

R Notebook

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This is an [R Markdown](#) Notebook. When you execute code within the notebook, the results appear beneath the code.

Try executing this chunk by clicking the *Run* button within the chunk or by placing your cursor inside it and pressing *Ctrl+Shift+Enter*.

```
library(ltm)

## Warning: package 'ltm' was built under R version 4.0.5
## Loading required package: MASS
## Loading required package: msm
## Warning: package 'msm' was built under R version 4.0.5
## Loading required package: polycor
## Warning: package 'polycor' was built under R version 4.0.5

library(mokken)

## Warning: package 'mokken' was built under R version 4.0.5
## Loading required package: poLCA
## Warning: package 'poLCA' was built under R version 4.0.5
## Loading required package: scatterplot3d

library(car)

## Warning: package 'car' was built under R version 4.0.5
## Loading required package: carData

## Error : 'format_warning' is not an exported object from 'namespace:cli'

## Warning: replacing previous import 'ellipsis::check_dots_unnamed' by
## 'rlang::check_dots_unnamed' when loading 'hms'

## Warning: replacing previous import 'ellipsis::check_dots_used' by
## 'rlang::check_dots_used' when loading 'hms'

## Warning: replacing previous import 'ellipsis::check_dots_empty' by
## 'rlang::check_dots_empty' when loading 'hms'
```

```

## Warning: replacing previous import 'ellipsis::check_dots_unnamed' by
## 'rlang::check_dots_unnamed' when loading 'tibble'

## Warning: replacing previous import 'ellipsis::check_dots_used' by
## 'rlang::check_dots_used' when loading 'tibble'

## Warning: replacing previous import 'ellipsis::check_dots_empty' by
## 'rlang::check_dots_empty' when loading 'tibble'

## Warning: replacing previous import 'ellipsis::check_dots_unnamed' by
## 'rlang::check_dots_unnamed' when loading 'pillar'

## Warning: replacing previous import 'ellipsis::check_dots_used' by
## 'rlang::check_dots_used' when loading 'pillar'

## Warning: replacing previous import 'ellipsis::check_dots_empty' by
## 'rlang::check_dots_empty' when loading 'pillar'

##
## Attaching package: 'car'

## The following object is masked from 'package:mokken':
##
##      recode

library(tidyverse)

## -- Attaching packages ----- tidyverse
1.3.0 --

## v ggplot2 3.3.5      v purrr  0.3.4
## v tibble  3.1.0      v dplyr  1.0.8
## v tidyr   1.1.3      v stringr 1.4.0
## v readr   1.4.0      v forcats 0.5.1

## Warning: package 'ggplot2' was built under R version 4.0.5

## Warning: package 'dplyr' was built under R version 4.0.5

## -- Conflicts -----
tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()     masks stats::lag()
## x dplyr::recode() masks car::recode(), mokken::recode()
## x dplyr::select() masks MASS::select()
## x purrr::some()   masks car::some()

library(latticeExtra)

## Warning: package 'latticeExtra' was built under R version 4.0.5

## Loading required package: lattice

```

```
##
## Attaching package: 'latticeExtra'

## The following object is masked from 'package:ggplot2':
##
##      layer

data <- read.csv("q1Throughq6andGender.csv", header=TRUE)
data<- data[c('Q1A', 'Q2A', 'Q3A', 'Q4A', 'Q5A', 'Q6A', 'gender')]
head(data)

##   Q1A Q2A Q3A Q4A Q5A Q6A gender
## 1   4   4   2   4   4   4      2
## 2   4   1   2   3   4   4      2
## 3   3   1   4   1   4   3      2
## 4   2   3   2   1   3   3      2
## 5   2   2   3   4   4   2      2
## 6   1   1   2   1   3   1      2

data$gender[data$gender==1]<-0
data$gender[data$gender==2]<-1
# converting the data to binary for dichotomous purposes
questions = c('Q1A', 'Q2A', 'Q3A', 'Q4A', 'Q5A', 'Q6A')

for (c in questions) {
  data[[c]] <- car::recode(data[[c]], "c(1, 2)='0';c(3, 4)='1'")
}

head(data)

##   Q1A Q2A Q3A Q4A Q5A Q6A gender
## 1   1   1   0   1   1   1      1
## 2   1   0   0   1   1   1      1
## 3   1   0   1   0   1   1      1
## 4   0   1   0   0   1   1      1
## 5   0   0   1   1   1   0      1
## 6   0   0   0   0   1   0      1

dat_base <-data[c('Q1A', 'Q2A', 'Q3A', 'Q4A', 'Q5A', 'Q6A')]
head(dat_base)

##   Q1A Q2A Q3A Q4A Q5A Q6A
## 1   1   1   0   1   1   1
## 2   1   0   0   1   1   1
## 3   1   0   1   0   1   1
## 4   0   1   0   0   1   1
## 5   0   0   1   1   1   0
## 6   0   0   0   0   1   0
```

```

datM <- data[data$gender == 0,]
datM <- datM[c('Q1A', 'Q2A', 'Q3A', 'Q4A', 'Q5A', 'Q6A')]
datF <- data[data$gender == 1,]
datF <- datF[c('Q1A', 'Q2A', 'Q3A', 'Q4A', 'Q5A', 'Q6A')]

```

#at first glance the mean of each question seems to be good because not everyone is answering one way to these wquestions
summary(dat_base)

```

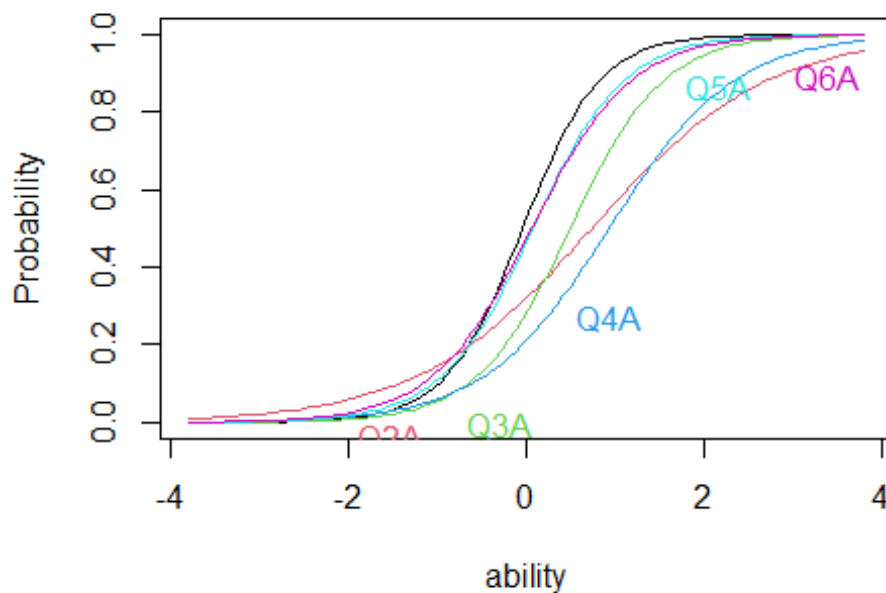
##           Q1A           Q2A           Q3A           Q4A
## Min.      :0.0000   Min.      :0.0000   Min.      :0.0000   Min.      :0.0000
## 1st Qu.:0.0000   1st Qu.:0.0000   1st Qu.:0.0000   1st Qu.:0.0000
## Median :1.0000   Median :0.0000   Median :0.0000   Median :0.0000
## Mean      :0.5117   Mean      :0.3491   Mean      :0.3558   Mean      :0.2738
## 3rd Qu.:1.0000   3rd Qu.:1.0000   3rd Qu.:1.0000   3rd Qu.:1.0000
## Max.      :1.0000   Max.      :1.0000   Max.      :1.0000   Max.      :1.0000
##           Q5A           Q6A
## Min.      :0.0000   Min.      :0.0000
## 1st Qu.:0.0000   1st Qu.:0.0000
## Median :0.0000   Median :0.0000
## Mean      :0.4769   Mean      :0.4817
## 3rd Qu.:1.0000   3rd Qu.:1.0000
## Max.      :1.0000   Max.      :1.0000

