 | 0.501 |
| n6 | 0.801 | |
| e1 | | 0.614 |
| e2 | | 0.537 |
| e3 | -0.294 | 0.199 |
| e4 | | 0.466 |
| e5 | 0.102 | 0.497 |
| e6 | -0.198 | 0.731 |

Factor Correlation matrix (2x2) Ψ :

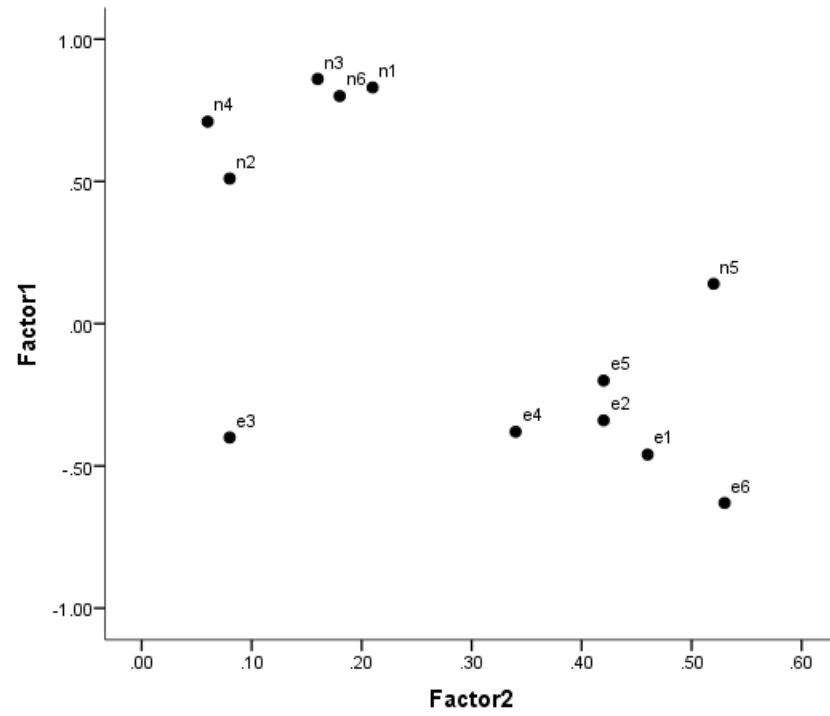
| | Factor1 | Factor2 |
|---------|---------|---------|
| Factor1 | 1.000 | -0.368 |
| Factor2 | -0.368 | 1.000 |

Diagonal covariance matrix (12x12) of residuals (Θ):

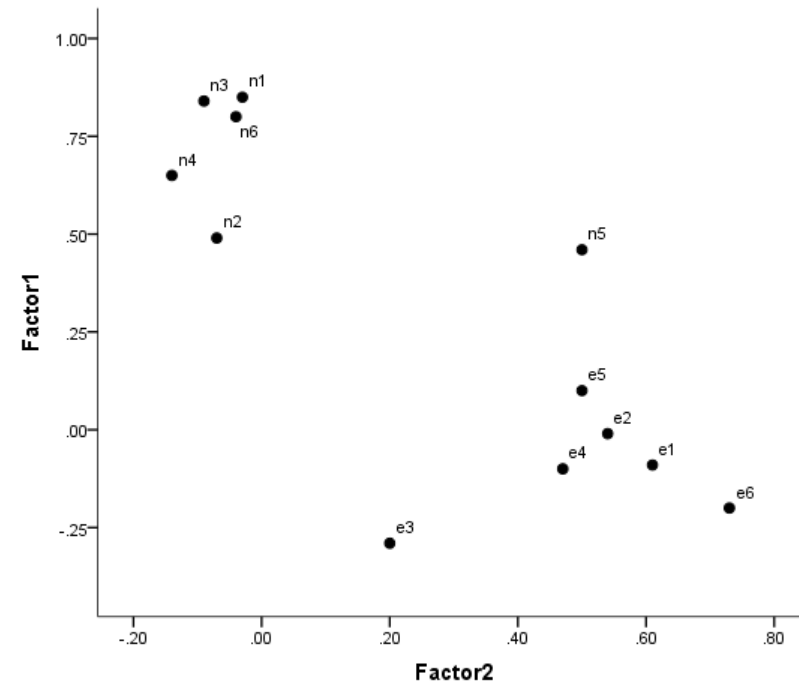
| | n1 | n2 | n3 | n4 | n5 | n6 | e1 | e2 | e3 | e4 | e5 | e6 |
|----|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|
| n1 | 0.259 | | | | | | | | | | | |
| n2 | | 0.735 | | | | | | | | | | |
| n3 | | | 0.237 | | | | | | | | | |
| n4 | | | | 0.496 | | | | | | | | |
| n5 | | | | | 0.705 | | | | | | | |
| n6 | | | | | | 0.330 | | | | | | |
| e1 | | | | | | | 0.574 | | | | | |
| e2 | | | | | | | | 0.707 | | | | |
| e3 | | | | | | | | | 0.831 | | | |
| e4 | | | | | | | | | | 0.738 | | |
| e5 | | | | | | | | | | | 0.780 | |
| e6 | | | | | | | | | | | | 0.321 |



unrotated



rotated



Goodness of fit of EFA 2 factor model.

