

SEM & Lavaan

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1. Fit a measurement model to your constructs at one time point. Try out the different types of scaling discussed in class. What changes what stays the same?

Measurement model was fit at timepoint 1. Under the marker method, parameter estimates were fixed at 1 for the first indicators (i.e., Animal_TW_Clusters_1 and Animal_Switches_1). Under the fixed factor method, in contrast, parameter estimates were fixed to 1 for the latent variables. In both cases, fit indices (e.g., logLikelihood, TLI, CFI, RMSEA) remain constant.

```
T1.mod <- '
    Semantic =~ Animal_TW_Clusters_1 + Food_TW_Clusters_1 + IQ_vocraw_1
    Phonemic =~ Animal_Switches_1 + Food_Switches_1 + IQ_mrraw_1
'
```

#Marker method

```
fit.marker <- cfa(T1.mod, data=mydata_wide)
summary(fit.marker, fit.measures = TRUE)
```

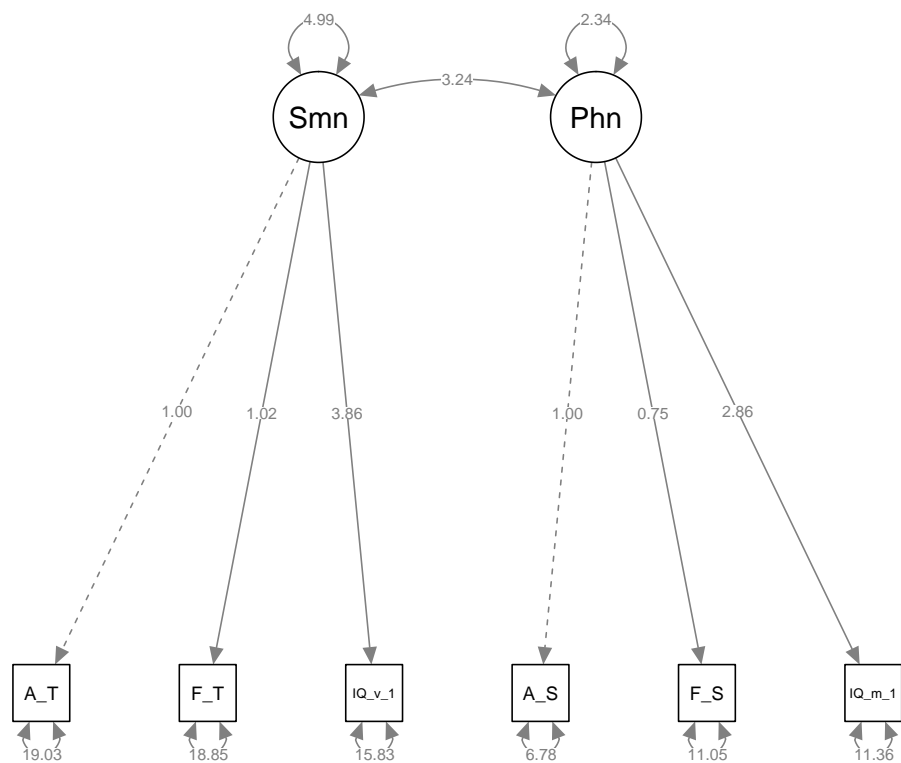
```
## lavaan (0.5-23.1097) converged normally after 53 iterations
##
##   Number of observations                67
##
##   Estimator                            ML
##   Minimum Function Test Statistic      22.669
##   Degrees of freedom                   8
##   P-value (Chi-square)                 0.004
##
## Model test baseline model:
##
##   Minimum Function Test Statistic      114.013
##   Degrees of freedom                   15
##   P-value                             0.000
##
## User model versus baseline model:
##
##   Comparative Fit Index (CFI)          0.852
##   Tucker-Lewis Index (TLI)            0.722
##
## Loglikelihood and Information Criteria:
##
##   Loglikelihood user model (H0)        -1161.512
##   Loglikelihood unrestricted model (H1) -1150.178
##
##   Number of free parameters            13
##   Akaike (AIC)                         2349.024
##   Bayesian (BIC)                       2377.685
##   Sample-size adjusted Bayesian (BIC)  2336.753
##
```

```

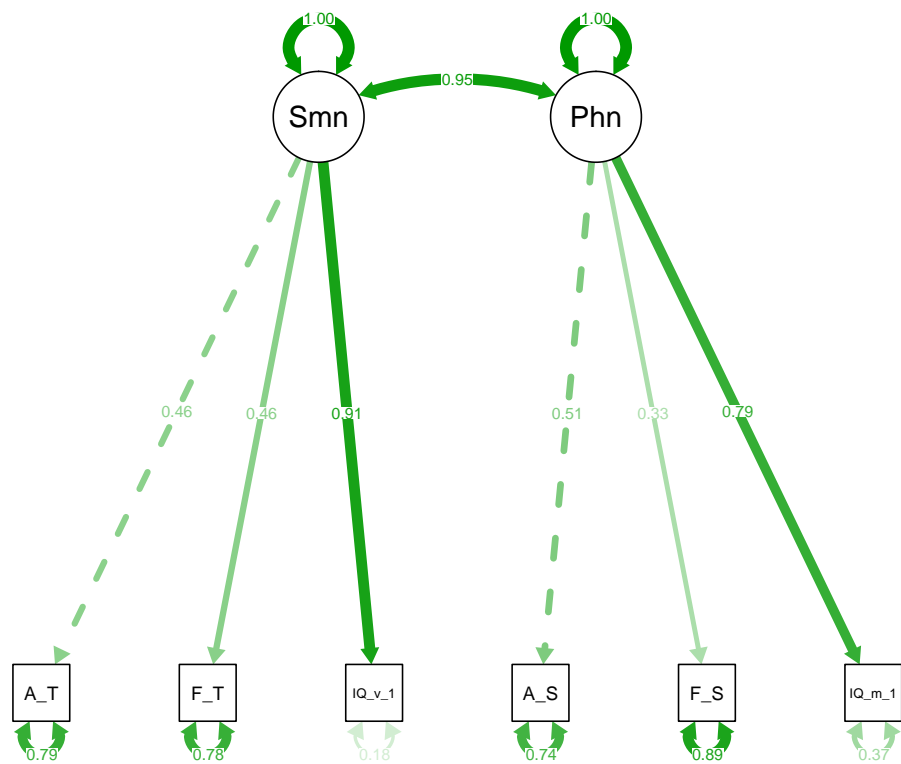
## Root Mean Square Error of Approximation:
##
##   RMSEA                                0.165
##   90 Percent Confidence Interval      0.087  0.248
##   P-value RMSEA <= 0.05              0.012
##
## Standardized Root Mean Square Residual:
##
##   SRMR                                0.091
##
## Parameter Estimates:
##
##   Information                        Expected
##   Standard Errors                    Standard
##
## Latent Variables:
##           Estimate  Std.Err  z-value  P(>|z|)
##   Semantic =~
##     Anml_TW_Clst_1    1.000
##     Fd_TW_Clst_1     1.019    0.366    2.786    0.005
##     IQ_vocraw_1      3.862    1.109    3.483    0.000
##   Phonemic =~
##     Animl_Swtchs_1    1.000
##     Food_Swtchs_1     0.755    0.338    2.231    0.026
##     IQ_mrraw_1       2.864    0.743    3.854    0.000
##
## Covariances:
##           Estimate  Std.Err  z-value  P(>|z|)
##   Semantic ~~
##     Phonemic         3.244    1.311    2.474    0.013
##
## Variances:
##           Estimate  Std.Err  z-value  P(>|z|)
##   .Anml_TW_Clst_1   19.027    3.439    5.533    0.000
##   .Fd_TW_Clst_1     18.855    3.416    5.519    0.000
##   .IQ_vocraw_1      15.830   11.974    1.322    0.186
##   .Animl_Swtchs_1    6.776    1.261    5.373    0.000
##   .Food_Swtchs_1    11.048    1.955    5.651    0.000
##   .IQ_mrraw_1       11.360    3.959    2.869    0.004
##   Semantic          4.991    2.726    1.831    0.067
##   Phonemic          2.343    1.154    2.030    0.042

```

```
semPaths(fit.marker, layout = "tree", whatLabels = "est")
```



```
semPaths(fit.marker, layout = "tree", what = "std")
```



```
#Fixed factor method
fit.fixed <- cfa(T1.mod, data=mydata_wide, std.lv = T)
summary(fit.fixed, fit.measures = TRUE)
```

