

# An introduction to Psychometric Theory with applications in R

## Structural Equation Modeling and applied scale construction

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## Outline

### Overview

- Measurement models

### Path algebra

- Wright's rules

### Observed-Observed

- As classic regression

### SEM

- With fixed variables

- latent variables

### Model comparison

- Causality?

lavaan as an alternative model package to sem

### Types of variables

### Confirmatory Factor Analysis

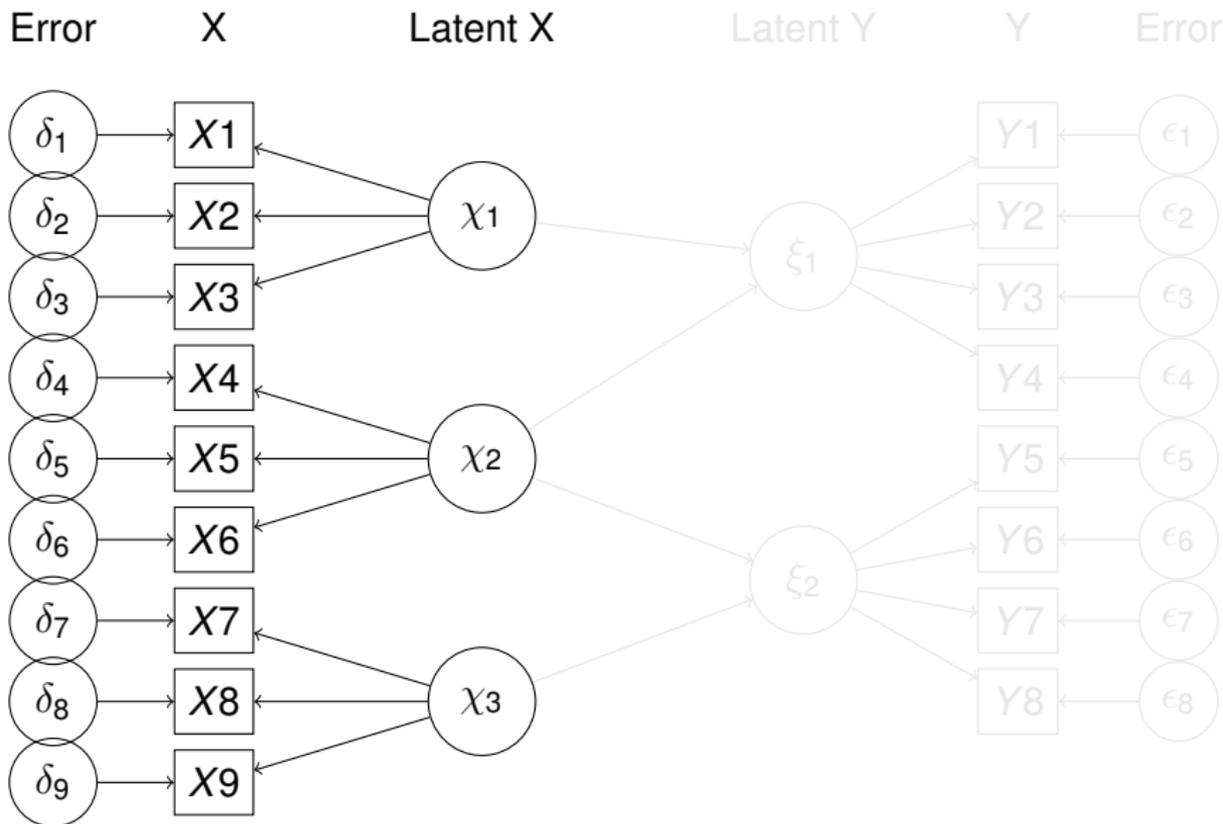
- Lavaan for CFA

### Measuring change

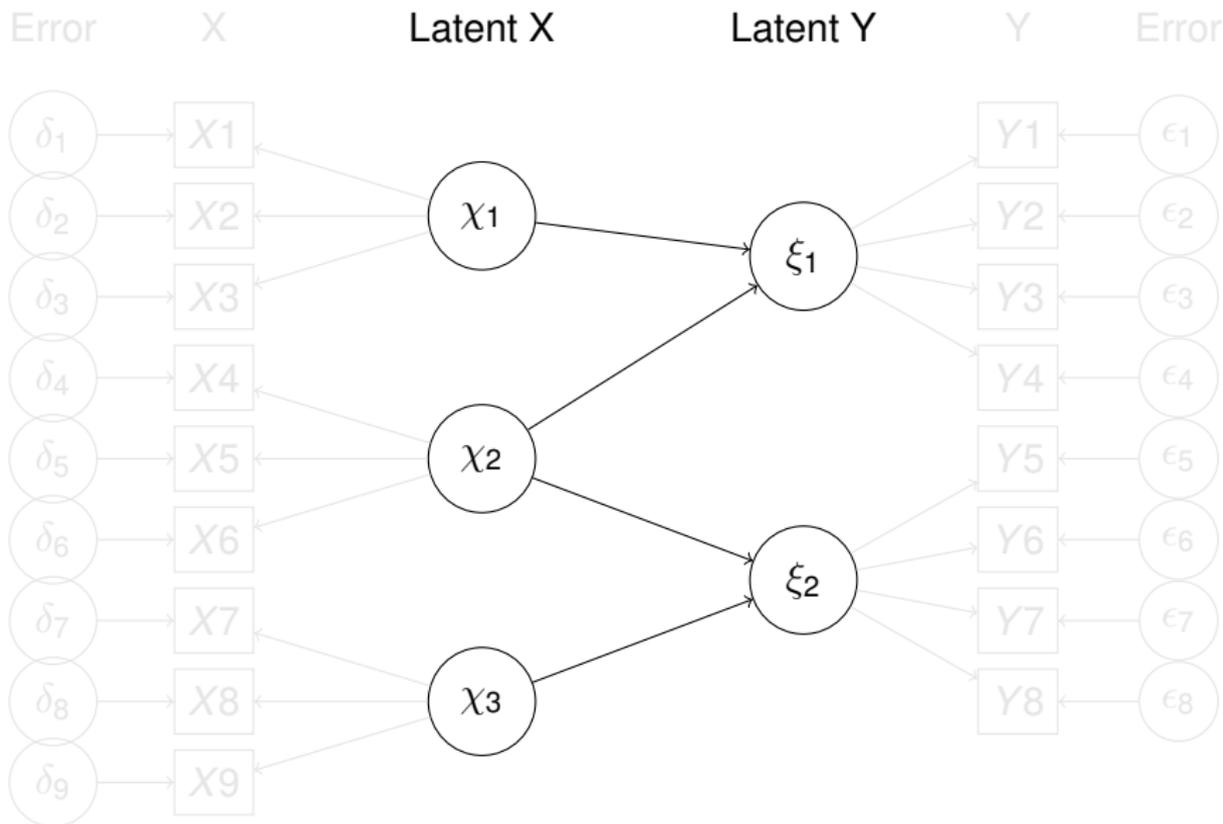
- create the data

Two time points invariant

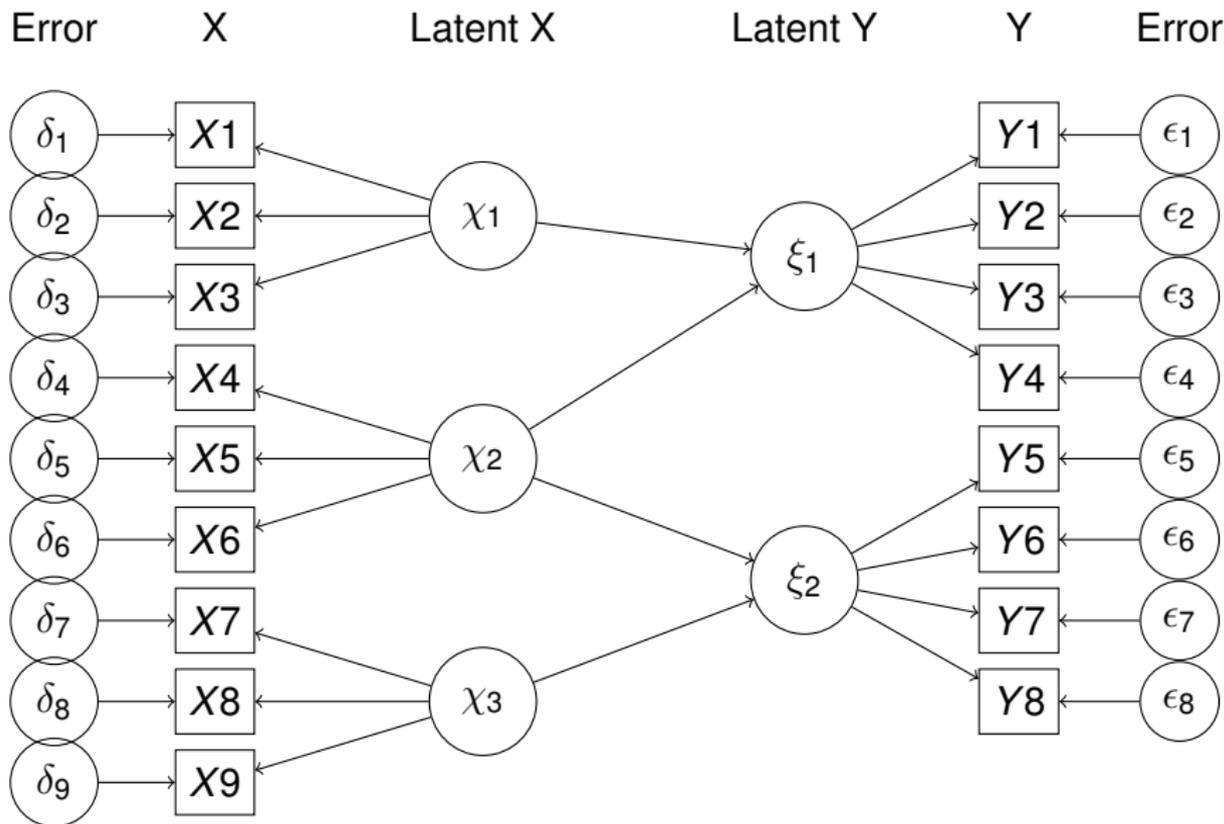
## Measurement: A latent variable approach.



## Theory



## Psychometric Theory: A conceptual Syllabus



## Two types of variables, three types of relationships

### 1. Variables

1.1 Observed Variables ( $X, Y$ )

1.2 Latent Variables ( $\xi, \eta, \zeta$ )

### 2. Three kinds of variance/covariances

2.1 Observed with Observed  $C_{xy}$  or  $\sigma_{xy}$

2.2 Observed with Latent  $\lambda$

2.3 Latent with Latent  $\phi$

### 3. Direction

- Bidirectional (correlation)
- Directional (regression)

## Observed Variables

X

$X_1$

$X_2$

$X_3$

$X_4$

$X_5$

$X_6$

Y

$Y_1$

$Y_2$

$Y_3$

$Y_4$

$Y_5$

$Y_6$

## Latent Variables

$\xi$

$\eta$

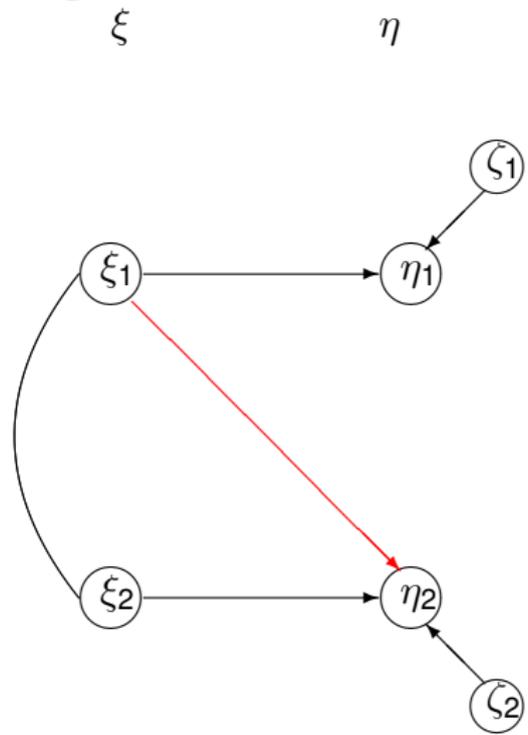
$\xi_1$

$\eta_1$

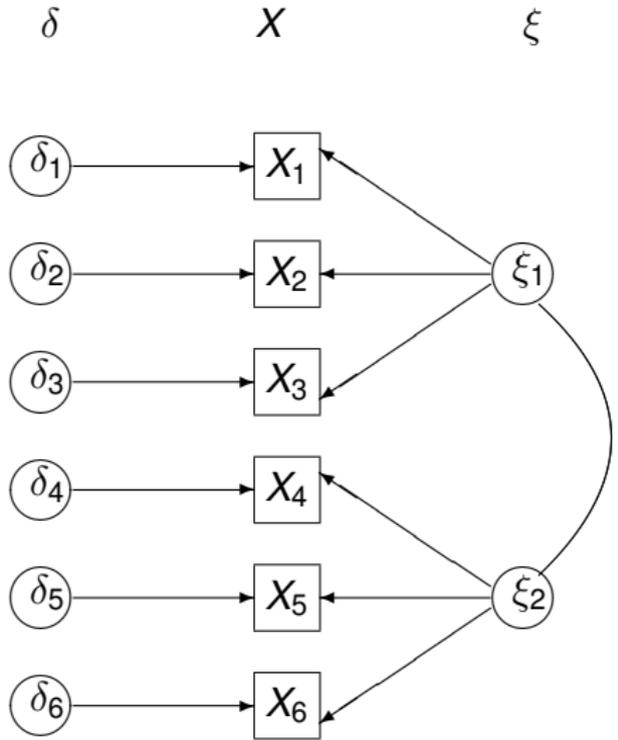
$\xi_2$

$\eta_2$

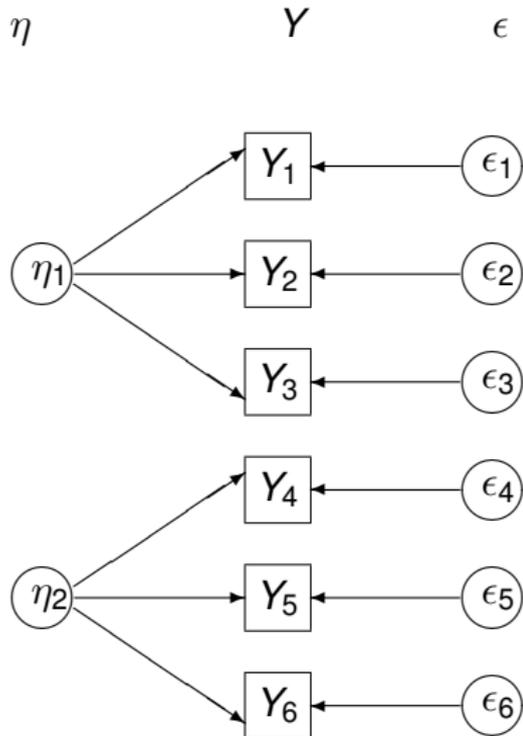
# Theory: A regression model of latent variables



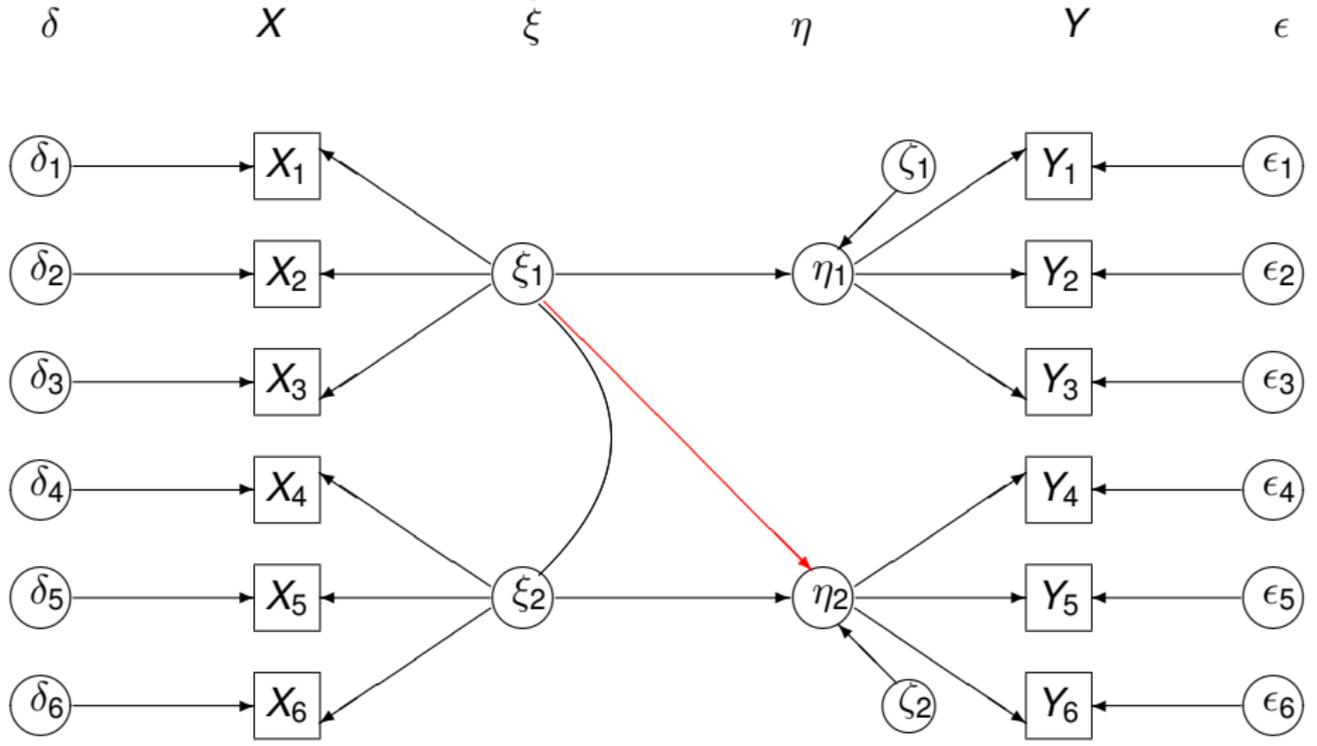
## A measurement model for X



## A measurement model for Y



## A complete structural model

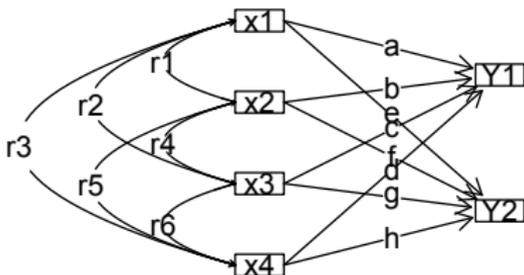


## Latent Variable Modeling

1. Requires measuring observed variables
  - Requires defining what is relevant and irrelevant to our theory.
  - Issues in quality of scale information, levels of measurement.
2. Formulating a measurement model of the data: estimating latent constructs
  - Perhaps based upon exploratory and then confirmatory factor analysis, definitely based upon theory.
  - Includes understanding the reliability of the measures.
3. Modeling the structure of the constructs
  - This is a combination of theory and fitting. Do the data fit the theory.
  - Comparison of models. Does one model fit better than alternative models?

