

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, missing = "ML")
summary(fit.marker, fit.measures = TRUE)
```

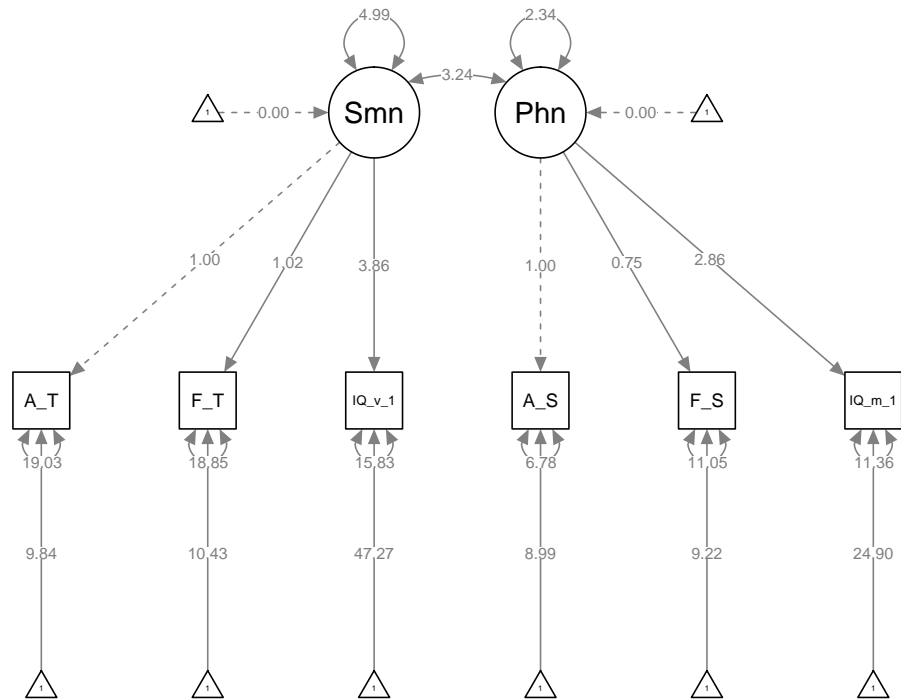
```
## lavaan (0.5-23.1097) converged normally after 53 iterations
##
##   Number of observations                    67
##
##   Number of missing patterns                1
##
##   Estimator                                ML
##   Minimum Function Test Statistic          22.669
##   Degrees of freedom                       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                19
##   Akaike (AIC)                            2361.024
##   Bayesian (BIC)                          2402.914
```

```

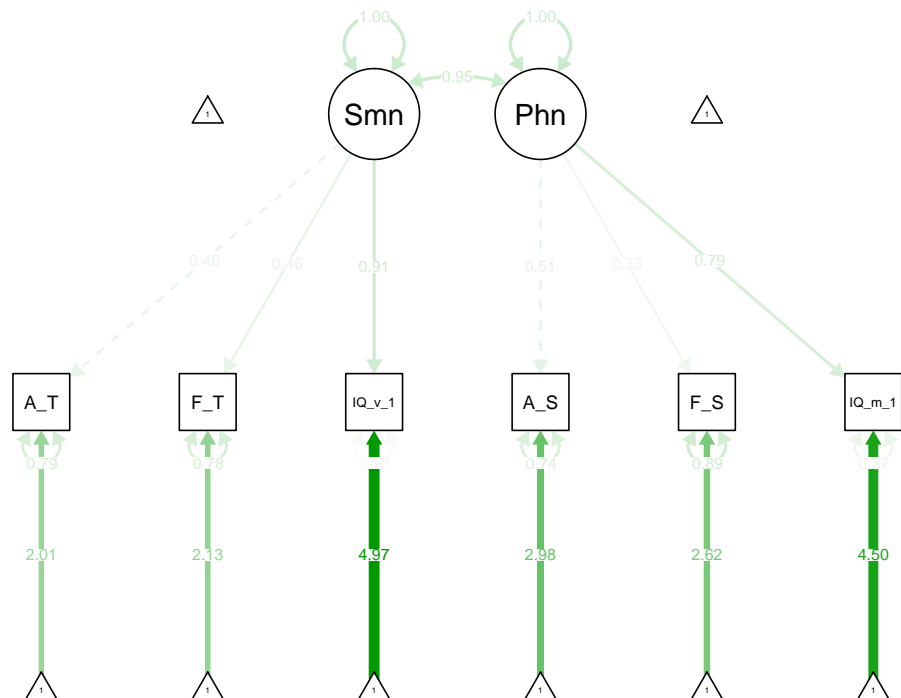
## Sample-size adjusted Bayesian (BIC) 2343.089
##
## Root Mean Square Error of Approximation:
##
## RMSEA 0.165
## 90 Percent Confidence Interval 0.087 0.248
## P-value RMSEA <= 0.05 0.012
##
## Standardized Root Mean Square Residual:
##
## SRMR 0.081
##
## Parameter Estimates:
##
## Information Observed
## 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 0.361 2.826 0.005
## IQ_vocraw_1 3.862 1.152 3.352 0.001
## Phonemic =~
## Animl_Swtchs_1 1.000
## Food_Swtchs_1 0.755 0.330 2.284 0.022
## IQ_mrraw_1 2.864 0.745 3.845 0.000
##
## Covariances:
## Estimate Std.Err z-value P(>|z|)
## Semantic ~~
## Phonemic 3.244 1.328 2.442 0.015
##
## Intercepts:
## Estimate Std.Err z-value P(>|z|)
## .Anml_TW_Clst_1 9.836 0.599 16.428 0.000
## .Fd_TW_Clst_1 10.433 0.599 17.418 0.000
## .IQ_vocraw_1 47.269 1.161 40.719 0.000
## .Animl_Swtchs_1 8.985 0.369 24.355 0.000
## .Food_Swtchs_1 9.224 0.430 21.456 0.000
## .IQ_mrraw_1 24.896 0.675 36.855 0.000
## Semantic 0.000
## Phonemic 0.000
##
## Variances:
## Estimate Std.Err z-value P(>|z|)
## .Anml_TW_Clst_1 19.027 3.487 5.457 0.000
## .Fd_TW_Clst_1 18.855 3.418 5.516 0.000
## .IQ_vocraw_1 15.830 11.157 1.419 0.156
## .Animl_Swtchs_1 6.776 1.273 5.323 0.000
## .Food_Swtchs_1 11.048 1.964 5.624 0.000
## .IQ_mrraw_1 11.360 3.529 3.219 0.001
## Semantic 4.991 2.786 1.791 0.073
## Phonemic 2.343 1.167 2.008 0.045

```

```
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, missing = "ML")
summary(fit.fixed, fit.measures = TRUE)
```

```
## lavaan (0.5-23.1097) converged normally after 48 iterations
##
```

```

##      Number of observations                67
##
##      Number of missing patterns           1
##
##      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            19
##      Akaike (AIC)                        2361.024
##      Bayesian (BIC)                      2402.914
##      Sample-size adjusted Bayesian (BIC) 2343.089
##
## 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                              0.081
##
## Parameter Estimates:
##
##      Information                        Observed
##      Standard Errors                    Standard
##
## Latent Variables:
##      Estimate Std.Err z-value P(>|z|)
##      Semantic =~
##      Anml_TW_Clst_1    2.234    0.624    3.583    0.000
##      Fd_TW_Clst_1      2.276    0.610    3.734    0.000
##      IQ_vocraw_1       8.629    1.088    7.928    0.000
##      Phonemic =~
##      Animl_Swtchs_1     1.531    0.381    4.015    0.000
##      Food_Swtchs_1     1.155    0.464    2.491    0.013

