

IRT in R

Doug Tommet

2/15/2018

Libraries

```
library(readr)
library(tidyr)
```

```
## Warning: package 'tidyr' was built under R version 3.4.2
```

```
library(tibble)
library(dplyr)
```

```
## Warning: package 'dplyr' was built under R version 3.4.2
```

```
##
## Attaching package: 'dplyr'
```

```
## The following objects are masked from 'package:stats':
##
##     filter, lag
```

```
## The following objects are masked from 'package:base':
##
##     intersect, setdiff, setequal, union
```

```
library(knitr)
```

```
## Warning: package 'knitr' was built under R version 3.4.3
```

```
library(MplusAutomation)
library(mirt)
```

```
## Warning: package 'mirt' was built under R version 3.4.3
```

```
## Loading required package: stats4
```

```
## Loading required package: lattice
```

```
library(lavaan)
```

```
## This is lavaan 0.5-23.1097
```

```
## lavaan is BETA software! Please report any bugs.
```

Read in the data

Read in the tester simulation data.

```
tester <- read_csv(file = "/Users/doulastommet/Dropbox/ADNIpsychometrics/PRIVATE/RNJ/tester.csv")
```

```
## Parsed with column specification:
## cols(
##   u1 = col_integer(),
##   u2 = col_integer(),
##   u3 = col_integer(),
##   u4 = col_integer(),
##   u5 = col_integer(),
##   u6 = col_integer(),
##   u7 = col_integer(),
##   u8 = col_integer(),
##   u9 = col_integer()
## )
```

Mplus - WLSMV theta parameterization

```

mod.mplus <- " f by u1* u2 u3 u4 u5 u6 u7 u8 u9;
              f @1; "
mod.wlsmv.t.body <- mplusObject(
  VARIABLE = "CATEGORICAL ARE all;",
  ANALYSIS = "estimator = wlsmv; parameterization = theta;",
  MODEL = mod.mplus,
  OUTPUT = "",
  SAVEDATA = "",
  usevariables = c("u1", "u2", "u3", "u4", "u5", "u6", "u7", "u8", "u9"),
  rdata = tester
)
mod.wlsmv.t.fit <- mplusModeler(mod.wlsmv.t.body,
                                   modelout = "mod.wlsmv.t.inp", run = TRUE,
                                   hashfilename = FALSE, writeData = 'always')

```

```
## Wrote model to: mod.wlsmv.t.inp
```

```
## Wrote data to: mod.wlsmv.t.dat
```

```

## Warning in prepareMplusData(df = data[i, , drop = FALSE], keepCols = object
## $usevariables, : The file 'mod.wlsmv.t.dat' currently exists and will be
## overwritten

```

```

## 
## Running model: mod.wlsmv.t.inp
## System command: cd "." && "/Applications/Mplus/mplus" "mod.wlsmv.t.inp"
## Reading model: mod.wlsmv.t.out

```

```

param <- readModels("mod.wlsmv.t.out")$parameters$unstandardized %>%
  as_tibble()

```

```
## Reading model: mod.wlsmv.t.out
```

```

param.wlsmv.theta <- param %>%
  rename(est_theta = est) %>%
  mutate(param = tolower(param))