```

#testing for monotonicity
#z1 implies that we have one latent variable that we are predicting
#IRT.param=TRUE is how we set this IRT Model to 2PL (difficulty and discrimination)
model<-ltm(dat_base~z1, IRT.param=TRUE)

the steeper the slope the more
discriminable an item is
plot(model, type="ICC", xlab='ability')

Item Characteristic Curves



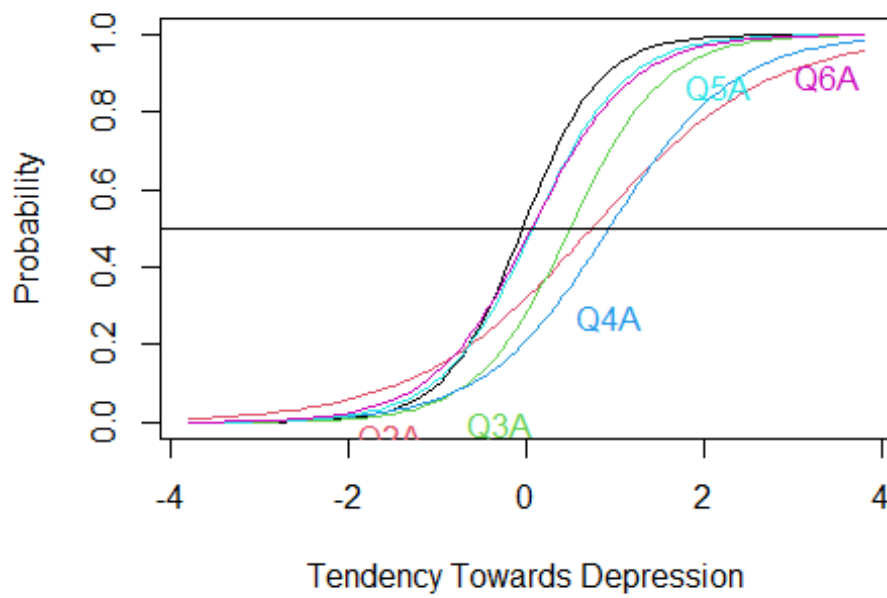
*#difficulty is a zcore of whatever the latent variable is (anxiety)
#discriminable is the slope or how good the question is at figuring a person out*

```
model<-ltm(dat_base~z1, IRT.param=TRUE)
print(coef(model))
```

```
##          Dffclt  Dscrmn
## Q1A -0.03346154 2.317070
## Q2A  0.74285394 1.016176
## Q3A  0.49802722 1.908842
## Q4A  0.93501859 1.423853
## Q5A  0.07939943 1.953463
## Q6A  0.06521985 1.790820
```

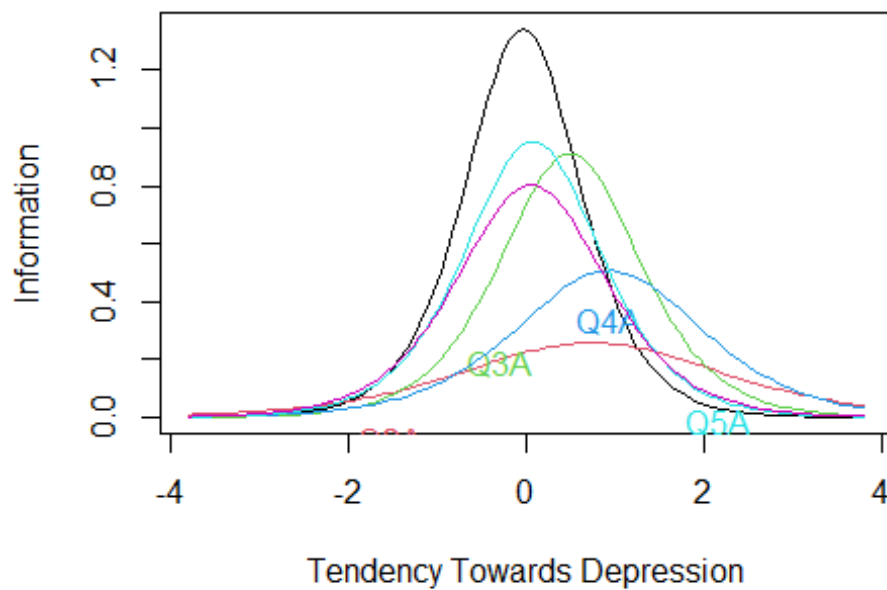
```
plot(model, type="ICC", items=c(1,2,3,4,5, 6), xlab='Tendency Towards
Depression')
abline(.5,0)
```

Item Characteristic Curves



```
plot(model, type="IIC", xlab='Tendency Towards Depression') #test information function
```

Item Information Curves



#gives you every combination of response patterns

```
factor.scores(model)
```

```
##
```

```
## Call:
```

```
## ltm(formula = dat_base ~ z1, IRT.param = TRUE)
```

```
##
```

```
## Scoring Method: Empirical Bayes
```

```
##
```

```
## Factor-Scores for observed response patterns:
```