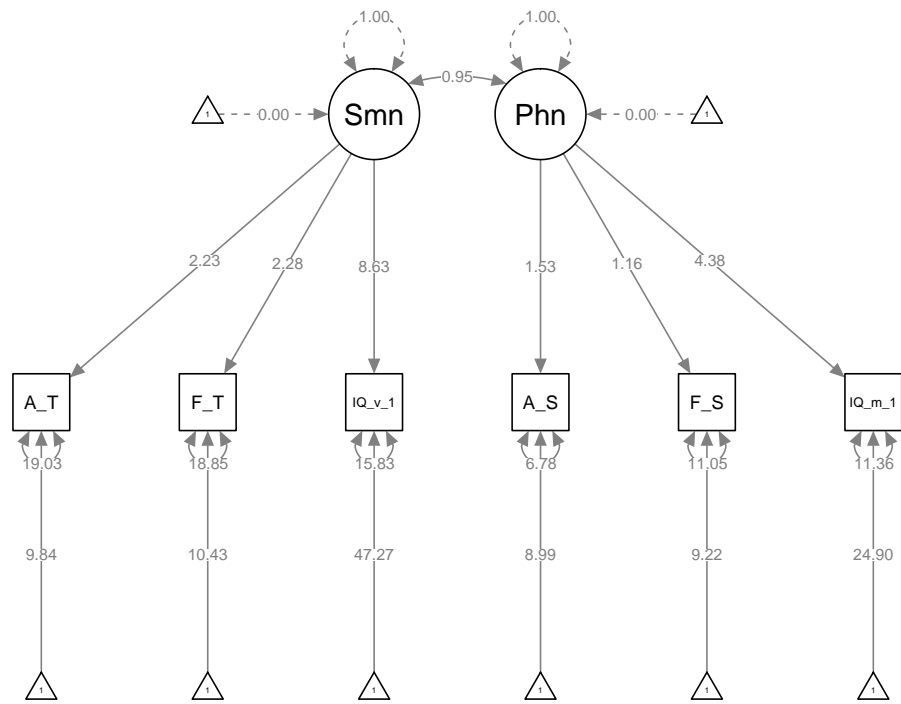
```

```

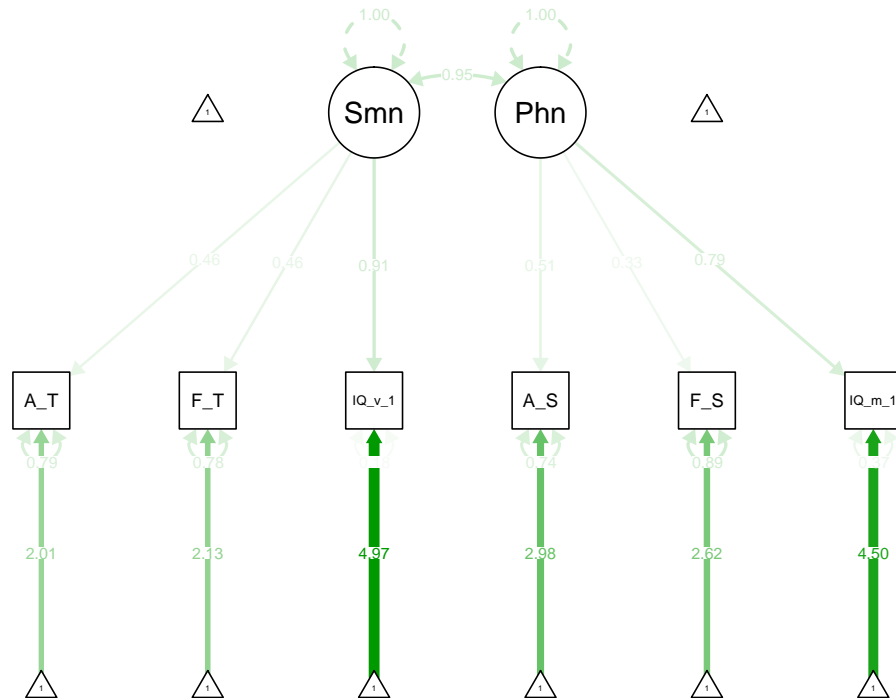
##      IQ_mrraw_1      4.383    0.652    6.724    0.000
##
## Covariances:
##              Estimate Std.Err  z-value  P(>|z|)
## Semantic ~~
## Phonemic      0.949    0.093   10.157    0.000
##
## Intercepts:
##              Estimate Std.Err  z-value  P(>|z|)
## .Anml_TW_Clst_1    9.836    0.599   16.428    0.000
## .Fd_TW_Clst_1     10.433    0.599   17.418    0.000
## .IQ_vocraw_1      47.269    1.161   40.719    0.000
## .Animl_Swtchs_1    8.985    0.369   24.355    0.000
## .Food_Swtchs_1     9.224    0.430   21.456    0.000
## .IQ_mrraw_1      24.896    0.675   36.856    0.000
## Semantic           0.000
## Phonemic           0.000
##
## Variances:
##              Estimate Std.Err  z-value  P(>|z|)
## .Anml_TW_Clst_1   19.027    3.487    5.457    0.000
## .Fd_TW_Clst_1     18.855    3.418    5.516    0.000
## .IQ_vocraw_1      15.830   11.157    1.419    0.156
## .Animl_Swtchs_1    6.776    1.273    5.323    0.000
## .Food_Swtchs_1    11.048    1.964    5.624    0.000
## .IQ_mrraw_1      11.360    3.529    3.219    0.001
## 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 = .081, 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 auto regressive; 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, indicators load strongly & significantly onto their respective latent constructs.

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 are insignificant ($p = .10$ and $.62$, respectively), likely due to the fact that their variability is accounted for by earlier timepoints. Likelihood ratio tests designate the longitudinal CFA model as the preferred model.

```

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

    ##correlated residuals across time
    Animal_TW_Clusters_1 ~~ Animal_TW_Clusters_2 + Animal_TW_Clusters_3
    Animal_TW_Clusters_2 ~~ Animal_TW_Clusters_3
    Food_TW_Clusters_1 ~~ Food_TW_Clusters_2 + Food_TW_Clusters_3
    Food_TW_Clusters_2 ~~ Food_TW_Clusters_3
    IQ_vocraw_1 ~~ IQ_vocraw_2 + IQ_vocraw_3
    IQ_vocraw_2 ~~ IQ_vocraw_3
'

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

```

```
## lavaan (0.5-23.1097) converged normally after 229 iterations
```

```

##
##   Number of observations                67
##
##   Number of missing patterns           4
##
##   Estimator                           ML
##   Minimum Function Test Statistic      26.593
##   Degrees of freedom                   15
##   P-value (Chi-square)                 0.032
##
## Model test baseline model:
##
##   Minimum Function Test Statistic      366.737
##   Degrees of freedom                   36
##   P-value                              0.000
##
## User model versus baseline model:
##
##   Comparative Fit Index (CFI)          0.965
##   Tucker-Lewis Index (TLI)            0.916
##
## Loglikelihood and Information Criteria:
##
##   Loglikelihood user model (H0)        -1749.532
##   Loglikelihood unrestricted model (H1) -1736.235
##
##   Number of free parameters            39
##   Akaike (AIC)                         3577.063
##   Bayesian (BIC)                       3663.046
##   Sample-size adjusted Bayesian (BIC)  3540.250
##
## Root Mean Square Error of Approximation:
##
##   RMSEA                                0.107
##   90 Percent Confidence Interval        0.031  0.173
##   P-value RMSEA <= 0.05                0.087