```

Mplus - WLSMV delta parameterization

```
mod.wlsmv.d.body <- mplusObject(
  VARIABLE = "CATEGORICAL ARE all;",
  ANALYSIS = "estimator = wlsmv; parameterization = delta;",
  MODEL = mod.mplus,
  OUTPUT = "",
  SAVEDATA = "",
  usevariables = c("u1", "u2", "u3", "u4", "u5", "u6", "u7", "u8", "u9"),
  rdata = tester
)
mod.wlsmv.d.fit <- mplusModeler(mod.wlsmv.d.body,
  modelout = "mod.wlsmv.d.inp", run = TRUE,
  hashfilename = FALSE, writeData = 'always')
```

```
## Wrote model to: mod.wlsmv.d.inp
```

```
## Wrote data to: mod.wlsmv.d.dat
```

```
## Warning in prepareMplusData(df = data[i, , drop = FALSE], keepCols = object
## $usevariables, : The file 'mod.wlsmv.d.dat' currently exists and will be
## overwritten
```

```
##
## Running model: mod.wlsmv.d.inp
## System command: cd "." && "/Applications/Mplus/mplus" "mod.wlsmv.d.inp"
## Reading model: mod.wlsmv.d.out
```

```
param <- readModels("mod.wlsmv.d.out")$parameters$unstandardized %>%
  as_tibble()
```

```
## Reading model: mod.wlsmv.d.out
```

```
param.wlsmv.delta <- param %>%
  rename(est_delta = est) %>%
  mutate(param = tolower(param))
```

Mplus - MLR

```
mod.mlr.body <- mplusObject(
  VARIABLE = "CATEGORICAL ARE all;",
  ANALYSIS = "estimator = mlr; ",
  MODEL = mod.mplus,
  OUTPUT = "",
  SAVEDATA = "",
  usevariables = c("u1", "u2", "u3", "u4", "u5", "u6", "u7", "u8", "u9"),
  rdata = tester
)
mod.mlr.fit <- mplusModeler(mod.mlr.body,
  modelout = "mod.mlr.inp", run = TRUE,
  hashfilename = FALSE, writeData = 'always')
```

Wrote model to: mod.mlr.inp

Wrote data to: mod.mlr.dat

Warning in prepareMplusData(df = data[i, , drop = FALSE], keepCols = object
\$usevariables, : The file 'mod.mlr.dat' currently exists and will be
overwritten

##
Running model: mod.mlr.inp
System command: cd "." && "/Applications/Mplus/mplus" "mod.mlr.inp"
Reading model: mod.mlr.out

No PROPORTION OF DATA PRESENT sections found within COVARIANCE COVERAGE OF DATA output.

```
param <- readModels("mod.mlr.out")$parameters$unstandardized %>%
  as_tibble()
```

Reading model: mod.mlr.out

No PROPORTION OF DATA PRESENT sections found within COVARIANCE COVERAGE OF DATA output.

```
param.mlr <- param %>%
  rename(est_mlr = est) %>%
  mutate(param = tolower(param))
```

Mplus - Bayes

```
mod.bayes.body <- mplusObject(
  VARIABLE = "CATEGORICAL ARE all;",
  ANALYSIS = "estimator = bayes; ",
  MODEL = mod.mplus,
  OUTPUT = "",
  SAVEDATA = "",
  usevariables = c("u1", "u2", "u3", "u4", "u5", "u6", "u7", "u8", "u9"),
  rdata = tester
)
# mod.bayes.fit <- mplusModeler(mod.bayes.body,
#                               modelout = "mod.bayes.inp", run = TRUE,
#                               hashfilename = FALSE, writeData = 'always')

param <- readModels("mod.bayes.out")$parameters$unstandardized %>%
  as_tibble()
```

Reading model: mod.bayes.out

```
param.bayes <- param %>%
  rename(est_bayes = est) %>%
  mutate(param = tolower(param))
```