##	Q1A	Q2A	Q3A	Q4A	Q5A	Q6A	Obs	Exp	z1	se.z1
## 1	0	0	0	0	0	0	8336	8061.059	-1.001	0.617
## 2	0	0	0	0	0	1	1421	1595.205	-0.472	0.486
## 3	0	0	0	0	1	0	1407	1406.395	-0.435	0.479
## 4	0	0	0	0	1	1	514	733.086	-0.067	0.435
## 5	0	0	0	1	0	0	478	592.888	-0.562	0.504
## 6	0	0	0	1	0	1	240	245.247	-0.169	0.443
## 7	0	0	0	1	1	0	216	228.727	-0.137	0.440
## 8	0	0	0	1	1	1	135	208.194	0.196	0.428
## 9	0	0	1	0	0	0	539	649.741	-0.445	0.481
## 10	0	0	1	0	0	1	204	332.363	-0.076	0.435
## 11	0	0	1	0	1	0	729	315.288	-0.045	0.433
## 12	0	0	1	0	1	1	390	343.448	0.285	0.431
## 13	0	0	1	1	0	0	78	104.066	-0.146	0.441
## 14	0	0	1	1	0	1	51	93.161	0.188	0.428
## 15	0	0	1	1	1	0	164	92.694	0.218	0.429
## 16	0	0	1	1	1	1	144	172.037	0.559	0.450
## 17	0	1	0	0	0	0	1325	1419.015	-0.671	0.529
## 18	0	1	0	0	0	1	496	484.756	-0.250	0.452
## 19	0	1	0	0	1	0	416	445.380	-0.218	0.448
## 20	0	1	0	0	1	1	224	347.805	0.121	0.428
## 21	0	1	0	1	0	0	262	163.498	-0.327	0.462
## 22	0	1	0	1	0	1	190	103.961	0.024	0.430
## 23	0	1	0	1	1	0	128	100.407	0.054	0.429
## 24	0	1	0	1	1	1	160	133.124	0.385	0.436
## 25	0	1	1	0	0	0	142	203.466	-0.226	0.449
## 26	0	1	1	0	0	1	98	156.215	0.113	0.428
## 27	0	1	1	0	1	0	312	153.332	0.143	0.428
## 28	0	1	1	0	1	1	180	243.723	0.478	0.442
## 29	0	1	1	1	0	0	43	45.249	0.046	0.429
## 30	0	1	1	1	0	1	57	59.007	0.376	0.435
## 31	0	1	1	1	1	0	152	60.781	0.407	0.437
## 32	0	1	1	1	1	1	193	168.497	0.777	0.477
## 33	1	0	0	0	0	0	1329	1496.169	-0.353	0.466
## 34	1	0	0	0	0	1	1389	905.690	0.001	0.431
## 35	1	0	0	0	1	0	650	871.137	0.031	0.430
## 36	1	0	0	0	1	1	1190	1103.485	0.362	0.434
## 37	1	0	0	1	0	0	206	274.704	-0.068	0.435
## 38	1	0	0	1	0	1	408	286.120	0.263	0.430

```

## 39  1  0  0  1  1  0 199 288.631 0.293 0.431
## 40  1  0  0  1  1  1 502 627.505 0.644 0.459
## 41  1  0  1  0  0  0 244 393.023 0.023 0.430
## 42  1  0  1  0  0  1 500 489.689 0.353 0.434
## 43  1  0  1  0  1  0 683 502.266 0.384 0.436
## 44  1  0  1  0  1  1 1800 1324.440 0.749 0.473
## 45  1  0  1  1  0  0  77 128.527 0.285 0.431
## 46  1  0  1  1  0  1 194 274.591 0.634 0.458
## 47  1  0  1  1  1  0 248 296.457 0.669 0.463
## 48  1  0  1  1  1  1 1224 1454.930 1.106 0.533
## 49  1  1  0  0  0  0 500 517.788 -0.146 0.441
## 50  1  1  0  0  0  1 663 463.626 0.188 0.428
## 51  1  1  0  0  1  0 337 461.311 0.218 0.429
## 52  1  1  0  0  1  1 716 856.360 0.560 0.450
## 53  1  1  0  1  0  0 202 130.238 0.121 0.428
## 54  1  1  0  1  0  1 410 197.776 0.455 0.441
## 55  1  1  0  1  1  0 234 206.667 0.486 0.443
## 56  1  1  0  1  1  1 742 679.121 0.873 0.492
## 57  1  1  1  0  0  0 139 206.199 0.210 0.428
## 58  1  1  1  0  0  1 271 376.292 0.551 0.449
## 59  1  1  1  0  1  0 381 400.139 0.584 0.453
## 60  1  1  1  0  1  1 1360 1625.403 0.995 0.513
## 61  1  1  1  1  0  0  78  91.141 0.478 0.442
## 62  1  1  1  1  0  1 238 293.882 0.862 0.490
## 63  1  1  1  1  1  0 380 330.550 0.901 0.496
## 64  1  1  1  1  1  1 2857 2759.328 1.431 0.601

```

#measures person ability for each pearson who responded a specific way
person.fit(model)

```

##
## Person-Fit Statistics and P-values
##
## Call:
## ltm(formula = dat_base ~ z1, IRT.param = TRUE)
##
## Alternative: Inconsistent response pattern under the estimated model
##

```

	Q1A	Q2A	Q3A	Q4A	Q5A	Q6A	L0	Lz	Pr(<Lz)
## 1	0	0	0	0	0	0	-0.6278	0.7994	0.788
## 2	0	0	0	0	0	1	-2.4153	0.4719	0.6815
## 3	0	0	0	0	1	0	-2.5445	0.4472	0.6726
## 4	0	0	0	0	1	1	-3.1919	0.6408	0.7392
## 5	0	0	0	1	0	0	-3.3937	-0.4043	0.343
## 6	0	0	0	1	0	1	-4.2938	-0.7237	0.2346
## 7	0	0	0	1	1	0	-4.3618	-0.7762	0.2188
## 8	0	0	0	1	1	1	-4.4135	-0.7438	0.2285
## 9	0	0	1	0	0	0	-3.3159	-0.1785	0.4292
## 10	0	0	1	0	0	1	-3.9836	-0.3027	0.381
## 11	0	0	1	0	1	0	-4.0336	-0.3324	0.3698