```

```

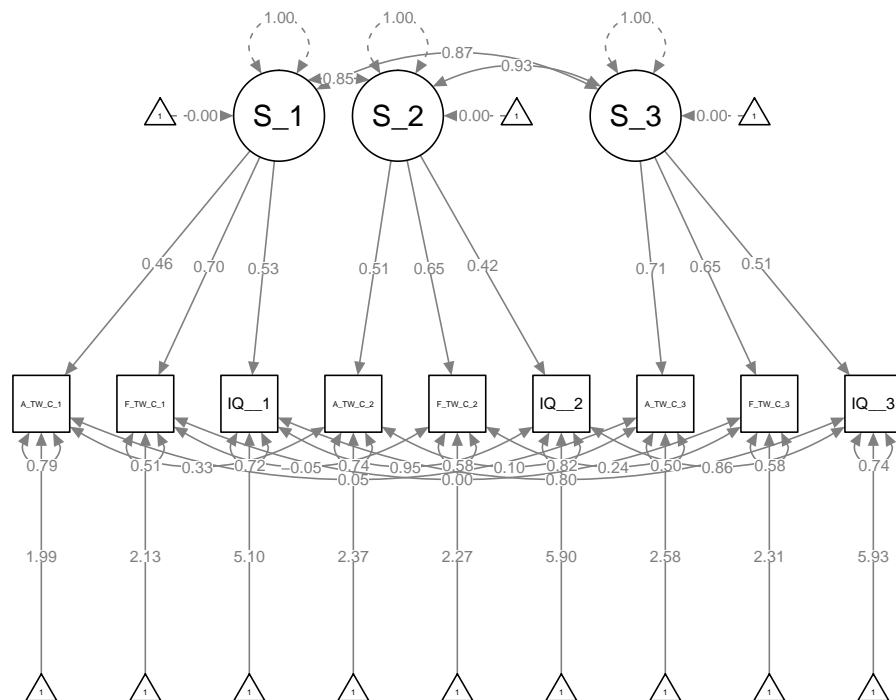
##
## Standardized Root Mean Square Residual:
##
##   SRMR                                0.071
##
## Parameter Estimates:
##
##   Information                                Observed
##   Standard Errors                            Standard
##
## Latent Variables:
##           Estimate  Std.Err  z-value  P(>|z|)
##   Semantic_1 =~
##     Anml_TW_Clst_1    2.276    0.713    3.192    0.001
##     Fd_TW_Clstrs_1    3.420    0.709    4.825    0.000
##     IQ_vocraw_1       4.949    1.423    3.479    0.001
##   Semantic_2 =~
##     Anml_TW_Clst_2    2.369    0.708    3.348    0.001
##     Fd_TW_Clstrs_2    3.303    0.834    3.960    0.000
##     IQ_vocraw_2       3.671    1.455    2.522    0.012
##   Semantic_3 =~
##     Anml_TW_Clst_3    3.190    0.658    4.847    0.000
##     Fd_TW_Clstrs_3    3.671    0.845    4.346    0.000
##     IQ_vocraw_3       4.518    1.445    3.127    0.002
##
## Covariances:
##           Estimate  Std.Err  z-value  P(>|z|)
##   .Animal_TW_Clusters_1 ~~
##     .Anml_TW_Clst_2      5.744    2.900    1.980    0.048
##     .Anml_TW_Clst_3      0.666    2.569    0.259    0.795
##   .Animal_TW_Clusters_2 ~~
##     .Anml_TW_Clst_3      1.224    2.525    0.485    0.628
##   .Food_TW_Clusters_1 ~~
##     .Fd_TW_Clstrs_2     -0.661    3.012   -0.219    0.826
##     .Fd_TW_Clstrs_3      0.029    3.234    0.009    0.993
##   .Food_TW_Clusters_2 ~~
##     .Fd_TW_Clstrs_3      4.065    3.980    1.021    0.307
##   .IQ_vocraw_1 ~~
##     .IQ_vocraw_2       58.266   12.100    4.815    0.000
##     .IQ_vocraw_3       48.365   11.610    4.166    0.000
##   .IQ_vocraw_2 ~~
##     .IQ_vocraw_3       51.882   11.751    4.415    0.000
##   Semantic_1 ~~
##     Semantic_2          0.849    0.092    9.231    0.000
##     Semantic_3          0.871    0.099    8.840    0.000
##   Semantic_2 ~~
##     Semantic_3          0.930    0.079   11.749    0.000
##
## Intercepts:
##           Estimate  Std.Err  z-value  P(>|z|)
##   .Anml_TW_Clst_1     9.836    0.603   16.301    0.000
##   .Fd_TW_Clstrs_1    10.433    0.599   17.406    0.000
##   .IQ_vocraw_1       47.269    1.133   41.732    0.000
##   .Anml_TW_Clst_2    10.955    0.565   19.388    0.000

```

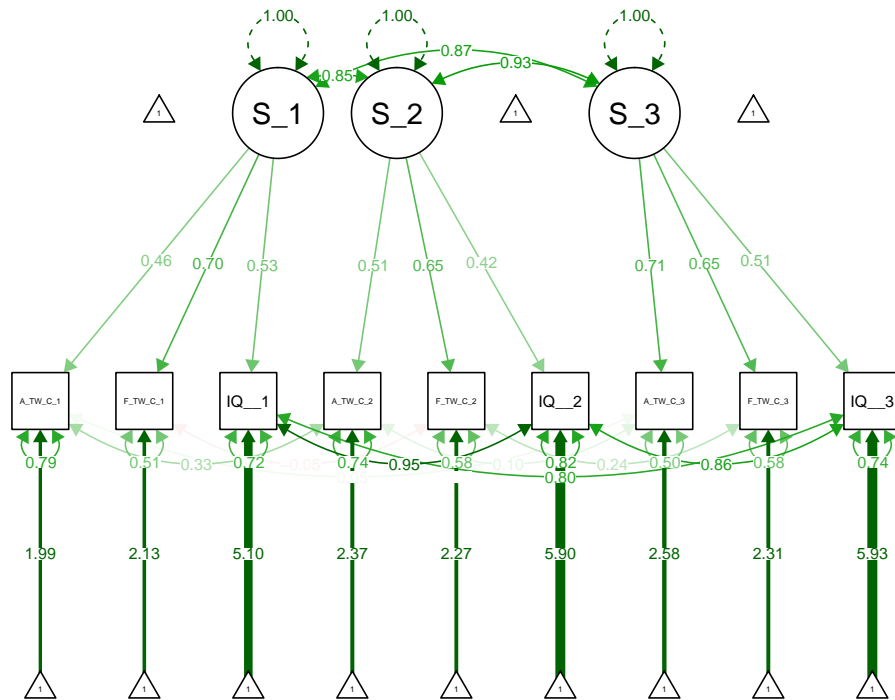


```
## .Fd_TW_Clstrs_2 11.567 0.623 18.562 0.000
## .IQ_vocraw_2 51.000 1.056 48.300 0.000
## .Anml_TW_Clst_3 11.674 0.559 20.898 0.000
## .Fd_TW_Clstrs_3 13.034 0.702 18.575 0.000
## .IQ_vocraw_3 52.971 1.093 48.468 0.000
## Semantic_1 0.000
## Semantic_2 0.000
## Semantic_3 0.000
##
## Variances:
## Estimate Std.Err z-value P(>|z|)
## .Anml_TW_Clst_1 19.212 3.902 4.923 0.000
## .Fd_TW_Clstrs_1 12.374 3.856 3.209 0.001
## .IQ_vocraw_1 61.463 13.218 4.650 0.000
## .Anml_TW_Clst_2 15.780 3.621 4.358 0.000
## .Fd_TW_Clstrs_2 15.110 4.970 3.040 0.002
## .IQ_vocraw_2 61.223 12.531 4.886 0.000
## .Anml_TW_Clst_3 10.267 3.276 3.134 0.002
## .Fd_TW_Clstrs_3 18.447 5.206 3.543 0.000
## .IQ_vocraw_3 59.299 12.906 4.595 0.000
## 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 =~ L1*Animal_TW_Clusters_1 + L2*Food_TW_Clusters_1 + L3*IQ_vocraw_1
  Semantic_2 =~ L1*Animal_TW_Clusters_2 + L2*Food_TW_Clusters_2 + L3*IQ_vocraw_2
  Semantic_3 =~ L1*Animal_TW_Clusters_3 + L2*Food_TW_Clusters_3 + L3*IQ_vocraw_3

  ##correlated residuals across time
  Animal_TW_Clusters_1 ~~ Animal_TW_Clusters_2 + Animal_TW_Clusters_3
  Animal_TW_Clusters_2 ~~ Animal_TW_Clusters_3
  Food_TW_Clusters_1 ~~ Food_TW_Clusters_2 + Food_TW_Clusters_3
  Food_TW_Clusters_2 ~~ Food_TW_Clusters_3
  IQ_vocraw_1 ~~ IQ_vocraw_2 + IQ_vocraw_3
  IQ_vocraw_2 ~~ IQ_vocraw_3

  ##directional regression paths
  Semantic_3 ~ Semantic_2
  Semantic_2 ~ Semantic_1

  ## free latent variances at later times (only set the scale once)
  Semantic_2 ~~ NA*Semantic_2
  Semantic_3 ~~ NA*Semantic_3
'
```

```
fit.auto <- sem(Auto.mod, data=mydata_wide, std.lv = T, missing = "ML")
summary(fit.auto, fit.measures = T)
```

```
## lavaan (0.5-23.1097) converged normally after 193 iterations
##
##   Number of observations              67
##
##   Number of missing patterns          4
##
##   Estimator                           ML
```

```

## Minimum Function Test Statistic          28.749
## Degrees of freedom                        20
## P-value (Chi-square)                     0.093
##
## Model test baseline model:
##
## Minimum Function Test Statistic          366.737
## Degrees of freedom                        36
## P-value                                  0.000
##
## User model versus baseline model:
##
## Comparative Fit Index (CFI)               0.974
## Tucker-Lewis Index (TLI)                 0.952
##
## Loglikelihood and Information Criteria:
##
## Loglikelihood user model (H0)             -1750.610
## Loglikelihood unrestricted model (H1)      -1736.235
##
## Number of free parameters                 34
## Akaike (AIC)                             3569.220
## Bayesian (BIC)                           3644.179
## Sample-size adjusted Bayesian (BIC)       3537.126
##
## Root Mean Square Error of Approximation:
##
## RMSEA                                     0.081
## 90 Percent Confidence Interval            0.000 0.142
## P-value RMSEA <= 0.05                    0.215
##
## Standardized Root Mean Square Residual:
##
## SRMR                                     0.064
##
## Parameter Estimates:
##
## Information                               Observed
## Standard Errors                           Standard
##
## Latent Variables:
##      Estimate Std.Err z-value P(>|z|)
## Semantic_1 =~
##   An_TW_C_1 (L1)    2.839   0.511   5.555   0.000
##   Fd_TW_C_1 (L2)    3.581   0.616   5.813   0.000
##   IQ_vcrw_1 (L3)    5.198   1.264   4.113   0.000
## Semantic_2 =~
##   An_TW_C_2 (L1)    2.839   0.511   5.555   0.000
##   Fd_TW_C_2 (L2)    3.581   0.616   5.813   0.000
##   IQ_vcrw_2 (L3)    5.198   1.264   4.113   0.000
## Semantic_3 =~
##   An_TW_C_3 (L1)    2.839   0.511   5.555   0.000
##   Fd_TW_C_3 (L2)    3.581   0.616   5.813   0.000
##   IQ_vcrw_3 (L3)    5.198   1.264   4.113   0.000