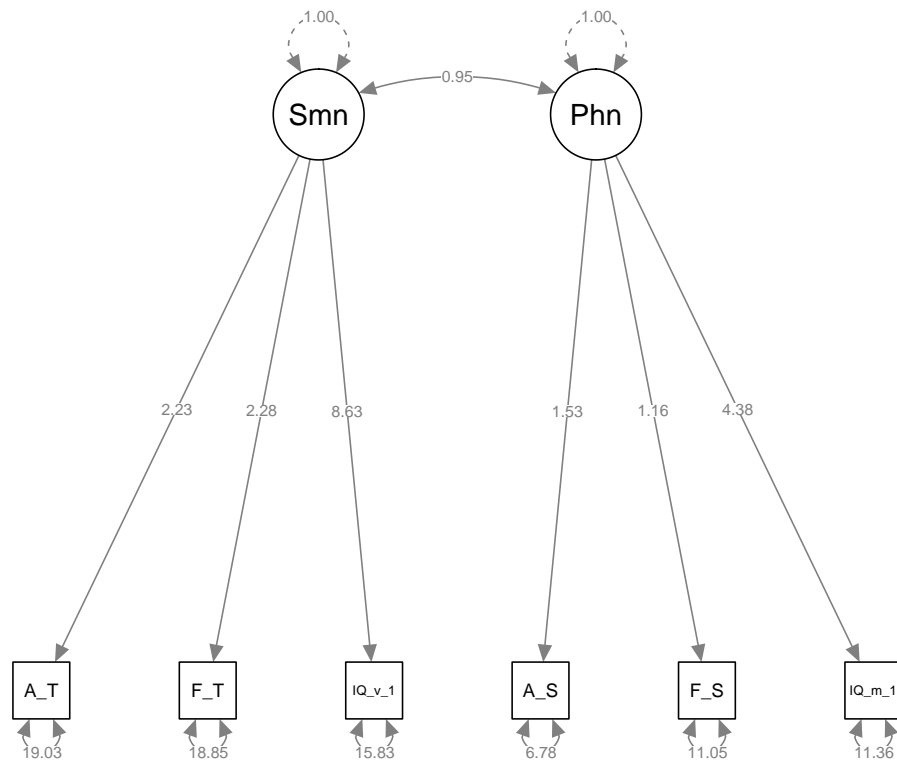
```

## lavaan (0.5-23.1097) converged normally after 48 iterations
##
##   Number of observations                67
##
##   Estimator                            ML
##   Minimum Function Test Statistic      22.669
##   Degrees of freedom                   8
##   P-value (Chi-square)                 0.004
##
## Model test baseline model:
##
##   Minimum Function Test Statistic      114.013
##   Degrees of freedom                   15
##   P-value                             0.000
##
## User model versus baseline model:
##
##   Comparative Fit Index (CFI)          0.852
##   Tucker-Lewis Index (TLI)           0.722
##
## Loglikelihood and Information Criteria:
##
##   Loglikelihood user model (H0)        -1161.512
##   Loglikelihood unrestricted model (H1) -1150.178
##
##   Number of free parameters            13
##   Akaike (AIC)                        2349.024
##   Bayesian (BIC)                      2377.685
##   Sample-size adjusted Bayesian (BIC)  2336.753
##
## Root Mean Square Error of Approximation:
##
##   RMSEA                                0.165
##   90 Percent Confidence Interval        0.087 0.248
##   P-value RMSEA <= 0.05                0.012
##
## Standardized Root Mean Square Residual:
##
##   SRMR                                0.091
##
## Parameter Estimates:
##
##   Information                          Expected
##   Standard Errors                      Standard
##
## Latent Variables:
##
##           Estimate  Std.Err  z-value  P(>|z|)
## Semantic =~
##   Anml_TW_Clst_1    2.234    0.610    3.661    0.000
##   Fd_TW_Clst_1      2.276    0.609    3.736    0.000
##   IQ_vocraw_1       8.629    1.117    7.723    0.000
## Phonemic =~
##   Animl_Swtchs_1    1.531    0.377    4.060    0.000
##   Food_Swtchs_1     1.155    0.456    2.531    0.011

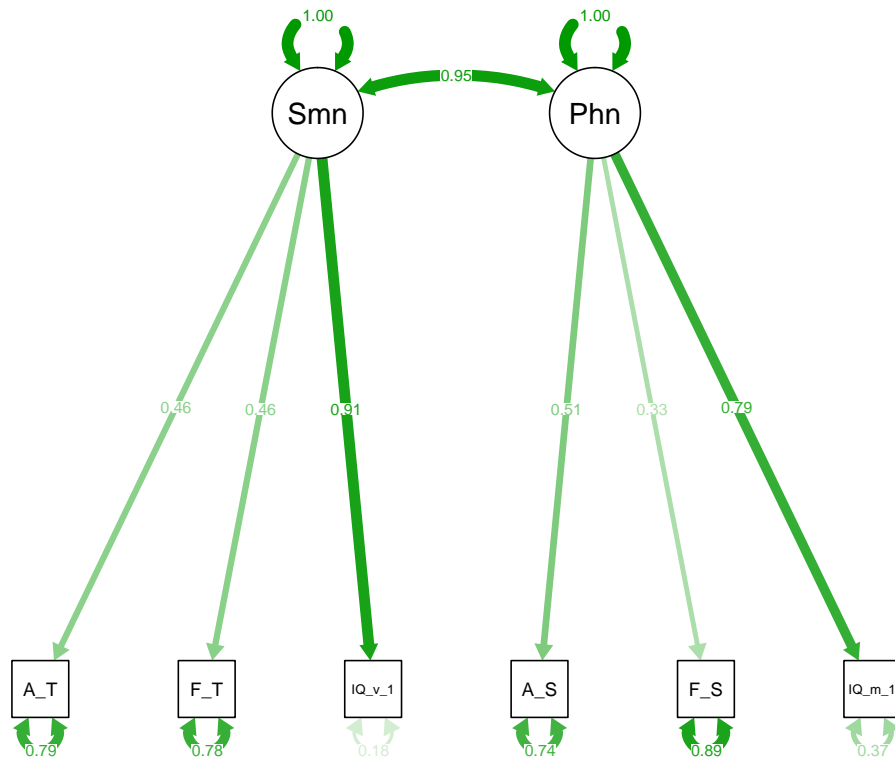
```

```
##      IQ_mrraw_1      4.383    0.683    6.416    0.000
##
## Covariances:
##              Estimate Std.Err  z-value  P(>|z|)
## Semantic ~~
## Phonemic      0.949    0.108    8.814    0.000
##
## Variances:
##              Estimate Std.Err  z-value  P(>|z|)
## .Anml_TW_Clst_1  19.027    3.439    5.533    0.000
## .Fd_TW_Clst_1    18.855    3.416    5.519    0.000
## .IQ_vocraw_1     15.830   11.974    1.322    0.186
## .Animl_Swtchs_1   6.776    1.261    5.373    0.000
## .Food_Swtchs_1   11.048    1.955    5.651    0.000
## .IQ_mrraw_1      11.360    3.959    2.869    0.004
## Semantic         1.000
## Phonemic         1.000
```

```
semPaths(fit.fixed, layout = "tree", whatLabels = "est")
```



```
semPaths(fit.fixed, layout = "tree", what = "std")
```



2. What do the fit statistics say about your latent variable? Good/bad? Is your latent variable Just identified/saturated, under identified or over identified?

- RMSEA = .165, SRMR = .091, TLI = .722, CFI = .852
- RMSEA & SRMR > .08 and TLI & CFI < .90, suggesting poor fit – that is, the latent variables are not effectively capturing commonalities among their indicator variables. This could be due to (1) high measurement error or (2) highly disparate indicators
- This model is over identified, as evidenced by the positive degrees of freedom (15).

3. Fit a longitudinal CFA model where you a) first correlate your latent factors across time and then b) a second model that predicts later times by a previous time (ie autoregressive; t1 -> t2 -> t3). What are your conclusions? How does one differ from the other?

For the longitudinal CFA model with correlated latent factors (Long.mod), I conclude that my latent factors are strongly correlated across time. Moreover, across all three timepoints, IQ verbal loads much more strongly onto the latent factors than Animal and Food total word counts, suggesting that IQ verbal may parsimoniously account for shared variance among indicators.

For the autoregressive CFA model (auto.mod), a similar picture emerges, suggesting that Semantic_1 is highly predictive of Semantic_2, which is highly predictive of Semantic_3. Of note, standardized variances for S_2 and S_3 latent factors are reduced for the autoregressive model compared to the correlated model (see path diagrams), likely due to the fact that S_2 and S_3 variances are accounted for by earlier timepoints.

```
Long.mod <- '
    Semantic_1 =~ Animal_TW_Clusters_1 + Food_TW_Clusters_1 + IQ_vocraw_1
```

```

        Semantic_2 =~ Animal_TW_Clusters_2 + Food_TW_Clusters_2 + IQ_vocraw_2
        Semantic_3 =~ Animal_TW_Clusters_3 + Food_TW_Clusters_3 + IQ_vocraw_3
,

fit.long <- cfa(Long.mod, data=mydata_wide, std.lv = T)
summary(fit.long)

```

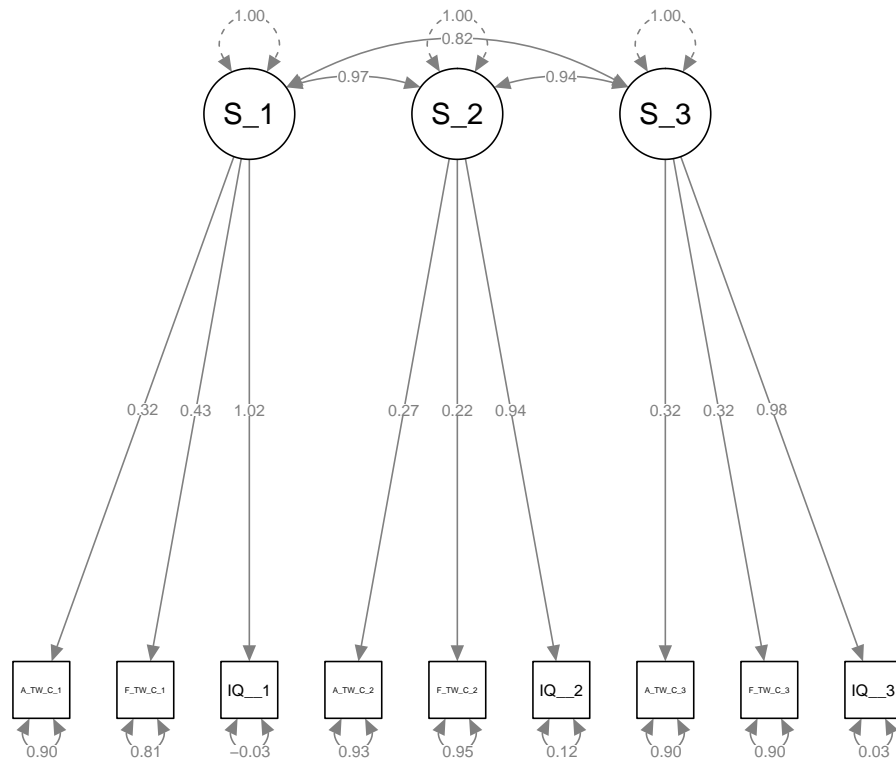
```
## lavaan (0.5-23.1097) converged normally after 103 iterations
```

```
##
##                                     Used      Total
##   Number of observations                64        67
##
##   Estimator                          ML
##   Minimum Function Test Statistic      91.983
##   Degrees of freedom                   24
##   P-value (Chi-square)                 0.000
##
## Parameter Estimates:
##
##   Information                        Expected
##   Standard Errors                   Standard
##
## Latent Variables:
##           Estimate  Std.Err  z-value  P(>|z|)
##   Semantic_1 =~
##     Anml_TW_Clst_1    1.508    0.565    2.669    0.008
##     Fd_TW_Clstrs_1    2.148    0.593    3.622    0.000
##     IQ_vocraw_1       9.609    0.917   10.479    0.000
##   Semantic_2 =~
##     Anml_TW_Clst_2    1.209    0.553    2.187    0.029
##     Fd_TW_Clstrs_2    1.107    0.610    1.814    0.070
##     IQ_vocraw_2       8.025    0.931    8.617    0.000
##   Semantic_3 =~
##     Anml_TW_Clst_3    1.422    0.560    2.539    0.011
##     Fd_TW_Clstrs_3    1.774    0.702    2.528    0.011
##     IQ_vocraw_3       8.726    1.024    8.523    0.000
##
## Covariances:
##           Estimate  Std.Err  z-value  P(>|z|)
##   Semantic_1 ~~
##     Semantic_2         0.971    0.072   13.413    0.000
##     Semantic_3         0.822    0.080   10.220    0.000
##   Semantic_2 ~~
##     Semantic_3         0.942    0.092   10.283    0.000
##
## Variances:
##           Estimate  Std.Err  z-value  P(>|z|)
##   .Anml_TW_Clst_1   19.557    3.452    5.665    0.000
##   .Fd_TW_Clstrs_1   19.929    3.534    5.639    0.000
##   .IQ_vocraw_1      -3.049    7.873   -0.387    0.699
##   .Anml_TW_Clst_2   18.646    3.299    5.652    0.000
##   .Fd_TW_Clstrs_2   23.199    4.101    5.656    0.000
##   .IQ_vocraw_2       8.726    7.815    1.117    0.264
##   .Anml_TW_Clst_3   18.073    3.214    5.623    0.000

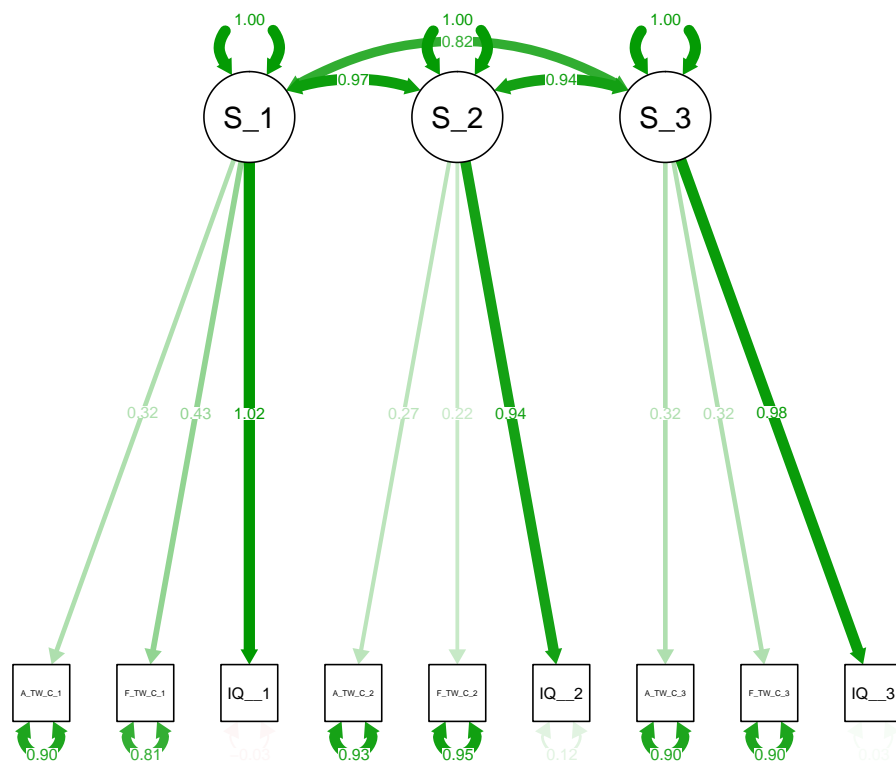
```

```
## .Fd_TW_Clstrs_3 28.428 5.055 5.623 0.000
## .IQ_vocraw_3 2.694 11.201 0.241 0.810
## Semantic_1 1.000
## Semantic_2 1.000
## Semantic_3 1.000
```

```
semPaths(fit.long, whatLabels = "std")
```



```
semPaths(fit.long, what = "std")
```

```
Auto.mod <- '
    Semantic_1 =~ Animal_TW_Clusters_1 + Food_TW_Clusters_1 + IQ_vocraw_1
    Semantic_2 =~ Animal_TW_Clusters_2 + Food_TW_Clusters_2 + IQ_vocraw_2
    Semantic_3 =~ Animal_TW_Clusters_3 + Food_TW_Clusters_3 + IQ_vocraw_3

    Semantic_3 ~ Semantic_2
    Semantic_2 ~ Semantic_1
'
```

```
fit.auto <- cfa(Auto.mod, data=mydata_wide, std.lv = T)
summary(fit.auto)
```

```
## lavaan (0.5-23.1097) converged normally after 81 iterations
```