# The generalized regression model

## Generalized regression model



$$\hat{Y} = \beta_x X + \epsilon$$

## The Kerchoff data set of educational outcomes

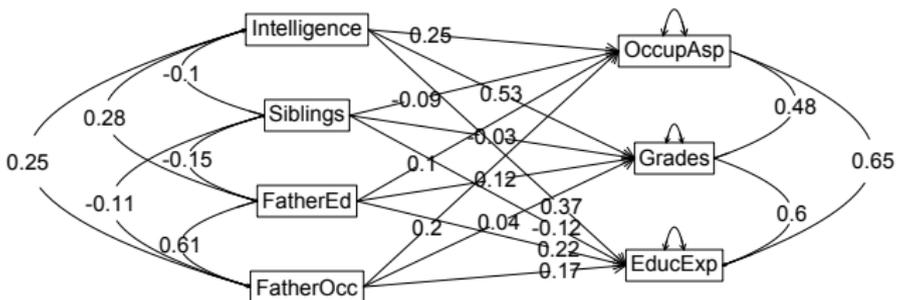
lowerMat (R.kerch)

	Int11	Sblng	FthrE	FthrO	Grads	EdcEx	OccpA
<b>Intelligence</b>	1.00						
<b>Siblings</b>	-0.10	1.00					
<b>FatherEd</b>	0.28	-0.15	1.00				
<b>FatherOcc</b>	0.25	-0.11	0.61	1.00			
<b>Grades</b>	0.57	-0.10	0.29	0.25	1.00		
<b>EducExp</b>	0.49	-0.21	0.45	0.41	0.60	1.00	
<b>OccupAsp</b>	0.34	-0.15	0.30	0.33	0.48	0.65	1.00

[Kerckhoff \(1974\)](#) (From LISREL manual, available in help page to mediate)

## The classic regression model

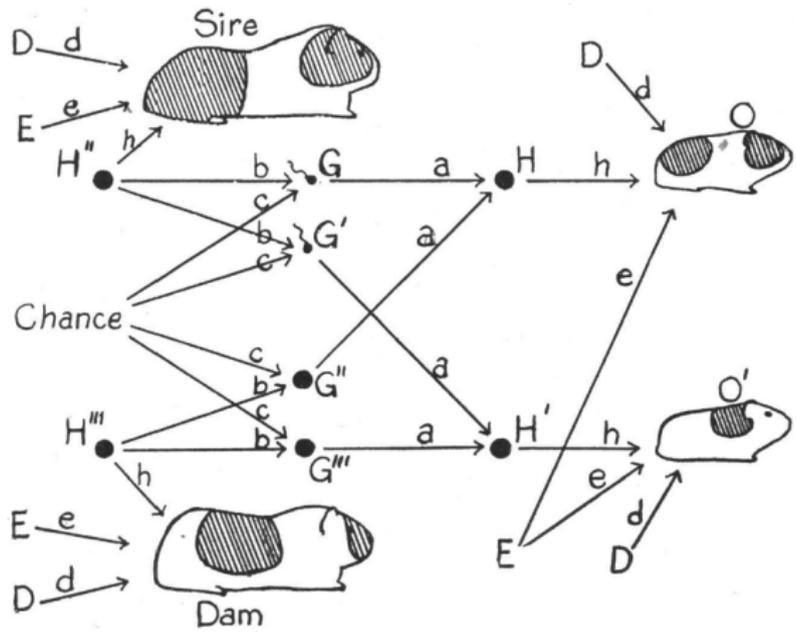
### Regression Models



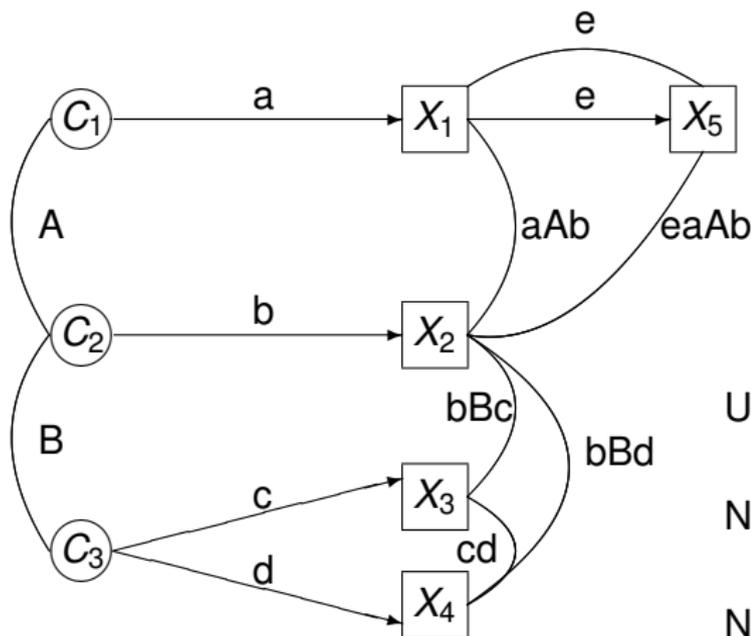
unweighted matrix correlation = 0.58

$$\hat{Y} = \beta_x X + \epsilon$$

## Wright's Path model of inheritance in the Guinea Pig (Wright, 1921)



## The basic rules of path analysis—think genetics



Up ... and over and down ...

No down and up

No double overs

Up ... and down ...

Parents cause children  
children do not cause parents

## Observed-observed

1. The [Kerckhoff \(1974\)](#) data set is used in the LISREL manual as an example of regression path models
  - “A sample of boys originally studied as ninth graders in 1969 was recontacted in 1974 to obtain information about high school performance and educational attainment.”
  - 767 twelfth grade males
  - A study of background, aspiration and educational attainment
2. Variables
  - Intelligence
  - Number of siblings
  - Fathers Education
  - Fathers’s occupation
  - Grades
  - Educational expectation
  - Occupational aspiration
3. Correlation matrix is available in the sem ([Fox, Nie & Byrnes, 2013](#)) package and the help page to mediate

## The Kerckhoff correlation matrix (N=767)

```
> R.kerch
  Intelligence Siblings FatherEd FatherOcc Grades EducExp OccupAsp
Intelligence    1.000    -0.100    0.277    0.250    0.572    0.489    0.335
Siblings        -0.100     1.000   -0.152   -0.108  -0.105  -0.213  -0.153
FatherEd         0.277    -0.152     1.000     0.611    0.294    0.446    0.303
FatherOcc        0.250    -0.108     0.611     1.000    0.248    0.410    0.331
Grades           0.572    -0.105     0.294     0.248     1.000    0.597    0.478
EducExp          0.489    -0.213     0.446     0.410    0.597     1.000    0.651
OccupAsp         0.335    -0.153     0.303     0.331    0.478    0.651     1.000
>
```

## Conceptual Kerckhoff model

### 1. Background variables

- Intelligence
- Number of siblings
- Fathers Education
- Fathers's occupation

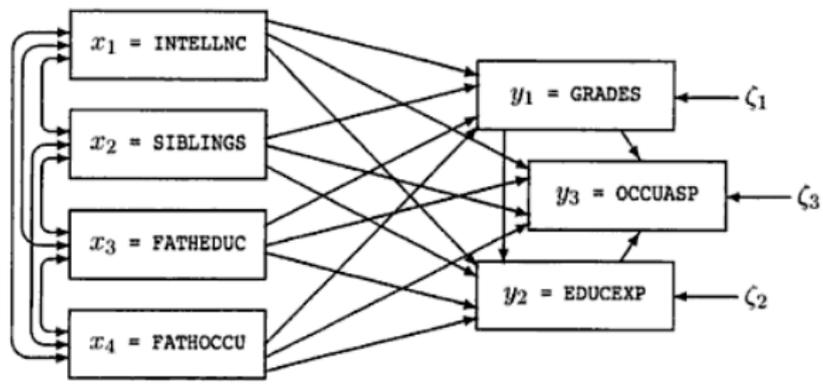
### 2. Intermediate variables

- Grades
- Educational expectation

### 3. Final outcomes

- Occupational aspiration

# The model (From the LISREL manual)



## Matrix regression

### 1. Most regression examples (and functions) use raw data

- $\hat{Y} = X\beta + \epsilon$
- $beta = (X'X)^{-1}X'Y$
- `lm(y ~ x , data = data)`

### 2. Regression is just solving the matrix equation

- $\beta = R^{-1}r_{xy}$
- `mat.regress(R,x,y)` (deprecated)
- `lmCor(y ~ x, data =R)` (recommended)

### 3. `setCor` will work on raw data as well.

## Simple regression 4 predictors, 3 criteria

### R code

```
model <- lmCor(Grades + EducExp + OccupAsp ~ Intelligence + Siblings +
              FatherEd + FatherOcc, data=R.kerch)
summary(model) #to give concise output
```

Multiple Regression from matrix input

cx

Beta weights

	Grades	EducExp	OccupAsp
Intelligence	0.53	0.37	0.25
Siblings	-0.03	-0.12	-0.09
FatherEd	0.12	0.22	0.10
FatherOcc	0.04	0.17	0.20

Multiple R

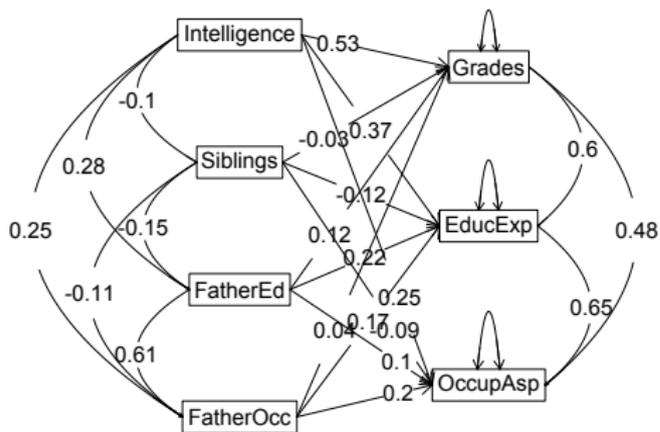
Grades	EducExp	OccupAsp
0.59	0.61	0.44

multiple R2

Grades	EducExp	OccupAsp
0.35	0.38	0.19

## The paths (From ImCor)

### Regression Models



unweighted matrix correlation = 0.58

## More complicated regression

```
> lmCor(y=6:7, x=1:5, data=R.kerch)
#easier for long names
```

**Call:** lmtCor(y = 6:7, x = 1:5, data = R.kerch)

Multiple Regression from **matrix** input

### Beta weights

	EducExp	OccupAsp
Intelligence	0.16	0.05
Siblings	-0.11	-0.08
FatherEd	0.17	0.05
FatherOcc	0.15	0.18
Grades	0.41	0.38

### Multiple R

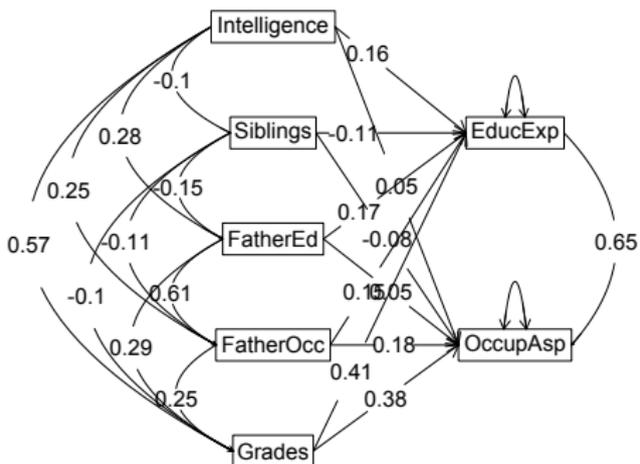
EducExp	OccupAsp
0.70	0.54

### multiple R2

EducExp	OccupAsp
0.48	0.29

## The paths (From setCor)

### Regression Models



unweighted matrix correlation = 0.64

## Predicting occupational aspirations from the intermediate set

```
lmCor( OccupAsp ~ Grades + EducExp , data=R.kerch)
```

**Call:** lmtCor(y = OccupAsp ~ Grades + EducExp, data = R.kerch)

Multiple Regression from **matrix** input

```
DV = OccupAsp
      slope VIF
Grades 0.14 1.55
EducExp 0.57 1.55
```

```
Multiple Regression
      R R2 Ruw R2uw
OccupAsp 0.66 0.44 0.63 0.4
```

## Predicting occupational aspirations from the entire set

```
> lmCor(y=7,x=1:6,data=R.kerch)
```

**Call:** `lmCor(y = 7, x = 1:6, data = R.kerch)`

Multiple Regression from **matrix** input

DV = OccupAsp

	slope	VIF
Intelligence	-0.04	1.57
Siblings	-0.02	1.05
FatherEd	-0.04	1.73
FatherOcc	0.10	1.66
Grades	0.16	1.85
EducExp	0.55	1.94

Multiple Regression

	R	R2	Ruw	R2uw
OccupAsp	0.67	0.44	0.57	0.32

## We can also examine mediation using regression (and the mediate function).

```
mediate(EducExp ~ FatherOcc + (Intelligence) + (Siblings) +
        (FatherEd), data=R.kerch, n.obs=767)
```

```
> mediate(6, x=4, m=1:3, data=R.kerch, n.obs=767)
```

The replication **data** matrices were simulated based upon the specified number of subjects and the observed correlation **matrix**.

### Mediation/Moderation Analysis

**Call:** mediate(y = EducExp ~ FatherOcc + (Intelligence) + (Siblings) + (FatherEd), data = R.kerch, n.obs = 767)

The DV (Y) was EducExp . The IV (X) was FatherOcc . The mediating variables were Intelligence Siblings FatherEd .