Test of the hypothesis that 2 factors are sufficient.
The chi square statistic is 289.76 on 43 degrees of
freedom. The p-value is **2.52e-38**

By this statistical criterion the model is judged to
be acceptable if the p-value is greater than the
chosen alpha (e.g. alpha=.05).

By the statistical criterion, we'd **reject** the model!

Part 4A: saturated model

```
ny=12      # number of indicators
ne=2      # expected number of common factors
varnames=colnames(datb2) # var names
```

fit the saturated model ###**# define the means and covariance matrix in OpenMx to obtain the saturated model loglikelihood**

```
Rsl=mxMatrix(type='Stand',nrow=ny,ncol=ny,free=TRUE,value=.05,
             lbound=-.9,ubound=.9,name='cor') # 12x12 correlation matrix
Sds1=mxMatrix(type='Diag',nrow=ny,ncol=ny,free=TRUE,value=5,name='sds') # 12x12 diagonal matrix (st devs)
Mean1=mxMatrix(type='Full',nrow=1,ncol=ny,free=TRUE,value=25,name='mean1') # 1x12 vector means
MkS1=mxAlgebra(expression=sds%*%cor%*%sds,name='Ssat1') # expected covariance matrix

satmodels1=mxModel('part1',Rsl, Sds1, Mean1,MkS1) # assemble the model
```

data + estimation function

```
satdats1=mxModel("part2",
  mxData( observed=datb2, type="raw"), # the data
  mxExpectationNormal( covariance="part1.Ssat1", means="part1.mean1",
    dimnames=varnames), # the fit function
  mxFitFunctionML()
) # data & expected cov/means
```

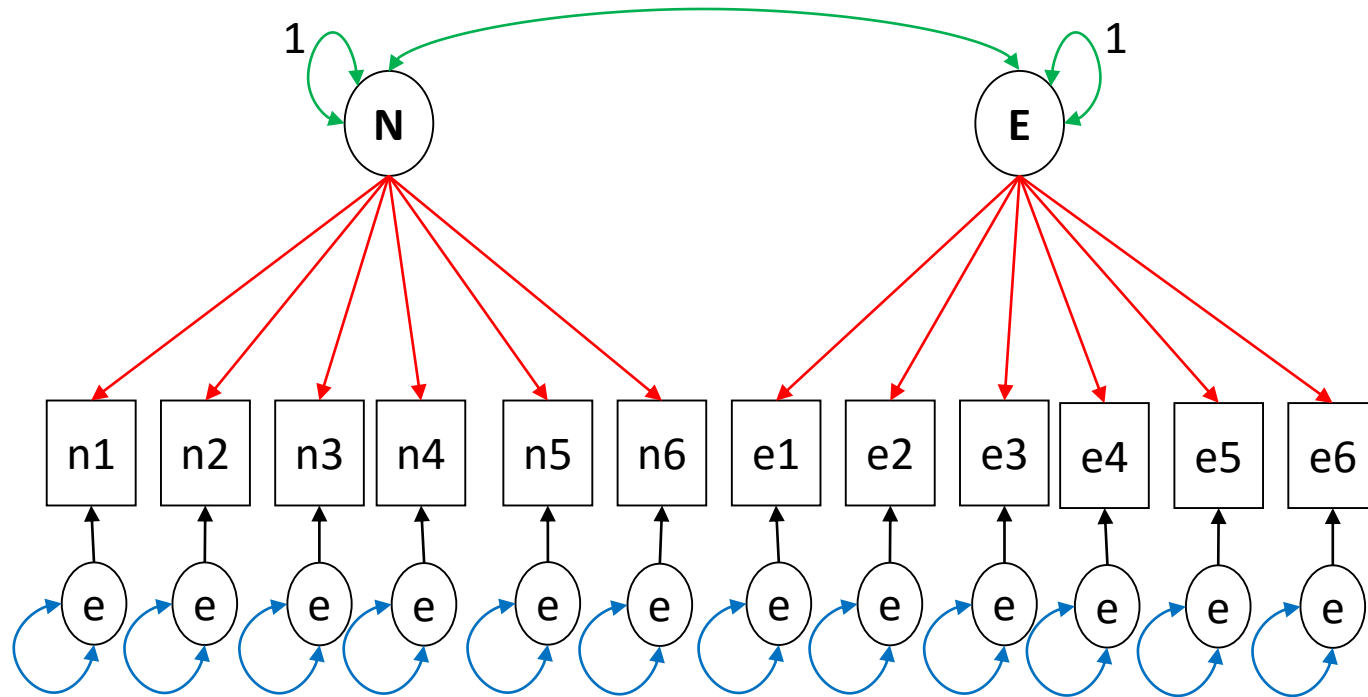
fit the saturated model....

```
Models1 <- mxModel("models1", satmodels1, satdats1,
  mxAlgebra(part2.objective, name="minus2loglikelihood"),
  mxFitFunctionAlgebra("minus2loglikelihood"))
Models1_out <- mxRun(Models1)
```

```
> summary(Models1_out)
```

```
observed statistics: 4332  
estimated parameters: 90  
degrees of freedom: 4242  
-2 log likelihood: 23578.09  
number of observations: 361
```