```

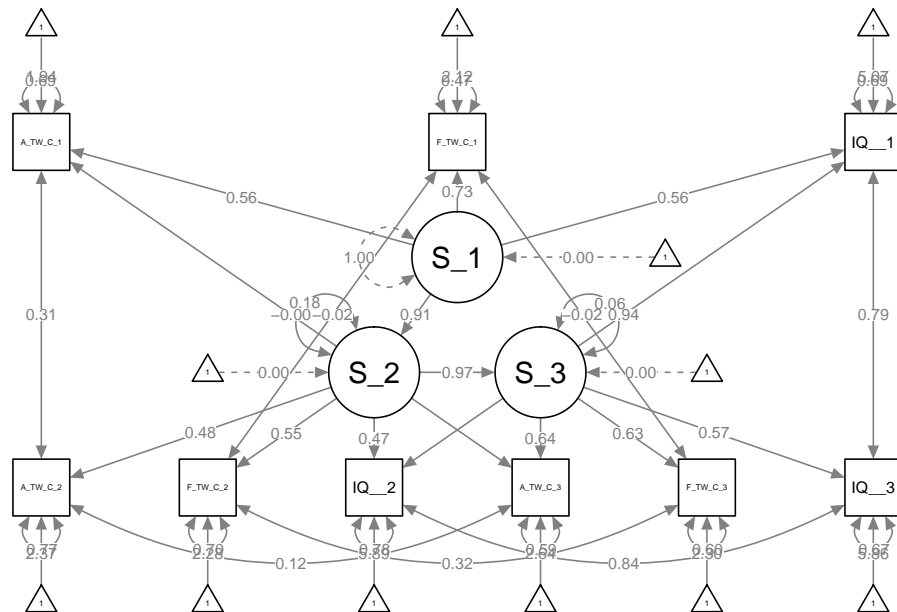
```

##
## Regressions:
##           Estimate Std.Err z-value P(>|z|)
## Semantic_3 ~
## Semantic_2      1.239   0.183   6.784   0.000
## Semantic_2 ~
## Semantic_1      0.705   0.100   7.046   0.000
##
## Covariances:
##           Estimate Std.Err z-value P(>|z|)
## .Animal_TW_Clusters_1 ~~
## .Anml_TW_Clst_2          5.269   2.728   1.932   0.053
## .Anml_TW_Clst_3         -0.028   2.461  -0.011   0.991
## .Animal_TW_Clusters_2 ~~
## .Anml_TW_Clst_3          1.621   2.397   0.677   0.499
## .Food_TW_Clusters_1 ~~
## .Fd_TW_Clstrs_2         -0.252   2.757  -0.091   0.927
## .Fd_TW_Clstrs_3         -0.280   3.114  -0.090   0.928
## .Food_TW_Clusters_2 ~~
## .Fd_TW_Clstrs_3          6.063   3.564   1.701   0.089
## .IQ_vocraw_1 ~~
## .IQ_vocraw_2          55.488  11.576   4.793   0.000
## .IQ_vocraw_3          45.660  10.958   4.167   0.000
## .IQ_vocraw_2 ~~
## .IQ_vocraw_3          47.970  10.928   4.390   0.000
##
## Intercepts:
##           Estimate Std.Err z-value P(>|z|)
## .Anml_TW_Clst_1      9.836   0.620  15.852   0.000
## .Fd_TW_Clstrs_1     10.433   0.602  17.317   0.000
## .IQ_vocraw_1       47.269   1.139  41.500   0.000
## .Anml_TW_Clst_2     10.955   0.566  19.371   0.000
## .Fd_TW_Clstrs_2     11.567   0.619  18.673   0.000
## .IQ_vocraw_2       51.000   1.058  48.215   0.000
## .Anml_TW_Clst_3     11.665   0.546  21.371   0.000
## .Fd_TW_Clstrs_3     13.028   0.703  18.535   0.000
## .IQ_vocraw_3       52.981   1.108  47.837   0.000
## Semantic_1          0.000
## .Semantic_2          0.000
## .Semantic_3          0.000
##
## Variances:
##           Estimate Std.Err z-value P(>|z|)
## .Semantic_2          0.109   0.065   1.669   0.095
## .Semantic_3          0.061   0.122   0.501   0.617
## .Anml_TW_Clst_1     17.732   3.718   4.769   0.000
## .Fd_TW_Clstrs_1     11.493   3.692   3.113   0.002
## .IQ_vocraw_1       59.903  12.746   4.700   0.000
## .Anml_TW_Clst_2     16.547   3.369   4.912   0.000
## .Fd_TW_Clstrs_2     17.943   3.957   4.535   0.000
## .IQ_vocraw_2       58.601  11.631   5.038   0.000
## .Anml_TW_Clst_3     11.501   2.977   3.863   0.000
## .Fd_TW_Clstrs_3     19.397   4.870   3.983   0.000
## .IQ_vocraw_3       55.096  12.059   4.569   0.000

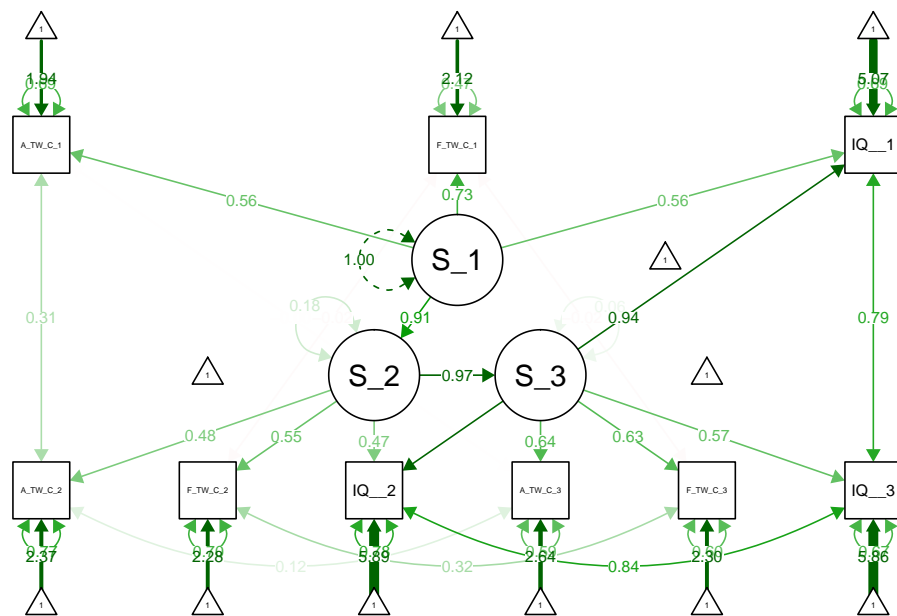
```

```
## Semantic_1 1.000
```

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



```
semPaths(fit.auto, layout = "tree", 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	15	3577.1	3663.0	26.593			
## fit.auto	20	3569.2	3644.2	28.749	2.1563	5	0.8271

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