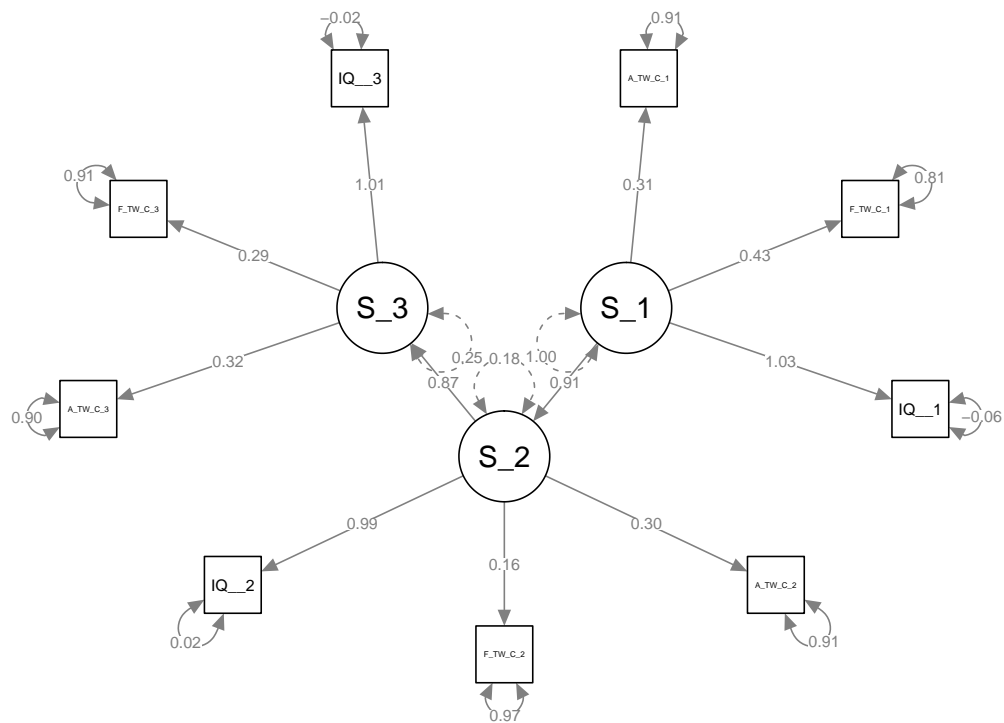
```
##
##                                     Used      Total
##   Number of observations                64        67
##
##   Estimator                          ML
##   Minimum Function Test Statistic    92.425
##   Degrees of freedom                  25
##   P-value (Chi-square)                0.000
##
## Parameter Estimates:
##
##   Information                        Expected
##   Standard Errors                    Standard
##
## Latent Variables:
##           Estimate Std.Err z-value P(>|z|)
##   Semantic_1 =~
```

```

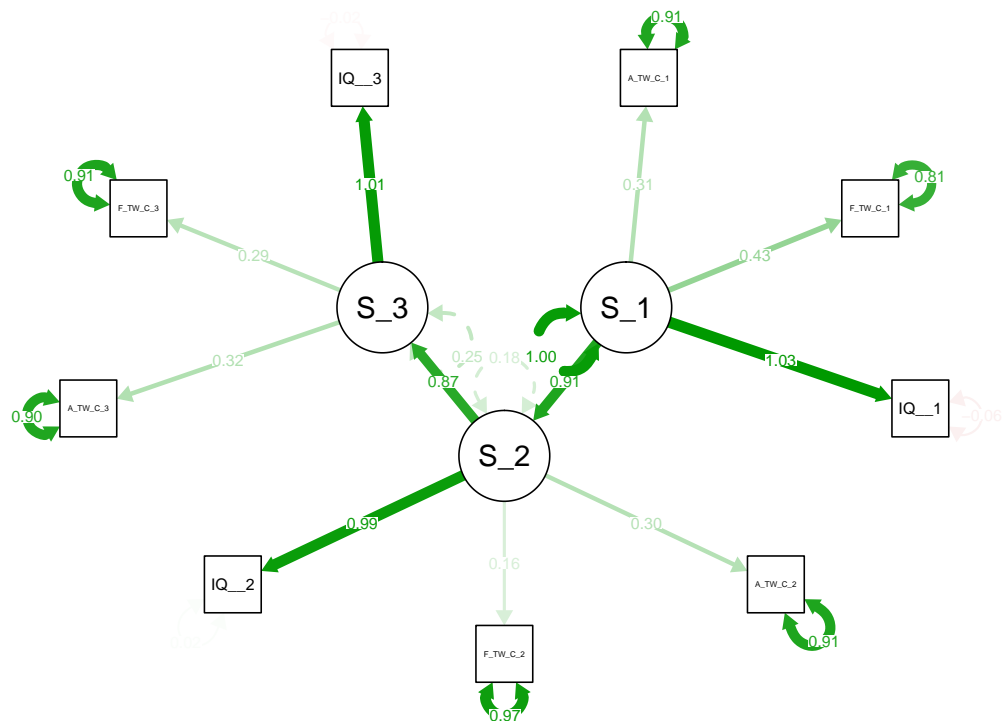
##      Anml_TW_Clst_1      1.428      0.559      2.552      0.011
##      Fd_TW_Clstrs_1      2.147      0.591      3.634      0.000
##      IQ_vocraw_1         9.733      0.916     10.628      0.000
##      Semantic_2 =~
##      Anml_TW_Clst_2      0.564      0.261      2.162      0.031
##      Fd_TW_Clstrs_2      0.328      0.269      1.222      0.222
##      IQ_vocraw_2         3.548      0.849      4.181      0.000
##      Semantic_3 =~
##      Anml_TW_Clst_3      0.700      0.289      2.424      0.015
##      Fd_TW_Clstrs_3      0.815      0.360      2.264      0.024
##      IQ_vocraw_3         4.451      1.479      3.010      0.003
##
## Regressions:
##              Estimate Std.Err  z-value  P(>|z|)
##      Semantic_3 ~
##      Semantic_2      0.734      0.300      2.446      0.014
##      Semantic_2 ~
##      Semantic_1      2.169      0.645      3.360      0.001
##
## Variances:
##              Estimate Std.Err  z-value  P(>|z|)
##      .Anml_TW_Clst_1  19.793      3.485      5.680      0.000
##      .Fd_TW_Clstrs_1  19.933      3.528      5.650      0.000
##      .IQ_vocraw_1     -5.452      8.398     -0.649      0.516
##      .Anml_TW_Clst_2  18.292      3.239      5.648      0.000
##      .Fd_TW_Clstrs_2  23.811      4.211      5.655      0.000
##      .IQ_vocraw_2      1.332      1.977      0.674      0.500
##      .Anml_TW_Clst_3  18.098      3.211      5.637      0.000
##      .Fd_TW_Clstrs_3  28.876      5.116      5.644      0.000
##      .IQ_vocraw_3     -1.773     12.578     -0.141      0.888
##      Semantic_1       1.000
##      .Semantic_2       1.000
##      .Semantic_3       1.000

```

```
semPaths(fit.auto, layout = "circle2", whatLabels = "std")
```



```
semPaths(fit.auto, layout = "circle2", what = "std")
```



```
anova(fit.long, fit.auto) #simpler model (fit.long) is preferred
```

```
## Chi Square Difference Test
```

```
##
```

```
##           Df AIC BIC  Chisq Chisq diff Df diff Pr(>Chisq)
```

```
## fit.long 24      91.983
```

```
## fit.auto 25          92.425    0.44184      1    0.5062
```