Total effect (c) of FatherOcc on EducExp = 0.41 S.E. = 0.03

t = 12.43 df= 762 with p = 1.9e-32

Direct effect (c') of FatherOcc on EducExp removing Intelligence Siblings

Indirect effect (ab) of FatherOcc on EducExp through Intelligence Siblings

Mean bootstrapped indirect effect = 0.26 with standard error = 0.03 Lower

R<sub>1</sub> = 0.61 R<sub>2</sub> = 0.38 F = 115.03 on 4 and 762 DF p-value : 9.89e-77

To see the longer output, specify short = FALSE in the print statement or

R code

summary(mod)

Call: mediate(y = EducExp ~ FatherOcc + (Intelligence) + (Siblings) + (FatherEd), data = R.kerch, n.obs = 767)

Total effect estimates (c)

	EducExp	se	t	df	Prob
FatherOcc	0.41	0.03	12.43	762	1.92e-32

Direct effect estimates (c')

	EducExp	se	t	df	Prob
FatherOcc	0.17	0.04	4.63	762	4.28e-06
Intelligence	0.37	0.03	12.45	762	1.64e-32
Siblings	-0.12	0.03	-4.27	762	2.17e-05
FatherEd	0.22	0.04	6.00	762	2.99e-09

R = 0.61 R2 = 0.38 F = 115.03 on 4 and 762 DF p-value: 9.89e-77

'a' effect estimates

	FatherOcc	se	t	df	Prob
Intelligence	0.25	0.04	7.14	765	2.15e-12
Siblings	-0.11	0.04	-3.00	765	2.75e-03
FatherEd	0.61	0.03	21.35	765	1.10e-79

'b' effect estimates

	EducExp	se	t	df	Prob
Intelligence	0.37	0.03	12.45	762	1.64e-32
Siblings	-0.12	0.03	-4.27	762	2.17e-05
FatherEd	0.22	0.04	6.00	762	2.99e-09

'ab' effect estimates

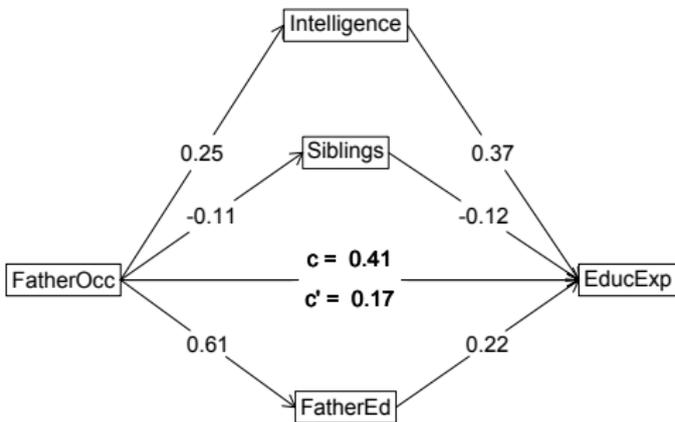
	EducExp	boot	sd	lower	upper
FatherOcc	0.24	0.22	0.03	0.17	0.27

'ab' effects estimates for each mediator

	Intelligence	boot	sd	lower	upper
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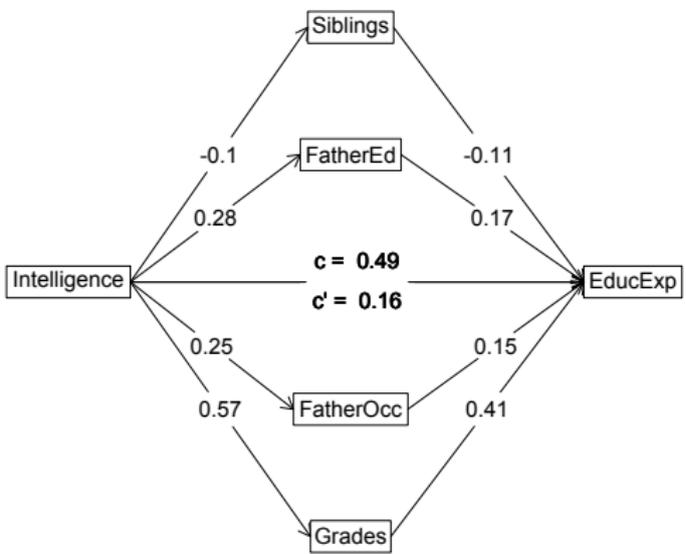
# The mediated paths (From mediate)

## Mediation model



## An alternative mediated model (From mediate)

Mediation model



## Modeling the Kerckhoff data using mediate

R code

```
model <- mediate(OccupAsp ~ Intelligence + Siblings +
  FatherEd + FatherOcc + (Grades) + (EducExp), data=R.kerch,n.o
  mediate.diagram(model, main="Kerckhoff model using mediate function")
```

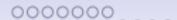
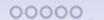
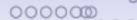
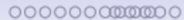
Direct effect estimates (traditional regression) (c')

	OccupAsp	se	t	df	Prob
Intercept	0.00	0.03	0.00	760	1.00e+00
Intelligence	-0.04	0.03	-1.16	760	2.46e-01
Siblings	-0.02	0.03	-0.68	760	4.98e-01
FatherEd	-0.04	0.04	-1.16	760	2.47e-01
FatherOcc	0.10	0.03	2.85	760	4.43e-03
Grades	0.16	0.04	4.29	760	2.06e-05
EducExp	0.55	0.04	14.60	760	1.01e-42

R = 0.67 R2 = 0.44 F = 100.9 on 6 and 760 DF p-value: 2.95e-93

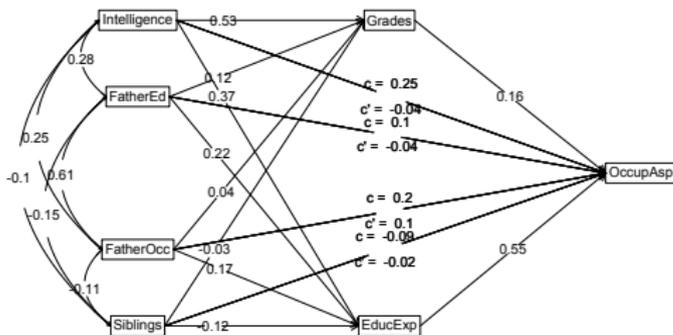
Total effect estimates (c)

	OccupAsp	se	t	df	Prob
Intercept	0.00	0.03	0.00	762	1.00e+00
Intelligence	0.25	0.03	7.29	762	7.63e-13
Siblings	-0.09	0.03	-2.78	762	5.60e-03
FatherEd	0.10	0.04	2.36	762	1.84e-02
FatherOcc	0.20	0.04	4.80	762	1.90e-06



## The full model (without one path)

Kerckhoff model using mediate function



## Can use sem functions (in either sem or lavaan) to estimate the mediation model

1. Treat all variables as observed (fixed)
  - Specify a limited number of paths rather than the full regression model
2. *sem* (Fox et al., 2013) commands
  - either in RAM (path) notation or
  - causal notation
3. *lavaan* (Rosseel, 2012) commands similar to causal notation of *sem* and the commercial programs Mplus and EQS
  - Output can mimic either MPlus or LISREL
  - Commands can be translated directly from *lavaan* to MPlus

## Kerckhoff-Kenny path analysis (modified from sem help page) to just predict the DVs)

```

model.kerch <- specifyModel()
  Intelligence -> Grades,      gam51
  Siblings -> Grades,         gam52
  FatherEd -> Grades,         gam53
  FatherOcc -> Grades,        gam54
  Intelligence -> EducExp,    gam61
  Siblings -> EducExp,        gam62
  FatherEd -> EducExp,        gam63
  FatherOcc -> EducExp,       gam64
  Intelligence -> OccupAsp,   gam71
  Siblings -> OccupAsp,       gam72
  FatherEd -> OccupAsp,       gam73
  FatherOcc -> OccupAsp,      gam74
  # Grades -> EducExp, beta65
  # Grades -> OccupAsp, beta75
  # EducExp -> OccupAsp, beta76

sem.kerch <- sem(model.kerch, R.kerch, 737, fixed.x=c('Intelligence', 'Siblings',
  'FatherEd', 'FatherOcc'))
summary(sem.kerch)

```

## Fixed path model – not modeling the DV correlations

Model Chisquare = 411.72 Df = 3 Pr(>Chisq) = 6.4133e-89  
 Chisquare (**null model**) = 1664.3 Df = 21  
 Goodness-of-fit **index** = 0.85747  
 Adjusted goodness-of-fit **index** = -0.33031  
 RMSEA **index** = 0.43024 90% CI: (0.39572, 0.46581)  
 Bentler-Bonnett NFI = 0.75262  
 Tucker-Lewis NNFI = -0.74103  
 Bentler CFI = 0.75128  
 SRMR = 0.099759  
 AIC = 441.72  
 AICc = 412.38  
 BIC = 510.76  
 CAIC = 388.91

### Normalized Residuals

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
0.000	0.000	0.000	0.956	0.000	10.100

### R-square for Endogenous Variables

Grades	EducExp	OccupAsp
0.3490	0.3765	0.1930

## With path coefficients of

### Parameter Estimates

	Estimate	Std Error	z value	Pr(> z )	
gam51	0.525902	0.031182	16.86530	8.0987e-64	Grades <--- Intelligence
gam52	-0.029942	0.030149	-0.99314	3.2064e-01	Grades <--- Siblings
gam53	0.118966	0.038259	3.10951	1.8740e-03	Grades <--- FatherEd
gam54	0.040603	0.037785	1.07456	2.8257e-01	Grades <--- FatherOcc
gam61	0.373339	0.030517	12.23376	2.0521e-34	EducExp <--- Intelligence
gam62	-0.123910	0.029506	-4.19954	2.6745e-05	EducExp <--- Siblings
gam63	0.220918	0.037442	5.90022	3.6302e-09	EducExp <--- FatherEd
gam64	0.168302	0.036979	4.55125	5.3328e-06	EducExp <--- FatherOcc
gam71	0.248827	0.034718	7.16718	7.6561e-13	OccupAsp <--- Intelligence
gam72	-0.091653	0.033567	-2.73047	6.3245e-03	OccupAsp <--- Siblings
gam73	0.098869	0.042596	2.32109	2.0282e-02	OccupAsp <--- FatherEd
gam74	0.198486	0.042069	4.71809	2.3807e-06	OccupAsp <--- FatherOcc
V[Grades]	0.650995	0.033935	19.18333	5.1010e-82	Grades <--> Grades
V[EducExp]	0.623511	0.032503	19.18333	5.1010e-82	EducExp <--> EducExp
V[OccupAsp]	0.806964	0.042066	19.18333	5.1010e-82	OccupAsp <--> OccupAsp

Note, we are not modeling the DV correlations so the residuals will be large

## Residuals from this model

```
> lowerMat(resid(sem.kerch))
```

	Intll	Sblng	FthrE	FthrO	Grads	EdcEx	OccpA
Intelligence	0.00						
Siblings	0.00	0.00					
FatherEd	0.00	0.00	0.00				
FatherOcc	0.00	0.00	0.00	0.00			
Grades	0.00	0.00	0.00	0.00	0.00		
EducExp	0.00	0.00	0.00	0.00	0.26	0.00	
OccupAsp	0.00	0.00	0.00	0.00	0.25	0.38	0.00

## fixed sem = regression

Compare the coefficients from this sem with the regression  $\beta$  values

```

round(sem.kerch$coeff,3)
  gam51      gam52      gam53      gam54      gam61      gam62      gam63
  gam64      gam71      gam72      gam73      gam74
  0.526      -0.030      0.119      0.041      0.373      -0.124      0.221
  0.168      0.249      -0.092      0.099      0.198
  V[Grades] V[EducExp] V[OccupAsp]
  0.651      0.624      0.807

sc.kk <- setCor(x=c(1:4),y=c(5:7),data=R.kerch)

round(as.vector(sc.kk$coefficients),3)
[1] 0.526 -0.030 0.119 0.041 0.373 -0.124 0.221 0.168 0.249 -0.092 0.099 0.198
    
```

## More complicated regression

1. Able to model the intercorrelations of the Y variables
  - This treats the Ys as both predictors and predicted
2. Able to let some of the Ys be part of the regression model

## Complete Kerckhoff-Kenny path analysis (taken from sem help page)

```

model.kerch1 <- specifyModel()
  Intelligence -> Grades,      gam51
  Siblings -> Grades,         gam52
  FatherEd -> Grades,         gam53
  FatherOcc -> Grades,        gam54
  Intelligence -> EducExp,    gam61
  Siblings -> EducExp,        gam62
  FatherEd -> EducExp,        gam63
  FatherOcc -> EducExp,       gam64
  Grades -> EducExp,          beta65
  Intelligence -> OccupAsp,   gam71
  Siblings -> OccupAsp,      gam72
  FatherEd -> OccupAsp,      gam73
  FatherOcc -> OccupAsp,     gam74
  Grades -> OccupAsp,        beta75
  EducExp -> OccupAsp,       beta76

sem.kerch1 <- sem(model.kerch1, R.kerch, 737, fixed.x=c('Intelligence', 'S
  'FatherEd', 'FatherOcc'))
summary(sem.kerch1)

```

## sem output

Model Chisquare = 3.2685e-13 Df = 0 Pr(>Chisq) = NA  
 Chisquare (**null model**) = 1664.3 Df = 21  
 Goodness-of-fit **index** = 1

AIC = 36  
 AICc = 0.95265  
 BIC = 118.85  
 CAIC = 3.2685e-13  
 Normalized Residuals

	Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
Normalized Residuals	-4.26e-15	-1.35e-15	0.00e+00	-4.17e-16	0.00e+00	1.49e-15

Parameter Estimates

	Estimate	Std Error	z value	Pr(> z )	
gam51	0.525902	0.031182	16.86530	8.0987e-64	Grades <--- Intelligence
gam52	-0.029942	0.030149	-0.99314	3.2064e-01	Grades <--- Siblings
gam53	0.118966	0.038259	3.10951	1.8740e-03	Grades <--- FatherEd
gam54	0.040603	0.037785	1.07456	2.8257e-01	Grades <--- FatherOcc
gam61	0.160270	0.032710	4.89979	9.5940e-07	EducExp <--- Intelligence
gam62	-0.111779	0.026876	-4.15899	3.1966e-05	EducExp <--- Siblings
gam63	0.172719	0.034306	5.03461	4.7882e-07	EducExp <--- FatherEd
gam64	0.151852	0.033688	4.50758	6.5571e-06	EducExp <--- FatherOcc
beta65	0.405150	0.032838	12.33799	5.6552e-35	EducExp <--- Grades
gam71	-0.039405	0.034500	-1.14215	2.5339e-01	OccupAsp <--- Intelligence
gam72	-0.018825	0.028222	-0.66700	5.0477e-01	OccupAsp <--- Siblings
gam73	-0.041333	0.036216	-1.14126	2.5376e-01	OccupAsp <--- FatherEd
gam74	0.099577	0.035446	2.80924	4.9658e-03	OccupAsp <--- FatherOcc
beta75	0.157912	0.037443	4.21738	2.4716e-05	OccupAsp <--- Grades
beta76	0.549593	0.038260	14.36486	8.5976e-47	OccupAsp <--- EducExp
V[Grades]	0.650995	0.033935	19.18333	5.1010e-82	Grades <--> Grades
V[EducExp]	0.516652	0.026932	19.18333	5.1010e-82	EducExp <--> EducExp
V[OccupAsp]	0.556617	0.029016	19.18333	5.1010e-82	OccupAsp <--> OccupAsp

## Note that these models give different path coefficients

```

> round(sem.kerch$coeff ,3)
gam51      gam52      gam53      gam54      gam61      gam62      gam63
gam64
  gam71      gam72      gam73
0.526      -0.030      0.119      0.041      0.373      -0.124      0.221
0.168
  0.249      -0.092      0.099
  gam74      V[Grades]  V[EducExp]  V[OccupAsp]
0.198      0.651      0.624      0.807
> round(sem.kerch1$coeff ,3)
gam51      gam52      gam53      gam54      gam61      gam62      gam63
gam64
  beta65      gam71      gam72
0.526      -0.030      0.119      0.041      0.160      -0.112      0.173
0.152
  0.405      -0.039      -0.019
  gam73      gam74      beta75      beta76      V[Grades]  V[EducExp]  V[OccupAsp]
-0.041      0.100      0.158      0.550      0.651      0.517      0.557
> round(sc.kk$coefficients ,2)
          Grades EducExp OccupAsp
Intelligence  0.53  0.37  0.25
Siblings     -0.03 -0.12 -0.09
FatherEd     0.12  0.22  0.10
FatherOcc    0.04  0.17  0.20

round(as.vector(sc.kk$coefficients),3)
[1] 0.526 -0.030 0.119 0.041 0.373 -0.124 0.221 0.168 0.249 -0.092 0.099 0.198

```

## A latent variable structural model

1. Taken from the LISREL User's reference guide
2. Data from Caslyn and Kenny (1977)
  - Self-concept of ability and perceived evaluation of others:  
Cause or effect of academic achievement
3. Variables
  - self concept
  - parental evaluation
  - teacher evaluation
  - friend evaluation
  - educational aspiration
  - college plans

## Caslyn and Kenny data

	self	<b>parent</b>	teacher	friend	edu_asp	college
self_concept	1.00	0.73	0.70	0.58	0.46	0.56
parental_ <b>eval</b>	0.73	1.00	0.68	0.61	0.43	0.52
teacher_ <b>eval</b>	0.70	0.68	1.00	0.57	0.40	0.48
friend_ <b>eval</b>	0.58	0.61	0.57	1.00	0.37	0.41
edu_aspir	0.46	0.43	0.40	0.37	1.00	0.72
college_plans	0.56	0.52	0.48	0.41	0.72	1.00

## Creating the model using `structure.diagram`

```
fx <- structure.list(6, list(c(1:4), c(5:6)), item.labels = rownames(ability),
                    f.labels=c("Ability", "Aspiration"))
mod.edu <- structure.diagram(fx, "r", title="Lisrel_example_3.2",
                            errors=TRUE, lr=FALSE, cex=.8)
```

`fx`

`fx`

	Ability	Aspiration
<code>self_concept</code>	"a1"	"0"
<code>parental_eval</code>	"a2"	"0"
<code>teacher_eval</code>	"a3"	"0"
<code>friend_eval</code>	"a4"	"0"
<code>edu_aspir</code>	"0"	"b5"
<code>college_plans</code>	"0"	"b6"

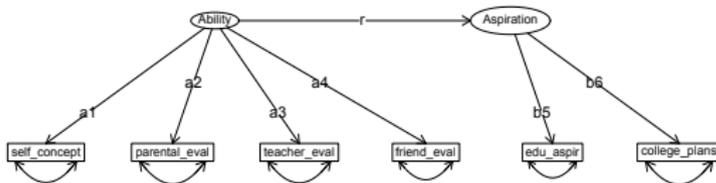
## The sem commands are in the mod.edu object

*mod.edu*

	<i>Path</i>	<i>Parameter</i>	<i>Value</i>
[1,]	"Ability ->self_concept"	"a1"	NA
[2,]	"Ability ->parental_eval"	"a2"	NA
[3,]	"Ability ->teacher_eval"	"a3"	NA
[4,]	"Ability ->friend_eval"	"a4"	NA
[5,]	"Aspiration ->edu_aspir"	"b5"	NA
[6,]	"Aspiration ->college_plans"	"b6"	NA
[7,]	"self_concept<->self_concept"	"x1e"	NA
[8,]	"parental_eval<->parental_eval"	"x2e"	NA
[9,]	"teacher_eval<->teacher_eval"	"x3e"	NA
[10,]	"friend_eval<->friend_eval"	"x4e"	NA
[11,]	"edu_aspir<->edu_aspir"	"x5e"	NA
[12,]	"college_plans<->college_plans"	"x6e"	NA
[13,]	"Aspiration<->Ability"	"rF2F1"	NA
[14,]	"Ability<->Ability"	NA	"1"
[15,]	"Aspiration<->Aspiration"	NA	"1"