$$\Sigma_y = \Lambda \Psi \Lambda^t + \Theta$$



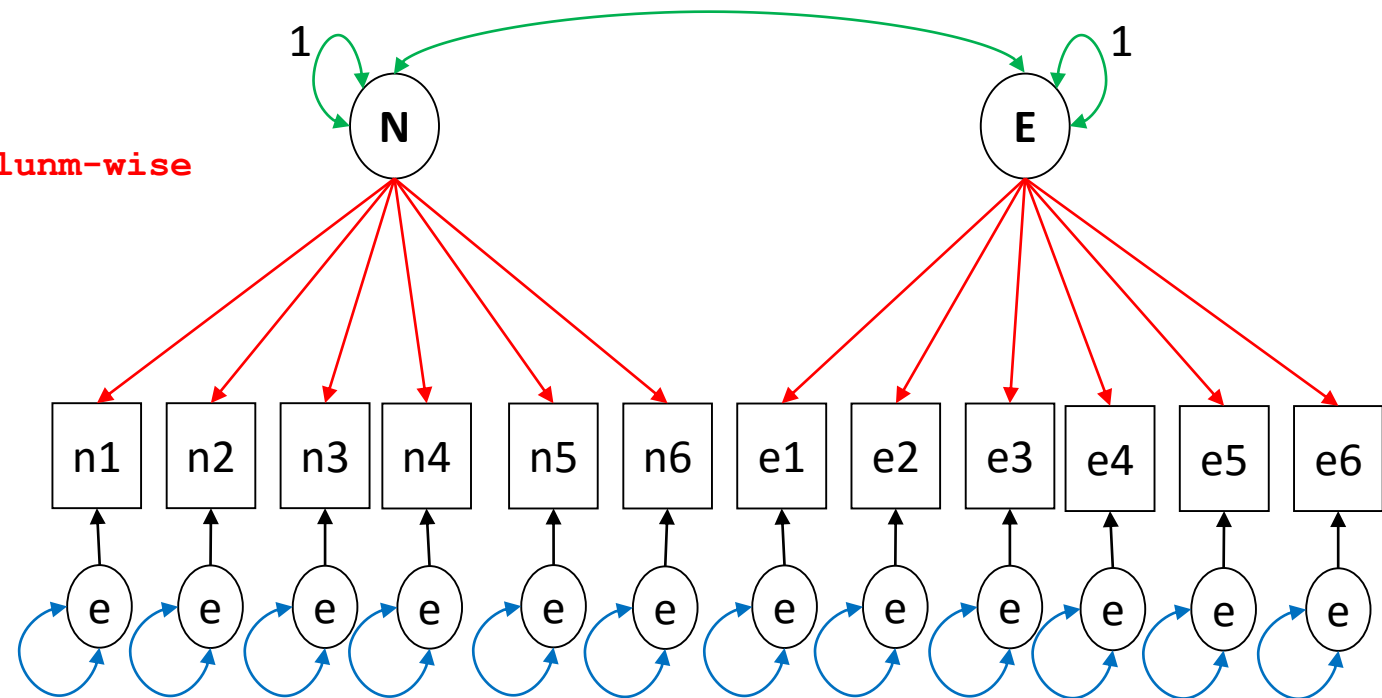
```

Ly=mxMatrix(type='Full',nrow=ny,ncol=ne,
  free=matrix(c(
    T,F,
    T,F,
    T,F,
    T,F,
    T,F,
    T,F,
    F,T,
    F,T,
    F,T,
    F,T,
    F,T,
    F,T),ny,ne,byrow=T),
  values=c(4,4,4,4,4,4,0,0,0,0,0,0,
           0,0,0,0,0,0,4,4,4,4,4,4), # read column-wise
  labels=matrix(c(
    'f1_1','f1_2',
    'f2_1','f2_2',
    'f3_1','f3_2',
    'f4_1','f4_2',
    'f5_1','f5_2',
    'f6_1','f6_2',
    'f7_1','f7_2',
    'f8_1','f8_2',
    'f9_1','f9_2',
    'f10_1','f10_2',
    'f11_1','f11_2',
    'f12_1','f12_2'),ny,ne,byrow=T),name='Ly'),