4. Fit a longitudinal growth model in SEM and in HLM. Compare and contrast the differences.

Estimates of *intercept* are similar (1) between fixed slope SEM & HLM models, (2) between random slope SEM & HLM models, and (3) between fixed-slope + covariate SEM & HLM models. Estimates of *slope* are similar (1) between fixed slope SEM & HLM models and (2) between random slope SEM & HLM models. Estimates of slope differ between SEM & HLM models when a covariate is added because the HLM slope can no longer be interpreted as the straight-forward rate of increase (y over x) between timepoints. As well, logLikelihood tests designated the fixed slope + covariate model as the preferred model regardless of whether SEM or HLM was used.

```
#HLM model
library(lme4)
library(car)
mod.HLM <- lmer(Sem_TotalCorrect ~ Timepoint + (1 | ID2), data = mydata) #fixed slope
summary(mod.HLM) #intercept = 31.06, slope = 3.85
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: Sem_TotalCorrect ~ Timepoint + (1 | ID2)
## Data: mydata
##
## REML criterion at convergence: 1405
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.20888 -0.57356 -0.05482  0.51023  2.96871
##
## Random effects:
## Groups Name Variance Std.Dev.
## ID2 (Intercept) 57.40 7.576
## Residual 38.03 6.167
## Number of obs: 200, groups: ID2, 67
##
## Fixed effects:
## Estimate Std. Error t value
## (Intercept) 31.0596 1.4786 21.005
## Timepoint 3.8471 0.5355 7.184
##
## Correlation of Fixed Effects:
## (Intr)
## Timepoint -0.722
car::Anova(mod.HLM)

## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: Sem_TotalCorrect
## Chisq Df Pr(>Chisq)
## Timepoint 51.614 1 6.757e-13 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
mod.HLM2 <- lmer(Sem_TotalCorrect ~ Timepoint + (Timepoint | ID2), data = mydata) #random slope
summary(mod.HLM2) #intercept = 31.07, slope = 3.84
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: Sem_TotalCorrect ~ Timepoint + (Timepoint | ID2)
## Data: mydata
##
## REML criterion at convergence: 1404
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.92900 -0.55230 -0.00624  0.49794  3.12399
##
## Random effects:
## Groups Name Variance Std.Dev. Corr
## ID2 (Intercept) 82.343 9.074
## Timepoint 4.664 2.160 -0.54
## Residual 33.485 5.787
## Number of obs: 200, groups: ID2, 67
##
## Fixed effects:
## Estimate Std. Error t value
## (Intercept) 31.0709 1.5494 20.054
## Timepoint 3.8386 0.5679 6.759
##
## Correlation of Fixed Effects:
## (Intr)
## Timepoint -0.752
```

```
car::Anova(mod.HLM2)
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: Sem_TotalCorrect
## Chisq Df Pr(>Chisq)
## Timepoint 45.685 1 1.389e-11 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
mod.HLM3 <- lmer(Sem_TotalCorrect ~ Timepoint + Age_at_time_of_testing + (1 | ID2),
data = mydata) #fixed slope, with covariate
summary(mod.HLM3) #intercept = 14.83, est. timepoint = 1.83, est. age = 1.55
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: Sem_TotalCorrect ~ Timepoint + Age_at_time_of_testing + (1 |
## ID2)
## Data: mydata
##
## REML criterion at convergence: 1379
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.30716 -0.60338 -0.01007  0.46251  3.13253
##
## Random effects:
```

```

## Groups      Name      Variance Std.Dev.
## ID2         (Intercept) 35.42    5.952
## Residual                37.79    6.147
## Number of obs: 200, groups: ID2, 67
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept)      14.8273      3.1802  4.662
## Timepoint         1.8291      0.6402  2.857
## Age_at_time_of_testing 1.5487      0.2738  5.655
##
## Correlation of Fixed Effects:
##              (Intr) Timpnt
## Timepoint      0.220
## Ag_t_tm_f_t   -0.904 -0.552
car::Anova(mod.HLM3)

## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: Sem_TotalCorrect
##              Chisq Df Pr(>Chisq)
## Timepoint      8.1641  1  0.004273 **
## Age_at_time_of_testing 31.9833  1  1.555e-08 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
anova(mod.HLM, mod.HLM2, mod.HLM3) #fixed slope, covariate model is preferred

## Data: mydata
## Models:
## mod.HLM: Sem_TotalCorrect ~ Timepoint + (1 | ID2)
## mod.HLM3: Sem_TotalCorrect ~ Timepoint + Age_at_time_of_testing + (1 |
## mod.HLM3: ID2)
## mod.HLM2: Sem_TotalCorrect ~ Timepoint + (Timepoint | ID2)
##              Df    AIC    BIC logLik deviance Chisq Chi Df Pr(>Chisq)
## mod.HLM      4 1415.5 1428.7 -703.74  1407.5
## mod.HLM3     5 1390.3 1406.8 -690.17  1380.3 27.144      1 1.888e-07 ***
## mod.HLM2     6 1418.5 1438.3 -703.27  1406.5  0.000      1      1
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
#Growth model
mod.SEM <- ' intercept =~ 1*Sem_TotalCorrect_1 + 1*Sem_TotalCorrect_2 + 1*Sem_TotalCorrect_3
            slope =~ 0*Sem_TotalCorrect_1 + 1*Sem_TotalCorrect_2 + 2*Sem_TotalCorrect_3
            slope ~~ 0*slope ' #fixed slope, no variance

mod.SEM.fixed <- growth(mod.SEM, missing = "ML", data = mydata_wide)
summary(mod.SEM.fixed) #intercept = 34.93, slope = 3.85

## lavaan (0.5-23.1097) converged normally after 68 iterations
##
##      Number of observations              67
##
##      Number of missing patterns          2
##

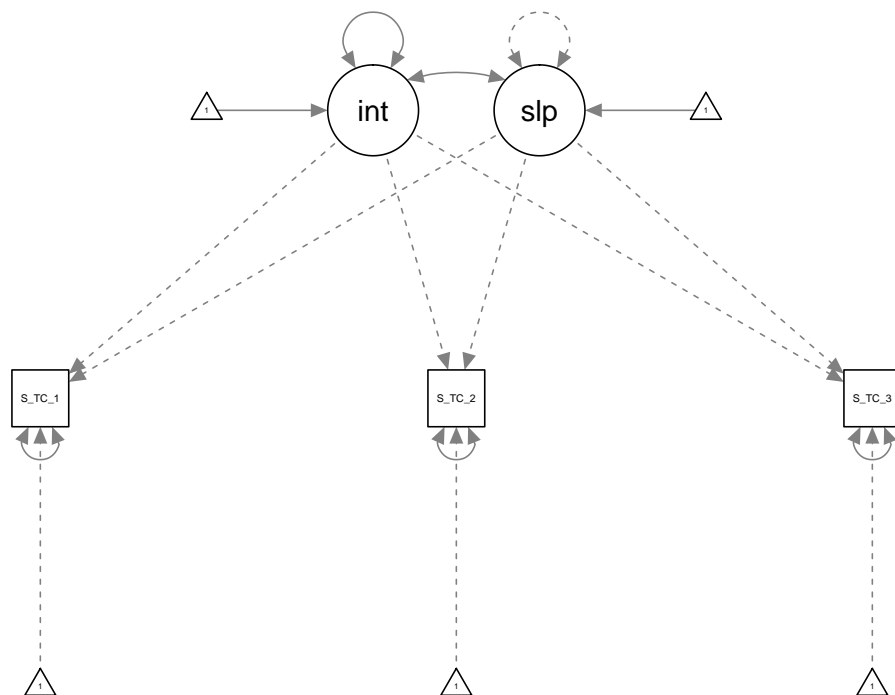
```

```

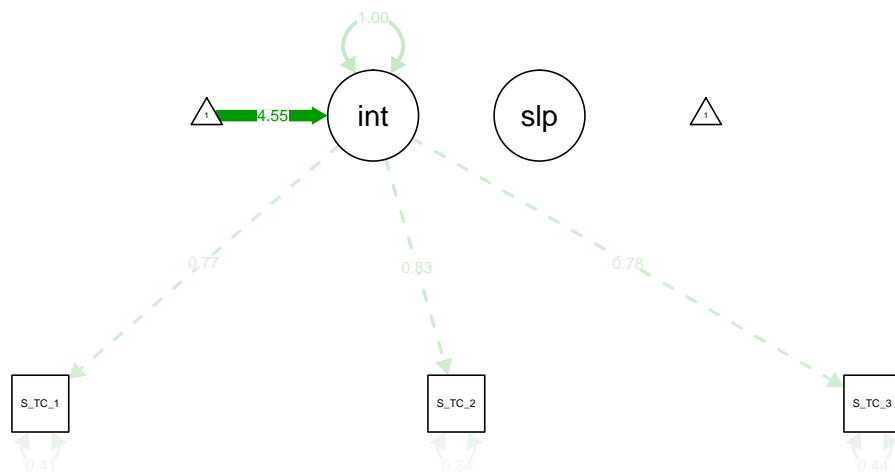
## Estimator ML
## Minimum Function Test Statistic 0.063
## Degrees of freedom 2
## P-value (Chi-square) 0.969
##
## Parameter Estimates:
##
## Information Observed
## Standard Errors Standard
##
## Latent Variables:
## Estimate Std.Err z-value P(>|z|)
## intercept =~
## Sem_TtlCrrct_1 1.000
## Sem_TtlCrrct_2 1.000
## Sem_TtlCrrct_3 1.000
## slope =~
## Sem_TtlCrrct_1 0.000
## Sem_TtlCrrct_2 1.000
## Sem_TtlCrrct_3 2.000
##
## Covariances:
## Estimate Std.Err z-value P(>|z|)
## intercept ~~
## slope -1.298 5.361 -0.242 0.809
##
## Intercepts:
## Estimate Std.Err z-value P(>|z|)
## .Sem_TtlCrrct_1 0.000
## .Sem_TtlCrrct_2 0.000
## .Sem_TtlCrrct_3 0.000
## intercept 34.925 1.172 29.805 0.000
## slope 3.845 0.564 6.815 0.000
##
## Variances:
## Estimate Std.Err z-value P(>|z|)
## slope 0.000
## .Sem_TtlCrrct_1 41.684 10.355 4.026 0.000
## .Sem_TtlCrrct_2 28.989 7.920 3.660 0.000
## .Sem_TtlCrrct_3 42.710 10.468 4.080 0.000
## intercept 58.911 16.124 3.654 0.000

```

```
semPaths(mod.SEM.fixed)
```



```
semPaths(mod.SEM.fixed, what = "std")
```



```

mod.SEM2 <- ' intercept =~ 1*Sem_TotalCorrect_1 + 1*Sem_TotalCorrect_2 + 1*Sem_TotalCorrect_3
              slope =~ 0*Sem_TotalCorrect_1 + 1*Sem_TotalCorrect_2 + 2*Sem_TotalCorrect_3 ' #random slope

mod.SEM.random <- growth(mod.SEM2, missing = "ML", data = mydata_wide)
summary(mod.SEM.random) #intercept = 34.93, slope = 3.85

```

```
## lavaan (0.5-23.1097) converged normally after 78 iterations
##
```

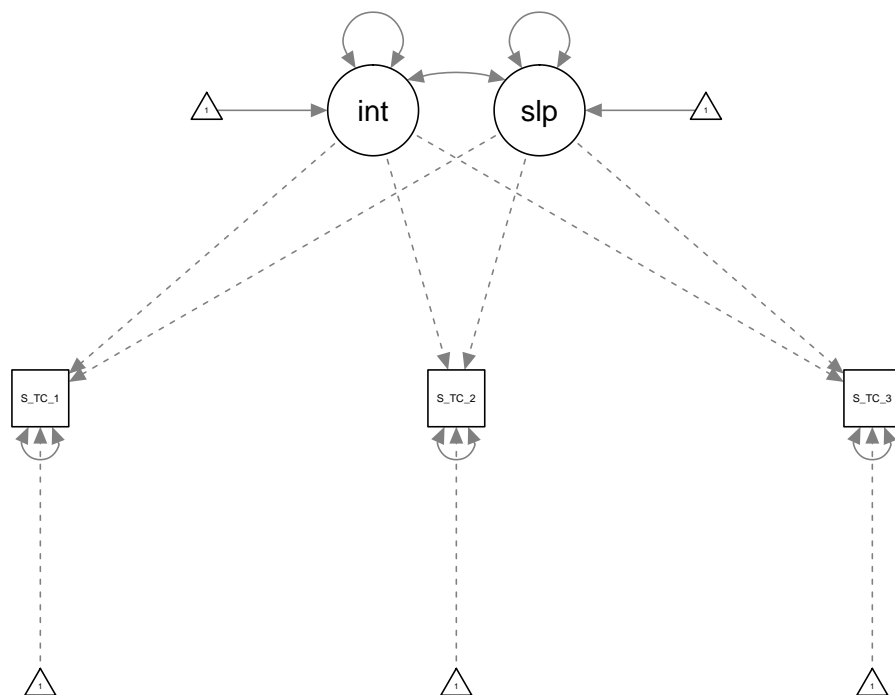


```

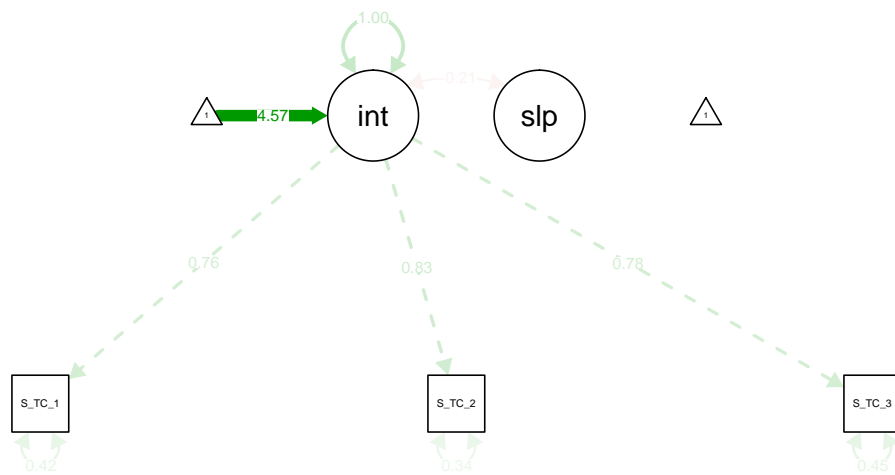
##      Number of observations                67
##
##      Number of missing patterns           2
##
##      Estimator                           ML
##      Minimum Function Test Statistic      0.062
##      Degrees of freedom                   1
##      P-value (Chi-square)                 0.803
##
## Parameter Estimates:
##
##      Information                          Observed
##      Standard Errors                      Standard
##
## Latent Variables:
##      Estimate Std.Err z-value P(>|z|)
##      intercept =~
##      Sem_TtlCrrct_1    1.000
##      Sem_TtlCrrct_2    1.000
##      Sem_TtlCrrct_3    1.000
##      slope =~
##      Sem_TtlCrrct_1    0.000
##      Sem_TtlCrrct_2    1.000
##      Sem_TtlCrrct_3    2.000
##
## Covariances:
##      Estimate Std.Err z-value P(>|z|)
##      intercept ~~
##      slope      -0.932   10.880   -0.086   0.932
##
## Intercepts:
##      Estimate Std.Err z-value P(>|z|)
##      .Sem_TtlCrrct_1    0.000
##      .Sem_TtlCrrct_2    0.000
##      .Sem_TtlCrrct_3    0.000
##      intercept    34.926    1.172   29.813    0.000
##      slope         3.846    0.564    6.824    0.000
##
## Variances:
##      Estimate Std.Err z-value P(>|z|)
##      .Sem_TtlCrrct_1   42.272   18.421    2.295    0.022
##      .Sem_TtlCrrct_2   28.836    8.842    3.261    0.001
##      .Sem_TtlCrrct_3   43.274   17.998    2.404    0.016
##      intercept    58.471   19.695    2.969    0.003
##      slope       -0.349    9.040   -0.039    0.969