## A model of the Caslyn-Kenny (1997) data set

### Structural model



```
ability <- as.matrix(ability) #sem requires matrix input
sem.edu <- sem(mod=mod.edu,S=ability,N=556)
summary(sem.edu)
```

Model Chisquare = 9.2557 Df = 8 Pr(>Chisq) = 0.32118  
 Chisquare (null model) = 1832 Df = 15  
 Goodness-of-fit index = 0.99443  
 Adjusted goodness-of-fit index = 0.98537  
 RMSEA index = 0.016817 90% CI: (NA, 0.054321)  
 Bentler-Bonnett NFI = 0.99495  
 Tucker-Lewis NNFI = 0.9987  
 Bentler CFI = 0.99931  
 SRMR = 0.012011  
 AIC = 35.256  
 AICc = 9.9273  
 BIC = 91.426  
 CAIC = -49.31

Normalized Residuals

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
-0.4410	-0.1870	0.0000	-0.0131	0.2110	0.5330

R-square for Endogenous Variables

self_concept	parental_eval	teacher_eval	friend_eval	edu_aspir	college_plans
0.7451	0.7213	0.6482	0.4834	0.6008	0.8629

## With parameter estimates

### Parameter Estimates

	Estimate	Std Error	z value	Pr(> z )	
a1	0.86320	0.035145	24.5612	3.2846e-133	self_concept <--- Ability
a2	0.84932	0.035450	23.9582	7.5937e-127	parental_eval <--- Ability
a3	0.80509	0.036405	22.1149	2.2725e-108	teacher_eval <--- Ability
a4	0.69527	0.038634	17.9964	2.0795e-72	friend_eval <--- Ability
b5	0.77508	0.040357	19.2058	3.3077e-82	edu_aspir <--- Aspiration
b6	0.92893	0.039410	23.5712	7.6153e-123	college_plans <--- Aspiration
x1e	0.25488	0.023367	10.9075	1.0617e-27	self_concept <--> self_concept
x2e	0.27865	0.024128	11.5491	7.4600e-31	parental_eval <--> parental_eval
x3e	0.35184	0.026919	13.0703	4.8660e-39	teacher_eval <--> teacher_eval
x4e	0.51660	0.034725	14.8768	4.6594e-50	friend_eval <--> friend_eval
x5e	0.39924	0.038196	10.4525	1.4266e-25	edu_aspir <--> edu_aspir
x6e	0.13709	0.043505	3.1511	1.6264e-03	college_plans <--> college_plans
rF2F1	0.66637	0.030954	21.5276	8.5783e-103	Ability <--> Aspiration

## Three competing models

1. Ability and aspirations are correlated
  - $r = .66$
2. Ability causes aspirations
  - $\text{beta} = .89$
3. Aspirations cause ability
  - $\text{beta} = .89$

## Create the model where ability lead to aspirations

```
phi <- phi.list(2,c(2))
```

```
phi
```

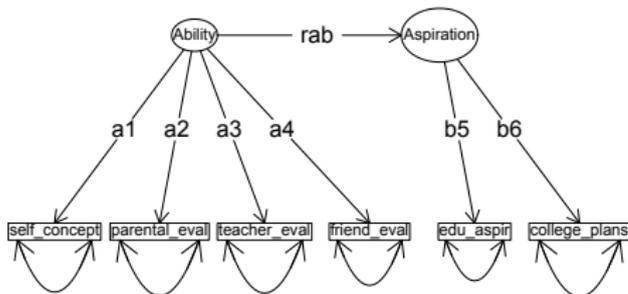
```
mod.edu <- structure.diagram(fx, phi, title="Aspiration_leads_to_ability",
                             errors=TRUE, lr=FALSE, cex=.7)
```

```
mod.edu
```

	Path	Parameter	Value
[1, ]	"Ability ->self_concept"	"a1"	NA
[2, ]	"Ability ->parental_eval"	"a2"	NA
[3, ]	"Ability ->teacher_eval"	"a3"	NA
[4, ]	"Ability ->friend_eval"	"a4"	NA
[5, ]	"Aspiration ->edu_aspir"	"b5"	NA
[6, ]	"Aspiration ->college_plans"	"b6"	NA
[7, ]	"self_concept<->self_concept"	"x1e"	NA
[8, ]	"parental_eval<->parental_eval"	"x2e"	NA
[9, ]	"teacher_eval<->teacher_eval"	"x3e"	NA
[10, ]	"friend_eval<->friend_eval"	"x4e"	NA
[11, ]	"edu_aspir<->edu_aspir"	"x5e"	NA
[12, ]	"college_plans<->college_plans"	"x6e"	NA
[13, ]	"Ability_->Aspiration"	"rF1F2"	NA
[14, ]	"Ability<->Ability"	NA	"1"
[15, ]	"Aspiration<->Aspiration"	NA	"1"

## Ability causes aspiration

Ability leads to Aspiration



## Fit statistics are identical

```
> sem.edu <- sem(mod=mod.edu, S=ability, N=556)
> summary(sem.edu)
```

```
summary(sem.edu)
```

```
Model Chi-square = 9.2557 Df = 8 Pr(>ChiSq) = 0.32118
Chi-square (null model) = 1832 Df = 15
Goodness-of-fit index = 0.99443
Adjusted goodness-of-fit index = 0.98537
RMSEA index = 0.016817 90% CI: (NA, 0.054321)
Bentler-Bonnett NFI = 0.99495
Tucker-Lewis NNFI = 0.9987
Bentler CFI = 0.99931
SRMR = 0.012011
AIC = 35.256
AICc = 9.9273
BIC = 91.426
CAIC = -49.31
```

Normalized Residuals

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
-0.4410	-0.1870	0.0000	-0.0131	0.2110	0.5330

**R-square for** Endogenous Variables

self_concept	parental_eval	teacher_eval	friend_eval	Aspiration	edu_aspir	college_plans
0.7451	0.7213	0.6482	0.4834	0.4440	0.6008	0.8629

## But the paths are different

### Parameter Estimates

	Estimate	Std Error	z value	Pr(> z )	
a1	0.86320	0.035145	24.5612	3.2848e-133	self_concept <--- Ability
a2	0.84932	0.035450	23.9582	7.5919e-127	parental_eval <--- Ability
a3	0.80509	0.036405	22.1149	2.2732e-108	teacher_eval <--- Ability
a4	0.69527	0.038634	17.9964	2.0794e-72	friend_eval <--- Ability
b5	0.57792	0.030630	18.8678	2.0977e-79	edu_aspir <--- Aspiration
b6	0.69263	0.037979	18.2370	2.6257e-74	college_plans <--- Aspiration
x1e	0.25488	0.023367	10.9075	1.0617e-27	self_concept <--> self_concept
x2e	0.27865	0.024128	11.5491	7.4610e-31	parental_eval <--> parental_eval
x3e	0.35184	0.026919	13.0703	4.8654e-39	teacher_eval <--> teacher_eval
x4e	0.51660	0.034725	14.8768	4.6595e-50	friend_eval <--> friend_eval
x5e	0.39924	0.038196	10.4525	1.4266e-25	edu_aspir <--> edu_aspir
x6e	0.13709	0.043505	3.1511	1.6264e-03	college_plans <--> college_plans
rF1F2	0.89371	0.074673	11.9683	5.2068e-33	Aspiration <--- Ability

<-----

Iterations = 30

## Let aspirations cause ability

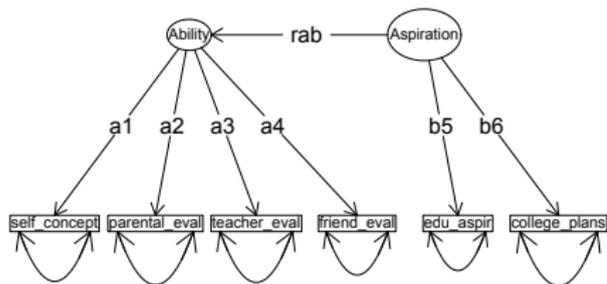
```
> phi[1,2] <- phi[2,1]
> phi[2,1] <- "0"
> mod.edu <- structure.diagram(fx, phi, main="Aspiration_leads_to_Ability", e
>mod.edu
```

```
> mod.edu
```

	Path	Parameter	Value
[1, ]	"Ability ->self_concept"	"a1"	NA
[2, ]	"Ability ->parental_eval"	"a2"	NA
[3, ]	"Ability ->teacher_eval"	"a3"	NA
[4, ]	"Ability ->friend_eval"	"a4"	NA
[5, ]	"Aspiration ->edu_aspir"	"b5"	NA
[6, ]	"Aspiration ->college_plans"	"b6"	NA
[7, ]	"self_concept<->self_concept"	"x1e"	NA
[8, ]	"parental_eval<->parental_eval"	"x2e"	NA
[9, ]	"teacher_eval<->teacher_eval"	"x3e"	NA
[10, ]	"friend_eval<->friend_eval"	"x4e"	NA
[11, ]	"edu_aspir<->edu_aspir"	"x5e"	NA
[12, ]	"college_plans<->college_plans"	"x6e"	NA
[13, ]	"Aspiration<- Ability"	"rF2F1"	NA
[14, ]	"Ability<->Ability"	NA	"1"
[15, ]	"Aspiration<->Aspiration"	NA	"1"

## Aspiration leads to ability

### Aspiration leads to Ability



## Fits are still identical

```
> sem.edu <- sem(mod=mod.edu, S=ability, N=556)
> summary(sem.edu)
```

```
Model Chi-square = 9.2557 Df = 8 Pr(>ChiSq) = 0.32118
Chi-square (null model) = 1832 Df = 15
Goodness-of-fit index = 0.99443
Adjusted goodness-of-fit index = 0.98537
RMSEA index = 0.016817 90% CI: (NA, 0.054321)
Bentler-Bonnett NFI = 0.99495
Tucker-Lewis NNFI = 0.9987
Bentler CFI = 0.99931
SRMR = 0.012011
AIC = 35.256
AICc = 9.9273
BIC = 91.426
CAIC = -49.31
```

### Normalized Residuals

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
-0.4410	-0.1870	0.0000	-0.0131	0.2110	0.5330

### R-square for Endogenous Variables

self_concept	parental_eval	teacher_eval	friend_eval	Aspiration
edu_aspir	college_plans			

## And the crucial coefficient is different

### Parameter Estimates

	Estimate	Std Error	z value	Pr(> z )	
a1	0.86320	0.035145	24.5612	3.2848e-133	self_concept <--- Ability
a2	0.84932	0.035450	23.9582	7.5919e-127	parental_eval <--- Ability
a3	0.80509	0.036405	22.1149	2.2732e-108	teacher_eval <--- Ability
a4	0.69527	0.038634	17.9964	2.0794e-72	friend_eval <--- Ability
b5	0.57792	0.030630	18.8678	2.0977e-79	edu_aspir <--- Aspiration
b6	0.69263	0.037979	18.2370	2.6257e-74	college_plans <--- Aspiration
x1e	0.25488	0.023367	10.9075	1.0617e-27	self_concept <--> self_concept
x2e	0.27865	0.024128	11.5491	7.4610e-31	parental_eval <--> parental_eval
x3e	0.35184	0.026919	13.0703	4.8654e-39	teacher_eval <--> teacher_eval
x4e	0.51660	0.034725	14.8768	4.6595e-50	friend_eval <--> friend_eval
x5e	0.39924	0.038196	10.4525	1.4266e-25	edu_aspir <--> edu_aspir
x6e	0.13709	0.043505	3.1511	1.6264e-03	college_plans <--> college_plans
rF2F1	0.89371	0.074673	11.9683	5.2068e-33	Aspiration <--- Ability

←-----

## Compare the three models

	correlated	asp	abil
a1	0.86	0.64	0.86
a2	0.85	0.63	0.85
a3	0.81	0.60	0.81
a4	0.70	0.52	0.70
b5	0.78	0.78	0.58
b6	0.93	0.93	0.69
x1e	0.25	0.25	0.25
x2e	0.28	0.28	0.28
x3e	0.35	0.35	0.35
x4e	0.52	0.52	0.52
x5e	0.40	0.40	0.40
x6e	0.14	0.14	0.14
rF2F1	0.67	0.89	0.89

1. Although fits were identical
2. Paths differ as a function of presumed influence
3. Which solution is correct?
4. Is this even possible to answer?

## lavaan syntax is perhaps easier (correlated latents)

R code

```
model <- "Ability =~ self_concept + parental_eval +
          teacher_eval + friend_eval
          Aspiration =~ edu_aspir + college_plans"
fit <- sem(model, sample.cov=casslyn, fixed.x=TRUE, sample.nobs=556,
           std.lv=TRUE, mimic="eqs") #note that the default is lavaan
```

lavaan (0.5-17) converged normally after 22 iterations

Number of observations	556
Estimator	ML
Minimum Function Test Statistic	9.256
Degrees of freedom	8
P-value (Chi-square)	0.321

Parameter estimates:

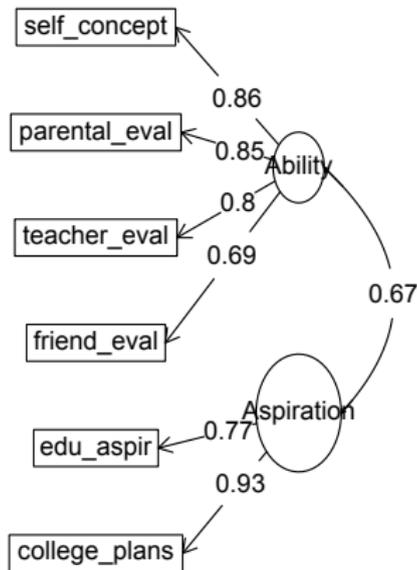
Information	Expected
Standard Errors	Standard

## lavaan output (continued)

	Estimate	Std.err	Z-value	P(> z )
<b>Latent variables:</b>				
<b>Ability =~</b>				
self_concept	0.863	0.035	24.561	0.000
parental_eval	0.849	0.035	23.958	0.000
teacher_eval	0.805	0.036	22.115	0.000
friend_eval	0.695	0.039	17.996	0.000
<b>Aspiration =~</b>				
edu_aspir	0.775	0.040	19.206	0.000
college_plans	0.929	0.039	23.571	0.000
<b>Covariances:</b>				
<b>Ability ~~</b>				
Aspiration	0.666	0.031	21.528	0.000
<b>Variances:</b>				
self_concept	0.255	0.023	10.907	0.000
parental_eval	0.279	0.024	11.549	0.000
teacher_eval	0.352	0.027	13.070	0.000
friend_eval	0.517	0.035	14.877	0.000
edu_aspir	0.399	0.038	10.453	0.000
college_plans	0.137	0.044	3.151	0.002
Ability	1.000			
Aspiration	1.000			

## A model of the Caslyn-Kenny (1997) data set using lavaan

### Confirmatory structure



## lavaan syntax for regressions

R code

```

model <- "Ability =~ self_concept + parental_eval +
          teacher_eval + friend_eval
          Aspiration =~ edu_aspir + college_plans
          Aspiration ~ Ability"
fit <- sem(model, sample.cov=casslyn, fixed.x=TRUE, sample.nobs=556,
           std.lv=TRUE, mimic="eqs") #note that the default is lavaan
    
```

lavaan (0.5-17) converged normally after 22 iterations

Number of observations	556
Estimator	ML
Minimum Function Test Statistic	9.256
Degrees of freedom	8
P-value (Chi-square)	0.321

Parameter estimates:

Information	Expected
Standard Errors	Standard

	Estimate	Std.err	Z-value	P(> z )
--	----------	---------	---------	---------

Latent variables:

Ability =~				
self_concept	0.863	0.035	24.561	0.000

## lavaan regression output, continued

	Estimate	Std.err	Z-value	P(> z )
<b>Latent variables:</b>				
<b>Ability =~</b>				
self_concept	0.863	0.035	24.561	0.000
parental_eval	0.849	0.035	23.958	0.000
teacher_eval	0.805	0.036	22.115	0.000
friend_eval	0.695	0.039	17.996	0.000
<b>Aspiration =~</b>				
edu_aspir	0.578	0.031	18.868	0.000
college_plans	0.693	0.038	18.237	0.000
<b>Regressions:</b>				
<b>Aspiration ~</b>				
Ability	0.894	0.075	11.968	0.000
<b>Variances:</b>				
self_concept	0.255	0.023	10.907	0.000
parental_eval	0.279	0.024	11.549	0.000
teacher_eval	0.352	0.027	13.070	0.000
friend_eval	0.517	0.035	14.877	0.000
edu_aspir	0.399	0.038	10.453	0.000
college_plans	0.137	0.044	3.151	0.002
Ability	1.000			
Aspiration	1.000			

## Implications of arrows

1. Need to fit alternative models
  - Create alternative plausible models
  - Create alternative implausible models (they will fit also).
2. Need to consider alternative representations
  - Try reversing arrows
3. Are there external variables (e.g., time) that allow one to choose between models?
4. Confirmation that a model fits does not confirm theoretical adequacy of the model.

## Two fundamentally different types of observed variables

1. Observed variables can be “reflective” of the latent variables. They are “effect indicators”.
  - Variables are caused by the latent variables.
  - Covariation between the variables are explained by the latent variables
2. Observed variables can be “causal indicators” or formative indicators that can directly effect the latent variable
  - Variables cause the “latent” variable
  - Covariation of the the observed variables is not modeled

## Formative indicators Bollen (2002)

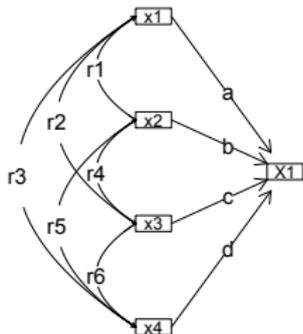
1. The correlational structure of formative indicators is independent of the loadings on a factor.
  - They are not locally independent
2. Examples of formative indicators Time spent in social interaction
  - Time spent with family, time spent with friends, time spent with coworkers.
  - These might in fact be negatively correlated even though total score is important.