```

Define factor loading matrix Λ

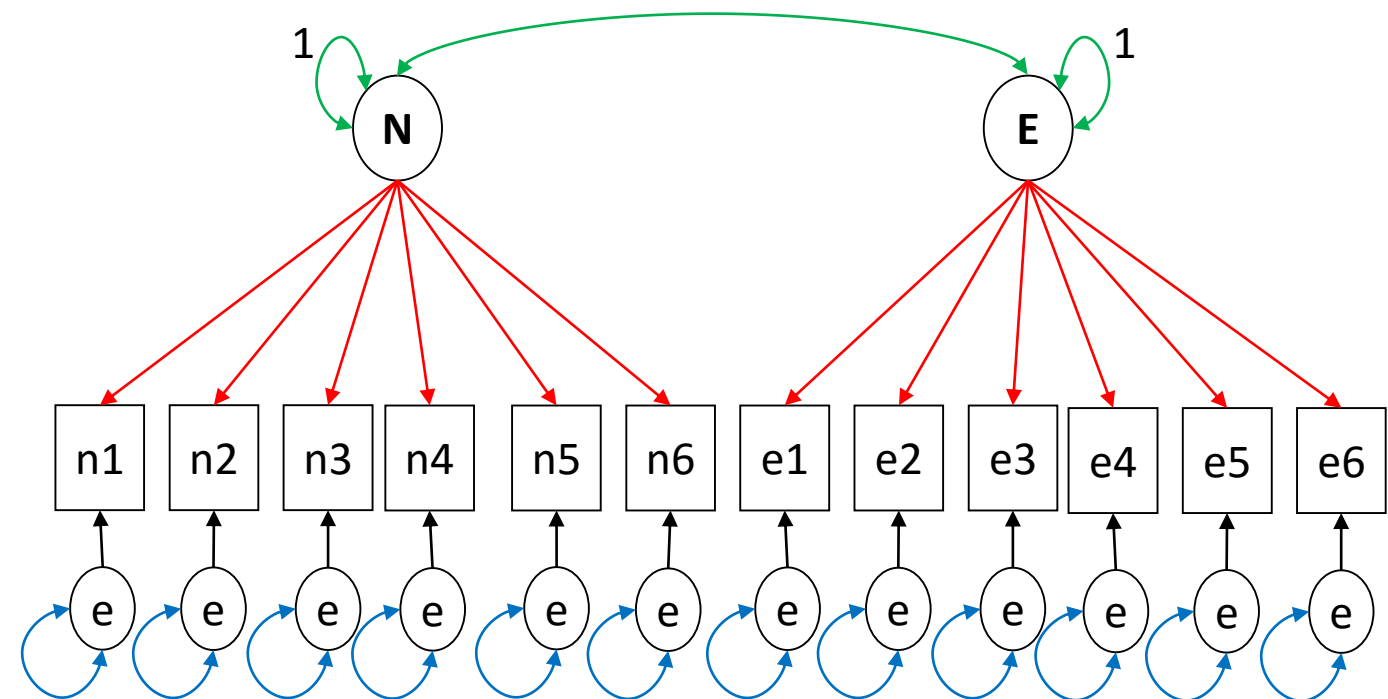
$$\Sigma_{\mathbf{y}} = \Lambda \Psi \Lambda^t + \Theta$$



Define covariance matrix of residuals 

```
Te=mxMatrix(type='Diag',nrow=ny,ncol=ny,
  labels=c('rn1','rn2','rn3','rn4','rn5','rn6',
    're1','re2','re3','re4','re5','re6'),
  free=TRUE,value=10,name='Te')
```

$$\Sigma_{\mathbf{y}} = \Lambda \Psi \Lambda^t + \Theta$$



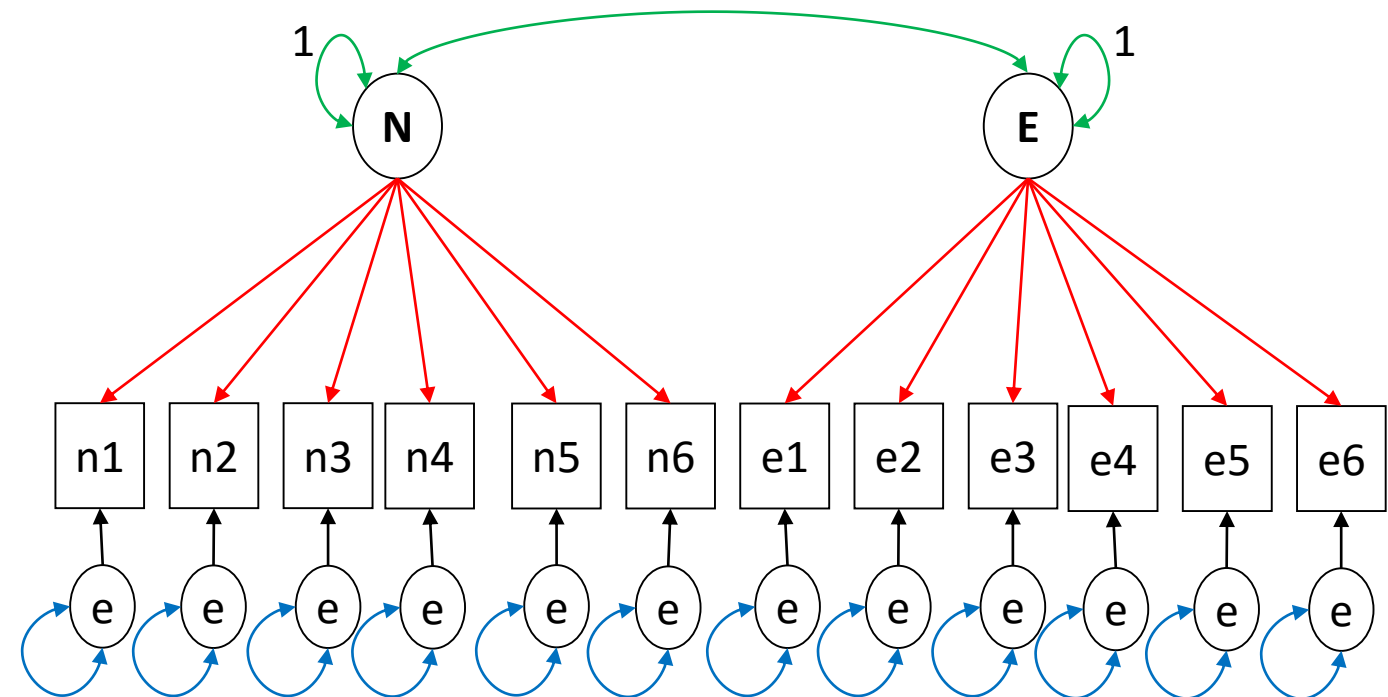
```
## latent correlation matrix
Ps=mxMatrix(type='Symm',nrow=ne,ncol=ne,
            free=c(FALSE,TRUE,FALSE),
            labels=c('v1_0','r12_0','v2_0'),
            values=c(1,.0,1),name='Ps')
```