Mirt

```
mod.mirt <- mirt(tester, 1)
```

```
##  
Iteration: 1, Log-Lik: -192023.608, Max-Change: 0.68539  
Iteration: 2, Log-Lik: -186726.777, Max-Change: 0.56274  
Iteration: 3, Log-Lik: -185487.476, Max-Change: 0.30051  
Iteration: 4, Log-Lik: -184966.068, Max-Change: 0.20538  
Iteration: 5, Log-Lik: -184734.757, Max-Change: 0.11239  
Iteration: 6, Log-Lik: -184619.378, Max-Change: 0.08752  
Iteration: 7, Log-Lik: -184550.889, Max-Change: 0.05454  
Iteration: 8, Log-Lik: -184487.880, Max-Change: 0.04380  
Iteration: 9, Log-Lik: -184458.065, Max-Change: 0.03829  
Iteration: 10, Log-Lik: -184434.485, Max-Change: 0.03671  
Iteration: 11, Log-Lik: -184419.066, Max-Change: 0.04024  
Iteration: 12, Log-Lik: -184405.511, Max-Change: 0.02249  
Iteration: 13, Log-Lik: -184390.744, Max-Change: 0.01721  
Iteration: 14, Log-Lik: -184381.806, Max-Change: 0.01177  
Iteration: 15, Log-Lik: -184376.337, Max-Change: 0.00897  
Iteration: 16, Log-Lik: -184372.953, Max-Change: 0.00695  
Iteration: 17, Log-Lik: -184370.890, Max-Change: 0.00593  
Iteration: 18, Log-Lik: -184369.581, Max-Change: 0.00458  
Iteration: 19, Log-Lik: -184367.471, Max-Change: 0.00063  
Iteration: 20, Log-Lik: -184367.448, Max-Change: 0.00045  
Iteration: 21, Log-Lik: -184367.437, Max-Change: 0.00037  
Iteration: 22, Log-Lik: -184367.412, Max-Change: 0.00007
```

```
param.mirt <- coef(mod.mirt, simplify=TRUE)$items %>%  
  as.data.frame() %>%  
  rownames_to_column("param") %>%  
  rename(a1_mirt = a1)
```

Mirt summary

```
summary(mod.mirt)
```

```
##      F1      h2
## u1  0.757  0.5727
## u2  0.532  0.2834
## u3  0.864  0.7472
## u4  0.857  0.7344
## u5  0.722  0.5206
## u6  0.301  0.0907
## u7  0.712  0.5063
## u8  0.495  0.2448
## u9  0.703  0.4935
##
## SS loadings:  4.194
## Proportion Var:  0.466
##
## Factor correlations:
##
##      F1
## F1   1
```

Mirt coef

```
coef(mod.mirt, simplify=TRUE)
```

```
## $items
##      a1      d g u
## u1 1.970 -4.380 0 1
## u2 1.070 -2.615 0 1
## u3 2.926 -3.549 0 1
## u4 2.830 -3.212 0 1
## u5 1.774 -2.070 0 1
## u6 0.538 -1.381 0 1
## u7 1.724 -1.632 0 1
## u8 0.969 -1.185 0 1
## u9 1.680 -1.250 0 1
##
## $means
## F1
## 0
##
## $cov
##      F1
## F1  1
```

Lavaan

```
mod.lavaan.body <- 'f == NA*u1 + u2 + u3 + u4 + u5 + u6 + u7 + u8 + u9
f--1*f
'
mod.lavaan.fit <- sem(mod.lavaan.body,
                       data=tester ,
                       ordered=c("u1", "u2", "u3", "u4", "u5", "u6", "u7", "u8", "u9"))

param.lavaan <- parameterestimates(mod.lavaan.fit) %>%
  rename(est_lavaan = est)
```