## Effect (reflective) indicators

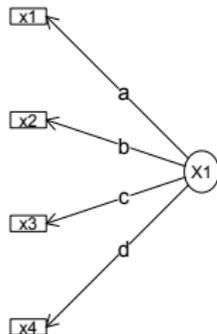
1. Test scores on various quantitative tests as effect indicators of trait
  - Feelings of self worth as effect indicators of self esteem
  - Ability items as indicators of ability
2. Correlational structure is a function of the path coefficients with latent variables
3. Variables are locally independent
  - (uncorrelated with each other when latent variable is partialled out)

## Formative vs. Effect variables as regression versus factors

Formative Variables



Effect Variables



## Confirmatory Factor Analysis as a way of testing factor models

1. Exploratory factor analysis (EFA) is a way of summarizing the relationships between variables
  - May have goodness of fit of factor model to the covariances but this is unusual
  - But all loadings are modeled
2. Confirmatory factor analysis (CFA) also models the data
  - But limits paths to specified paths, sets other to zero
  - Goodness of fit statistics are always produced
3. Blending CFA with regression is a full Structural Equation Model

## Using sim.structural to create models

```
fx <-matrix(c( .9,.8,.6,rep(0,4),.6,.8,-.7),ncol=2)
fy <- matrix(c(.6,.5,.4),ncol=1)
rownames(fx) <- c("V","Q","A","nach","Anx")
rownames(fy) <- c("gpa","Pre","MA")
Phi <-matrix( c(1,0,.7,.0,1,.6,.7,.6,1),ncol=3)
gre.gpa <- sim.structural(fx,Phi,fy,n=1000)
gre.gpa
```

```
> fx
> fy
Call: sim.structural(fx = fx, Phi = Phi, fy = fy)
> Phi
```

\$model (Population correlation matrix)

	V	Q	A	nach	Anx	gpa
Pre	MA					
V	1.00	0.72	0.54	0.00	0.00	0.38
0.32	0.25					
Q	0.72	1.00	0.48	0.00	0.00	0.34
0.28	0.22					
A	0.54	0.48	1.00	0.48	-0.42	0.47
0.39	0.31					
nach	0.00	0.00	0.48	1.00	-0.56	0.29
0.24	0.19					
Anx	0.00	0.00	-0.42	-0.56	1.00	
1.00	-0.25	-0.21	-0.17			
gpa	0.38	0.34	0.47	0.29	-0.25	1.00
0.30	0.24					
Pre	0.32	0.28	0.39	0.24	-0.21	0.30
1.00	0.20					
MA	0.25	0.22	0.31	0.19	-0.17	0.24
0.20	1.00					

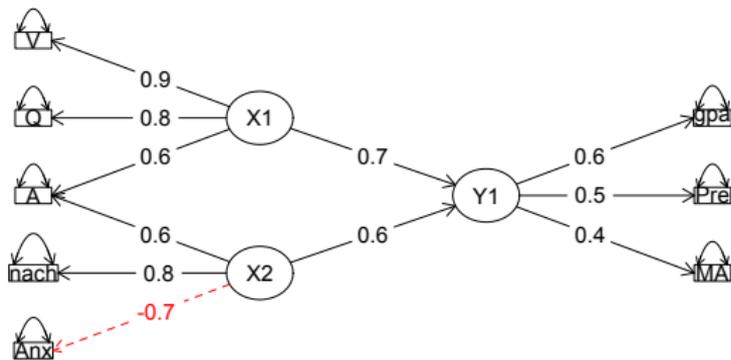
```
> fx
      [,1] [,2]
V      0.9  0.0
Q      0.8  0.0
A      0.6  0.6
nach   0.0  0.8
Anx    0.0 -0.7
> Phi
      [,1] [,2] [,3]
[1,]  1.0  0.0  0.7
[2,]  0.0  1.0  0.6
[3,]  0.7  0.6  1.0
> fy
      [,1]
gpa  0.6
Pre  0.5
MA   0.4
>
```

\$reliability (population reliability)

## Create a figure (and write sem code for the sem package)

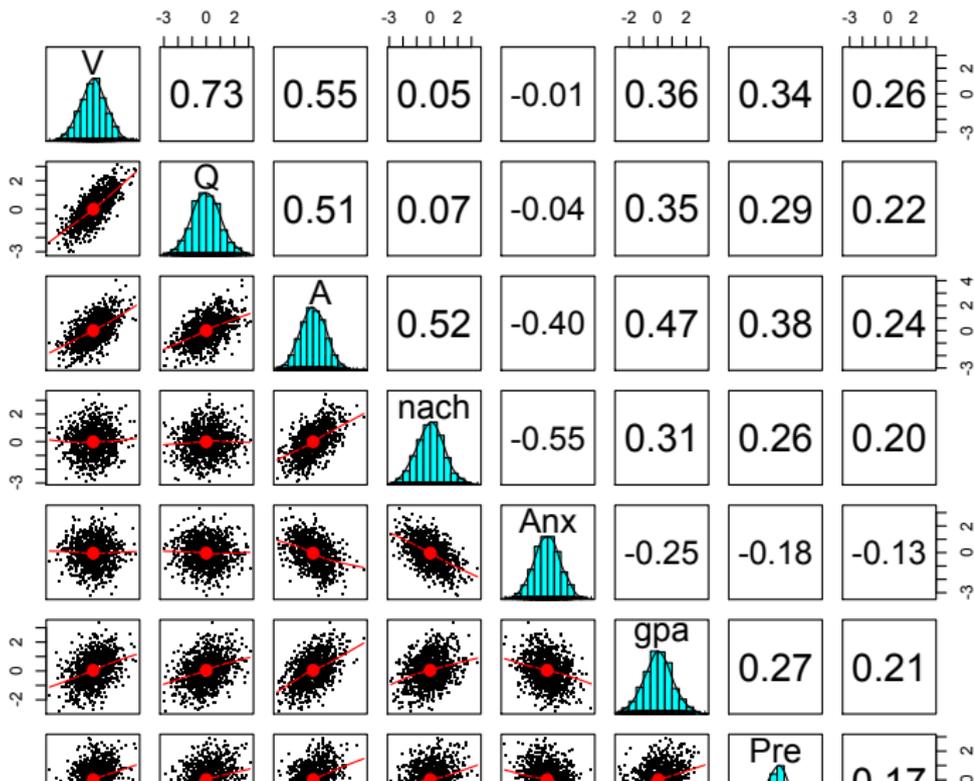
```
mod4 <- structure.diagram(fx,Phi,fy,errors=TRUE,e.size=.3)
```

Structural model



## Show the SPLOM

```
pairs.panels(gre.gpa$observed,pch=".")
```



## Show the sem code – This uses RAM path notation for all arrows

```
> mod4
```

	Path	Parameter	Value
[1,]	"X1->V"	"F1V"	NA
[2,]	"X1->Q"	"F1Q"	NA
[3,]	"X1->A"	"F1A"	NA
[4,]	"X2->A"	"F2A"	NA
[5,]	"X2->nach"	"F2nach"	NA
[6,]	"X2->Anx"	"F2Anx"	NA
[7,]	"V<->V"	"x1e"	NA
[8,]	"Q<->Q"	"x2e"	NA
[9,]	"A<->A"	"x3e"	NA
[10,]	"nach<->nach"	"x4e"	NA
[11,]	"Anx<->Anx"	"x5e"	NA
[12,]	"Y1->gpa"	"Fygpa"	NA
[13,]	"Y1->Pre"	"FyPre"	NA
[14,]	"Y1->MA"	"FyMA"	NA
[15,]	"gpa<->gpa"	"y1e"	NA
[16,]	"Pre<->Pre"	"y2e"	NA
[17,]	"MA<->MA"	"y3e"	NA
[18,]	"X2<->X1"	"rF2F1"	NA
[19,]	"X1->Y1"	"rX1Y1"	NA
[20,]	"X2->Y1"	"rX2Y1"	NA
[21,]	"X1<->X1"	NA	"1"
[22,]	"X2<->X2"	NA	"1"
[23,]	"Y1<->Y1"	NA	"1"

```
attr(,"class")
[1] "mod"
```

## Can we recover the structure using a confirmatory factor model?

### R code

```

model <- 'Ability =~ V + Q + A
          Motivation =~ A + Anx
          Performance =~ gpa + MA + Pre'
fit <- sem(model, gre.gpa$observed, std.lv=TRUE)
summary(fit)
lavaan.diagram(fit)
    
```

## lavaan output

lavaan (0.5-17) converged normally after 28 iterations

Number of observations	1000
Estimator	ML
Minimum Function Test Statistic	12.116
Degrees of freedom	10
P-value (Chi-square)	0.277

Parameter estimates:

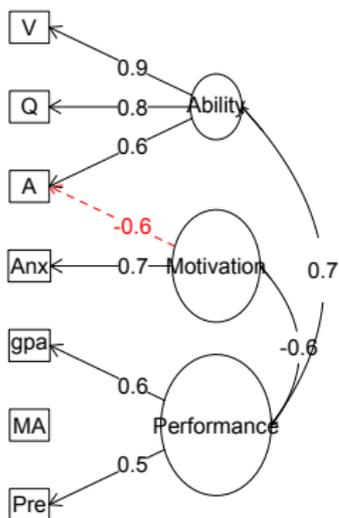
	Information		Expected		Standard
	Standard Errors		Standard		
	Estimate	Std.err	Z-value	P(> z )	
<b>Latent variables:</b>					
<b>Ability =~</b>					
V	0.924	0.029	32.165	0.000	
Q	0.830	0.029	28.263	0.000	
A	0.614	0.033	18.540	0.000	
<b>Motivation =~</b>					
A	-0.593	0.038	-15.642	0.000	
Anx	0.660	0.044	14.990	0.000	
<b>Performance =~</b>					
gpa	0.570	0.036	16.001	0.000	
MA	0.341	0.035	9.637	0.000	
Pre	0.483	0.035	13.678	0.000	

Covariances:

<b>Ability ~~</b>					
Motivation	-0.033	0.051	-0.654	0.513	
Performance	0.748	0.039	19.386	0.000	
<b>Motivation ~~</b>					

## Create a figure from the lavaan output)

### Confirmatory structure



## But the data can also be modeled as full SEM model

### R code

```

model <- 'Ability =~ V + Q + A
          Motivation =~ A + Anx
          Performance =~ gpa + MA + Pre
          Performance ~ Ability + Motivation'

fit <- sem(model, gre.gpa$observed, std.lv=TRUE)
summary(fit)
standardizedSolution(fit)
lavaan.diagram(fit)
    
```

## lavaan solution for the regression model

standardizedSolution(fit)

	lhs	op	rhs	est.	std	se	z	pvalue
1	Ability	==	V	0.899	0.028		32.165	0.000
2	Ability	==	Q	0.810	0.029		28.263	0.000
3	Ability	==	A	0.593	0.032		18.540	0.000
4	Motivation	==	A	-0.573	0.037		-15.642	0.000
5	Motivation	==	Anx	0.672	0.045		14.990	0.000
6	Performance	==	gpa	0.579	0.198		2.932	0.003
7	Performance	==	MA	0.340	0.117		2.891	0.004
8	Performance	==	Pre	0.484	0.165		2.941	0.003
9	Performance	~	Ability	0.729	0.252		2.889	0.004
10	Performance	~	Motivation	-0.575	0.209		-2.749	0.006
11	V	~~	V	0.193	0.026		7.444	0.000
12	Q	~~	Q	0.344	0.025		13.704	0.000
13	A	~~	A	0.297	0.034		8.625	0.000
14	Anx	~~	Anx	0.548	0.053		10.301	0.000
15	gpa	~~	gpa	0.664	0.039		17.001	0.000
16	MA	~~	MA	0.885	0.041		21.435	0.000
17	Pre	~~	Pre	0.766	0.039		19.660	0.000
18	Ability	~~	Ability	1.000	NA		NA	NA
19	Motivation	~~	Motivation	1.000	NA		NA	NA
20	Performance	~~	Performance	0.110	NA		NA	NA
21	Ability	~~	Motivation	-0.033	0.051		-0.654	0.513

## Measuring structure at two (or more) time points

1. Is the structure the same
  - Structural Invariance (is the graph the same)
  - Measurement invariance (are the loadings the same)
  - Strong measurement invariance (are the item intercepts the same?)
  - Measuring change
2. Do the means change (is there growth)
  - This is the means of the latent trait, not the means of the items
3. Do the latent traits correlate across two or more occasions?
  - Just two occasions, can not separate trait from state effects
  - With  $> 2$  occasions, can examine trait and state effects
4. Compare several different simulations

## Create some basic data and add in some change

```
> set.seed(42)
> fx <- matrix(c(.8,.7,.6,rep(0,6),.8,.7,.6),ncol=2)
> fx
```

```
      [,1] [,2]
[1,]  0.8  0.0
[2,]  0.7  0.0
[3,]  0.6  0.0
[4,]  0.0  0.8
[5,]  0.0  0.7
[6,]  0.0  0.6
```

```
> Phi <- matrix(c(1,.6,.6,1),ncol=2)
> Phi
```

```
      [,1] [,2]
[1,]  1.0  0.6
[2,]  0.6  1.0
```

```
> x.model <- sim(fx=fx,Phi=Phi,mu=c(0,1),n=250)
```

```
> x <- x.model$observed
```

```
> structure.diagram(fx,Phi,lr=FALSE,e.size=.3,main="A_basic_two_time_model")
```

```
> describe(x,skew=FALSE)
```

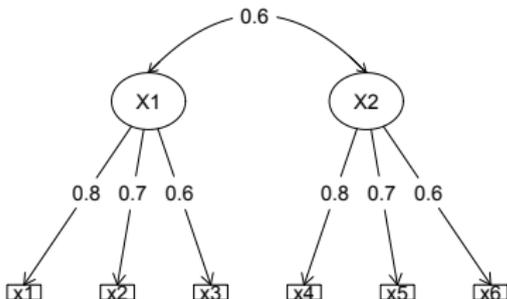
```
> pairs.panels(x,pch=". ")
```

	vars	n	mean	sd	median	trimmed	mad	min
<b>max range</b>			skew	kurtosis	se			
1	1	250	0.01	0.97	0.06	0.01	0.92	-2.56 2.68
5.24			0.05	0.03	0.06			
2	2	250	0.07	1.04	0.05	0.08	0.98	-2.76 3.07
5.83			-0.05	-0.04	0.07			
3	3	250	0.00	0.98	-0.03	-0.04	0.97	-2.62 3.90
6.52			0.53	0.79	0.06			
4	4	250	0.83	0.95	0.85	0.84	1.01	2.04 3.64

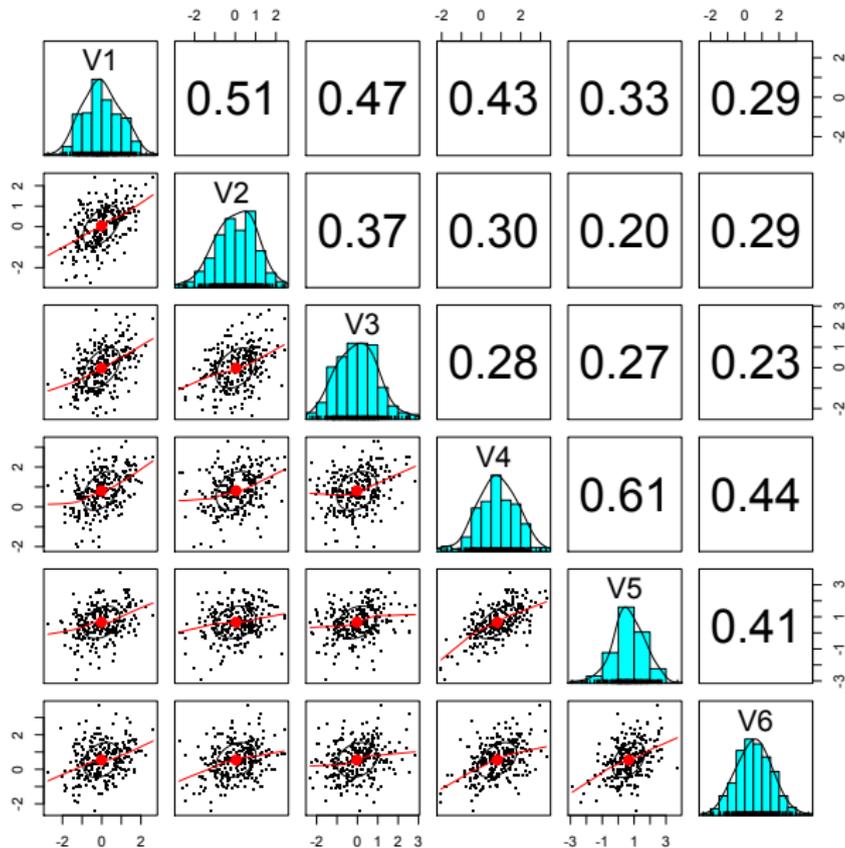
1. Set the random seed
2. Create a one factor structure
3. Put in some change
4. Show the structural model
5. Describe it
6. Show the splom (with small pch)

# A basic two occasion trait model

## A basic two time model



## Splom of 6 basic variables



## Multiple models

1. Ignore time, are the data congeneric?
  - Are they all measures of the same thing?
  - This is a one factor model
2. Include time, do we recover a correlation across time?
  - Try a two factor model
  - Plot the resulting structure

