

Master in *Statistical Sciences*
Lab 2 *Latent variable models*

Exercise 1
Attitude towards abortion

The data contain responses given by 410 individuals to four out of seven items concerning attitude to abortion. A small number of individual (379) did not answer to some of the questions and this data set contains only the complete cases.

1. Load and display the *dataframe* **Abortion** (library **ltm**).
2. Use the function **descript** to summarize the principal characteristics of the data set dataset. In particular evaluate:
 - which one among the 4 items shows the lowest degree of agreement among the sampled individuals;
 - what is the distribution of the *total score* in the sample;
 - if there are problems in some pairs of items ;

```
library(ltm)
data(Abortion)
?Abortion
dim(Abortion)
head(Abortion)
?descript
dsc <- descript(Abortion)
dsc$perc
dsc$items
dsc$pw.ass
```

2. Use the function **ltm** to fit **the 2 IRT model** to the data. Comment the results. In particular describe :
 - 2.1 Which item shows the highest positive attitude towards the abortion by the interviewed individuals
 - 2.2 Which item more discriminate among individuals in favour and against the abortion
 - 2.3 Which item less discriminate among individuals in favour and against the abortion
 - 2.4 The interpretation of the latent variable

```
?ltm
m1<-ltm(Abortion ~ z1)
```

```

m1.rip<-ltm(Abortion~ z1,IRT.param=FALSE)
summary(m1)
summary(m1.rip)

```

- 2.5 Verify the relation between the parameter estimates in the two parameterizations.

```

-summary(m1)$coefficients[1,1]*summary(m1)$coefficients[5,1]
summary(m1.rip)$coefficients[1,1]

```

- 4.3 Compute the standardized discriminant parameters and the probability of positive response of the median individual

```

alpha<-m1.rip$coeff[,2]
stalpha<-alpha/sqrt(1+alpha^2)
coef(m1.rip,prob=TRUE,order=TRUE)

```

6. Represent the characteristic curves of the four items for the estimated model. Comment the results.

```

plot(m1, legend = TRUE, cx = "bottomright", xlab="Attitude toward
abortion", lwd = 3, cex.main = 1.5, cex.lab = 1.3, cex = 1.1)

```

7. Based on the obtained results evaluate if the model has a good fit to the data.

```

E<-fitted(m1)[,5]
O<- m1$patterns$obs
cbind(m1$pattern$X,m1$pattern$obs,E)
Chisq<-sum((E-O)^2/E)
DOF<-14-2*4-1
pvalueC<-1-pchisq(Chisq,DOF)
LR<-2*sum(O*log(O/E))
pvalueLR<-1-pchisq(LR,DOF)

```

8. Use the function `margins` to compute the values of the two-way and three-way chi-square residuals and evaluate if the model has a local goodness of fit to the data.

```

?margins
margins(m1)
margins(m1,type="three-way",nprint=2)
margins(m1,type="three-way",nprint=3)

```

9. Use the function `factor.scores` to estimate the values of the latent variable for each individual.

```

?factor.scores
fs<-factor.scores(m1,method="EAP")
fs
Comp<-factor.scores(m1,method="Component")
Comp

```

```
resp.pattern<-fs$score.dat[,1:4]
total.score<-apply(resp.pattern,1,sum)
total.score
Cp<- Comp$score.dat[,7] tab<-cbind(fs$score.dat,Cp,total.score)
round(tab,3)
round(tab[order(total.score),],3)
```

10. Estimate the abilities of the *response pattern* that do not occur in the dataset.

```
fitted(m1,resp.pattern=rbind(c(1,0,0,1),c(1,0,1,0)))
```

Exercise 2

Law School Admission Test

The LSAT is a classical example in educational testing for measuring ability traits. This test was designed to measure a single latent ability scale. This LSAT example is a part of a data set given in Bock and Lieberman (1970).

1. Load and display the *dataframe* LSAT (library `ltm`).
2. Use the function `descript` to summarize the principal characteristics of the data set. In particular evaluate:
 - which one among the 5 items is the most difficult among the candidates;
 - what is the distribution of the *total score* in the sample;
 - if there are problems in some pairs of items `item` ;
 - which item has the highest/smallest proportion of correct responses given the same value of the *total score*.

```
dsc <- descript(LSAT)
dsc$perc
dsc$items
dsc$pw.ass
plot(dsc)
```

3. Use the function `rasch` to fit the **constrained Rasch model** (same discrimination parameter equal to 1 for all the `items`).

```
?rasch
m1<-rasch(LSAT, IRT.param=TRUE,constraint = cbind(ncol(LSAT) + 1,
1))
m1.rip<-rasch(LSAT, IRT.param=FALSE,constraint = cbind(ncol(LSAT)
+ 1, 1))
```

- 3.1 Display and comment the parameter estimates in the latent trait parameterization.

```
summary(m1.rip)
```

- 3.2 Compute and comment the estimates of the correct probability of the five items by the median individual

```
coef(m1.rip,prob=TRUE,order=TRUE)
```

- 3.3 Use the function `GoF.rasch` to estimate the *bootstrap p – value* of the X^2 statistic and establish if the model has an overall goodness of fit to the data.

```
pval.boot<-GoF.rasch(m1,B=199)
pval.boot$Tobs
pval.boot
```

- 3.4 Use the function `margins` to compute the values of the two-way and three-way chi-square residuals and evaluate if the model has a local goodness of fit to the data.

```
margins(m1)
margins(m1,type="three-way",nprint=2)
margins(m1,type="three-way",nprint=3)
```

4. Use the function `rasch` to fit **unconstrained Rasch model** ($a \neq 1$).

```
m2<-rasch(LSAT)
m2.rip<-rasch(LSAT,IRT.param=FALSE)
```

- 4.1 Compare the new estimates with the previous ones: what is the effect of removal the constrain on the parameters?

```
summary(m2.rip)
```

- 4.2 Verify the relation between the parameter estimates in the two parameterizations.

```
summary(m2)$coefficients[1,1]*summary(m2)$coefficients[6,1]
summary(m2.rip)$coefficients[1,1]
```

- 4.3 Establish if the discrimination parameter can be considered not equal to 1.

```
anova(m1,m2)
```

- 4.4 Verify if the model has problems of local of fit (two and three way margins).

```
margins(m2)
margins(m2,type="three-way",nprint=2)
```

- 4.5 Use the *AIC* e *BIC* criteria for the selection between the two Rasch models.

5. Use the function `ltm` to fit **the 2 IRT model** to the data.

```
m3<-ltm(LSAT ~z1)
m3.rip<-ltm(LSAT~z1,IRT.param=FALSE)
```

- 5.1 Compare the estimates of the discrimination parameters of the items.

```
summary(m3.rip)
```

- 5.2 Evaluate if there is a significant difference between this model and the previous one in terms of goodness of fit.

```
anova(m2,m3)
```

6. Represent the characteristic curves of the five items `item` for the selected model. Comment the results.

```
plot(m2, legend = TRUE, cx = "bottomright", lwd = 3, cex.main = 1.5,
cex.lab = 1.3, cex = 1.1)
```

7. Use the function `factor.scores` to estimate the candidates of the Law School according to the selected model.

```
fs<-factor.scores(m2,method="EAP")
fs
plot(fs$score.dat$z1)
resp.pattern<-fs$score.dat[,1:5]
total.score<-apply(resp.pattern,1,sum)
total.score
round(fs$score.dat[order(total.score),],3)
```

8. Estimate the abilities of the *response pattern* that do not occur in the dataset.

```
factor.scores(m2,resp.pattern=rbind(c(0,1,1,0,0),c(0,1,0,1,0)))
```