```

```
semPaths(mod.SEM.random)
```



```
semPaths(mod.SEM.random, what = "std")
```



```
mod.SEM3 <- ' intercept =~ 1*Sem_TotalCorrect_1 + 1*Sem_TotalCorrect_2 + 1*Sem_TotalCorrect_3
slope =~ 0*Sem_TotalCorrect_1 + 1*Sem_TotalCorrect_2 + 2*Sem_TotalCorrect_3
slope ~~ 0*slope

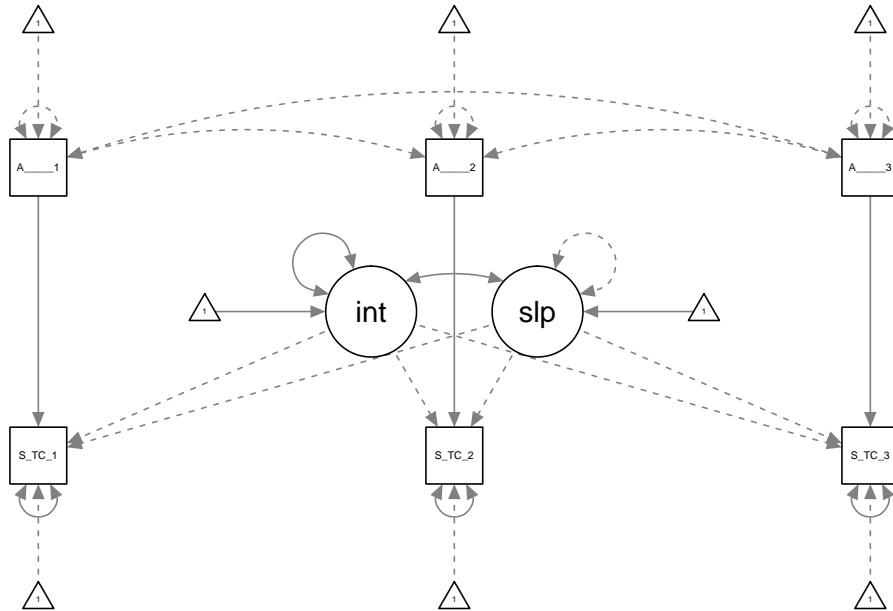
Sem_TotalCorrect_1 ~ Age_at_time_of_testing_1
Sem_TotalCorrect_2 ~ Age_at_time_of_testing_2
Sem_TotalCorrect_3 ~ Age_at_time_of_testing_3 ' #fixed slope, with covariate
```

```
mod.SEM.cov <- growth(mod.SEM3, missing = "ML", data = mydata_wide)
summary(mod.SEM.cov) #intercept = 12.38, slope = 6.65
```

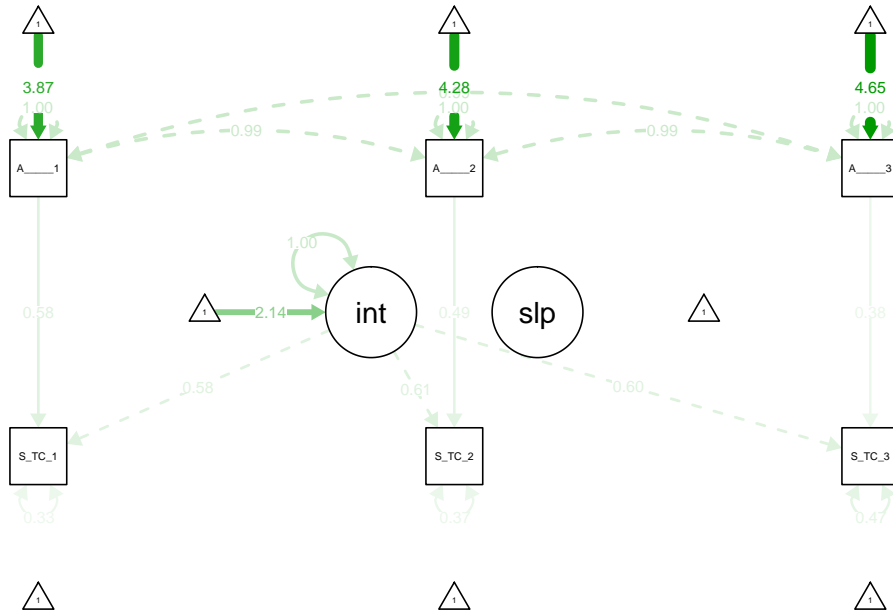
```
## lavaan (0.5-23.1097) converged normally after 67 iterations
##
##   Number of observations                    67
##
##   Number of missing patterns                2
##
##   Estimator                                ML
##   Minimum Function Test Statistic          20.725
##   Degrees of freedom                       8
##   P-value (Chi-square)                     0.008
##
## Parameter Estimates:
##
##   Information                                Observed
##   Standard Errors                          Standard
##
## Latent Variables:
##           Estimate Std.Err z-value P(>|z|)
##   intercept =~
##     Sem_TtlCrrct_1    1.000
##     Sem_TtlCrrct_2    1.000
##     Sem_TtlCrrct_3    1.000
##   slope =~
##     Sem_TtlCrrct_1    0.000
##     Sem_TtlCrrct_2    1.000
##     Sem_TtlCrrct_3    2.000
##
## Regressions:
##           Estimate Std.Err z-value P(>|z|)
##   Sem_TotalCorrect_1 ~
##     Ag_t_tm_f_ts_1    1.904    0.316    6.027    0.000
##   Sem_TotalCorrect_2 ~
##     Ag_t_tm_f_ts_2    1.517    0.279    5.445    0.000
##   Sem_TotalCorrect_3 ~
##     Ag_t_tm_f_ts_3    1.173    0.336    3.487    0.000
##
## Covariances:
##           Estimate Std.Err z-value P(>|z|)
##   intercept ~~
##     slope              0.659    4.249    0.155    0.877
##
## Intercepts:
##           Estimate Std.Err z-value P(>|z|)
##     .Sem_TtlCrrct_1    0.000
##     .Sem_TtlCrrct_2    0.000
##     .Sem_TtlCrrct_3    0.000
##     intercept          12.377    3.863    3.204    0.001
##     slope                6.652    2.490    2.672    0.008
##
## Variances:
##           Estimate Std.Err z-value P(>|z|)
```

```
##      slope          0.000
##      .Sem_TtlCrrct_1 32.435    8.948    3.625    0.000
##      .Sem_TtlCrrct_2 32.778    8.019    4.087    0.000
##      .Sem_TtlCrrct_3 44.160   10.815    4.083    0.000
##      intercept     33.307   11.097    3.001    0.003
```

```
semPaths(mod.SEM.cov)
```



```
semPaths(mod.SEM.cov, what = "std")
```



```
anova(mod.SEM.fixed, mod.SEM.random, mod.SEM.cov) #fixed slope, covariate model is preferred
```

```
## Chi Square Difference Test
```

```
##
##      Df AIC BIC   Chisq Chisq diff Df diff Pr(>Chisq)
## mod.SEM.random  1      0.0620
```

```
## mod.SEM.fixed    2          0.0635    0.0015    1    0.96914
## mod.SEM.cov      8          20.7247    20.6613    6    0.00211 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

5. Constrain the residual variances to be equal. Does this change the fit of your model?

Constraining the residual variances does not significantly change model fit. LogLikelihood tests indicate that a simpler model, where residual variances are allowed to vary, is preferred to a more complex model where they are constrained to be equal.

```
mod.SEM4 <- ' intercept =~ 1*Sem_TotalCorrect_1 + 1*Sem_TotalCorrect_2 + 1*Sem_TotalCorrect_3
              slope =~ 0*Sem_TotalCorrect_1 + 1*Sem_TotalCorrect_2 + 2*Sem_TotalCorrect_3

              Sem_TotalCorrect_1 ~ Age_at_time_of_testing_1
              Sem_TotalCorrect_2 ~ Age_at_time_of_testing_2
              Sem_TotalCorrect_3 ~ Age_at_time_of_testing_3