Define covariance matrix of factors Ψ

$$\Sigma_y = \Lambda \Psi \Lambda^t + \Theta$$

NOTE: scaling of the common factors by fixing the variances to equal 1.

Ψ is a correlation matrix!



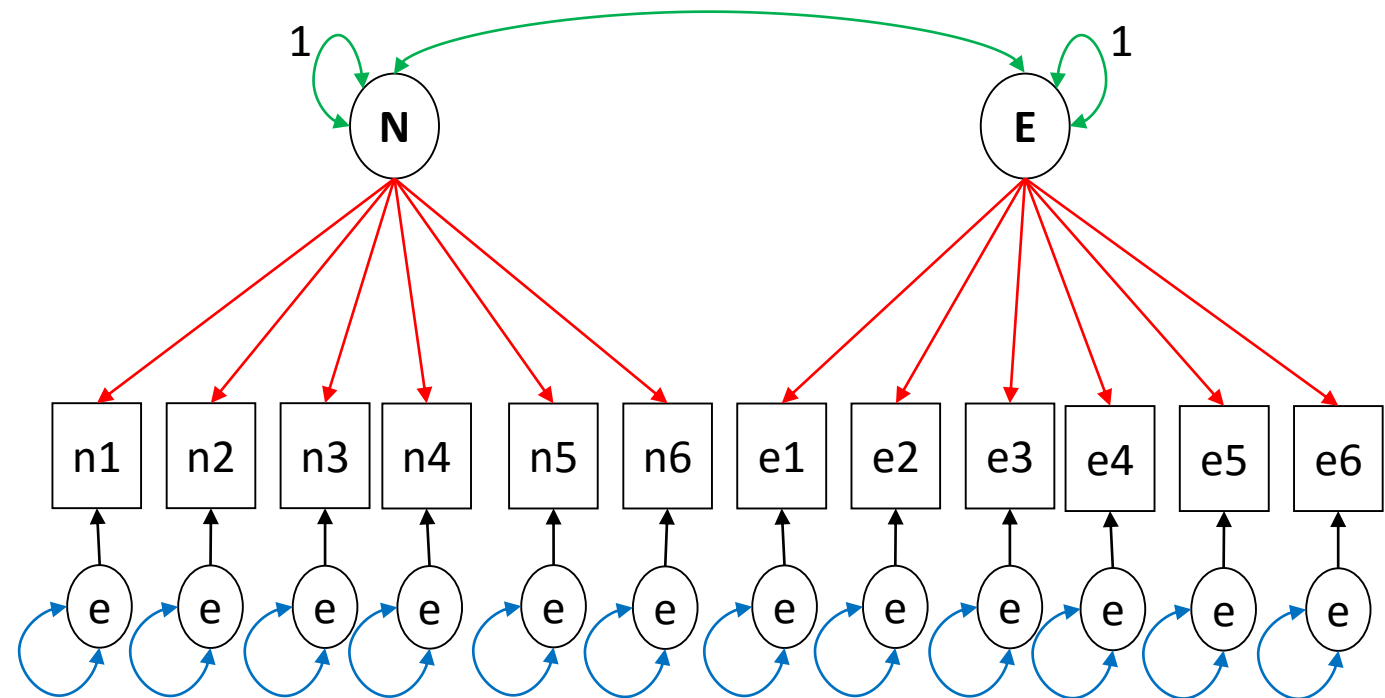
```
# means
Tau=mxMatrix(type='Full',nrow=1,ncol=ny,free=TRUE,value=25,
             labels=c('mn1','mn2','mn3','mn4','mn5','mn6',
                    'me1','me2','me3','me4','me5','me6'),
             name='Means')
```

$$\Sigma_y = \Lambda \Psi \Lambda^t + \Theta$$

```
MKS=mxAlgebra(expression=Ly %*% (Ps) %*%t(Ly) + Te %*%t(Te), name='Sigma'),
MKM=mxAlgebra(expression=Means, name='means')
```

.... assemble the model and run

```
CFM1_out = mxRun(cfamodel1)
```



```
> mxCompare (Models1_out, CFM1_out)
```