Lavaan summary

```
summary(mod.lavaan.fit, fit.measures=TRUE, standardized=TRUE)
```

```
## lavaan (0.5-23.1097) converged normally after 11 iterations
##
## Number of observations 50000
##
## Estimator DWLS Robust
## Minimum Function Test Statistic 12.004 18.568
## Degrees of freedom 27 27
## P-value (Chi-square) 0.994 0.885
## Scaling correction factor 0.657
## Shift parameter 0.308
##      for simple second-order correction (Mplus variant)
##
## Model test baseline model:
##
## Minimum Function Test Statistic 131325.879 101198.182
## Degrees of freedom 36 36
## P-value 0.000 0.000
##
## User model versus baseline model:
##
## Comparative Fit Index (CFI) 1.000 1.000
## Tucker-Lewis Index (TLI) 1.000 1.000
##
## Robust Comparative Fit Index (CFI) NA
## Robust Tucker-Lewis Index (TLI) NA
##
## Root Mean Square Error of Approximation:
##
## RMSEA 0.000 0.000
## 90 Percent Confidence Interval 0.000 0.000 0.000 0.002
## P-value RMSEA <= 0.05 1.000 1.000
##
## Robust RMSEA NA
## 90 Percent Confidence Interval 0.000 NA
##
## Standardized Root Mean Square Residual:
##
## SRMR 0.005 0.005
##
## Weighted Root Mean Square Residual:
##
## WRMR 0.516 0.516
##
## Parameter Estimates:
##
## Information Expected
## Standard Errors Robust.sem
##
## Latent Variables:
##             Estimate Std.Err z-value P(>|z|) Std.lv Std.all
## f ==
## u1        0.703  0.008  91.099  0.000  0.703  0.703
## u2        0.496  0.008  62.419  0.000  0.496  0.496
## u3        0.853  0.004 206.122  0.000  0.853  0.853
## u4        0.846  0.004 206.358  0.000  0.846  0.846
## u5        0.711  0.005 138.736  0.000  0.711  0.711
## u6        0.296  0.007  39.734  0.000  0.296  0.296
## u7        0.707  0.005 142.783  0.000  0.707  0.707
## u8        0.496  0.006  78.923  0.000  0.496  0.496
## u9        0.701  0.005 143.558  0.000  0.701  0.701
```

```

## Intercepts:
##             Estimate Std.Err z-value P(>|z|) Std.lv Std.all
##   .u1        0.000
##   .u2        0.000
##   .u3        0.000
##   .u4        0.000
##   .u5        0.000
##   .u6        0.000
##   .u7        0.000
##   .u8        0.000
##   .u9        0.000
##   f         0.000
##
## Thresholds:
##             Estimate Std.Err z-value P(>|z|) Std.lv Std.all
##   u1|t1     1.645  0.009 174.042  0.000  1.645  1.645
##   u2|t1     1.283  0.008 167.700  0.000  1.283  1.283
##   u3|t1     1.036  0.007 151.276  0.000  1.036  1.036
##   u4|t1     0.961  0.007 144.354  0.000  0.961  0.961
##   u5|t1     0.835  0.006 130.952  0.000  0.835  0.835
##   u6|t1     0.793  0.006 126.001  0.000  0.793  0.793
##   u7|t1     0.669  0.006 109.995  0.000  0.669  0.669
##   u8|t1     0.614  0.006 102.201  0.000  0.614  0.614
##   u9|t1     0.520  0.006  88.309  0.000  0.520  0.520
##
## Variances:
##             Estimate Std.Err z-value P(>|z|) Std.lv Std.all
##   f          1.000
##   .u1        0.506
##   .u2        0.754
##   .u3        0.273
##   .u4        0.285
##   .u5        0.494
##   .u6        0.913
##   .u7        0.501
##   .u8        0.754
##   .u9        0.509
##
## Scales y*:
##             Estimate Std.Err z-value P(>|z|) Std.lv Std.all
##   u1        1.000
##   u2        1.000
##   u3        1.000
##   u4        1.000
##   u5        1.000
##   u6        1.000
##   u7        1.000
##   u8        1.000
##   u9        1.000