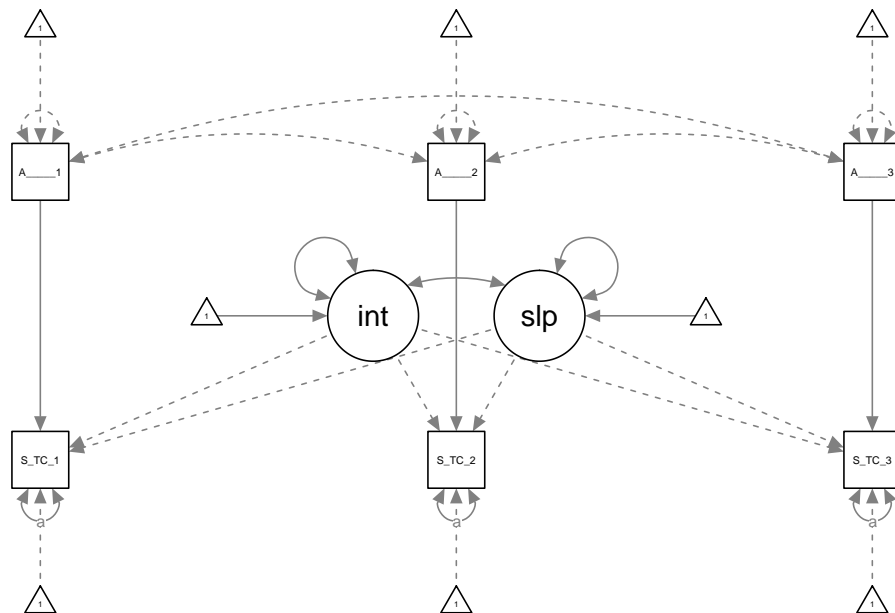
              Sem_TotalCorrect_1 ~~ a*Sem_TotalCorrect_1
              Sem_TotalCorrect_2 ~~ a*Sem_TotalCorrect_2
              Sem_TotalCorrect_3 ~~ a*Sem_TotalCorrect_3 ' #random slope, with covariate, residual vari

mod.SEM.cov2 <- growth(mod.SEM4, missing = "ML", data = mydata_wide)
summary(mod.SEM.cov2) #intercept = 12.53, slope = 6.23
```

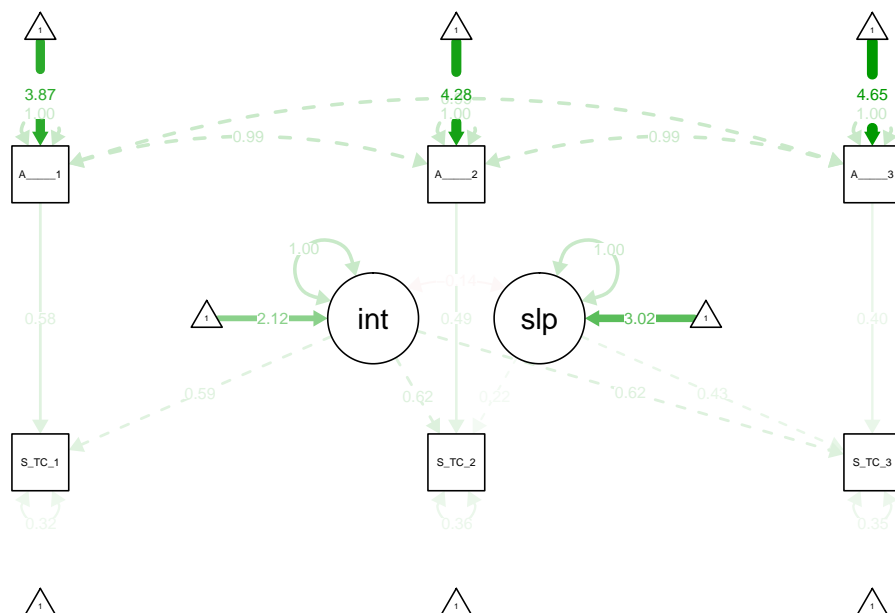
```
## lavaan (0.5-23.1097) converged normally after 86 iterations
##
##   Number of observations              67
##
##   Number of missing patterns          2
##
##   Estimator                          ML
##   Minimum Function Test Statistic     20.579
##   Degrees of freedom                  9
##   P-value (Chi-square)                0.015
##
## Parameter Estimates:
##
##   Information                        Observed
##   Standard Errors                    Standard
##
## Latent Variables:
##           Estimate Std.Err z-value P(>|z|)
## intercept =~
##   Sem_TtlCrrct_1    1.000
##   Sem_TtlCrrct_2    1.000
##   Sem_TtlCrrct_3    1.000
## slope =~
##   Sem_TtlCrrct_1    0.000
##   Sem_TtlCrrct_2    1.000
##   Sem_TtlCrrct_3    2.000
##
## Regressions:
```

```
##               Estimate Std.Err z-value P(>|z|)
## Sem_TotalCorrect_1 ~
##   Ag_t_tm_f_ts_1      1.892   0.316   5.991   0.000
## Sem_TotalCorrect_2 ~
##   Ag_t_tm_f_ts_2      1.535   0.277   5.544   0.000
## Sem_TotalCorrect_3 ~
##   Ag_t_tm_f_ts_3      1.218   0.335   3.631   0.000
##
## Covariances:
##               Estimate Std.Err z-value P(>|z|)
## intercept ~~
##   slope          -1.747   5.666  -0.308   0.758
##
## Intercepts:
##               Estimate Std.Err z-value P(>|z|)
## .Sem_TtlCrrct_1      0.000
## .Sem_TtlCrrct_2      0.000
## .Sem_TtlCrrct_3      0.000
## intercept          12.534   3.858   3.249   0.001
## slope              6.234   2.513   2.481   0.013
##
## Variances:
##               Estimate Std.Err z-value P(>|z|)
## .Sm_TtlCr_1 (a)      32.259   5.580   5.781   0.000
## .Sm_TtlCr_2 (a)      32.259   5.580   5.781   0.000
## .Sm_TtlCr_3 (a)      32.259   5.580   5.781   0.000
## intercept          34.943  11.672   2.994   0.003
## slope              4.275   4.506   0.949   0.343
```

```
semPaths(mod.SEM.cov2)
```



```
semPaths(mod.SEM.cov2, what = "std")
```



```
anova(mod.SEM.cov, mod.SEM.cov2) #simpler model (mod.SEM.cov) is preferred
```

```
## Chi Square Difference Test
```

```
##
```

```
##           Df AIC BIC   Chisq Chisq diff Df diff Pr(>Chisq)
```

```
## mod.SEM.cov    8      20.725
```

```
## mod.SEM.cov2   9      20.579   -0.14545      1      1
```

6. Constrain your slope to be fixed, not random. How does this change your model?

Constraining slopes to be fixed does not significantly change my model ($p = .97$).

```
anova(mod.SEM.fixed, mod.SEM.random) #see problem #4 for model specifics
```

```
## Chi Square Difference Test
```

```
##
```

```
##           Df AIC BIC   Chisq Chisq diff Df diff Pr(>Chisq)
```

```
## mod.SEM.random  1      0.0620
```

```
## mod.SEM.fixed   2      0.0635  0.0014966      1      0.9691
```

7. Change the time metric in your SEM growth model. How does that change your estimates? Does it change your fit statistics?

I changed my time metric such that the intercept was centered at TP 3 rather than TP1. This increased the intercept (which makes sense, given age-related change) but did not change model fit.

I also changed my time metric such that “duration” between TP3 & TP2 > “duration” between TP2 & TP1. This had little impact on the intercept, but decreased the slope and improved model fit.

```
# mydata_wide$time_1 <- 0
# mydata_wide$time_2 <- as.numeric(mydata_wide$Age_at_time_of_testing_2) -
#   as.numeric(mydata_wide$Age_at_time_of_testing_1)
# mydata_wide$time_3 <- as.numeric(mydata_wide$Age_at_time_of_testing_3) -
```

```

# as.numeric(mydata_wide$Age_at_time_of_testing_1)

mod.SEM5 <- ' intercept =~ 1*Sem_TotalCorrect_1 + 1*Sem_TotalCorrect_2 + 1*Sem_TotalCorrect_3
            slope =~ -2*Sem_TotalCorrect_1 + -1*Sem_TotalCorrect_2 + 0*Sem_TotalCorrect_3 '

mod.SEM.time <- growth(mod.SEM5, missing = "ML", data = mydata_wide)
summary(mod.SEM.time)

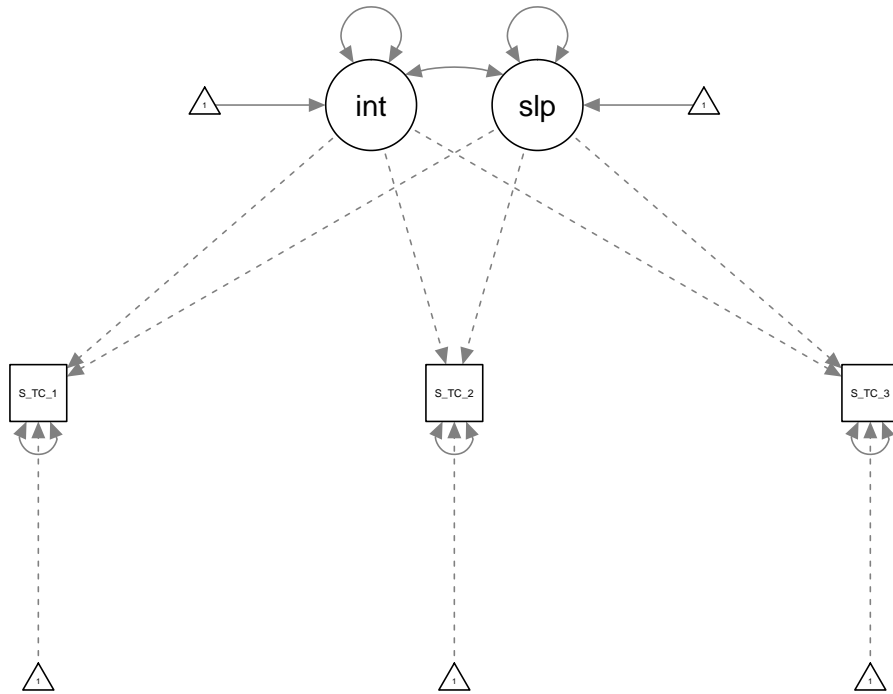
## lavaan (0.5-23.1097) converged normally after 93 iterations
##
## Number of observations                    67
##
## Number of missing patterns                2
##
## Estimator                                ML
## Minimum Function Test Statistic           0.062
## Degrees of freedom                        1
## P-value (Chi-square)                      0.803
##
## Parameter Estimates:
##
## Information                                Observed
## Standard Errors                           Standard
##
## Latent Variables:
##           Estimate Std.Err z-value P(>|z|)
## intercept =~
##   Sem_TtlCrrct_1    1.000
##   Sem_TtlCrrct_2    1.000
##   Sem_TtlCrrct_3    1.000
## slope =~
##   Sem_TtlCrrct_1   -2.000
##   Sem_TtlCrrct_2   -1.000
##   Sem_TtlCrrct_3    0.000
##
## Covariances:
##           Estimate Std.Err z-value P(>|z|)
## intercept ~~
##   slope           -1.630   10.117   -0.161   0.872
##
## Intercepts:
##           Estimate Std.Err z-value P(>|z|)
## .Sem_TtlCrrct_1    0.000
## .Sem_TtlCrrct_2    0.000
## .Sem_TtlCrrct_3    0.000
## intercept         42.618    1.146   37.181   0.000
## slope              3.846    0.564    6.824   0.000
##
## Variances:
##           Estimate Std.Err z-value P(>|z|)
## .Sem_TtlCrrct_1   42.272   18.421    2.295   0.022
## .Sem_TtlCrrct_2   28.836    8.842    3.261   0.001
## .Sem_TtlCrrct_3   43.274   17.998    2.404   0.016
## intercept         53.347   18.762    2.843   0.004

```

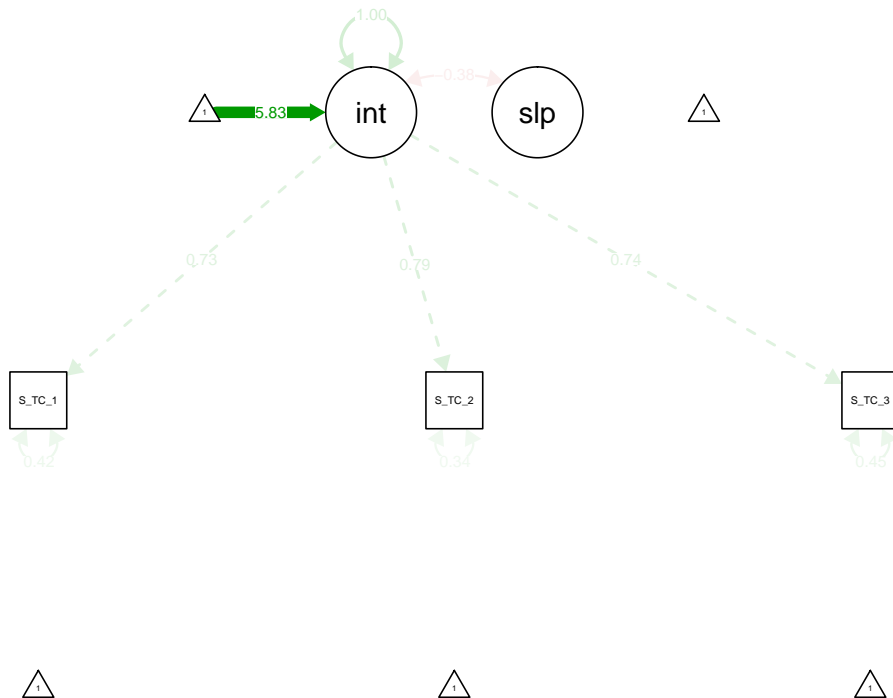