## 12	0	0	1	0	1	1	-3.8947	0.0272	0.5109
## 13	0	0	1	1	0	0	-5.1498	-1.6241	0.0522
## 14	0	0	1	1	0	1	-5.2192	-1.9399	0.0262
## 15	0	0	1	1	1	0	-5.2185	-1.9413	0.0261
## 16	0	0	1	1	1	1	-4.5118	-0.9074	0.1821
## 17	0	1	0	0	0	0	-2.4993	0.0646	0.5257
## 18	0	1	0	0	0	1	-3.6148	-0.1411	0.4439
## 19	0	1	0	0	1	0	-3.6989	-0.1744	0.4308
## 20	0	1	0	0	1	1	-3.9136	-0.0347	0.4862
## 21	0	1	0	1	0	0	-4.7015	-1.1988	0.1153
## 22	0	1	0	1	0	1	-5.1354	-1.7229	0.0425
## 23	0	1	0	1	1	0	-5.1662	-1.7881	0.0369
## 24	0	1	0	1	1	1	-4.8190	-1.3039	0.0961
## 25	0	1	1	0	0	0	-4.4825	-0.9480	0.1716
## 26	0	1	1	0	0	1	-4.7153	-1.1862	0.1178
## 27	0	1	1	0	1	0	-4.7290	-1.2099	0.1132
## 28	0	1	1	0	1	1	-4.1888	-0.4750	0.3174
## 29	0	1	1	1	0	0	-5.9644	-2.8564	0.0021
## 30	0	1	1	1	0	1	-5.6347	-2.4455	0.0072
## 31	0	1	1	1	1	0	-5.5972	-2.3556	0.0092
## 32	0	1	1	1	1	1	-4.4445	-0.9540	0.17
## 33	1	0	0	0	0	0	-2.4869	0.6690	0.7483
## 34	1	0	0	0	0	1	-2.9734	1.0677	0.8572
## 35	1	0	0	0	1	0	-3.0086	1.0923	0.8626
## 36	1	0	0	0	1	1	-2.7098	1.6659	0.9521
## 37	1	0	0	1	0	0	-4.1735	-0.5188	0.302
## 38	1	0	0	1	0	1	-4.0822	-0.2466	0.4026
## 39	1	0	0	1	1	0	-4.0669	-0.2279	0.4099
## 40	1	0	0	1	1	1	-3.1873	0.4953	0.6898
## 41	1	0	1	0	0	0	-3.8056	0.0258	0.5103
## 42	1	0	1	0	0	1	-3.5243	0.5242	0.6999
## 43	1	0	1	0	1	0	-3.4914	0.5382	0.7048
## 44	1	0	1	0	1	1	-2.3957	1.0960	0.8635
## 45	1	0	1	1	0	0	-4.8777	-1.4218	0.0775
## 46	1	0	1	1	0	1	-4.0174	-0.3965	0.3459
## 47	1	0	1	1	1	0	-3.9273	-0.3347	0.3689
## 48	1	0	1	1	1	1	-2.0851	0.6663	0.7474
## 49	1	1	0	0	0	0	-3.5453	0.0927	0.5369
## 50	1	1	0	0	0	1	-3.6145	0.4408	0.6703
## 51	1	1	0	0	1	0	-3.6137	0.4504	0.6738
## 52	1	1	0	0	1	1	-2.9067	0.9804	0.8366
## 53	1	1	0	1	0	0	-4.8960	-1.4469	0.074
## 54	1	1	0	1	0	1	-4.4045	-0.7430	0.2287
## 55	1	1	0	1	1	0	-4.3514	-0.6852	0.2466
## 56	1	1	0	1	1	1	-3.0008	0.2821	0.6111
## 57	1	1	1	0	0	0	-4.4206	-0.7529	0.2258
## 58	1	1	1	0	0	1	-3.7321	0.0208	0.5083
## 59	1	1	1	0	1	0	-3.6594	0.0632	0.5252
## 60	1	1	1	0	1	1	-2.0536	0.8908	0.8135
## 61	1	1	1	1	0	0	-5.1726	-1.7290	0.0419

```

## 62    1    1    1    1    0    1 -3.8444 -0.4702  0.3191
## 63    1    1    1    1    1    0 -3.7044 -0.3910  0.3479
## 64    1    1    1    1    1    1 -1.1452  0.8861  0.8122

item.fit(model)

##
## Item-Fit Statistics and P-values
##
## Call:
## ltm(formula = dat_base ~ z1, IRT.param = TRUE)
##
## Alternative: Items do not fit the model
## Ability Categories: 10
##
##           X^2 Pr(>X^2)
## Q1A 3789.449 <0.0001
## Q2A 9243.511 <0.0001
## Q3A 1704.668 <0.0001
## Q4A 3804.557 <0.0001
## Q5A 2943.172 <0.0001
## Q6A 1302.295 <0.0001

library(glue)

##
## Attaching package: 'glue'

## The following object is masked from 'package:dplyr':
##
##      collapse

# graphing both genders ICC
modelM<-ltm(datM~z1, IRT.param=TRUE)
modelF<-ltm(datF~z1, IRT.param=TRUE)

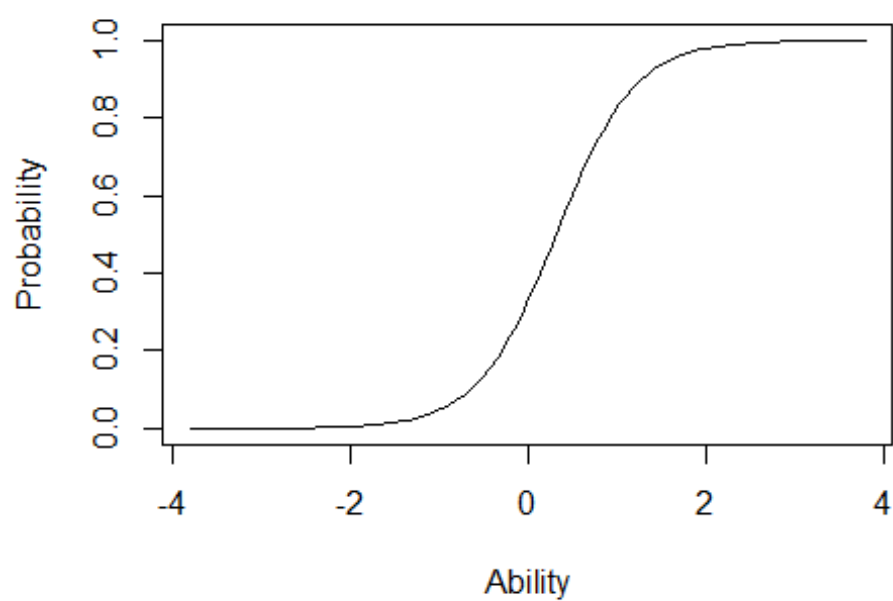
for (i in 1:6) {
  g<- ggplot()
  p1 = as.data.frame(plot(modelM,type="ICC",item = {i}))
  p2= as.data.frame(plot(modelF,type="ICC", item = {i}))

  # p1<- as.data.frame(pm)
  # p2 <- as.data.frame(pf)

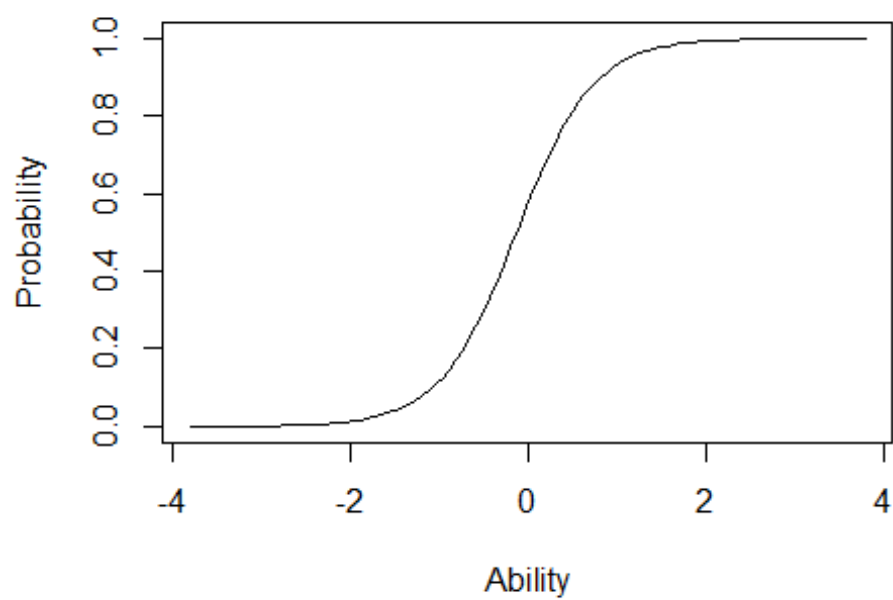
  g<- g+ geom_line(aes(x=p1$z, y=p1$V2, color = 'Male', size = .5)) +
  geom_line(aes(x=p2$z, y=p2$V2, color = 'Female', size=.5)) +
  labs(title=glue("Item ",{i}) ,
        x ="Probability", y = "Ability")
  print(g)
}

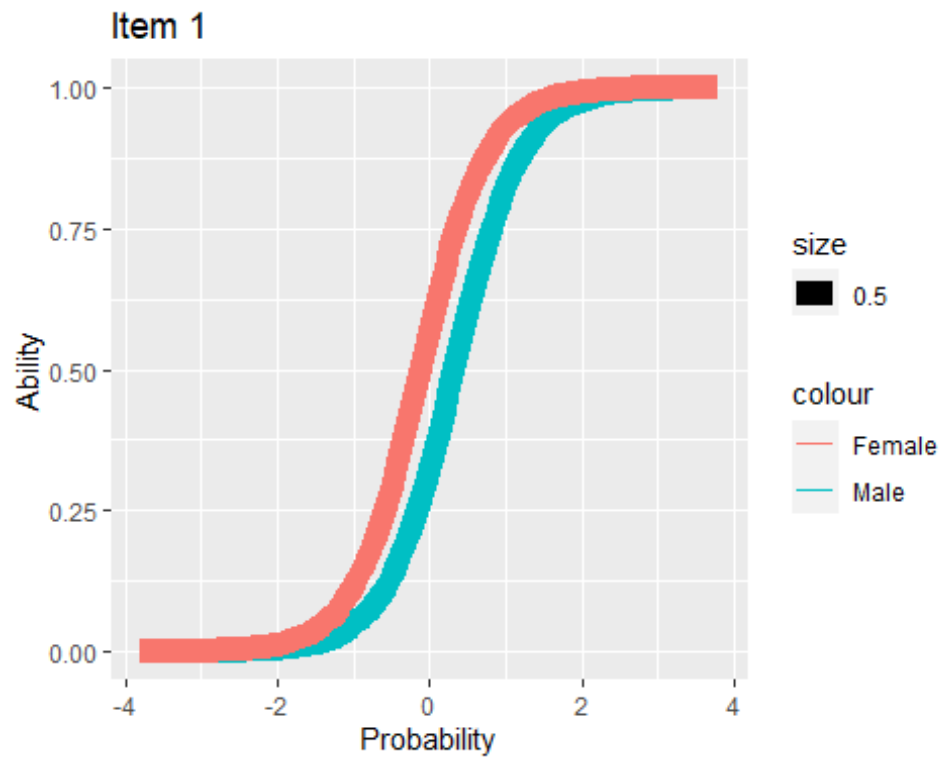
```