```

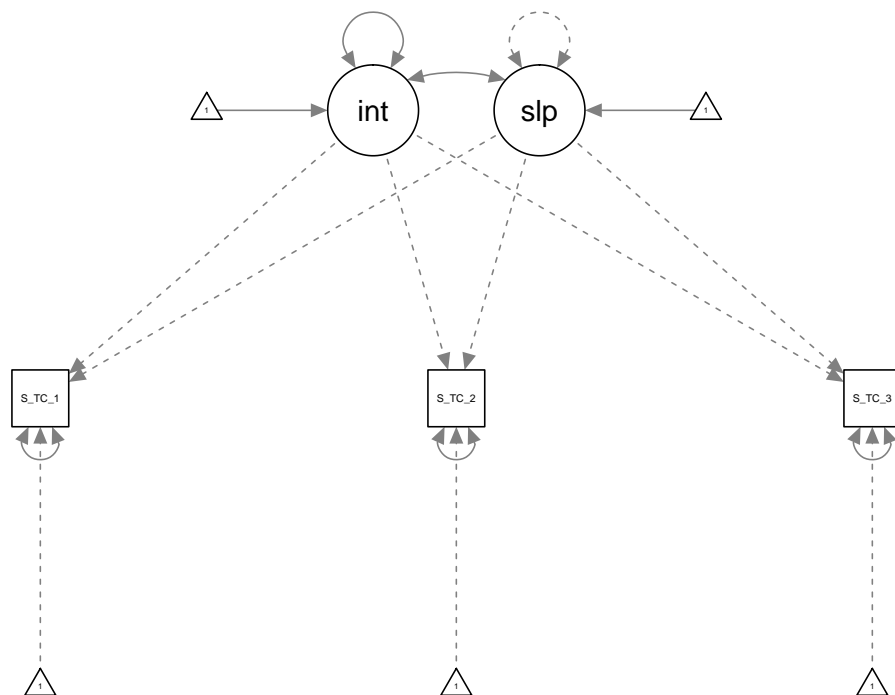


```

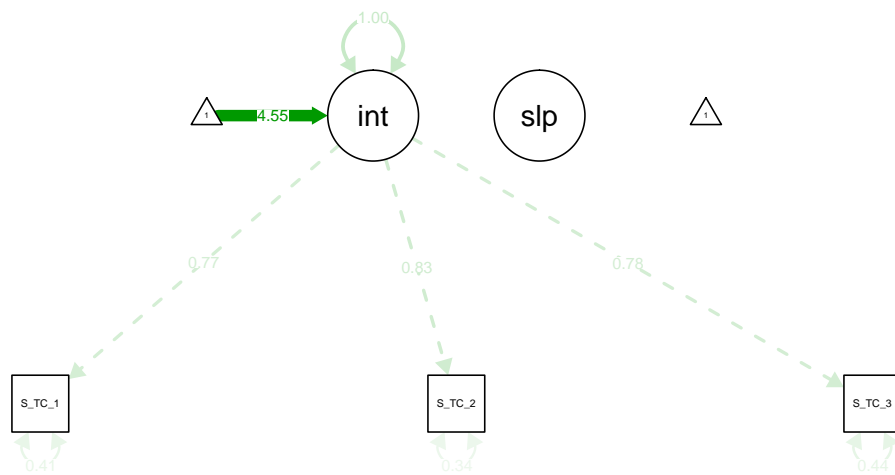
##
## Parameter Estimates:
##
## Information                                Observed
## Standard Errors                            Standard
##
## Latent Variables:
##           Estimate  Std.Err  z-value  P(>|z|)
## intercept =~
##   Sem_TtlCrrct_1    1.000
##   Sem_TtlCrrct_2    1.000
##   Sem_TtlCrrct_3    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

```

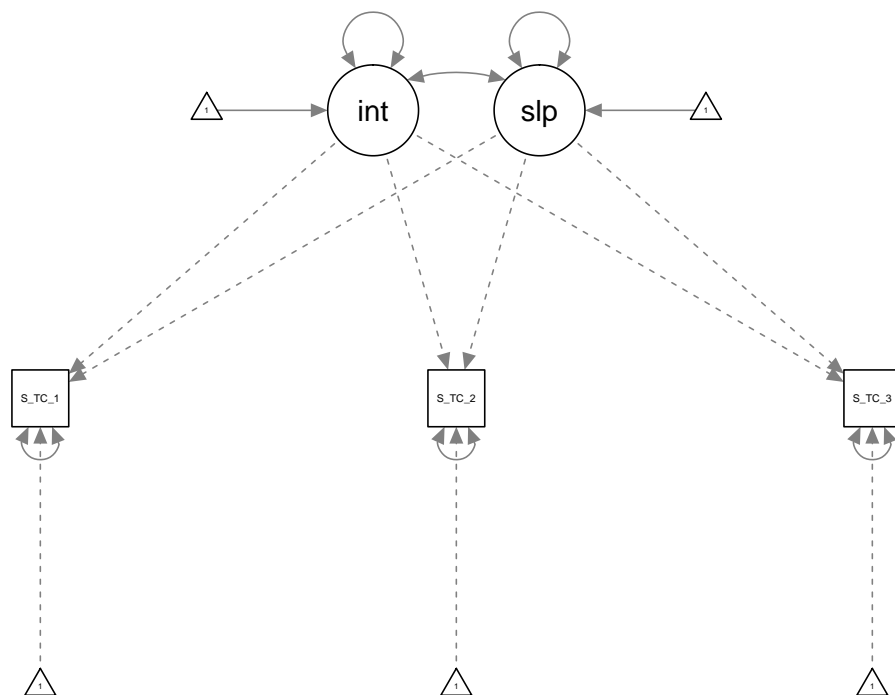
```
## lavaan (0.5-23.1097) converged normally after 78 iterations
##
```

```

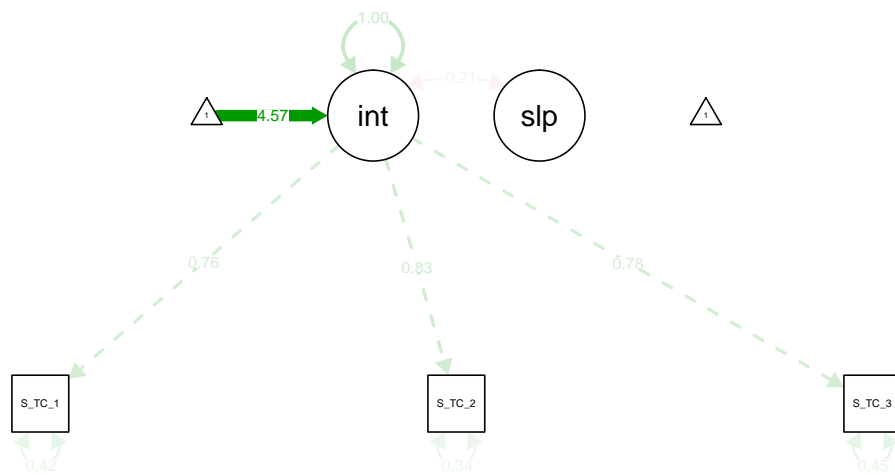
##      Number of observations                67
##
##      Number of missing patterns           2
##
##      Estimator                           ML
##      Minimum Function Test Statistic      0.062
##      Degrees of freedom                   1
##      P-value (Chi-square)                 0.803
##
## 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

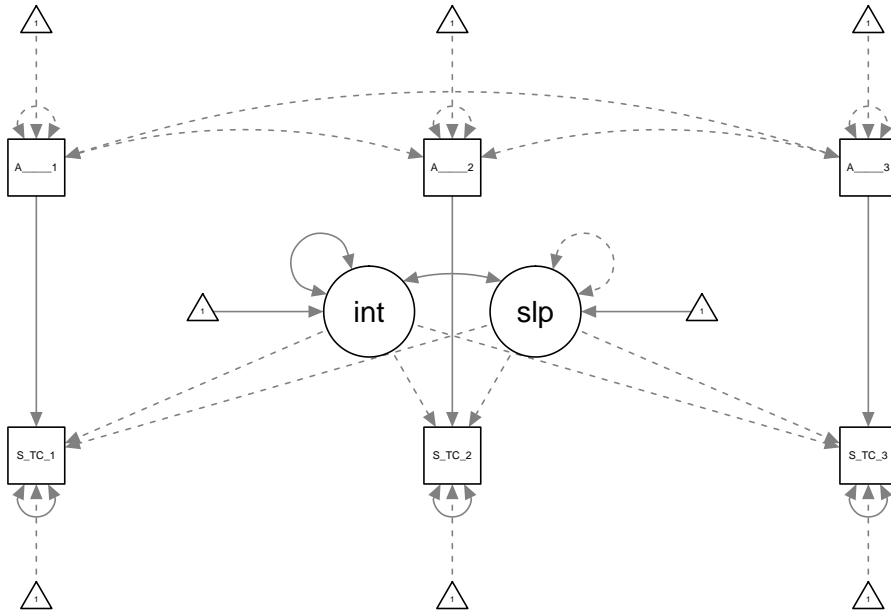
Sem_TotalCorrect_1 ~ Age_at_time_of_testing_1
Sem_TotalCorrect_2 ~ Age_at_time_of_testing_2
Sem_TotalCorrect_3 ~ Age_at_time_of_testing_3 ' #fixed slope, with covariate
```