```
##      slope      -0.349    9.040   -0.039    0.969
```

```
semPaths(mod.SEM.time)
```



```
semPaths(mod.SEM.time, what = "std")
```



```
anova(mod.SEM.random, mod.SEM.time)
```

```
## Chi Square Difference Test
```

```
##
```

```
##      Df AIC BIC Chisq  Chisq diff Df diff Pr(>Chisq)
```

```

## mod.SEM.random 1          0.062
## mod.SEM.time 1          0.062 -2.3803e-13          0          1
mod.SEM6 <- ' intercept =~ 1*Sem_TotalCorrect_1 + 1*Sem_TotalCorrect_2 + 1*Sem_TotalCorrect_3
              slope =~ 0*Sem_TotalCorrect_1 + 1*Sem_TotalCorrect_2 + 6*Sem_TotalCorrect_3 '

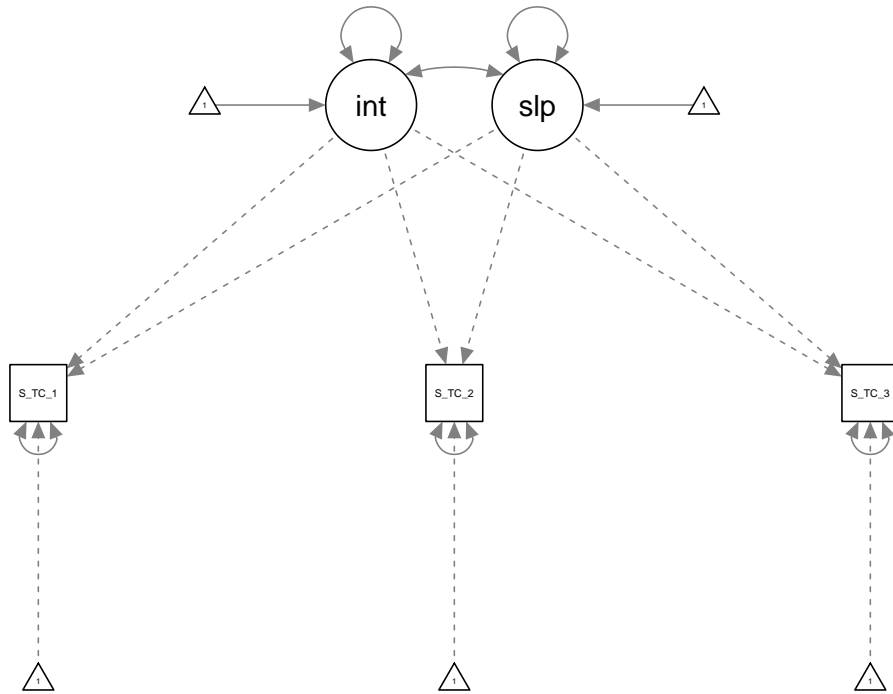
mod.SEM.time2 <- growth(mod.SEM6, missing = "ML", data = mydata_wide)
summary(mod.SEM.time2)

## lavaan (0.5-23.1097) converged normally after 89 iterations
##
## Number of observations          67
##
## Number of missing patterns      2
##
## Estimator                      ML
## Minimum Function Test Statistic 8.210
## Degrees of freedom              1
## P-value (Chi-square)            0.004
##
## Parameter Estimates:
##
## Information                      Observed
## Standard Errors                  Standard
##
## Latent Variables:
##           Estimate Std.Err z-value P(>|z|)
## intercept =~
##   Sem_TtlCrrct_1    1.000
##   Sem_TtlCrrct_2    1.000
##   Sem_TtlCrrct_3    1.000
## slope =~
##   Sem_TtlCrrct_1    0.000
##   Sem_TtlCrrct_2    1.000
##   Sem_TtlCrrct_3    6.000
##
## Covariances:
##           Estimate Std.Err z-value P(>|z|)
## intercept ~~
##   slope              0.411    2.280    0.180    0.857
##
## Intercepts:
##           Estimate Std.Err z-value P(>|z|)
##   .Sem_TtlCrrct_1    0.000
##   .Sem_TtlCrrct_2    0.000
##   .Sem_TtlCrrct_3    0.000
##   intercept          36.518    1.155    31.627    0.000
##   slope              1.067    0.184    5.794    0.000
##
## Variances:
##           Estimate Std.Err z-value P(>|z|)
##   .Sem_TtlCrrct_1    48.629    14.123    3.443    0.001
##   .Sem_TtlCrrct_2    31.574    10.548    2.994    0.003
##   .Sem_TtlCrrct_3    56.364    63.287    0.891    0.373
##   intercept          54.935    14.286    3.845    0.000

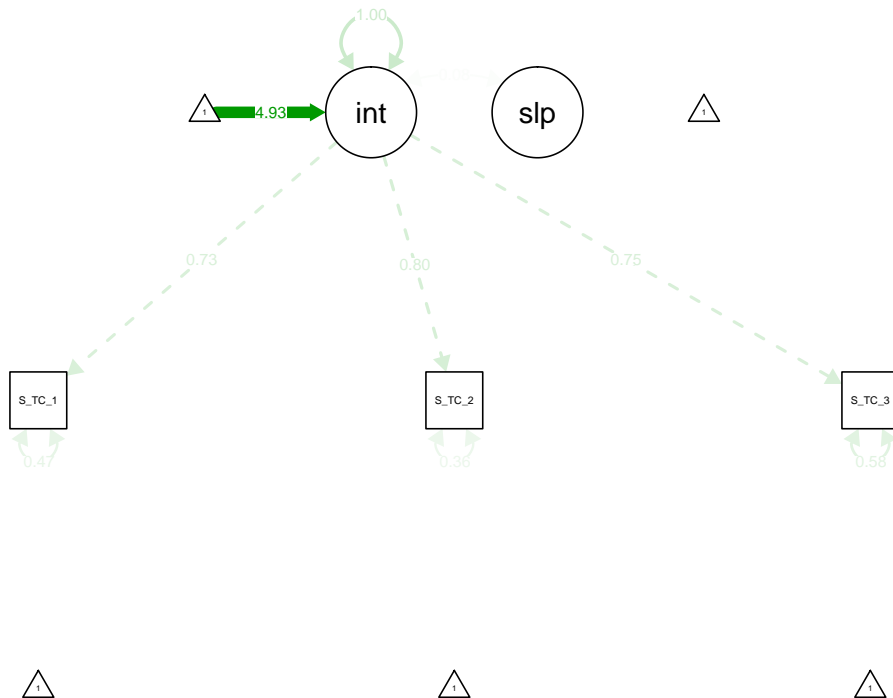
```

```
##      slope      -0.535    2.126   -0.252    0.801
```

```
semPaths(mod.SEM.time2)
```



```
semPaths(mod.SEM.time2, what = "std")
```



```
anova(mod.SEM.random, mod.SEM.time2)
```

```
## Chi Square Difference Test
```

```
##
```

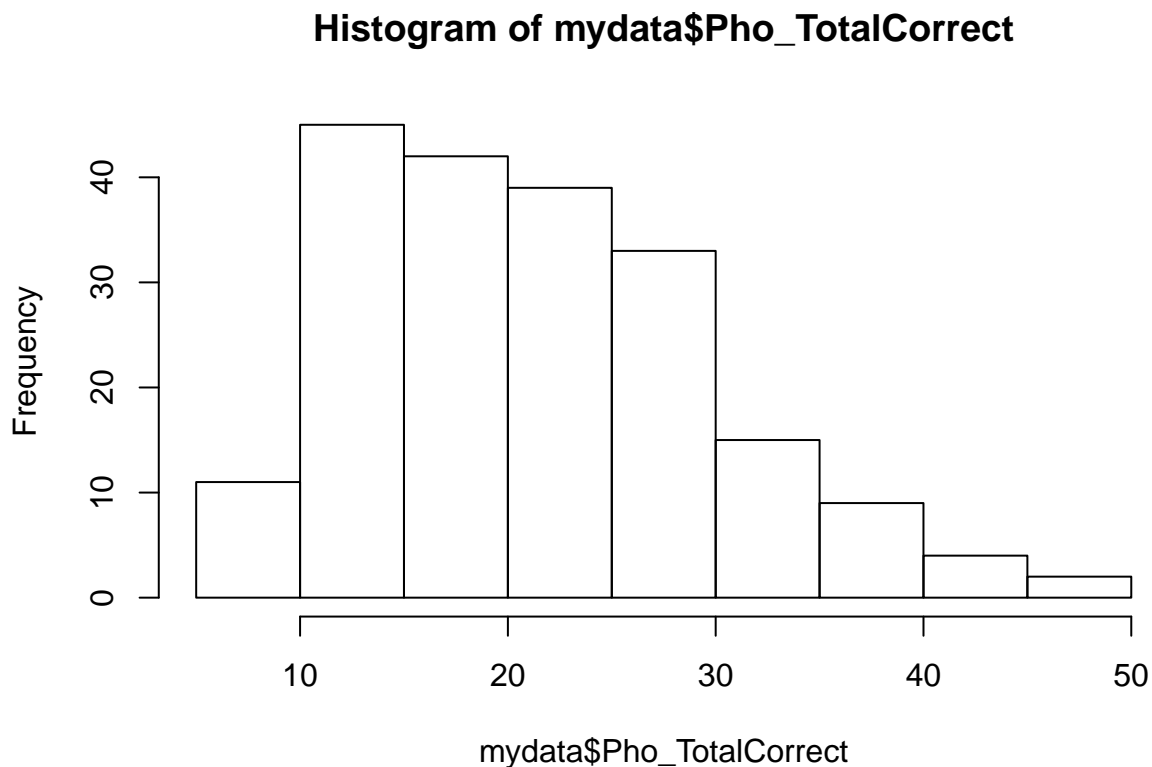
```
##      Df AIC BIC  Chisq Chisq diff Df diff Pr(>Chisq)
```

```
## mod.SEM.random 1          0.0620
## mod.SEM.time2   1          8.2098      8.1478      0 < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

8. Try a different type of estimation (see lavaan tutorial for details). How does that change your model?

Despite the fact that `Pho_TotalCorrect` is non-normal, changing the estimator has no effect on model fit, intercept, or slope.

```
#What if we want to try to model non-normal data?
hist(mydata$Pho_TotalCorrect)
```



```
mod.SEM7 <- ' intercept =~ 1*Pho_TotalCorrect_1 + 1*Pho_TotalCorrect_2 + 1*Pho_TotalCorrect_3
              slope =~ -2*Pho_TotalCorrect_1 + -1*Pho_TotalCorrect_2 + 0*Pho_TotalCorrect_3 '

mod.SEM.MLM <- growth(mod.SEM7, estimator = "MLM", data = mydata_wide)
mod.SEM.ML <- growth(mod.SEM7, estimator = "ML", data = mydata_wide)

summary(mod.SEM.MLM)
```