Item Characteristic Curves

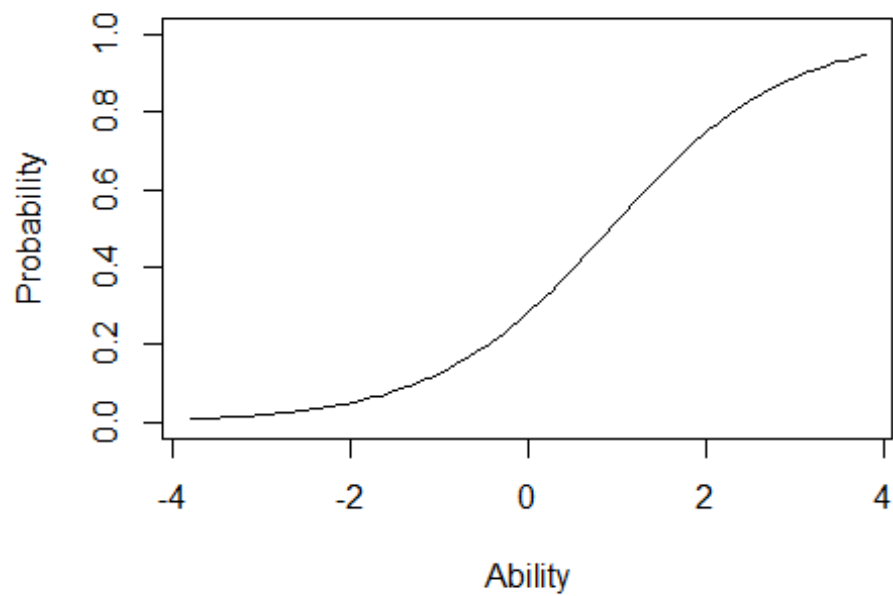


Item Characteristic Curves

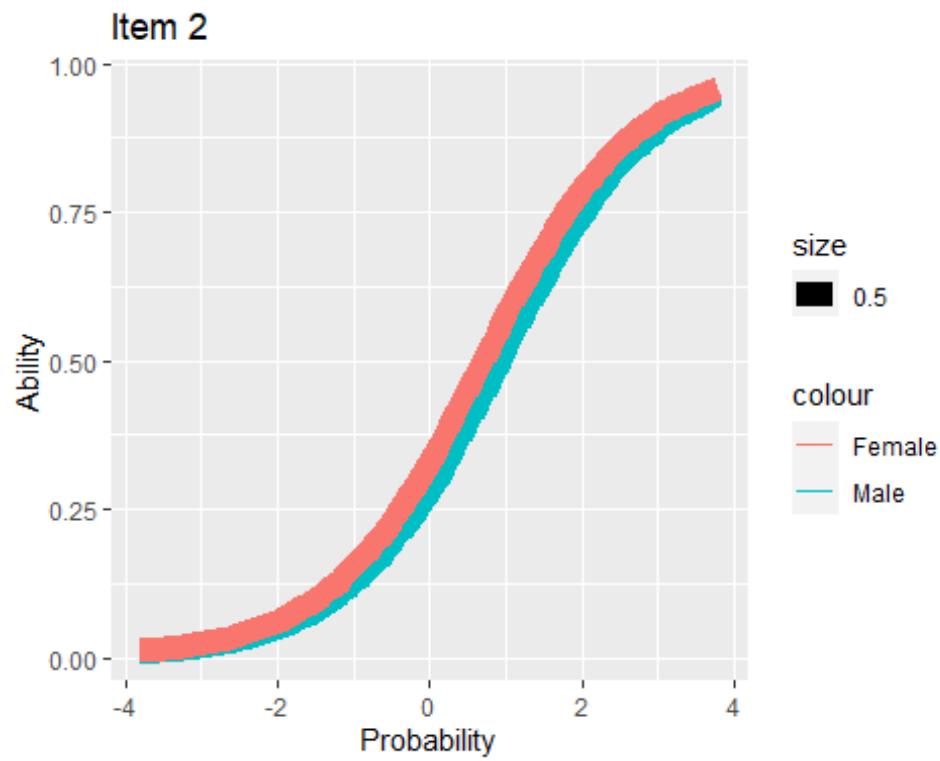
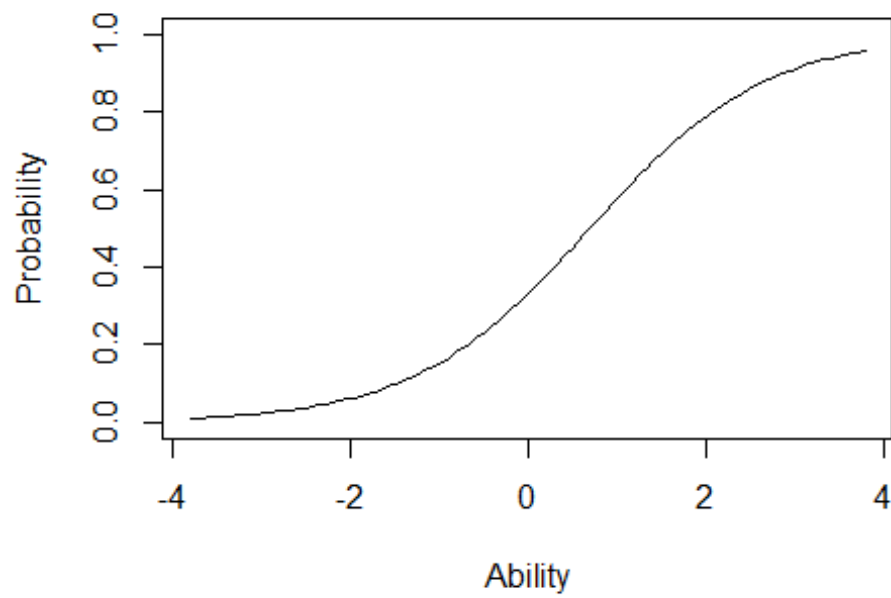