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

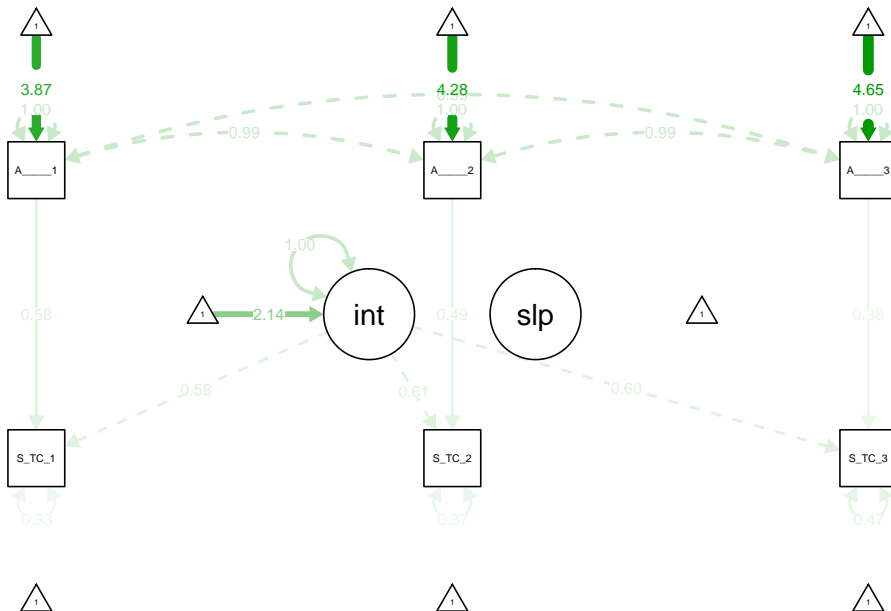
```
## lavaan (0.5-23.1097) converged normally after 67 iterations
##
##   Number of observations                67
##
##   Number of missing patterns            2
##
##   Estimator                            ML
##   Minimum Function Test Statistic      20.725
##   Degrees of freedom                    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

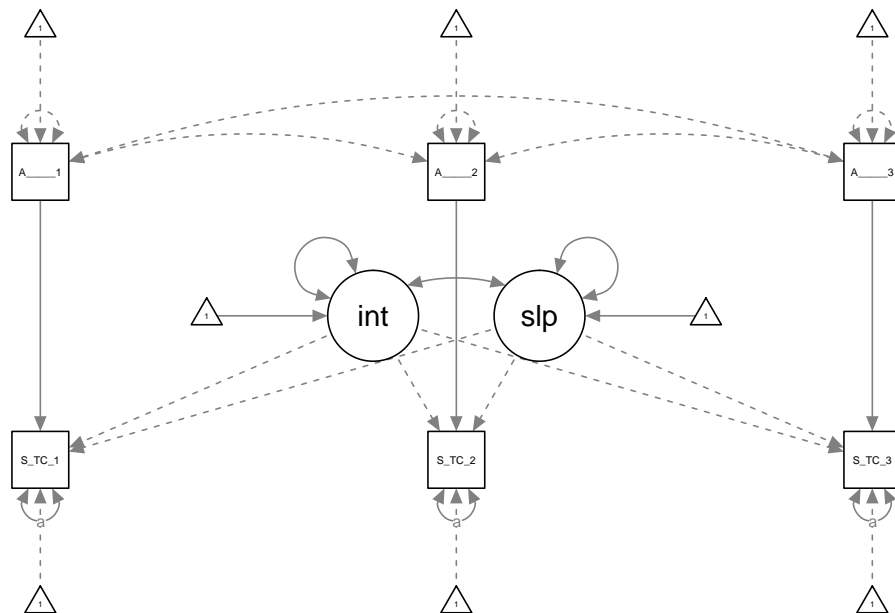
              Sem_TotalCorrect_1 ~~ a*Sem_TotalCorrect_1
              Sem_TotalCorrect_2 ~~ a*Sem_TotalCorrect_2
              Sem_TotalCorrect_3 ~~ a*Sem_TotalCorrect_3 ' #random slope, with covariate, residual vari

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

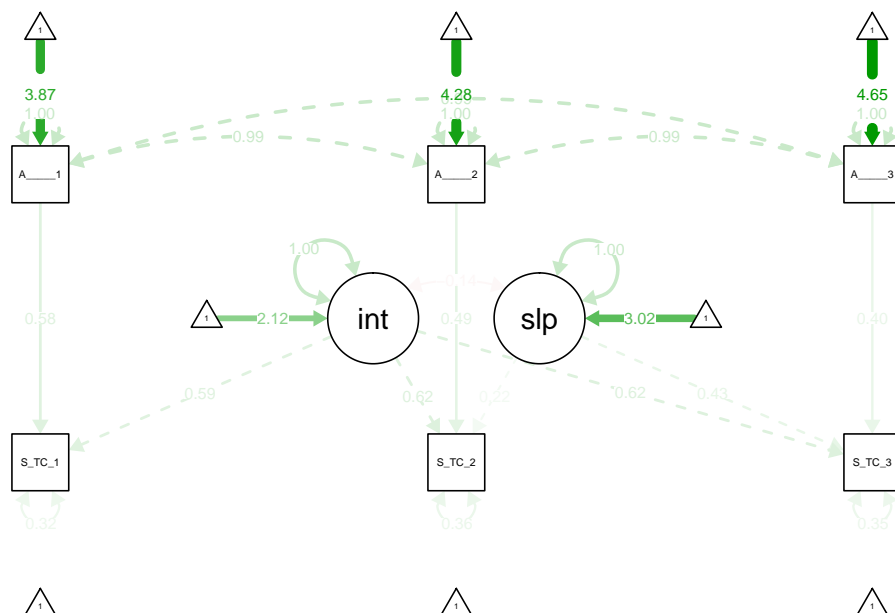
```
## lavaan (0.5-23.1097) converged normally after 86 iterations
##
##   Number of observations              67
##
##   Number of missing patterns          2
##
##   Estimator                           ML
##   Minimum Function Test Statistic      20.579
##   Degrees of freedom                   9
##   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 '

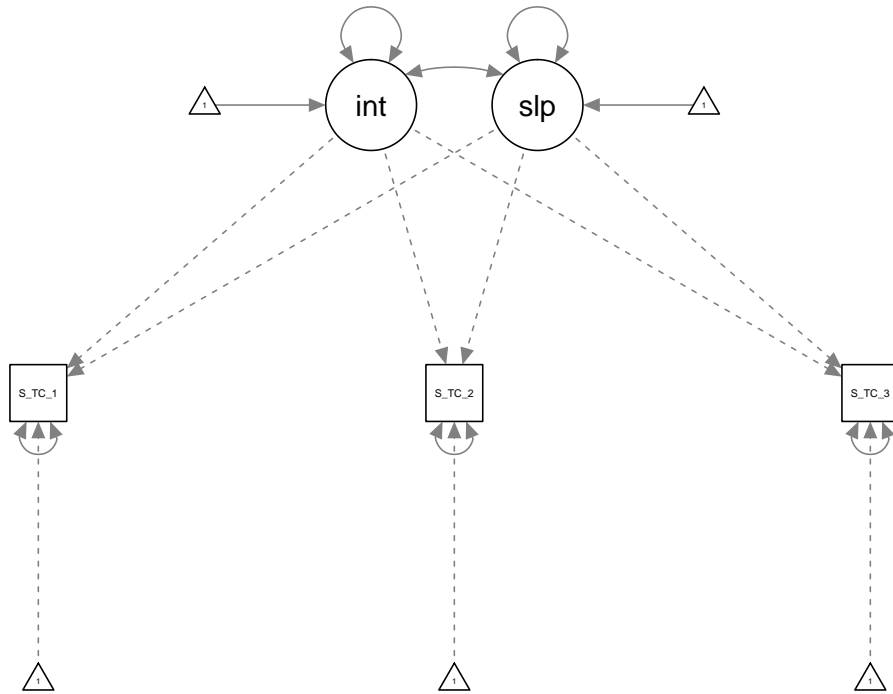
mod.SEM.time <- growth(mod.SEM5, missing = "ML", data = mydata_wide)
summary(mod.SEM.time)