```

Results

Merging the slope results together.

```

param.truth <- tibble(param = c("u1", "u2", "u3", "u4", "u5", "u6", "u7", "u8", "u9"),
                      truth.slope = c("1.000", "0.577", "1.614", "1.614", "1.000", "0.314", "1.000", "0.577", "1.000"))

param.wlsmv.theta.sub <- param.wlsmv.theta %>%
  filter(paramHeader=="F.BY") %>%
  select(param, est_theta) %>%
  rename(slope_theta = est_theta)

param.wlsmv.delta.sub <- param.wlsmv.delta %>%
  filter(paramHeader=="F.BY") %>%
  select(param, est_delta) %>%
  rename(slope_delta = est_delta)

param.mlrv.sub <- param.mlrv %>%
  filter(paramHeader=="F.BY") %>%
  select(param, est_mlrv) %>%
  rename(slope_mlrv = est_mlrv)

param.bayes.sub <- param.bayes %>%
  filter(paramHeader=="F.BY") %>%
  select(param, est_bayes) %>%
  rename(slope_bayes = est_bayes)

param.mirt.sub <- param.mirt %>%
  select(param, al_mirt) %>%
  rename(slope_mirt = al_mirt)

param.lavaan.sub <- param.lavaan %>%
  filter(op=="-") %>%
  rename(param = rhs) %>%
  select(param, est_lavaan) %>%
  rename(slope_lavaan = est_lavaan)

param.table.slope <- param.truth %>%
  left_join(param.wlsmv.theta.sub, by = "param") %>%
  left_join(param.wlsmv.delta.sub, by = "param") %>%
  left_join(param.mlrv.sub, by = "param") %>%
  left_join(param.bayes.sub, by = "param") %>%
  left_join(param.mirt.sub, by = "param") %>%
  left_join(param.lavaan.sub, by = "param")

```

```

param.truth <- tibble(param = c("u1", "u2", "u3", "u4", "u5", "u6", "u7", "u8", "u9"),
                      truth.thresh = c("2.327", "2.563", "1.219", "1.123", "1.190", "2.574", "0.954", "1.226", "0.742"))

param.wlsmv.theta.sub <- param.wlsmv.theta %>%
  filter(paramHeader=="Thresholds") %>%
  separate(param, c("param", "threshold"), sep = "\\$") %>%
  select(param, est_theta) %>%
  rename(thres_theta = est_theta)

param.wlsmv.delta.sub <- param.wlsmv.delta %>%
  filter(paramHeader=="Thresholds") %>%
  separate(param, c("param", "threshold"), sep = "\\$") %>%
  select(param, est_delta) %>%
  rename(thres_delta = est_delta)

param.mlrv.sub <- param.mlrv %>%
  filter(paramHeader=="Thresholds") %>%

```

```

filter(paramHeader == "thresholds", ~
separate(param, c("param", "threshold"), sep = "\\$") %>%
select(param, est_mlr) %>%
rename(thres_mlr = est_mlr)

param.bayes.sub <- param.bayes %>%
filter(paramHeader == "Thresholds") %>%
separate(param, c("param", "threshold"), sep = "\\$") %>%
select(param, est_bayes) %>%
rename(thres_bayes = est_bayes)

param.mirt.sub <- param.mirt %>%
select(param, d) %>%
rename(d_mirt = d)

param.lavaan.sub <- param.lavaan %>%
filter(op == "|") %>%
rename(param = lhs) %>%
select(param, est_lavaan) %>%
rename(thres_lavaan = est_lavaan)

param.table.threshold <- param.truth %>%
left_join(param.wlsmv.theta.sub, by = "param") %>%
left_join(param.wlsmv.delta.sub, by = "param") %>%
left_join(param.mlr.sub, by = "param") %>%
left_join(param.bayes.sub, by = "param") %>%
left_join(param.mirt.sub, by = "param") %>%
left_join(param.lavaan.sub, by = "param")

```

```

param.table <- param.table.slope %>%
left_join(param.table.threshold, by = "param")

```

Slope estimate table

These are the slope estimates, not standardized, under various estimator and parameterization options.

```
param.table %>%
  select(param, truth.slope, starts_with("slope_")) %>%
  kable(digits = 3)
```

param	truth.slope	slope_theta	slope_delta	slope_mlr	slope_bayes	slope_wls
u1	1.000	0.987	0.703	1.969	0.990	0.990
u2	0.577	0.571	0.496	1.070	0.564	0.564
u3	1.614	1.632	0.853	2.926	1.642	1.642
u4	1.614	1.585	0.846	2.830	1.591	1.591
u5	1.000	1.012	0.711	1.774	1.007	1.007
u6	0.314	0.310	0.296	0.538	0.311	0.311
u7	1.000	0.999	0.707	1.724	1.000	1.000
u8	0.577	0.571	0.496	0.969	0.572	0.572
u9	1.000	0.982	0.701	1.681	0.978	0.978