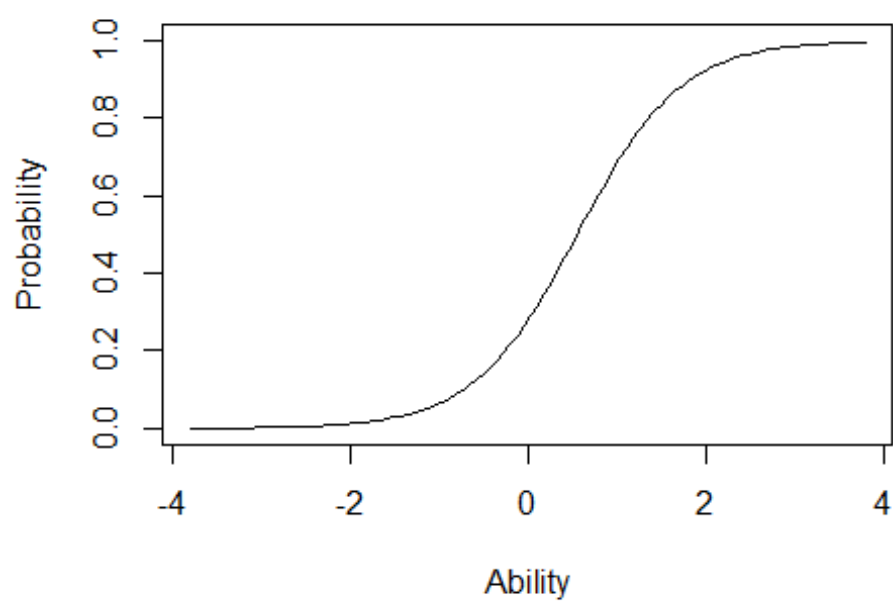
Item Characteristic Curves



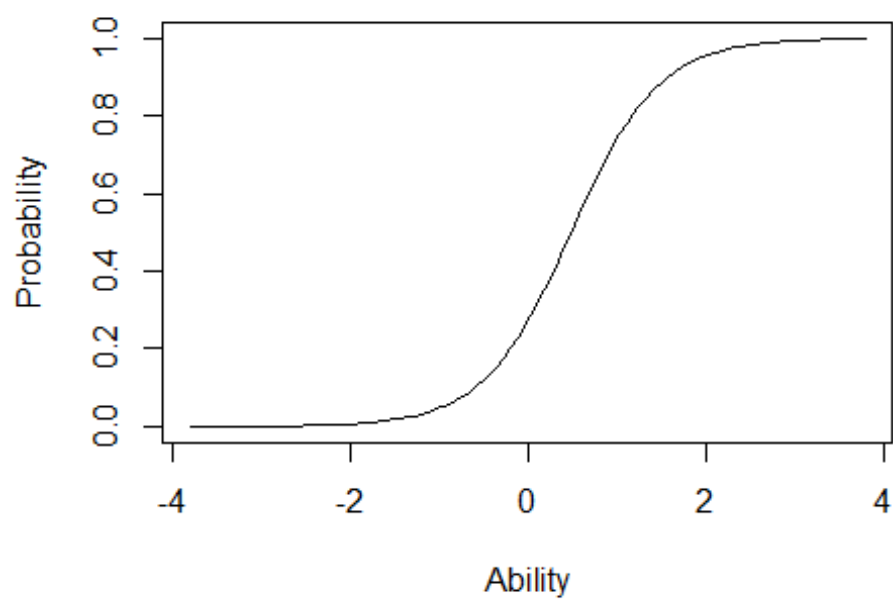
Item Characteristic Curves

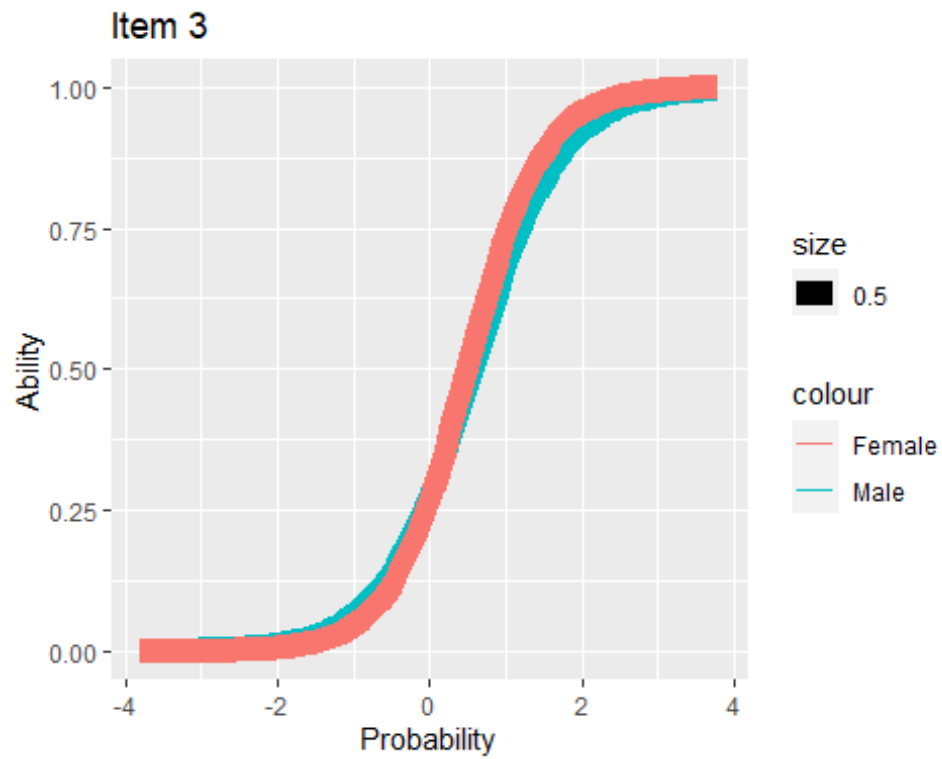


Item Characteristic Curves

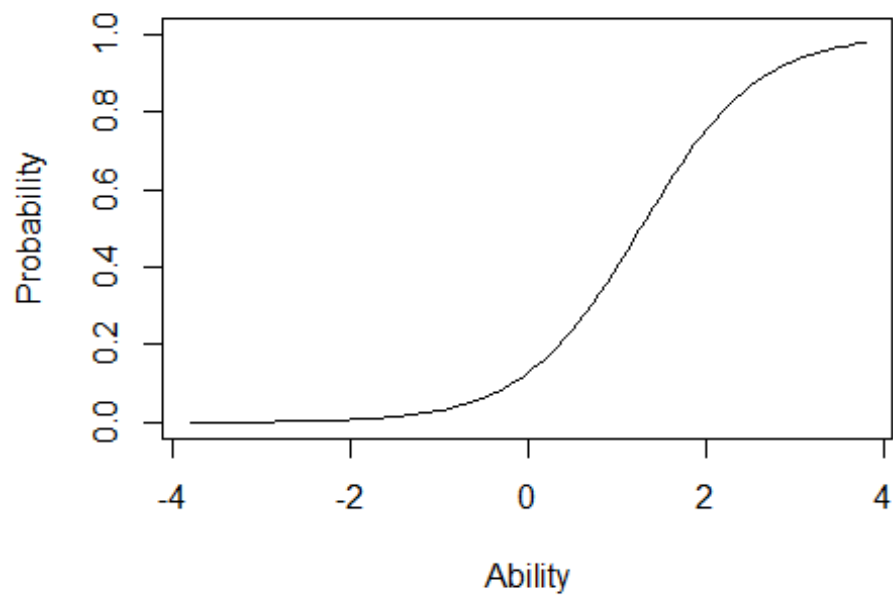


Item Characteristic Curves

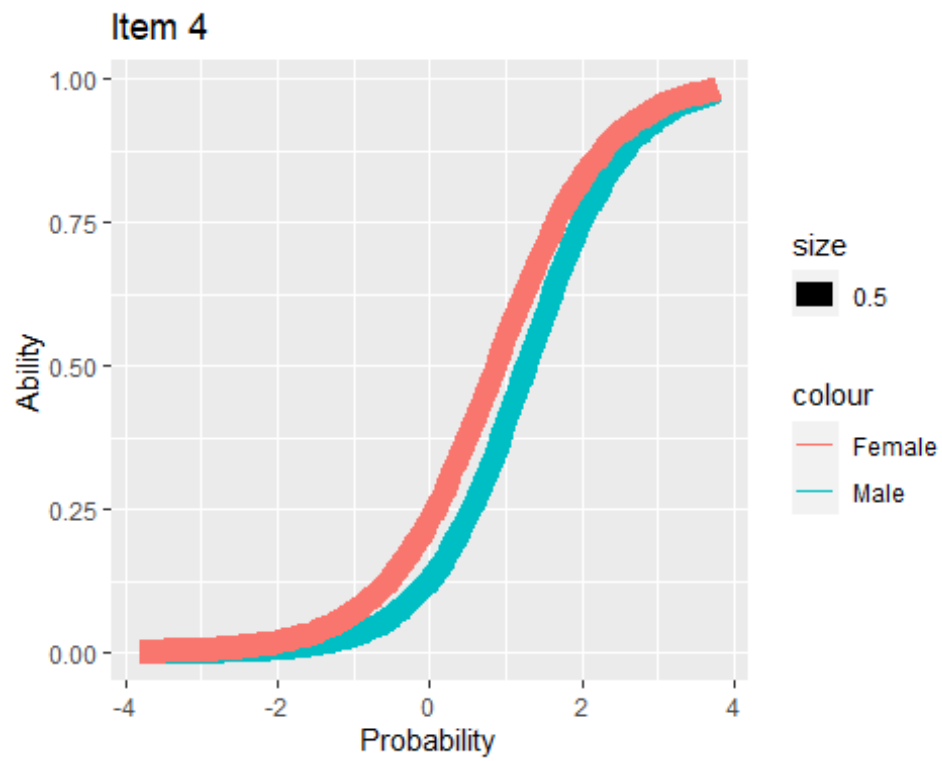
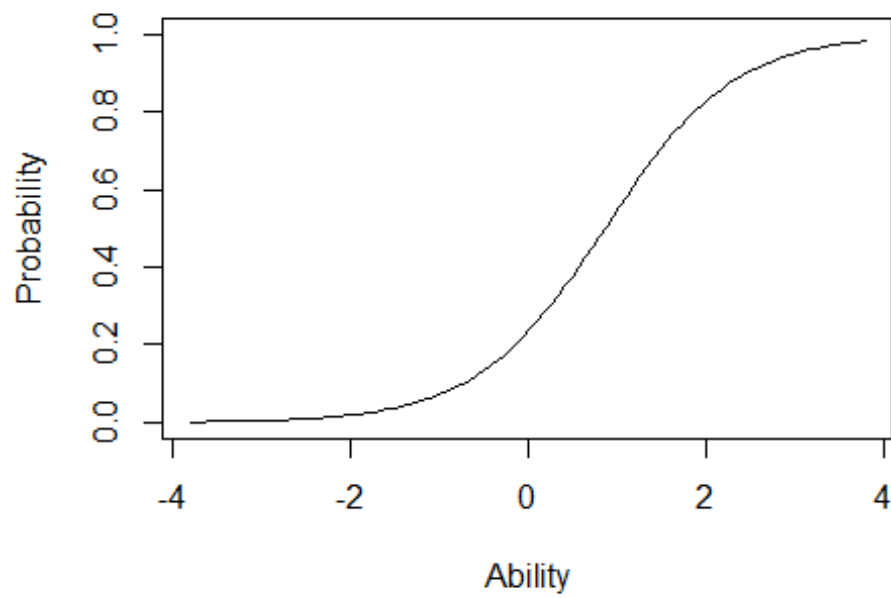