## 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

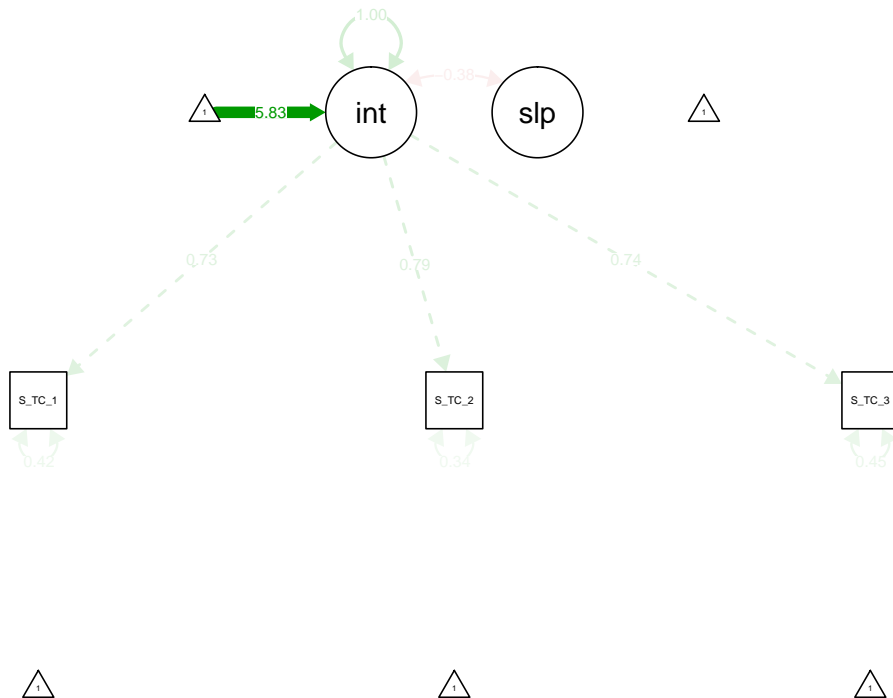
```

```
##      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 '

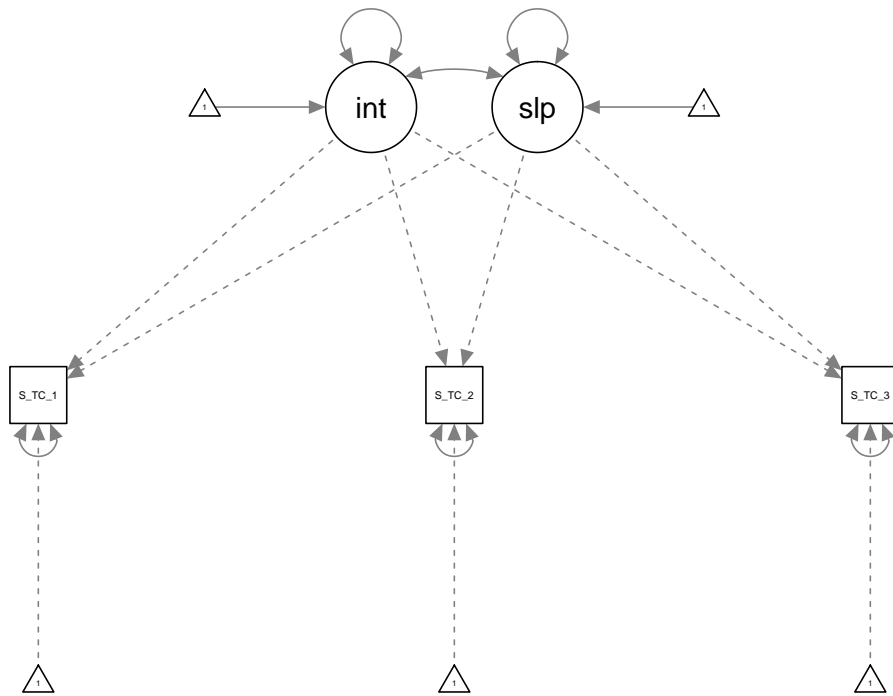
mod.SEM.time2 <- growth(mod.SEM6, missing = "ML", data = mydata_wide)
summary(mod.SEM.time2)

## 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

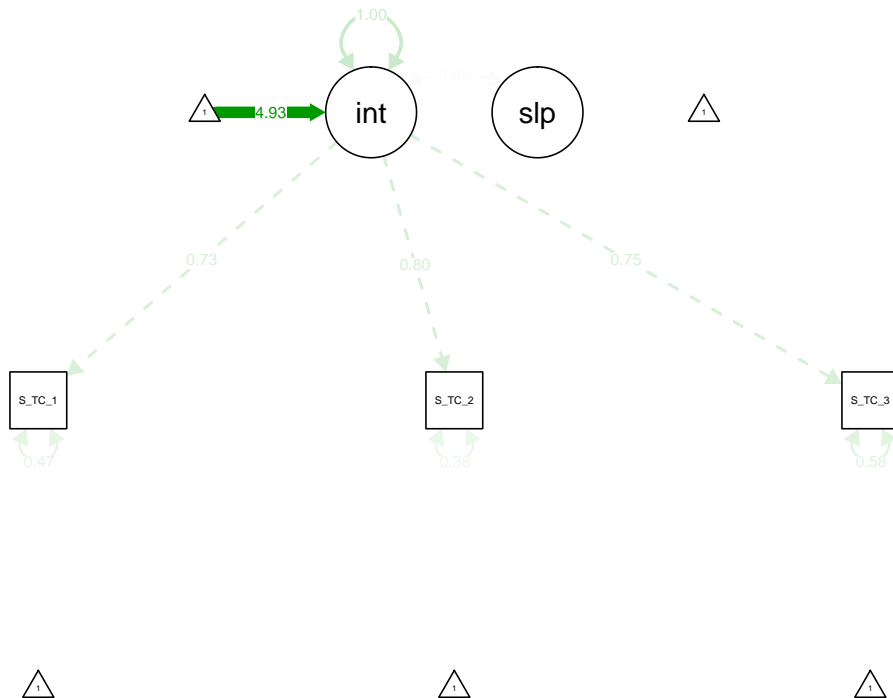
```

```
##      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
```

```
##
```

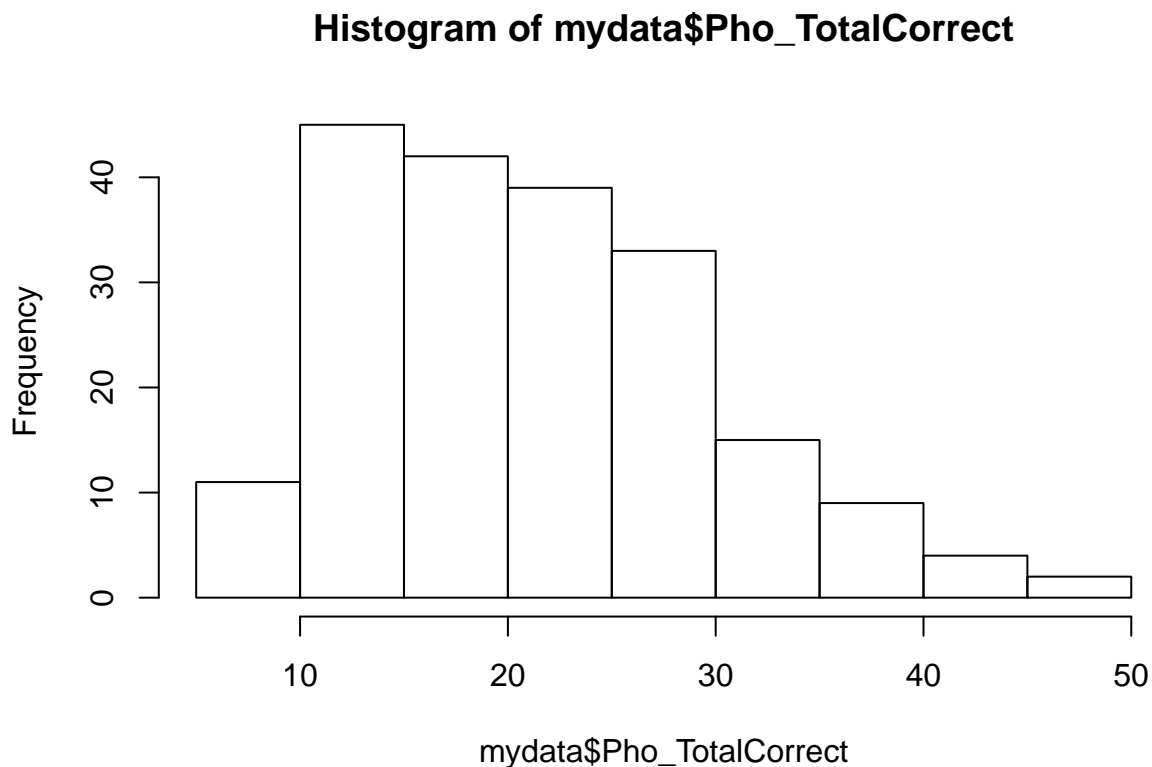
```
##      Df AIC BIC  Chisq Chisq diff Df diff Pr(>Chisq)
```