2PL slope estimate table

These are the slope estimates converted to 2PL a-parameters using **math**.

```
param.table <- param.table %>%
  mutate(a_theta = slope_theta,
        a_delta = slope_delta/sqrt(1-slope_delta^2),
        a_mlr = slope_mlr/1.7,
        a_bayes = slope_bayes,
        a_mirt = slope_mirt/1.7,
        a_lavaan = slope_lavaan/sqrt(1-slope_lavaan^2))
```

```
param.table %>%
  select(param, truth.slope, starts_with("a_")) %>%
  kable(digits = 3)
```

param	truth.slope	a_theta	a_delta	a_mlr	a_bayes	a_mirt	a_lavaan
u1	1.000	0.987	0.988	1.158	0.990	1.159	0.987
u2	0.577	0.571	0.571	0.629	0.564	0.630	0.571
u3	1.614	1.632	1.634	1.721	1.642	1.721	1.632
u4	1.614	1.585	1.587	1.665	1.591	1.665	1.585
u5	1.000	1.012	1.011	1.044	1.007	1.043	1.012
u6	0.314	0.310	0.310	0.316	0.311	0.316	0.309
u7	1.000	0.999	1.000	1.014	1.000	1.014	0.999
u8	0.577	0.571	0.571	0.570	0.572	0.570	0.571
u9	1.000	0.982	0.983	0.989	0.978	0.988	0.982

Threshold estimate table

These are threshold estimates, not standardized, under various estimator and parameterization options.

```
param.table %>%
  select(param, starts_with("thres_"), d_mirt) %>%
  kable(digits = 3)
```

param	thres_theta	thres_delta	thres_mir	thres_bayes	thres_lavaan	d_mirt
u1	2.312	1.645	4.378	2.317	1.645	
u2	1.477	1.283	2.615	1.473	1.283	
u3	1.982	1.036	3.548	1.990	1.036	
u4	1.800	0.961	3.211	1.805	0.961	
u5	1.188	0.835	2.069	1.185	0.835	
u6	0.830	0.793	1.381	0.831	0.793	
u7	0.946	0.669	1.632	0.947	0.669	
u8	0.707	0.614	1.185	0.708	0.614	
u9	0.729	0.520	1.250	0.728	0.520	

2PL location estimate

These are the threshold estimates converted to 2PL b-parameters using **math**.

```
param.table <- param.table %>%
  mutate(b_theta = thres_theta/slope_theta,
        b_delta = thres_delta/slope_delta,
        b_mlr = thres_mlr/slope_mlr,
        b_bayes = thres_bayes/slope_bayes,
        b_lavaan = thres_lavaan/slope_lavaan,
        b_mirt = -1 * d_mirt/slope_mirt)
```

```
param.table %>%
  select(param, truth.thresh, starts_with("b_")) %>%
  kable(digits = 3)
```

param	truth.thresh	b_theta	b_delta	b_mlr	b_bayes	b_lavaan	b_mirt
u1	2.327	2.342	2.340	2.223	2.340	2.342	2.223
u2	2.563	2.587	2.587	2.444	2.612	2.587	2.444
u3	1.219	1.214	1.215	1.213	1.212	1.215	1.213
u4	1.123	1.136	1.136	1.135	1.135	1.136	1.135
u5	1.190	1.174	1.174	1.166	1.177	1.174	1.167
u6	2.574	2.677	2.679	2.567	2.672	2.683	2.569
u7	0.954	0.947	0.946	0.947	0.947	0.947	0.947
u8	1.226	1.238	1.238	1.223	1.238	1.238	1.223
u9	0.742	0.742	0.742	0.744	0.744	0.742	0.744