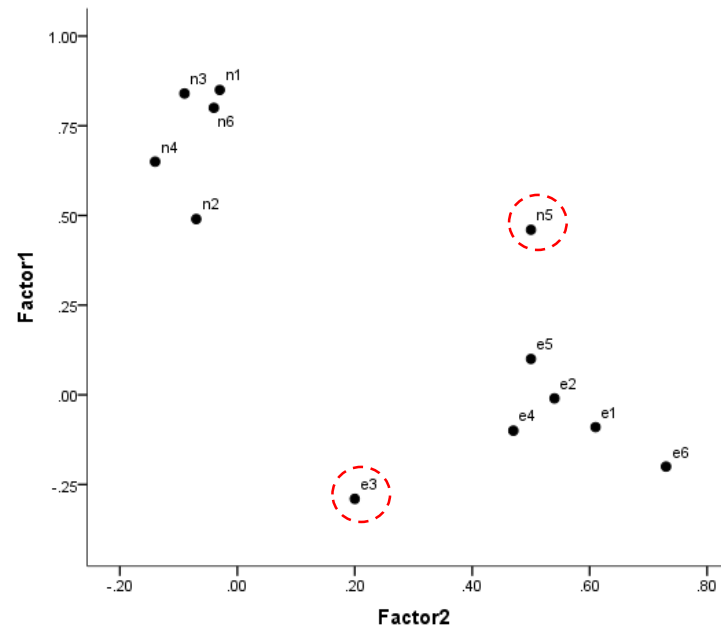
| | base | comparison | ep | minus2LL | df | AIC | diffLL | diffdf | p |
|---|---------|------------|----|----------|------|----------|----------|--------|--------------|
| 1 | models1 | <NA> | 90 | 23578.09 | 4242 | 15094.09 | NA | NA | NA |
| 2 | models1 | CFM1 | 37 | 23975.63 | 4295 | 15385.63 | 397.5483 | 53 | 2.785297e-54 |

This model does not fit (but we already know this from the EFA results).

```
> print(round(fals1pr$loadings[,1:2],4))
```

| | Factor1 | Factor2 |
|----|----------------|---------------|
| n1 | 0.8515 | -0.0252 |
| n2 | 0.4858 | -0.0682 |
| n3 | 0.8377 | -0.0870 |
| n4 | 0.6465 | -0.1404 |
| n5 | 0.4638 | <u>0.5005</u> |
| n6 | 0.8014 | -0.0444 |
| e1 | -0.0907 | 0.6142 |
| e2 | -0.0126 | 0.5367 |
| e3 | <u>-0.2943</u> | 0.1988 |
| e4 | -0.0996 | 0.4664 |
| e5 | 0.1024 | 0.4973 |
| e6 | -0.1978 | 0.7308 |