```
## 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)
```

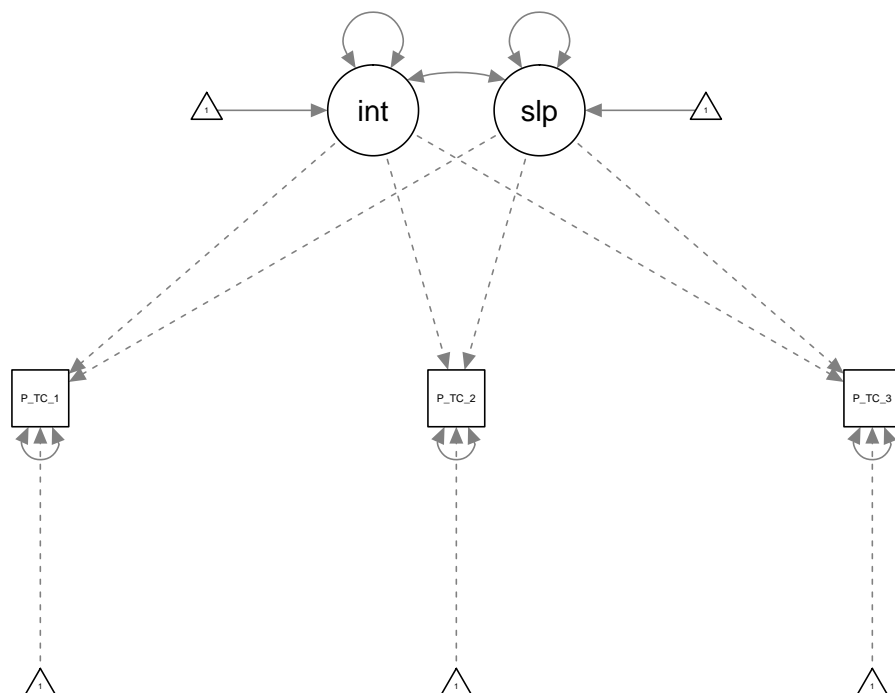
```
## lavaan (0.5-23.1097) converged normally after 80 iterations
##
##                               Used      Total
## Number of observations           66       67
##
## Estimator                      ML       Robust
## Minimum Function Test Statistic 4.151    4.420
## Degrees of freedom         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

```

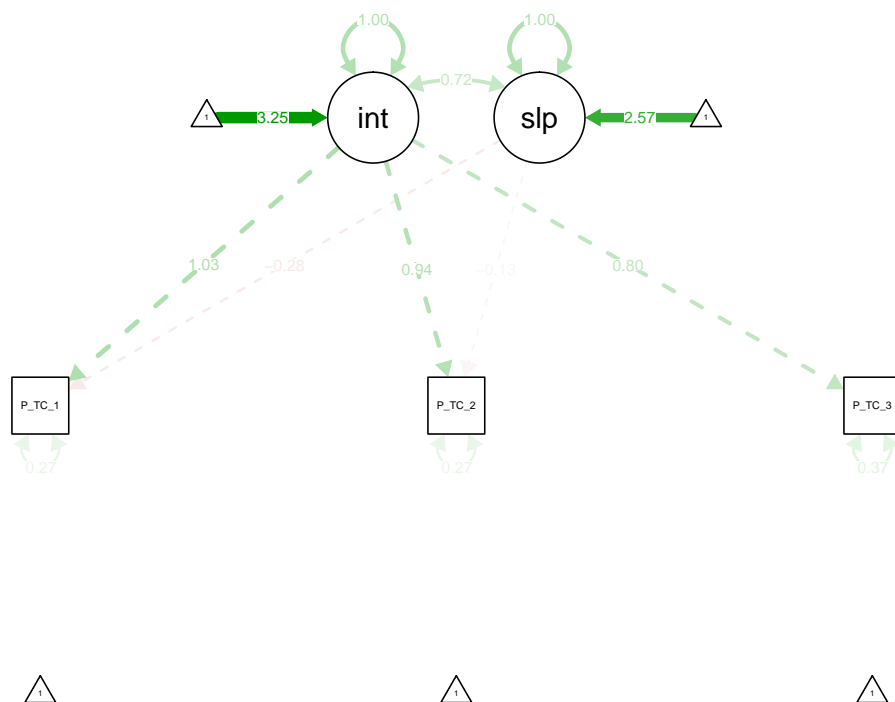
##      P-value (Chi-square)                0.042      0.036
##      Scaling correction factor            0.939
##      for the Satorra-Bentler correction
##
## Parameter Estimates:
##
##      Information                Expected
##      Standard Errors            Robust.sem
##
## Latent Variables:
##      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
```

```
##
```

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

```
##
```

```
##           Estimator                               ML
```

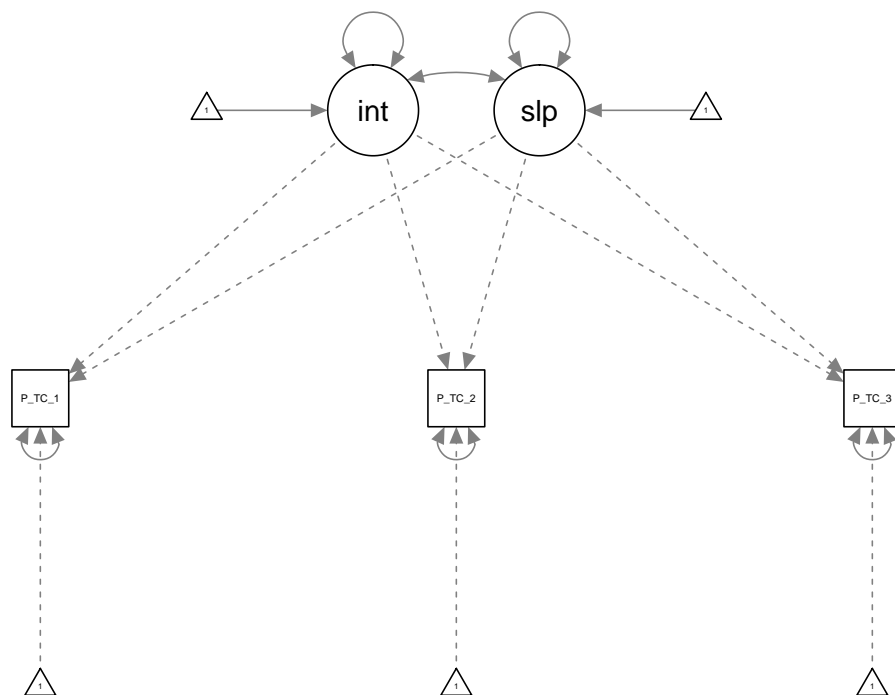


```

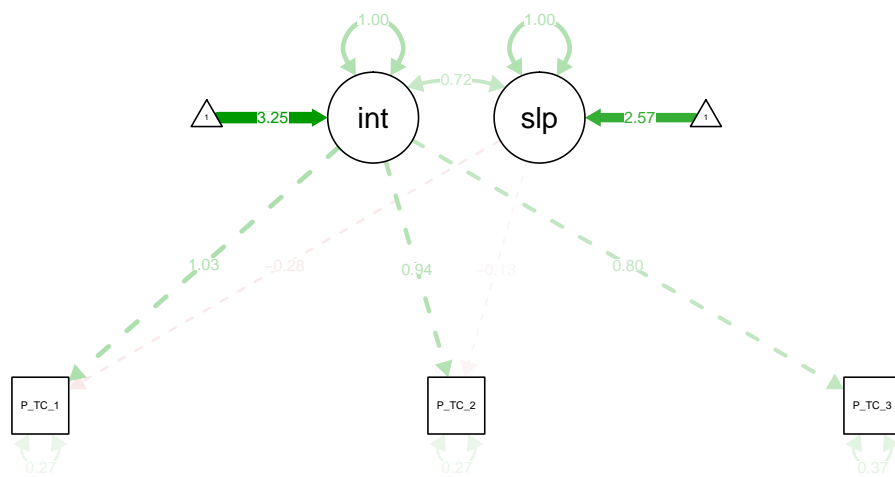
## Minimum Function Test Statistic          4.151
## Degrees of freedom                      1
## P-value (Chi-square)                    0.042
##
## Parameter Estimates:
##
## Information                               Expected
## Standard Errors                         Standard
##
## Latent Variables:
##      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)