## A one factor model

> fa(x)

```
Factor Analysis using method = minres
Call: fa(r = x)
Standardized loadings (pattern matrix) based upon
  MR1 h2 u2 com
1 0.76 0.57 0.43 1
2 0.65 0.42 0.58 1
3 0.56 0.31 0.69 1
4 0.64 0.41 0.59 1
5 0.50 0.25 0.75 1
6 0.50 0.25 0.75 1

                MR1
SS loadings      2.21
Proportion Var 0.37
```

Mean item complexity = 1  
 Test of the hypothesis that 1 factor is sufficient

The degrees of freedom for the null model are 15

and the objective function was 1.57  
 with Chi Square of 387.63

The degrees of freedom for the model are 9  
 and the objective function was 0.28

The root mean square of the residuals (RMSR) is 0.09

The df corrected root mean square of the residuals is 0.12

correlation matrix

The harmonic number of observations is 250  
 with the empirical chi square 62.96 with prob < 3.6e-10

The total number of observations was 250  
 with MLE Chi Square = 67.96 with prob < 3.8e-11

Tucker Lewis Index of factoring reliability = 0.736

RMSEA index = 0.164

and the 90 % confidence intervals are 0.127 0.199

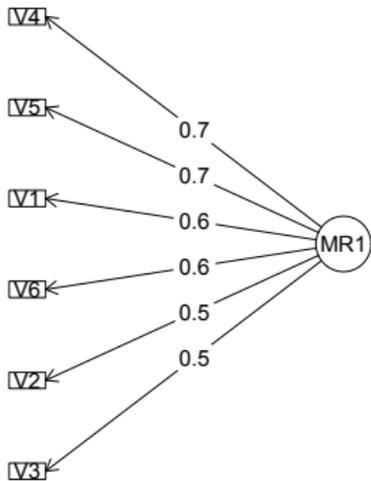
BIC = 18.27

Fit based upon off diagonal values = 0.94  
 Measures of factor score adequacy

Correlation of scores with factors 0.89

## Fit a one factor model to the data `fa.diagram(fa(x),sort=FALSE)`

A one factor model



## Try a two factor solution

```
> fa(x,2)
```

Factor Analysis using method = minres

Call: fa(r = x, nfact = 2)

Standardized loadings (pattern matrix) based upon correlation matrix

	MR1	MR2	h2	u2	com
1	0.88	-0.01	0.76	0.24	1.0
2	0.66	0.02	0.45	0.55	1.0
3	0.58	0.00	0.33	0.67	1.0
4	-0.01	0.87	0.75	0.25	1.0
5	-0.01	0.60	0.36	0.64	1.0
6	0.10	0.48	0.29	0.71	1.1

	MR1	MR2
SS loadings	1.58	1.37
Proportion Var	0.26	0.23
Cumulative Var	0.26	0.49
Proportion Explained	0.53	0.47
Cumulative Proportion	0.53	1.00

With factor correlations of

	MR1	MR2
MR1	1.00	0.58
MR2	0.58	1.00

Measures of factor score adequacy

MR2

Correlation of scores with factors

0.91 0.90

Multiple R square of scores with factors

0.83 0.81

Test of the hypothesis that 2 factors are sufficient

The degrees of freedom for the null model are

1.57 with Chi Square of 387.63

The degrees of freedom for the model are 4

and the objective function was

0.01

The root mean square of the residuals (RMSR) is

0.02

The df corrected root mean square of the residuals

0.03

The harmonic number of observations is 250

with the empirical chi square

1.97 with prob < 0.74

The total number of observations was 250

with MLE Chi Square = 2.51

with prob < 0.64

Tucker Lewis Index of factoring reliability =

1.015

RMSEA index = 0

and the 90 % confidence intervals are

NA 0.077

BIC = -19.58

Fit based upon off diagonal values = 1

## Compare the two solutions

Factor Analysis using method = minres

**Call:** fa(r = x)

Factor Analysis using method = minres

**Call:** fa(r = x)

Standardized loadings (pattern **matrix**) based upon

	MR1	h2	u2	com
1	0.76	0.57	0.43	1
2	0.65	0.42	0.58	1
3	0.56	0.31	0.69	1
4	0.64	0.41	0.59	1
5	0.50	0.25	0.75	1
6	0.50	0.25	0.75	1

	MR1
SS loadings	2.21
Proportion Var	0.37

Factor Analysis using method = minres

Factor Analysis using method = minres

**Call:** fa(r = x, nfactors = 2)

Standardized loadings (pattern **matrix**) based upon

	MR1	MR2	h2	u2	com
1	0.88	-0.01	0.76	0.24	1.0
2	0.66	0.02	0.45	0.55	1.0
3	0.58	0.00	0.33	0.67	1.0
4	-0.01	0.87	0.75	0.25	1.0
5	-0.01	0.60	0.36	0.64	1.0
6	0.10	0.48	0.29	0.71	1.1

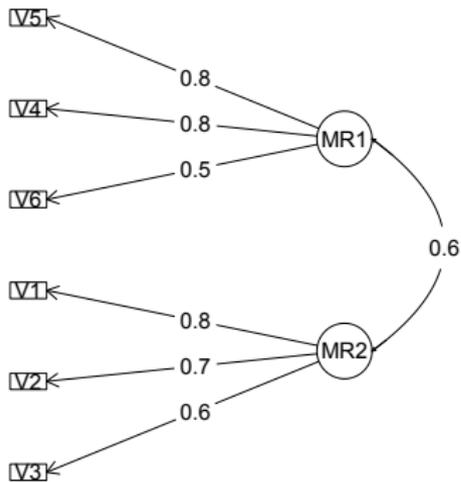
	MR1	MR2
SS loadings	1.58	1.37
Proportion Var	0.26	0.23
Cumulative Var	0.26	0.49
Proportion Explained	0.53	0.47
Cumulative Proportion	0.53	1.00

With **factor** correlations of

	MR1	MR2
MR1	1.00	0.58
MR2	0.58	1.00

# EFA two factor solution

## Factor Analysis



## Now try three different sems (using lavaan)

1. Two correlated factors, free loadings
2. Two correlated factors, equal loadings across occasions
3. Two correlated factors, all loadings equal

## A simple sem with a standardized solution using lavaan

R code

```
#factor model
mod2f <- 'F1 =~ V1 + V2 + V3
          F2 =~ V4 + V5 + V6
          #correlation between factors
          F1 ~~F2'# now fit it and summarize it
fit <- sem(mod2f, data=x,std.lv=TRUE)
summary(fit, fit.measures=TRUE)
standardizedSolution(fit)
```

	lhs	op	rhs	est.	std	se	z	pvalue
1	F1	==	V1	0.865	0.062	13.921	0	
2	F1	==	V2	0.681	0.063	10.756	0	
3	F1	==	V3	0.580	0.064	9.000	0	
4	F2	==	V4	0.838	0.067	12.599	0	
5	F2	==	V5	0.607	0.066	9.142	0	
6	F2	==	V6	0.558	0.067	8.348	0	
7	F1	~~	F2	0.602	0.061	9.887	0	
8	V1	~~	V1	0.251	0.068	3.713	0	
9	V2	~~	V2	0.536	0.064	8.443	0	
10	V3	~~	V3	0.664	0.068	9.763	0	
11	V4	~~	V4	0.297	0.077	3.885	0	
12	V5	~~	V5	0.631	0.070	9.036	0	
13	V6	~~	V6	0.689	0.072	9.609	0	
14	F1	~~	F1	1.000	NA	NA	NA	
15	F2	~~	F2	1.000	NA	NA	NA	

## lavaan fit statistics

lavaan (0.5–17) converged normally after 16 iterations

Number of observations  
250

Estimator

ML

Minimum Function Test Statistic  
3.986

Degrees of freedom  
8

P-value (Chi-square)  
0.858

Model test baseline **model**:

Minimum Function Test Statistic  
393.670

Degrees of freedom  
15

P-value  
0.000

User **model** versus baseline **model**:

Comparative Fit Index (CFI)  
1.000

Tucker-Lewis Index (TLI)  
1.020

Loglikelihood and Information Criteria:

Root Mean Square Error of Approximation:

RMSEA

0.000

90 Percent Confidence Interval

0.000 0.040

P-value RMSEA <= 0.05  
0.972

Standardized Root Mean Square Residual:

SRMR

0.017

Parameter estimates:

Information

Expected

Standard Errors

Standard

Z-value	P(> z )	Estimate	Std. err
Latent variables:			
F1 =~			
		V1	0.840 0.060
13.921	0.000	V2	0.703 0.065
10.756	0.000	V3	0.568 0.063
9.000	0.000		

F1 =~

V1 0.840 0.060

V2 0.703 0.065

V3 0.568 0.063

9.000 0.000

## Create two new models with some equality constraints

The first sets the loadings for 1-3 equal to those of 4-6, the second says they are all equal

R code

```
mod2fa <- 'F1 =~ a*V1 + b*V2 + c*V3
          F2 =~ a*V4 + b*V5 + c*V6
          F1 ~~F2'
fit2a <- sem(mod2fa,data=x.df,std.lv=TRUE)
summary(fit2a,fit.measures=TRUE)

mod2fe <- 'F1 =~ a*V1 + a*V2 + a*V3
          F2 =~ a*V4 + a*V5 + a*V6
          F1 ~~F2'
fit2e <- sem(mod2fe,data=x.df,std.lv=TRUE)
summary(fit2e,fit.measures=TRUE)
```

## Equal across time

lavaan (0.5-17) converged normally after 15 iterations

Number of observations  
250

Estimator  
ML

Minimum Function Test Statistic  
5.338

Degrees of freedom  
11

P-value (Chi-square)  
0.914

Model test baseline **model**:

Minimum Function Test Statistic  
393.670

Degrees of freedom  
15

P-value  
0.000

User **model** versus baseline **model**:

Comparative Fit Index (CFI)  
1.000

Tucker-Lewis Index (TLI)  
1.020

Loglikelihood and Information Criteria:

Root Mean Square Error of Approximation:

RMSEA  
0.000

90 Percent Confidence Interval  
0.000 0.025

P-value RMSEA <= 0.05  
0.991

Standardized Root Mean Square Residual:

SRMR  
0.035

Parameter estimates:

Information  
Expected  
Standard Errors  
Standard

	Z-value	P(> z )	Estimate	Std. err
Latent variables:				
F1 =~				
V1 (a)	17.711	0.000	0.817	0.046
V2 (b)	13.630	0.000	0.662	0.049
V3 (c)	11.949	0.000	0.549	0.046

## All loadings equal

```
> summary(fit2e, fit.measures=TRUE)
lavaan (0.5-17) converged normally after
12 iterations
```

```
Number of observations
250

Estimator
ML
Minimum Function Test Statistic
26.140
Degrees of freedom
13
P-value (Chi-square)
0.016
```

```
Model test baseline model:

Minimum Function Test Statistic
393.670
Degrees of freedom
15
P-value
0.000
```

```
User model versus baseline model:

Comparative Fit Index (CFI)
0.965
Tucker-Lewis Index (TLI)
0.960
```

Root Mean Square Error of Approximation:

```
RMSEA
0.064
90 Percent Confidence Interval
0.026 0.099
P-value RMSEA <= 0.05
0.235
```

Standardized Root Mean Square Residual:

```
SRMR
0.084
```

Parameter estimates:

```
Information
Expected
Standard Errors
Standard
```

	Z-value	P(> z )	Estimate	Std. err
Latent variables:				
F1 =~				
V1 (a)	20.964	0.000	0.686	0.033
V2 (a)	20.964	0.000	0.686	0.033
V3 (a)	20.964	0.000	0.686	0.033

## Do the models differ?

These are nested models, and we can compare their  $\chi^2$  values.

R code

```
anova (fit, fit2a)
anova (fit2a, 2e)
```

### Chi Square Difference Test

	Df	AIC	BIC	Chisq	Chisq diff	Df diff	Pr(>Chisq)
fit	8	3838.9	3884.6	3.9861			
fit2a	11	3834.2	3869.4	5.3380	1.3519	3	0.7169

### hi Square Difference Test

	Df	AIC	BIC	Chisq	Chisq diff	Df diff	Pr(>Chisq)
fit2a	11	3834.2	3869.4	5.338			
fit2e	13	3851.0	3879.2	26.140	20.802	2	3.04e-05 ***

## Measurement Invariance: Does a test measure the same thing

1. Across groups
  - Different schools
  - Different groups (e.g., ethnicity, age, gender)
2. Across time
  - Is today's measure the same as next year's measure?
3. Types of invariance
  - Configural: Are the arrows the same
  - Weak invariance: Are the loadings the same across groups
  - Strong invariance: Loadings and intercepts are equal across groups
  - Super strong: Loadings, intercepts and means are equal across groups

## Consider the Holzinger Swineford data set

1. 9 ability measures from two schools
  - 145 from Grant-White
  - 156 from Pasteur
2. Are the factor structures the same across schools
  - Although lavaan does this in one call, lets do it part by part
  - over all factor structure
  - factors with schools
  - constrain factors to have the same loadings, etc.

## First, some descriptive statistics

*describe (HolzingerSwineford1939 , skew=FALSE)*

	var	n	mean	sd	median	trimmed	mad	min	max	range	se
id	1	301	176.55	105.94	163.00	176.78	140.85	1.00	351.00	350.00	6.11
sex	2	301	1.51	0.50	2.00	1.52	0.00	1.00	2.00	1.00	0.03
ageyr	3	301	13.00	1.05	13.00	12.89	1.48	11.00	16.00	5.00	0.06
agemo	4	301	5.38	3.45	5.00	5.32	4.45	0.00	11.00	11.00	0.20
school*	5	301	1.52	0.50	2.00	1.52	0.00	1.00	2.00	1.00	0.03
grade	6	300	7.48	0.50	7.00	7.47	0.00	7.00	8.00	1.00	0.03
x1	7	301	4.94	1.17	5.00	4.96	1.24	0.67	8.50	7.83	0.07
x2	8	301	6.09	1.18	6.00	6.02	1.11	2.25	9.25	7.00	0.07
x3	9	301	2.25	1.13	2.12	2.20	1.30	0.25	4.50	4.25	0.07
x4	10	301	3.06	1.16	3.00	3.02	0.99	0.00	6.33	6.33	0.07
x5	11	301	4.34	1.29	4.50	4.40	1.48	1.00	7.00	6.00	0.07
x6	12	301	2.19	1.10	2.00	2.09	1.06	0.14	6.14	6.00	0.06
x7	13	301	4.19	1.09	4.09	4.16	1.10	1.30	7.43	6.13	0.06
x8	14	301	5.53	1.01	5.50	5.49	0.96	3.05	10.00	6.95	0.06
x9	15	301	5.37	1.01	5.42	5.37	0.99	2.78	9.25	6.47	0.06

## describeBy each group

```
> describeBy(HolzingerSwineford1939, group=HolzingerSwineford1939$school, sk
```

group: Grant-White

	var	n	mean	sd	median	trimmed	mad	min	max	range
<b>se</b>										
sex	2	145	1.50	0.50	2.00	1.50	0.00	1.00	2.00	1.00 0.00
ageyr	3	145	12.72	0.97	13.00	12.67	1.48	11.00	16.00	5.00 0.00
...										
grade	6	144	7.45	0.50	7.00	7.44	0.00	7.00	8.00	1.00 0.00
x1	7	145	4.93	1.15	5.00	4.96	1.24	1.83	8.50	6.67 0.10
x2	8	145	6.20	1.11	6.25	6.14	1.11	2.25	9.25	7.00 0.00
...										
x8	14	145	5.49	1.05	5.50	5.45	0.89	3.05	10.00	6.95 0.00
x9	15	145	5.33	1.03	5.31	5.33	1.15	3.11	9.25	6.14 0.00

group: Pasteur

	var	n	mean	sd	median	trimmed	mad	min	max	range
<b>se</b>										
sex	2	156	1.53	0.50	2.00	1.53	0.00	1.00	2.00	1.00 0.04
ageyr	3	156	13.25	1.06	13.00	13.15	1.48	12.00	16.00	4.00 0.09
...										
grade	6	156	7.50	0.50	7.50	7.50	0.74	7.00	8.00	1.00 0.04
x1	7	156	4.94	1.19	5.00	4.97	1.24	0.67	7.50	6.83 0.09
x2	8	156	5.98	1.23	5.75	5.89	1.11	3.50	9.25	5.75 0.10

## EFA for both groups

by(HolzingerSwineford1939[,7:15],HolzingerSwineford1939[,5],fa,nfactors

HolzingerSwineford1939[, 5]: Grant-White  
 Factor Analysis using method = minres

Call: FUN(r = data[x, , drop = FALSE], nfactors = 3, standardized loadings (pattern matrix) based upon

	MR1	MR2	MR3	h2	u2
x1	0.09	0.06	0.64	0.50	0.50
x2	0.02	-0.03	0.51	0.26	0.74
x3	0.11	-0.03	0.64	0.47	0.53
x4	0.86	-0.04	0.04	0.76	0.24
x5	0.82	0.09	-0.03	0.70	0.30
x6	0.81	-0.04	0.05	0.68	0.32
x7	0.14	0.78	-0.19	0.60	0.40
x8	-0.11	0.79	0.18	0.69	0.31
x9	0.08	0.46	0.40	0.54	0.46

	MR1	MR2	MR3
SS loadings	2.23	1.53	1.44
Proportion Var	0.25	0.17	0.16
Cumulative Var	0.25	0.42	0.58
Proportion Explained	0.43	0.29	0.28
Cumulative Proportion	0.43	0.72	1.00

With factor correlations of

	MR1	MR2	MR3
MR1	1.00	0.25	0.41
MR2	0.25	1.00	0.31
MR3	0.41	0.31	1.00

HolzingerSwineford1939[, 5]: Pasteur  
 Factor Analysis using method = minres

Call: FUN(r = data[x, , drop = FALSE], nfactors = 3, standardized loadings (pattern matrix) based upon