Λ



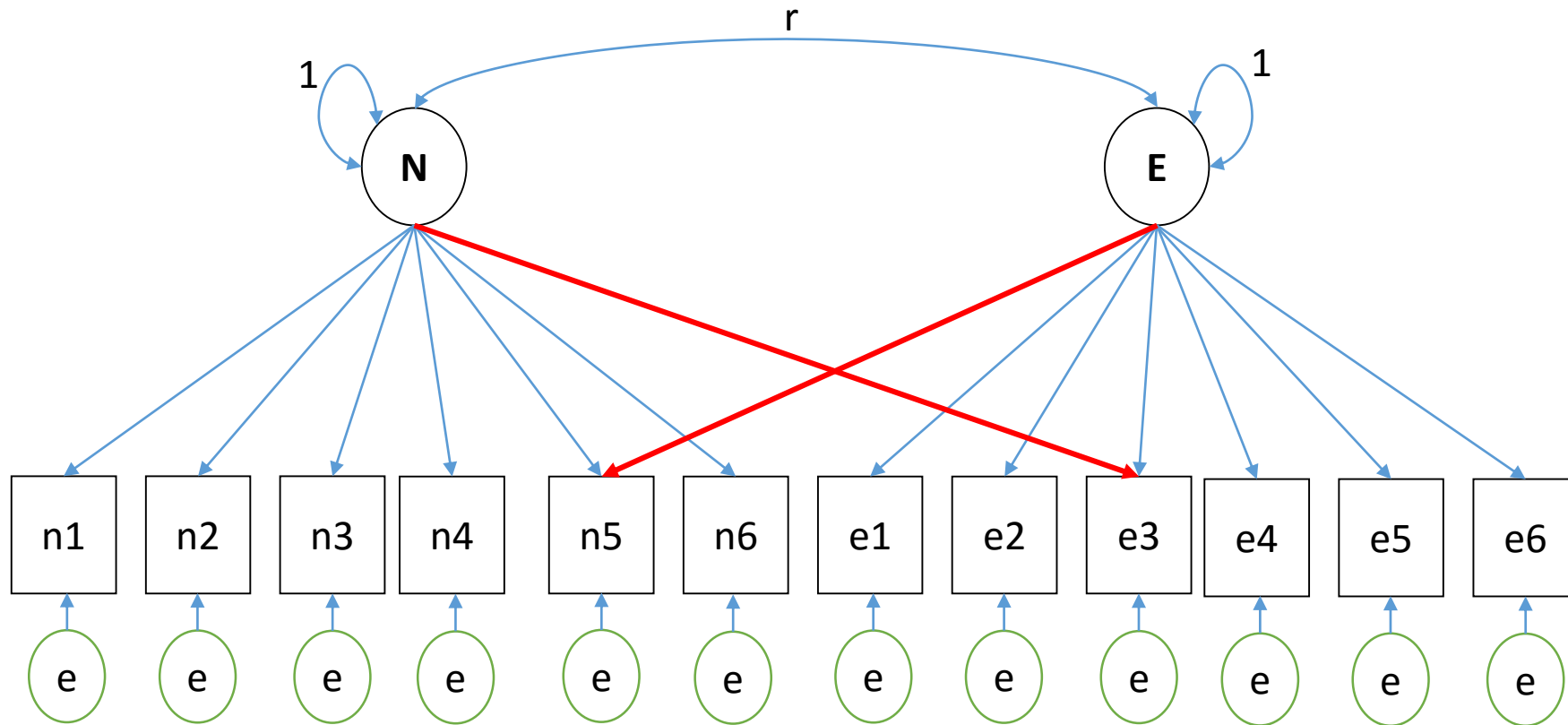
```
> round(est_Ly,3)
```

| | [,1] | [,2] |
|-------|--------------|--------------|
| [1,] | 4.923 | 0.000 |
| [2,] | 2.383 | 0.000 |
| [3,] | 5.197 | 0.000 |
| [4,] | 2.993 | 0.000 |
| [5,] | 0.889 | <u>0.000</u> |
| [6,] | 3.785 | 0.000 |
| [7,] | 0.000 | 2.565 |
| [8,] | 0.000 | 2.225 |
| [9,] | <u>0.000</u> | 1.822 |
| [10,] | 0.000 | 1.873 |
| [11,] | 0.000 | 1.576 |
| [12,] | 0.000 | 3.691 |

Λ

To do: free the cross loadings Ly[5,2] and Ly[9,1]

To do: free the cross loadings $\text{Ly}[5,2]$ and $\text{Ly}[9,1]$



Test whether the cross loadings are unequal zero using the likelihood ratio test

| base | comparison | ep | minus2LL | df | AIC | diffLL | diffdf | p |
|------|------------|------|----------|----------|------|----------|----------|----------------|
| 1 | CFM1 | <NA> | 39 | 23891.96 | 4293 | 15305.96 | NA | NA |
| 2 | CFM1 | CFM1 | 37 | 23975.63 | 4295 | 15385.63 | 83.67033 | 2 6.779838e-19 |

Given $\alpha=.05$, we reject the hypothesis $\text{Ly}[5,2] = \text{Ly}[9,1] = 0$
 Therefore either one or both are not equal to zero.

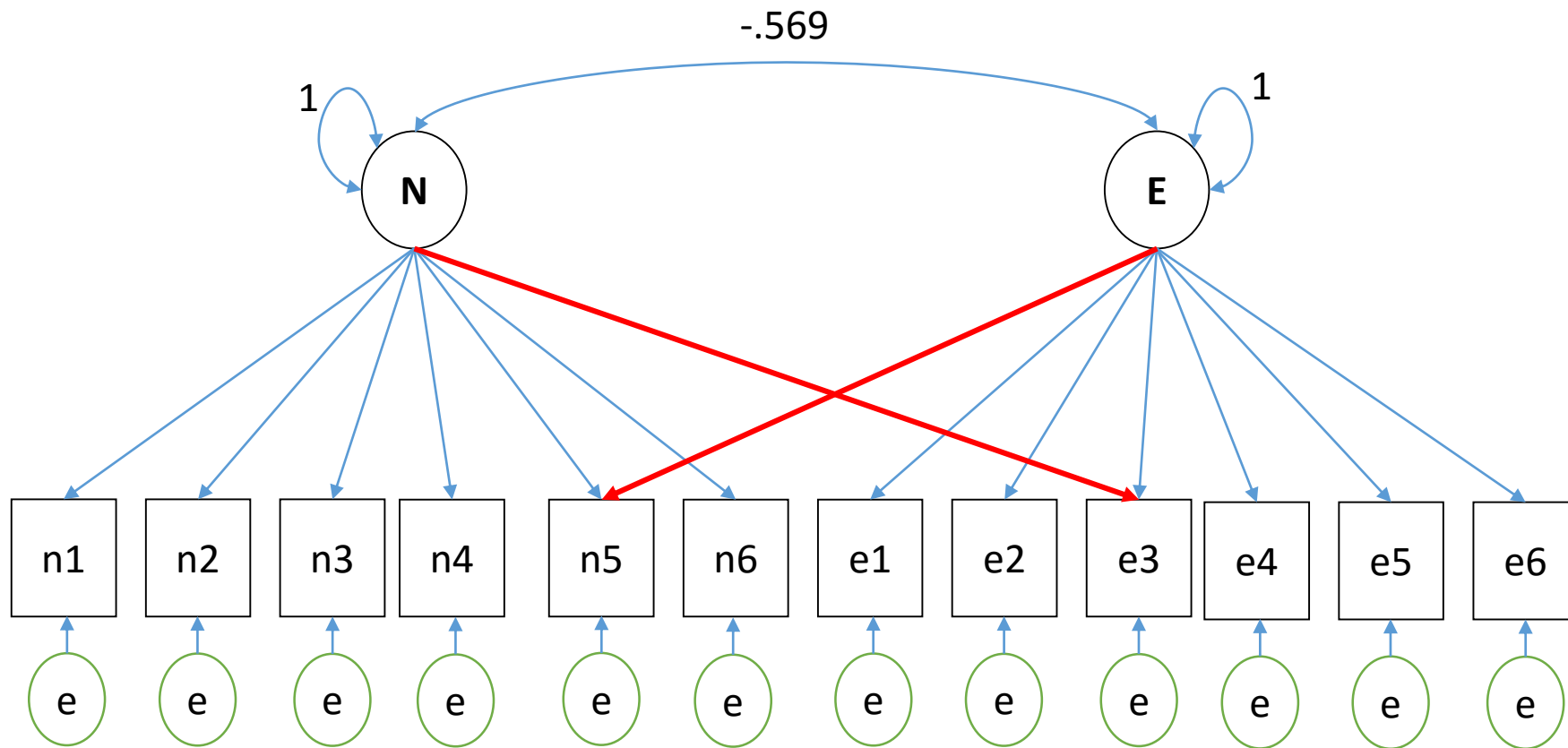
```
print(round(est_Ly,2))
```

```

      [,1] [,2]
[1,]  4.88 0.00
[2,]  2.38 0.00
[3,]  5.20 0.00
[4,]  3.02 0.00
[5,]  2.30 2.32
[6,]  3.80 0.00
[7,]  0.00 2.50
[8,]  0.00 2.23
[9,] -1.47 0.89
[10,] 0.00 1.85
[11,] 0.00 1.74
[12,] 0.00 3.73