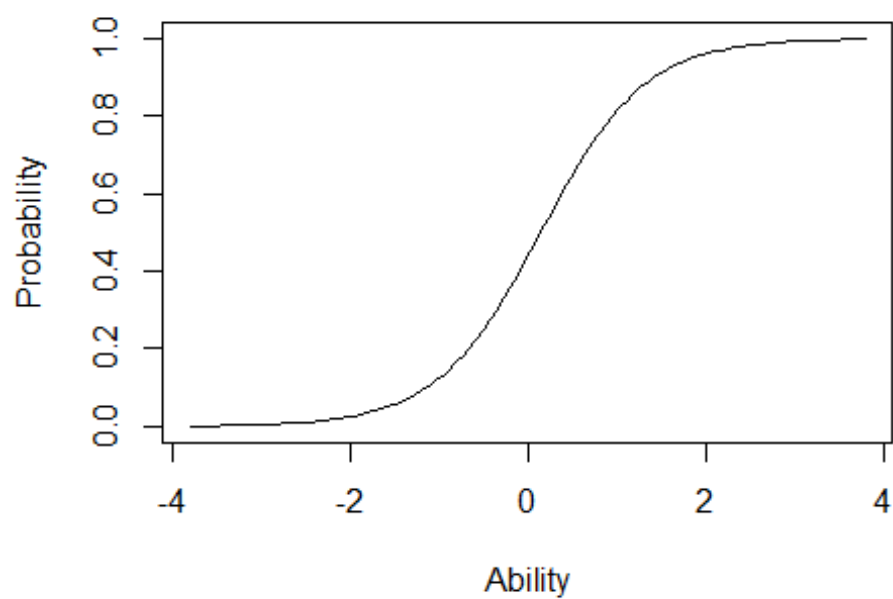
Item Characteristic Curves



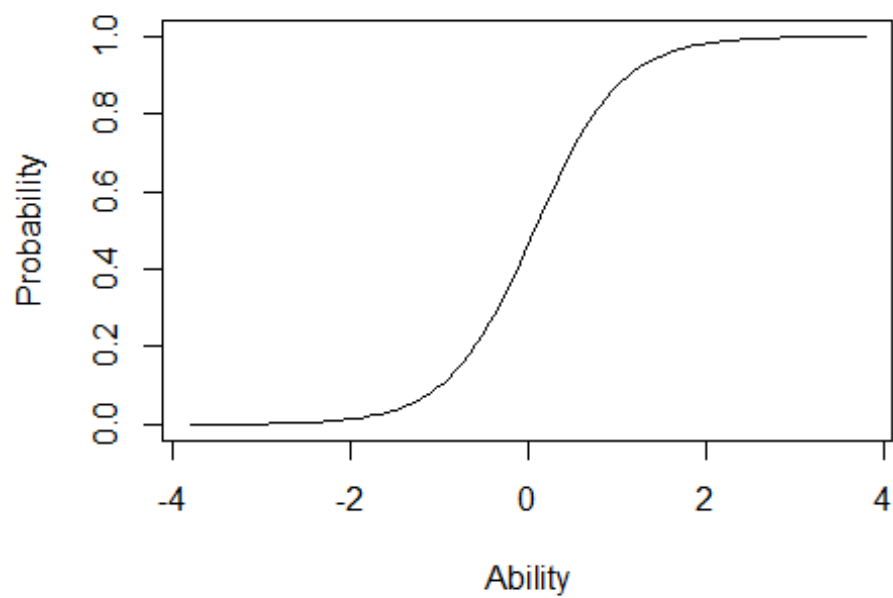
Item Characteristic Curves

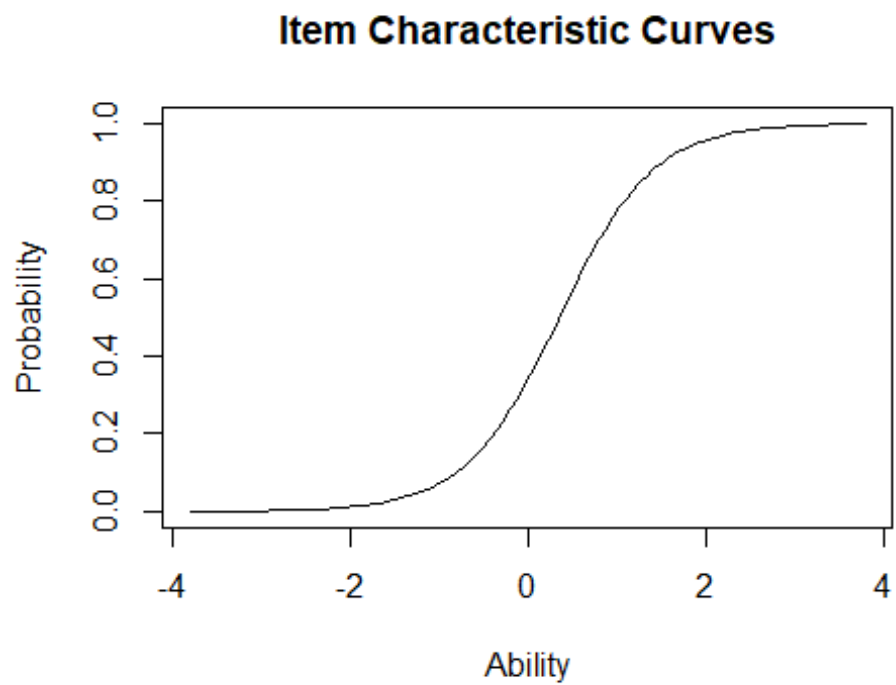
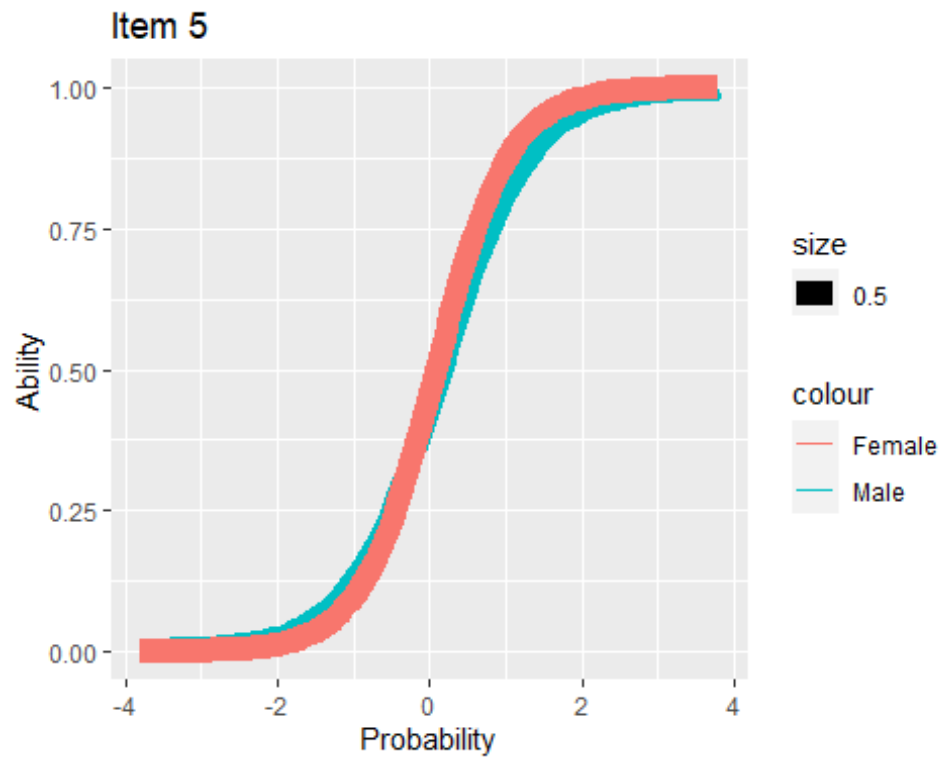


Item Characteristic Curves

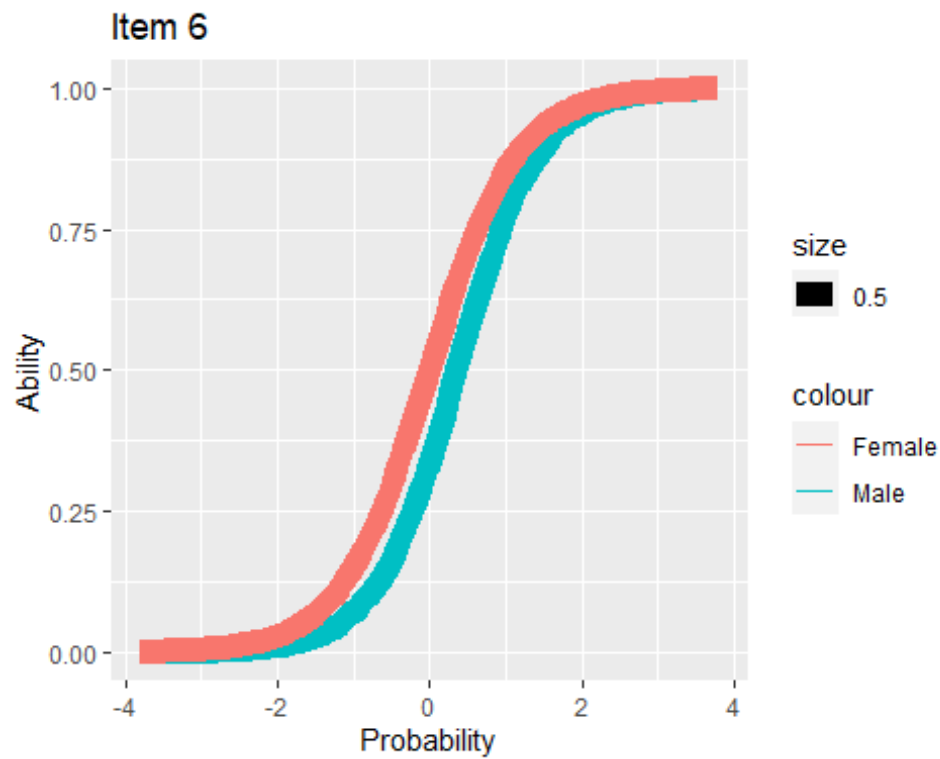
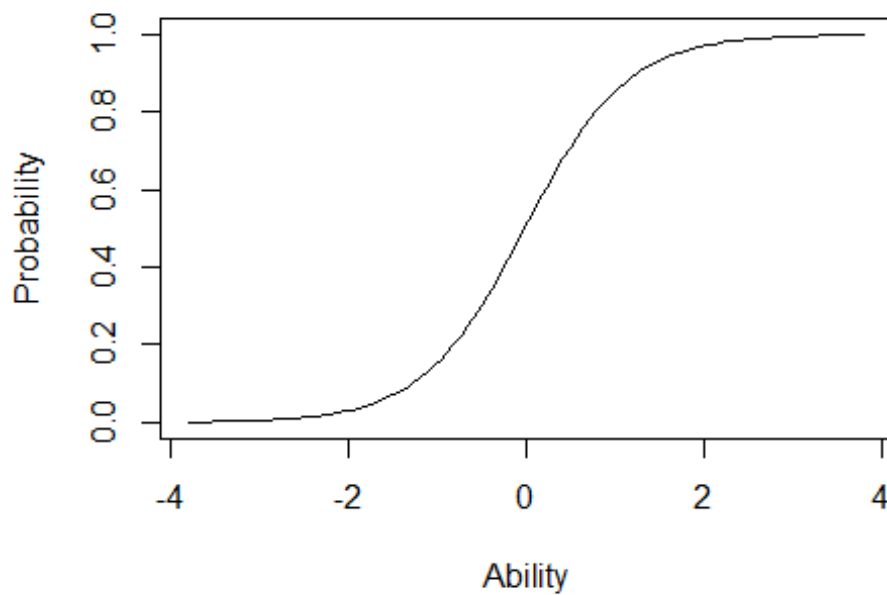


Item Characteristic Curves





Item Characteristic Curves



```
mod1p1<-rasch(dat_base)
mod2p1<-ltm(dat_base~z1, IRT.param=TRUE)
anova(mod1p1, mod2p1)
```

```
##  
## Likelihood Ratio Table  
##           AIC      BIC   log.Lik      LRT df p.value  
## mod1pl 279537.2 279597.4 -139761.6  
## mod2pl 277908.5 278011.5 -138942.2 1638.79  5  <0.001  
  
# anova(mod2pl, mod3pl)  
# anova(mod2pl, mod1pl)
```

Add a new chunk by clicking the *Insert Chunk* button on the toolbar or by pressing *Ctrl+Alt+I*.

When you save the notebook, an HTML file containing the code and output will be saved alongside it (click the *Preview* button or press *Ctrl+Shift+K* to preview the HTML file).

The preview shows you a rendered HTML copy of the contents of the editor. Consequently, unlike *Knit*, *Preview* does not run any R code chunks. Instead, the output of the chunk when it was last run in the editor is displayed.