	MR1	MR2	MR3	h2	u2
x1	0.27	0.59	0.00	0.51	0.49
x2	0.03	0.49	-0.16	0.25	0.75
x3	-0.08	0.73	0.01	0.50	0.50
x4	0.80	0.02	0.06	0.68	0.32
x5	0.92	-0.07	-0.06	0.79	0.21
x6	0.78	0.13	0.06	0.70	0.30
x7	0.06	-0.14	0.71	0.52	0.48
x8	-0.02	0.13	0.60	0.39	0.61
x9	-0.02	0.37	0.40	0.34	0.66

	MR1	MR2	MR3
SS loadings	2.22	1.36	1.10
Proportion Var	0.25	0.15	0.12
Cumulative Var	0.25	0.40	0.52
Proportion Explained	0.48	0.29	0.23
Cumulative Proportion	0.48	0.77	1.00

With factor correlations of

	MR1	MR2	MR3
MR1	1.00	0.27	0.26
MR2	0.27	1.00	0.14
MR3	0.26	0.14	1.00

## How similar are the solutions: factor congruence

Factor congruence is the cosine of the angle between two vectors:

$$\text{Congruence} = (\text{diag}(X'X))^{-.5} Y' X (\text{diag}((Y'Y)))^{-.5}$$

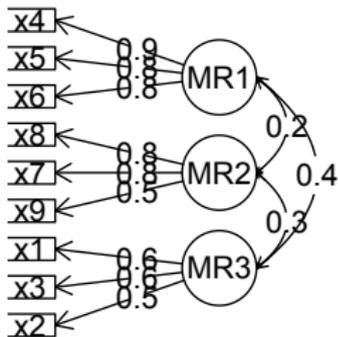
```
> f3.pasteur <- fa(HolzingerSwineford1939[1:156,7:15],3)
> f3.grant <- fa(HolzingerSwineford1939[157:301,7:15],3)
> factor.congruence(f3.pasteur, f3.grant)
```

	MR1	MR2	MR3
MR1	0.97	0.03	0.10
MR2	0.12	0.11	0.99
MR3	0.08	0.97	0.05

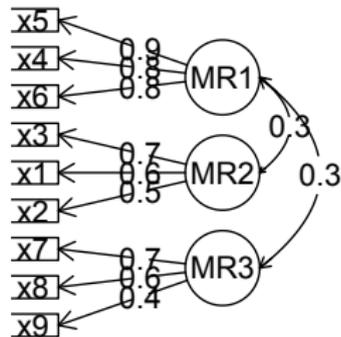
```
cross <- t(y) %*% x
sumsx <- sqrt(1/diag(t(x) %*% x))
sumsy <- sqrt(1/diag(t(y) %*% y))
```

## Do they look alike?

### Factor Analysis



### Factor Analysis



## Test if the model fits the combined data

```
HS.model <- 'visual =~ x1 + x2 + x3
            textual =~ x4 + x5 + x6
            speed =~ x7 + x8 + x9'
```

```
fit <- cfa(HS.model, data=HolzingerSwineford1939, std.lv=TRUE)
summary(fit, fit.measures=TRUE)
```

lavaan (0.4-14) converged normally after 41 iterations

Number of observations	301
Estimator	ML
Minimum Function Chi-square	85.306
Degrees of freedom	24
P-value	0.000

Chi-square test baseline **model**:

Minimum Function Chi-square	918.852
Degrees of freedom	36
P-value	0.000

Full **model** versus baseline **model**:

Comparative Fit Index (CFI)	0.931
Tucker-Lewis Index (TLI)	0.896

Loglikelihood and Information Criteria:

Loglikelihood user <b>model</b> (H0)	-3737.745
Loglikelihood unrestricted <b>model</b> (H1)	-3695.092

## With values of

Number of free parameters	21
Akaike (AIC)	7517.490
Bayesian (BIC)	7595.339
Sample-size adjusted Bayesian (BIC)	7528.739

### Root Mean Square Error of Approximation:

RMSEA	0.092
90 Percent Confidence Interval	0.071 0.114
P-value RMSEA <= 0.05	0.001

### Standardized Root Mean Square Residual:

SRMR	0.065
------	-------

### Parameter estimates:

Information	Expected			
Standard Errors	Standard			
Estimate	Std. err	Z-value	P(> z )	
<b>Latent variables:</b>				
visual =~				
x1	0.900	0.081	11.127	0.000
x2	0.498	0.077	6.429	0.000
x3	0.656	0.074	8.817	0.000
textual =~				
x4	0.990	0.057	17.474	0.000
x5	1.102	0.063	17.576	0.000
x6	0.917	0.054	17.082	0.000
speed =~				
x7	0.619	0.070	8.903	0.000
x8	0.731	0.066	11.090	0.000

## Now do it for both groups one analysis

```
> fit2 <- cfa(HW.model, data=HolzingerSwineford1939, group="school", std.lv=TRUE)
> summary(fit2)
```

Number of observations per group	
Pasteur	156
Grant-White	145
Estimator	ML
Minimum Function Chi-square	115.851
Degrees of freedom	48
P-value	0.000

Chi-square for each group:

Pasteur	64.309
Grant-White	51.542

Parameter estimates:

Information	Expected
Standard Errors	Standard

## Results for Pasteur

Group 1 [Pasteur]:

	Estimate	Std. err	Z-value	P(> z )
<b>Latent variables:</b>				
visual =~				
x1	1.047	0.132	7.934	0.000
x2	0.412	0.110	3.753	0.000
x3	0.597	0.108	5.525	0.000
textual =~				
x4	0.946	0.079	11.927	0.000
x5	1.119	0.089	12.604	0.000
x6	0.827	0.068	12.230	0.000
speed =~				
x7	0.591	0.106	5.557	0.000
x8	0.665	0.102	6.531	0.000
x9	0.545	0.097	5.596	0.000
<b>Covariances:</b>				
visual ~~				
textual	0.484	0.086	5.600	0.000
speed	0.299	0.109	2.755	0.006
textual ~~				
speed	0.325	0.100	3.256	0.001
<b>Intercepts:</b>				
x1	4.941	0.095	52.249	0.000
x2	5.984	0.098	60.949	0.000
x3	2.487	0.093	26.778	0.000
x4	2.823	0.092	30.689	0.000
x5	3.995	0.105	38.183	0.000
x6	1.922	0.079	24.321	0.000
x7	4.432	0.087	51.181	0.000
x8	5.563	0.078	71.214	0.000

## Results for Grant-White

### Latent variables:

visual =~

x1	0.777	0.103	7.525	0.000
x2	0.572	0.101	5.642	0.000
x3	0.719	0.093	7.711	0.000

textual =~

x4	0.971	0.079	12.355	0.000
x5	0.961	0.083	11.630	0.000
x6	0.935	0.081	11.572	0.000

speed =~

x7	0.679	0.087	7.819	0.000
x8	0.833	0.087	9.568	0.000
x9	0.719	0.086	8.357	0.000

### Covariances:

visual ~~

textual	0.541	0.085	6.355	0.000
speed	0.523	0.094	5.562	0.000

textual ~~

speed	0.336	0.091	3.674	0.000
-------	-------	-------	-------	-------

### Intercepts:

x1	4.930	0.095	51.696	0.000
x2	6.200	0.092	67.416	0.000
x3	1.996	0.086	23.195	0.000
x4	3.317	0.093	35.625	0.000
x5	4.712	0.096	48.986	0.000
x6	2.469	0.094	26.277	0.000
x7	3.921	0.086	45.819	0.000
x8	5.488	0.087	63.174	0.000
x9	5.327	0.085	62.571	0.000

visual

0.000

textual

0.000

## Constrain the loadings to be the same

```
> fit2e <- cfa(HW.model, data=HolzingerSwineford1939, group="school", std.lv=TRUE, group.equal="loadings")
> summary(fit2e)
```

lavaan (0.4-14) converged normally after 30 iterations

Number of observations per group	
Pasteur	156
Grant-White	145
Estimator	ML
Minimum Function Chi-square	127.834
Degrees of freedom	57
P-value	0.000

Chi-square for each group:

Pasteur	71.064
Grant-White	56.770

Parameter estimates:

Information	Expected
Standard Errors	Standard

## With parameter values of

Group 1 [Pasteur]:

	Estimate	Std. err	Z-value	P(> z )
Latent variables:				
visual =~				
x1	0.866	0.078	11.149	0.000
x2	0.523	0.076	6.916	0.000
x3	0.683	0.071	9.689	0.000
textual =~				
x4	0.954	0.056	17.002	0.000
x5	1.033	0.061	17.012	0.000
x6	0.870	0.052	16.750	0.000
speed =~				
x7	0.630	0.066	9.500	0.000
x8	0.752	0.065	11.586	0.000
x9	0.650	0.064	10.205	0.000
Covariances:				
visual ~~				
textual	0.485	0.087	5.555	0.000
speed	0.341	0.109	3.126	0.002
textual ~~				
speed	0.336	0.094	3.590	0.000
Intercepts:				
x1	4.941	0.092	53.661	0.000
x2	5.984	0.099	60.420	0.000
x3	2.487	0.093	26.734	0.000
x4	2.823	0.093	30.400	0.000
x5	3.995	0.100	39.756	0.000
x6	1.922	0.081	23.732	0.000
x7	4.432	0.089	49.903	0.000

## Now, compare the fit of the two group model to the one with equal parameters

```
> anova(fit2 , fit2e)
```

### Chi Square Difference Test

	Df	AIC	BIC	Chisq	Chisq	diff	Df	diff	Pr(>Chisq)
fit2	48	7484.4	7706.8	115.85					
fit2e	57	7478.4	7667.4	127.83	11.982		9		0.2143

## But, another way to specify the fit2e model – without making the latent variables standardized

```
fit2e <- cfa(HW.model, data=HolzingerSwinefeld1939, group="school", group.equal=c("loadings"))
```

Number of observations per group

Pasteur	156
Grant-White	145

Estimator	ML
Minimum Function Chi-square	124.044
Degrees of freedom	54
P-value	0.000

Chi-square for each group:

Pasteur	68.825
Grant-White	55.219

Parameter estimates:

Information	Expected
Standard Errors	Standard

Group 1 [Pasteur]:

	Estimate	Std. err	Z-value	P(> z )
Latent variables:				
visual =~				
x1	1.000			
x2	0.599	0.100	5.979	0.000
x3	0.784	0.108	7.267	0.000
textual =~				
x4	1.000			
x5	1.083	0.067	16.049	0.000
x6	0.912	0.058	15.785	0.000
speed =~				
x7	1.000			

## Test fit by comparing models

```
> anova(fit2 , fit2e)
```

Chi Square Difference Test

	Df	AIC	BIC	Chisq	Chisq	<b>diff</b>	Df	<b>diff</b>	Pr(>Chisq)
fit2	48	7484.4	7706.8	115.85					
fit2e	54	7480.6	7680.8	124.04	8.1922		6		0.2244

## Continue this logic, of successive tests with more constraints, but do it automatically

```
mi <- measurementInvariance(HW.model, data=HolzingerSwinefeld1939, group="school")
```

Measurement invariance tests:

Model 1: configural invariance:

chisq	df	pvalue	cfi	rmsea	bic
115.851	48.000	0.000	0.923	0.097	7706.822

Model 2: weak invariance (**equal** loadings):

chisq	df	pvalue	cfi	rmsea	bic
124.044	54.000	0.000	0.921	0.093	7680.771

[Model 1 versus **model 2**]

delta.chisq	delta.df	delta.p.value	delta.cfi
8.192	6.000	0.224	0.002

Model 3: strong invariance (**equal** loadings + intercepts):

chisq	df	pvalue	cfi	rmsea	bic
164.103	60.000	0.000	0.882	0.107	7686.588

[Model 1 versus **model 3**]

delta.chisq	delta.df	delta.p.value	delta.cfi
48.251	12.000	0.000	0.041

[Model 2 versus **model 3**]

delta.chisq	delta.df	delta.p.value	delta.cfi
40.059	6.000	0.000	0.038

Model 4: **equal** loadings + intercepts + means:

chisq	df	pvalue	cfi	rmsea	bic
204.605	63.000	0.000	0.840	0.122	7709.969

[Model 1 versus **model 4**]

delta.chisq	delta.df	delta.p.value	delta.cfi
88.754	15.000	0.000	0.083

[Model 3 versus **model 4**]

delta.chisq	delta.df	delta.p.value	delta.cfi
40.502	3.000	0.000	0.042

## Create a data set with non-invariant factor loadings

```

> set.seed(42)
> fx <- matrix(c(.8,.7,.6, rep(0,6), .6,.7,.8), ncol=2)
> fx
      [,1] [,2]
[1,]  0.8  0.0
[2,]  0.7  0.0
[3,]  0.6  0.0
[4,]  0.0  0.6
[5,]  0.0  0.7
[6,]  0.0  0.8
> Phi <- matrix(c(1,.6,.6,1), ncol=2)
> Phi
> set.seed(42)
> x.model <- sim(fx=fx, Phi=Phi, mu=c(0,1), n=250)
> x <- x.model$observed
> structure.diagram(fx, Phi, lr=FALSE, e.size=.3, main="A_basic_two_time_mod")
> describe(x, skew=FALSE)

```

	var	n	mean	sd	median	trimmed	mad	min	max	range	se
V1	1	250	0.02	0.99	0.01	0.03	1.02	-2.64	2.76	5.40	0.06
V2	2	250	-0.02	0.96	-0.01	-0.03	0.94	-2.66	3.12	5.78	0.06
V3	3	250	0.02	0.97	-0.06	0.01	0.95	-2.74	2.41	5.15	0.06
V4	4	250	0.61	0.99	0.59	0.65	0.99	-3.26	3.15	6.41	0.06

## One factor model

```
> f1n <- fa(x)
> f1n
```

Factor Analysis using method = minres

Call: fa(r = x)

Standardized loadings (pattern matrix) based upon

	MR1	h2	u2
V1	0.59	0.35	0.65
V2	0.56	0.31	0.69
V3	0.48	0.23	0.77
V4	0.54	0.29	0.71
V5	0.69	0.48	0.52
V6	0.72	0.52	0.48

	MR1
SS loadings	2.18
Proportion Var	0.36

Test of the hypothesis that 1 factor is sufficient.

The degrees of freedom for the null model are 15

and the objective function was

1.58

with Chi Square of 389.85

The degrees of freedom for the model are 9

and the objective function was

0.33

The root mean square of the residuals (RMSR) is 0.07

The df corrected root mean square of the residuals

is 0.13

The number of observations was 250

with Chi Square = 79.93 with prob < 1.7e-13

Tucker Lewis Index of factoring reliability = 0.684

RMSEA index = 0.179

and the 90 % confidence intervals are

0.143 0.214

BIC = 30.24

Fit based upon off diagonal values = 0.93

Measures of factor score adequacy

Correlation of scores with factors

0.89

Multiple R square of scores with factors

0.79

Minimum correlation of possible factor scores

0.57

MR

## Two factor model

```
> f2n <- fa(x,2)
> f2n
```

Factor Analysis using method = minres  
**Call:** fa(r = x, nfactors = 2)  
 Standardized loadings (pattern **matrix**) based upon correlation **matrix**

	MR1	MR2	h2	u2
V1	-0.01	0.79	0.62	0.38
V2	-0.01	0.73	0.53	0.47
V3	0.15	0.40	0.25	0.75
V4	0.51	0.07	0.30	0.70
V5	0.79	-0.03	0.60	0.40
V6	0.80	0.02	0.65	0.35

	MR1	MR2
SS loadings	1.58	1.37
Proportion Var	0.26	0.23
Cumulative Var	0.26	0.49
Proportion Explained	0.53	0.47
Cumulative Proportion	0.53	1.00

With **factor** correlations of

	MR1	MR2
MR1	1.00	0.54
MR2	0.54	1.00

Correlation of scores with factors  
 0.90 0.88  
 Multiple **R** square of scores with factors  
 0.80 0.77  
 Minimum correlation of possible **factor** scores

Test of the hypothesis that 2 factors are sufficient  
 The degrees of freedom **for** the **null model** are 15  
 and the objective **function** was 1.58  
 with Chi Square of 389.85  
 The degrees of freedom **for** the **model** are 4  
 and the objective **function** was 0.02  
 The root **mean square** of the **residuals** (RMSR) is 0.01  
 The **df** corrected root **mean square** of the **residuals** is 0.04  
 The number of observations was 250 with Chi Square = 4.7  
 with prob < 0.32  
 Tucker Lewis Index of factoring reliability = 0.993  
 RMSEA **index** = 0.028  
 and the 90 % confidence intervals are NA 0.102  
 BIC = -17.39  
 Fit based upon **off** diagonal values = 1  
 Measures of **factor** score adequacy

## Compare the two solutions

Factor Analysis using method = minres

**Call:** fa(r = x)

Standardized loadings (pattern **matrix**)  
based upon correlation **matrix**

	MR1	h2	u2
V1	0.59	0.35	0.65
V2	0.56	0.31	0.69
V3	0.48	0.23	0.77
V4	0.54	0.29	0.71
V5	0.69	0.48	0.52
V6	0.72	0.52	0.48

	MR1
SS loadings	2.18
Proportion Var	0.36

Factor Analysis using method = minres

**Call:** fa(r = x, nfactors = 2)

Standardized loadings (pattern **matrix**)  
based upon correlation **matrix**

	MR1	MR2	h2	u2
V1	-0.01	0.79	0.62	0.38
V2	-0.01	0.73	0.53	0.47
V3	0.15	0.40	0.25	0.75
V4	0.51	0.07	0.30	0.70
V5	0.79	-0.03	0.60	0.40
V6	0.80	0.02	0.65	0.35