```

Inspect the correlation matrix Ψ

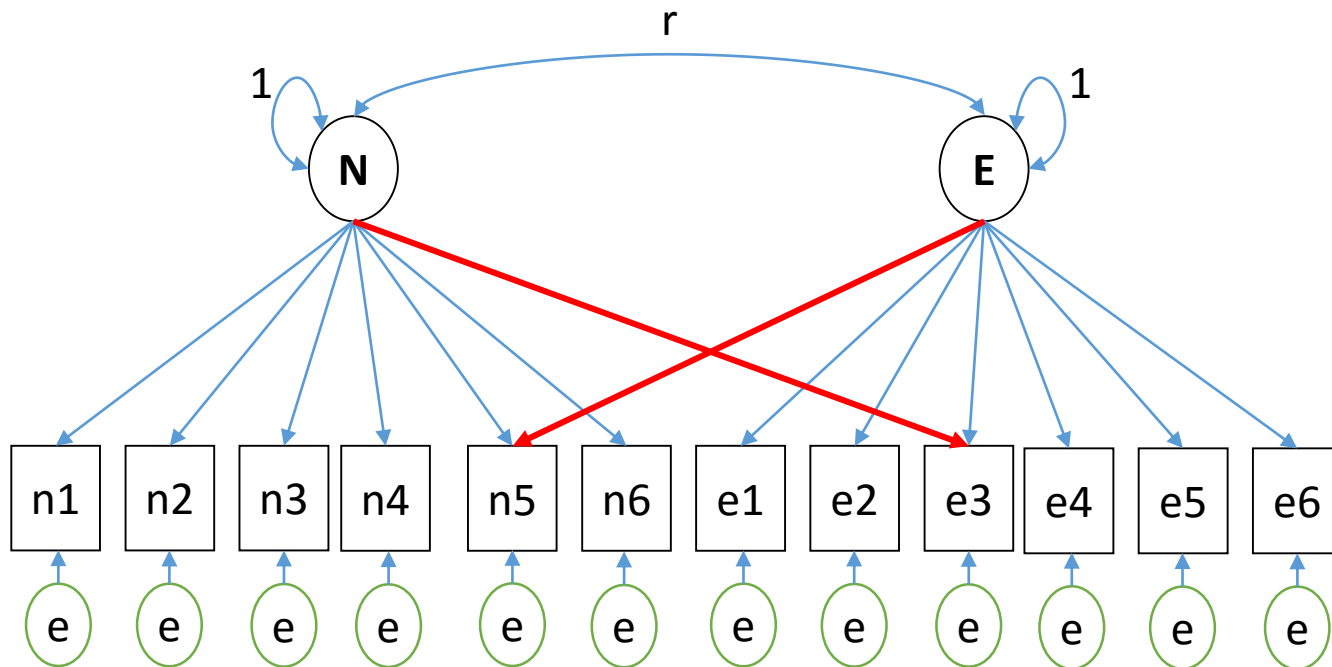


```
1.000 -0.569
-0.569 1.000
```

Reliability of the indicators

```
> round(diag(Sfit1)/diag(Sfit),2)
[1] 0.74 0.26 0.77 0.50 0.28 0.66 0.42 0.25 0.17 0.26 0.16 0.74
```

```
est_Ly=mxEval(CFAModel1.Ly,CFM2_out)
est_Ps=mxEval(CFAModel1.Ps,CFM2_out)
est_Te=mxEval(CFAModel1.Te,CFM2_out)
est_Te=est_Te^2
Sfit1=est_Ly%*%est_Ps%*%t(est_Ly)
Sfit=Sfit1+est_Te
rel=diag(Sfit1)/diag(Sfit)
print(round(rel,3))
```



Variance of n1 due to N divided by the total variance of n1:

$$23.77 / 32.24 = .74.$$

The common factor N explains 74% of the variance in item 1.