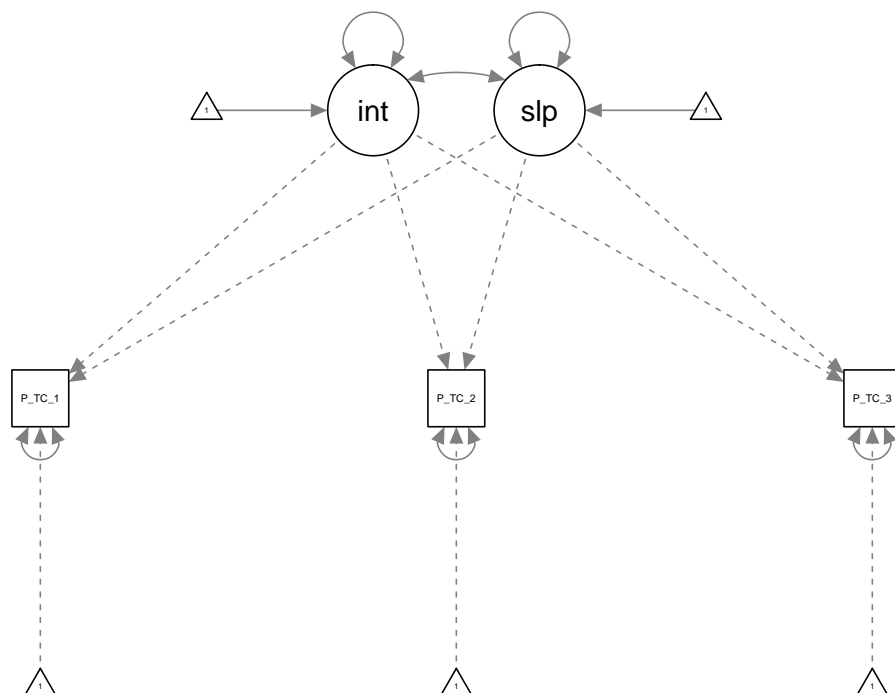
```
## lavaan (0.5-23.1097) converged normally after 80 iterations
##
##                               Used      Total
##   Number of observations           66       67
##
##   Estimator                      ML      Robust
##   Minimum Function Test Statistic    4.151    4.420
##   Degrees of freedom                 1         1
```

```

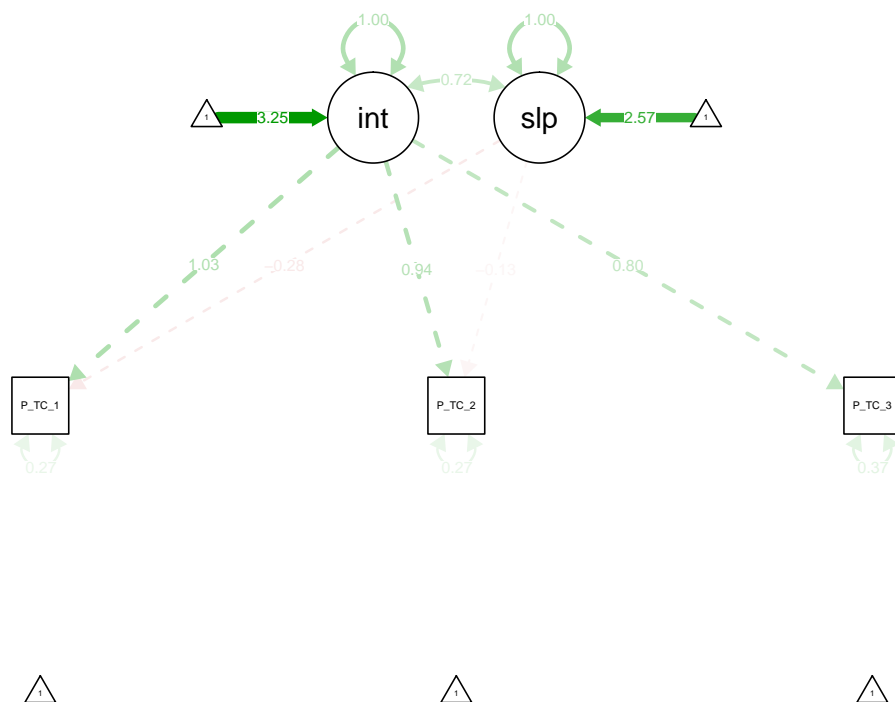
##      P-value (Chi-square)                0.042      0.036
##      Scaling correction factor            0.939
##      for the Satorra-Bentler correction
##
## Parameter Estimates:
##
##      Information                Expected
##      Standard Errors            Robust.sem
##
## Latent Variables:
##      Estimate  Std.Err  z-value  P(>|z|)
##      intercept =~
##      Pho_TtlCrrct_1    1.000
##      Pho_TtlCrrct_2    1.000
##      Pho_TtlCrrct_3    1.000
##      slope =~
##      Pho_TtlCrrct_1   -2.000
##      Pho_TtlCrrct_2   -1.000
##      Pho_TtlCrrct_3    0.000
##
## Covariances:
##      Estimate  Std.Err  z-value  P(>|z|)
##      intercept ~~
##      slope      5.513    6.633    0.831    0.406
##
## Intercepts:
##      Estimate  Std.Err  z-value  P(>|z|)
##      .Pho_TtlCrrct_1    0.000
##      .Pho_TtlCrrct_2    0.000
##      .Pho_TtlCrrct_3    0.000
##      intercept    24.238    1.101    22.023    0.000
##      slope        2.622    0.429    6.113    0.000
##
## Variances:
##      Estimate  Std.Err  z-value  P(>|z|)
##      .Pho_TtlCrrct_1   14.290    9.002    1.588    0.112
##      .Pho_TtlCrrct_2   17.235    5.583    3.087    0.002
##      .Pho_TtlCrrct_3   32.117   12.723    2.524    0.012
##      intercept    55.557   15.868    3.501    0.000
##      slope        1.043    5.083    0.205    0.837

```

`semPaths(mod.SEM.MLM)`



```
semPaths(mod.SEM.MLM, what = "std")
```



```
summary(mod.SEM.ML)
```

```
## lavaan (0.5-23.1097) converged normally after 80 iterations
```

```
##
```

```
##
##           Used      Total
## Number of observations      66      67
```

```
##
```

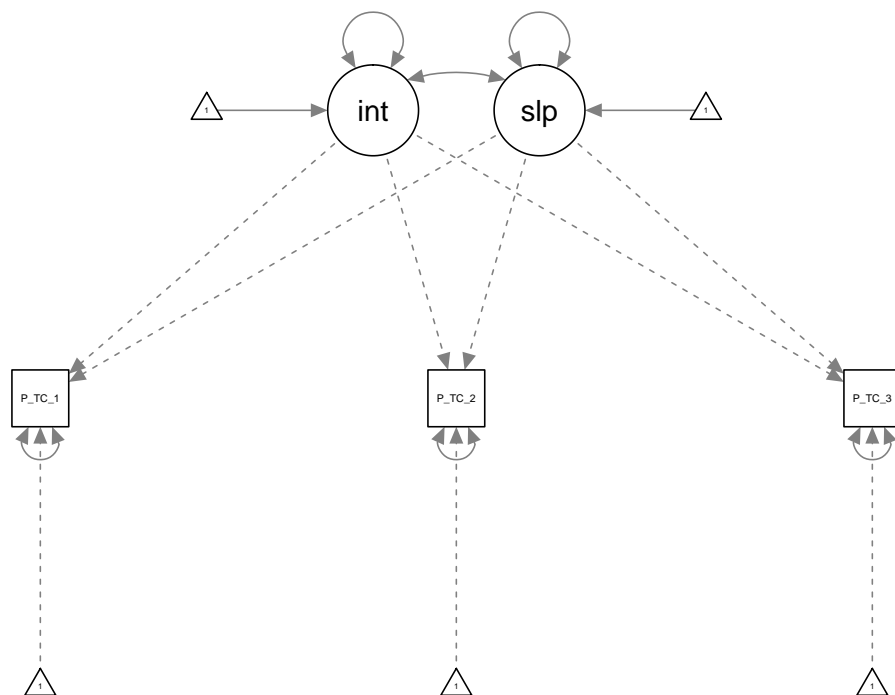
```
## Estimator              ML
```

```

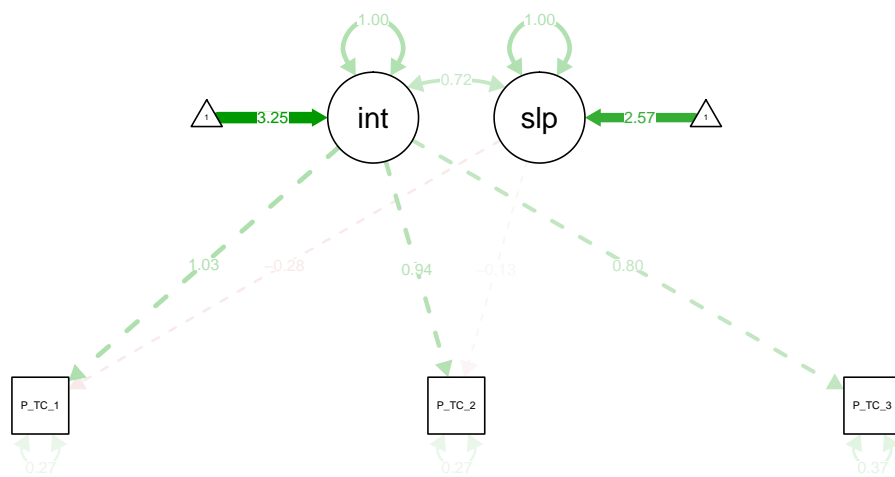
## Minimum Function Test Statistic          4.151
## Degrees of freedom                      1
## P-value (Chi-square)                    0.042
##
## Parameter Estimates:
##
## Information                               Expected
## Standard Errors                          Standard
##
## Latent Variables:
##      Estimate Std.Err z-value P(>|z|)
## intercept =~
##   Pho_TtlCrrct_1    1.000
##   Pho_TtlCrrct_2    1.000
##   Pho_TtlCrrct_3    1.000
## slope =~
##   Pho_TtlCrrct_1   -2.000
##   Pho_TtlCrrct_2   -1.000
##   Pho_TtlCrrct_3    0.000
##
## Covariances:
##      Estimate Std.Err z-value P(>|z|)
## intercept ~~
##   slope        5.513    6.612    0.834    0.404
##
## Intercepts:
##      Estimate Std.Err z-value P(>|z|)
##   .Pho_TtlCrrct_1    0.000
##   .Pho_TtlCrrct_2    0.000
##   .Pho_TtlCrrct_3    0.000
##   intercept    24.238    1.092   22.192    0.000
##   slope         2.622    0.426    6.159    0.000
##
## Variances:
##      Estimate Std.Err z-value P(>|z|)
##   .Pho_TtlCrrct_1   14.290    9.691    1.475    0.140
##   .Pho_TtlCrrct_2   17.235    5.652    3.050    0.002
##   .Pho_TtlCrrct_3   32.117   12.994    2.472    0.013
##   intercept    55.557   15.203    3.654    0.000
##   slope         1.043    5.427    0.192    0.848

```

`semPaths(mod.SEM.ML)`



```
semPaths(mod.SEM.ML, what = "std")
```



```
anova(mod.SEM.MLM, mod.SEM.ML)
```

```
## Chi Square Difference Test
```

```
##
```

```
##           Df AIC BIC  Chisq Chisq diff Df diff Pr(>Chisq)
```

```
## mod.SEM.MLM  1      4.1505
```

```
## mod.SEM.ML   1      4.1505      0      0      1
```


9. Provide semplots for each of the models

Incorporated throughout code (above)