	MR1	MR2
SS loadings	1.58	1.37
Proportion Var	0.26	0.23
Cumulative Var	0.26	0.49
Proportion Explained	0.53	0.47
Cumulative Proportion	0.53	1.00

With **factor** correlations of

	MR1	MR2
MR1	1.00	0.54
MR2	0.54	1.00

## CFA 2 correlated factors

```
> y.df <- data.frame(y)
> fitn <- sem(mod2f, data=y.df, std.lv=TRUE)
> summary(fitn, fit.measures=TRUE)
```

lavaan (0.4-14) converged normally after 17 iterations

Number of observations  
250

Estimator

ML

Minimum Function Chi-square  
2.552

Degrees of freedom  
8

P-value  
0.959

Chi-square test baseline **model**:

Minimum Function Chi-square  
341.309

Degrees of freedom  
15

P-value  
0.000

Full **model** versus baseline **model**:

Comparative Fit Index (CFI)  
1.000

Tucker-Lewis Index (TLI)  
1.031

Root Mean Square Error of Approximation:

RMSEA  
0.000

90 Percent Confidence Interval  
0.000 0.000

P-value RMSEA <= 0.05  
0.994

Standardized Root Mean Square Residual:

SRMR  
0.014

Parameter estimates:

Information

Expected  
Standard Errors  
Standard

Z-value	P(> z )	Estimate	Std. err
Latent variables:			
F1 =~			
V1		0.750	0.063
11.880	0.000		
V2		0.725	0.065
11.160	0.000		
V3		0.621	0.066
9.410	0.000		
F2 =~			
V4		0.705	0.074
11.160	0.000		

## Are the factors measurement invariant? Well, maybe.

```

> summary(fitn2a, fit.measures=TRUE)

lavaan (0.4-14) converged normally after 14 iterations

  Number of observations
250

  Estimator
ML
  Minimum Function Chi-square
8.093
  Degrees of freedom
11
  P-value
0.705

Chi-square test baseline model:

  Minimum Function Chi-square
341.309
  Degrees of freedom
15
  P-value
0.000

Full model versus baseline model:

  Comparative Fit Index (CFI)
1.000
  Tucker-Lewis Index (TLI)

```

Root Mean Square Error of Approximation:

```

RMSEA
0.000
90 Percent Confidence Interval
0.000 0.051
P-value RMSEA <= 0.05
0.946

```

Standardized Root Mean Square Residual:

```

SRMR
0.049

```

Parameter estimates:

Information		Estimate	Std. err
Expected Standard Errors			
Z-value	P(> z )		
Latent variables:			
F1 =~			
V1	(a)	0.686	0.050
13.774	0.000		
V2	(b)	0.675	0.049
13.834	0.000		
V3	(c)	0.657	0.048
13.677	0.000		
F2 =~			
	( )	0.000	0.000

## Create the original data set again

```
> fx <- matrix(c(.8,.7,.6,rep(0,6),.8,.7,.6),ncol=2)
> x.model <- sim(fx=fx,Phi=Phi,mu=c(0,1),n=250)
> x <- x.model$observed
> x.df <- data.frame(x)
```

```
> describe(x,skew=FALSE)
```

	var	n	mean	sd	median	trimmed	mad	min	max	range	se
V1	1	250	0.02	0.99	-0.01	-0.01	1.04	-2.38	3.26	5.64	0.06
V2	2	250	0.13	1.07	0.03	0.11	1.08	-2.53	3.65	6.19	0.07
V3	3	250	0.03	1.03	0.05	0.02	0.97	-2.68	3.12	5.80	0.07
V4	4	250	0.87	1.04	0.90	0.85	1.03	-1.63	3.61	5.24	0.07
V5	5	250	0.73	0.97	0.76	0.75	0.90	-2.36	3.29	5.65	0.06
V6	6	250	0.65	1.00	0.67	0.64	1.03	-2.58	3.17	5.75	0.06

## Model change in the items

```

mod2fc <- 'F1_~_a*V1_+_b*_V2_+_c*V3
          F2_~_a*_V4_+_b*V5_+c*_V6
          #correlation_between_factors
          F2_~_F1_' #the regression
fit2c <- sem(mod2fc, data=x.df, meanstructure=TRUE)
summary(fit2c, fit.measures=TRUE)

```

## Correlated factors

Root Mean Square Error of Approximation:

RMSEA  
 0.016  
 90 Percent Confidence Interval  
 0.000 0.071  
 P-value RMSEA <= 0.05  
 0.793

lavaan (0.4-14) converged normally after 23 iterations

Number of observations  
 250

Estimator  
 ML  
 Minimum Function Chi-square  
 10.616  
 Degrees of freedom  
 10  
 P-value  
 0.388

Standardized Root Mean Square Residual:

SRMR  
 0.032

Parameter estimates:

Information  
 Expected  
 Standard Errors  
 Standard

Chi-square test baseline **model**:

Minimum Function Chi-square  
 406.795  
 Degrees of freedom  
 15  
 P-value  
 0.000

Z-value P(>|z|) Estimate Std. err

Latent variables:

F1 =~  
 V1 (a) 1.000  
 V2 (b) 0.925 0.076  
 V3 (c) 0.816 0.072  
 12.243 0.000  
 11.386 0.000

Full **model** versus baseline **model**:

F2 =~  
 V4 (a) 1.000  
 V5 (b) 0.925 0.076

Comparative Fit Index (CFI)

## With the intercepts

**Intercepts :**

V1		0.021	0.063
0.332	0.740		
V2		0.135	0.066
2.033	0.042		
V3		0.032	0.066
0.482	0.630		
V4		0.865	0.065
13.294	0.000		
V5		0.729	0.062
11.696	0.000		
V6		0.649	0.063
10.369	0.000		
F1		0.000	
F2		0.000	

**Variances :**

V1	0.383	0.060
V2	0.576	0.068
V3	0.670	0.071
V4	0.486	0.065
V5	0.481	0.061
V6	0.596	0.065
F1	0.613	0.088
F2	0.319	0.063

## Try fitting a moments model

```

mod2fc <- 'F1_~_a*V1_+_b*_V2_+_c*V3
           F2_~_a*_V4_+_b*V5_+_c*_V6
           means_=_~_F1_+_F2_+_1*V1_+_1*V2+_1*V3+_1*V4+_1*V5+_1*V6
           #correlation_between_factors

           F2_~_F1_' #the regression
fit2c <- sem(mod2fc, data=x.df, meanstructure=TRUE)
summary(fit2c, fit.measures=TRUE)

```

# Using the moments matrix

## Modeling change using the moments matrix

1. McArdle (2009) Latent variable modeling of differences and changes with longitudinal data. *Annual Review of Psychology*, 60, 577-605
  - Use moments rather than covariances

## Create the model to be fit in sem

```

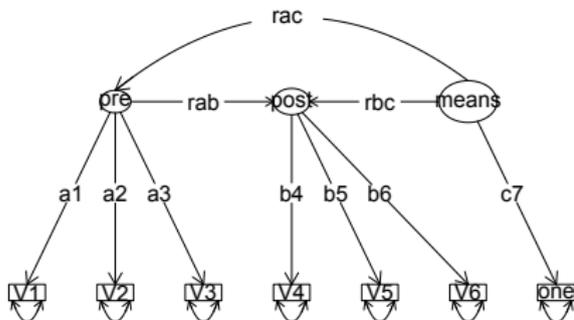
fxg
  pre  post means
V1 "a1" "0" "0"
V2 "a2" "0" "0"
V3 "a3" "0" "0"
V4 "0"  "b4" "0"
V5 "0"  "b5" "0"
V6 "0"  "b6" "0"
one "0"  "0"  "c7"
> phi
  F1  F2  F3
F1 "1"  "0" "rac"
F2 "rab" "1" "rbc"
F3 "0"  "0" "1"

mod.mom1 <- structure.diagram(fog, phi, errors=TRUE)

```

# Modeling the means in a moments matrix

## Structural model



## the basic path model, with some editing

mod.mom1

	Path	Parameter	Value	
[1, ]	"pre->V1"	"a1"	NA	
[2, ]	"pre->V2"	"a2"	NA	
[3, ]	"pre->V3"	"a3"	NA	
[4, ]	"post->V4"	"b4"	NA	
[5, ]	"post->V5"	"b5"	NA	
[6, ]	"post->V6"	"b6"	NA	
[7, ]	"means->one"	"c7"	NA	
[8, ]	"V1<->V1"	"x1e"	NA	
[9, ]	"V2<->V2"	"x2e"	NA	
[10, ]	"V3<->V3"	"x3e"	NA	
[11, ]	"V4<->V4"	"x4e"	NA	
[12, ]	"V5<->V5"	"x5e"	NA	
[13, ]	"V6<->V6"	"x6e"	NA	
[14, ]	"one<->one"	NA	"1"	<-- edited
[15, ]	"pre_>post"	"rF1F2"	NA	
[16, ]	"means->pre"	"rF3F1"	NA	<- edited
[17, ]	"means->post"	"rF3F2"	NA	<- edited
[18, ]	"pre<->pre"	NA	"1"	
[19, ]	"post<->post"	NA	"1"	
[20, ]	"means<->means"	NA	"1"	
<b>attr</b> (, "class")				
[1]	"mod"			

## sem output

### Parameter Estimates

	Estimate	Std Error	z value	Pr(> z )
a1	0.830111	0.069823	11.8888	
1.3536e-32 V1 <--- pre				
a2	0.881776	0.076506	11.5256	
9.7982e-31 V2 <--- pre				
a3	0.698963	0.074824	9.3415	
9.5005e-21 V3 <--- pre				
TRUE 0.664838 0.062395 10.6553				
1.6468e-26 V4 <--- post				
b5	0.554498	0.053510	10.3626	
3.6687e-25 V5 <--- post				
b6	0.510270	0.051499	9.9084	
3.8265e-23 V6 <--- post				
c7	1.118034	0.050000		
22.3607 9.5054e-111 one <--- means				
x1e	0.525613	0.078258	6.7164	
1.8623e-11 V1 <--> V1				
x2e	0.673405	0.093381	7.2114	
5.5387e-13 V2 <--> V2				
x3e	0.836504	0.090362	9.2572	
2.0983e-20 V3 <--> V3				
x4e	0.567286	0.081353	6.9732	
3.0990e-12 V4 <--> V4				
x5e	0.628041	0.073463	8.5490	
1.2412e-17 V5 <--> V5				
x6e	0.750522	0.080300	9.3465	
9.0592e-21 V6 <--> V6				
rF1F2	0.893184	0.142998	6.2461	
4.2076e-10 post <--- pre				
rF3F1	0.086583	0.073172	1.1833	
2.3669e-01 pre <--- means				

> sem.mom1 <- sem(mod.mom, MomMat, N=250, raw=TRUE)

> summary(sem.mom1)

Model fit to raw moment **matrix**.

Model Chisquare = 12.077 Df =

12

Pr(>Chisq) = 0.43954

AIC = 44.077

AICc = 14.411

BIC = 100.42

CAIC = -66.181

Normalized Residuals

Min.	1st Qu.	Median	Max.
-1.2500	-0.1150	0.0000	-0.0103
0.0972	0.9730		

## Traits and States and time

1. With just two time points, traits and states are confounded
  - Is the correlation a trait like stability
  - or does the state dissipate slowly?
2. With  $> 2$  time points we can distinguish states and traits
  - States should have an autocorrelation component
  - Traits should be consistent across time
3. Consider the simplex structure of 4 time points
  - Clean within time factor structure
  - Simplex across time points

## A factor simplex

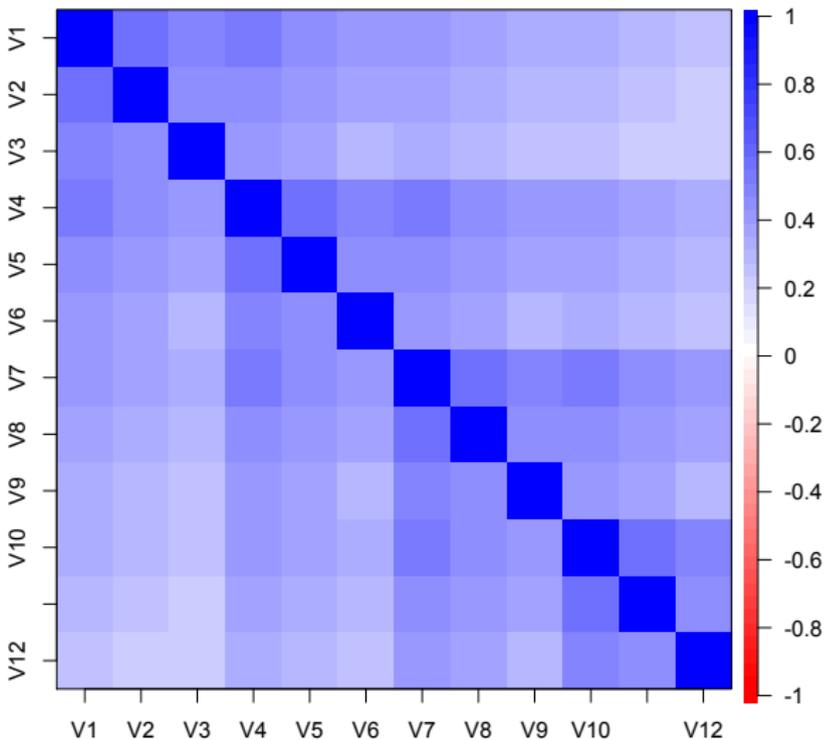
```
simp <- sim()
```

```
$model (Population correlation matrix)
```

	V1	V2	V3	V4	V5	V6	V7	V8	V9	V10	V11	V12
V1	1.00	0.56	0.48	0.51	0.45	0.38	0.41	0.36	0.31	0.33	0.29	0.25
V2	0.56	1.00	0.42	0.45	0.39	0.34	0.36	0.31	0.27	0.29	0.25	0.22
V3	0.48	0.42	1.00	0.38	0.34	0.29	0.31	0.27	0.23	0.25	0.22	0.18
V4	0.51	0.45	0.38	1.00	0.56	0.48	0.51	0.45	0.38	0.41	0.36	0.31
V5	0.45	0.39	0.34	0.56	1.00	0.42	0.45	0.39	0.34	0.36	0.31	0.27
V6	0.38	0.34	0.29	0.48	0.42	1.00	0.38	0.34	0.29	0.31	0.27	0.23
V7	0.41	0.36	0.31	0.51	0.45	0.38	1.00	0.56	0.48	0.51	0.45	0.38
V8	0.36	0.31	0.27	0.45	0.39	0.34	0.56	1.00	0.42	0.45	0.39	0.34
V9	0.31	0.27	0.23	0.38	0.34	0.29	0.48	0.42	1.00	0.38	0.34	0.29
V10	0.33	0.29	0.25	0.41	0.36	0.31	0.51	0.45	0.38	1.00	0.56	0.48
V11	0.29	0.25	0.22	0.36	0.31	0.27	0.45	0.39	0.34	0.56	1.00	0.42
V12	0.25	0.22	0.18	0.31	0.27	0.23	0.38	0.34	0.29	0.48	0.42	1.00

# A simplex

## Correlation plot



## Factor structure of a simplex

```
> fsimp <- fa(simp$model)
> fsimp
```

Factor Analysis using method = minres

**Call:** fa(r = simp\$model)

Standardized loadings (pattern **matrix**) based upon correlation **matrix**

	MR1	h2	u2
V1	0.64	0.41	0.59
V2	0.57	0.33	0.67
V3	0.49	0.24	0.76
V4	0.73	0.54	0.46
V5	0.65	0.42	0.58
V6	0.56	0.31	0.69
V7	0.73	0.54	0.46
V8	0.65	0.42	0.58
V9	0.56	0.31	0.69
V10	0.64	0.41	0.59
V11	0.57	0.33	0.67
V12	0.49	0.24	0.76

	MR1
SS loadings	4.50
Proportion Var	0.38

## Factors over time

```
> fsimp4 <- fa(simp$model, 4)
> fsimp4
```

Factor Analysis using method = minres

**Call:** fa(r = simp\$model, nfactors = 4)

Standardized loadings (pattern **matrix**)  
based upon correlation **matrix**

	MR3	MR1	MR2	MR4	h2	u2
V1	0.8	0.0	0.0	0.0	0.64	0.36
V2	0.7	0.0	0.0	0.0	0.49	0.51
V3	0.6	0.0	0.0	0.0	0.36	0.64
V4	0.0	0.0	0.0	0.8	0.64	0.36
V5	0.0	0.0	0.0	0.7	0.49	0.51
V6	0.0	0.0	0.0	0.6	0.36	0.64
V7	0.0	0.8	0.0	0.0	0.64	0.36
V8	0.0	0.7	0.0	0.0	0.49	0.51
V9	0.0	0.6	0.0	0.0	0.36	0.64
V10	0.0	0.0	0.8	0.0	0.64	0.36
V11	0.0	0.0	0.7	0.0	0.49	0.51
V12	0.0	0.0	0.6	0.0	0.36	0.64

	MR3	MR1	MR2	MR4
SS loadings	1.49	1.49	1.49	1.49
Proportion Var	0.12	0.12	0.12	0.12
Cumulative Var	0.12	0.25	0.37	0.50
Proportion Explained	0.25	0.25	0.25	0.25
Cumulative Proportion	0.25	0.50	0.75	1.00

With **factor** correlations of

	MR3	MR1	MR2	MR4
MR3	1.00	0.64	0.51	0.80
MR1	0.64	1.00	0.80	0.80
MR2	0.51	0.80	1.00	0.64
MR4	0.80	0.80	0.64	1.00

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