

Heatmap for patterns of association in log-linear models

Below I provide code and step-by-step explanations to produce a heatmap for log odds-ratios from log-linear models. I exemplify the implementation in R using data on intergenerational class mobility for England, France and Sweden.

Steps:

1. Load the following packages for data manipulation (tidyverse,modelr,reshape2), log-linear model estimation (vcdExtra,logmult) and plotting (cowplot). Install previously if you do not have them.

```
library("tidyverse")
library("modelr")
library("reshape2")
library("vcdExtra")
library("logmult")
library("cowplot")
```

2. Input the contingency table and turn it into a data frame. I use the dataset erikson from the package gnm, a dependency of package logmult. This is a cross-classification of subject's occupational status (destination) and his father's occupational status (origin) across 3 countries.

```
# Inpute data and create contingency table as data.frame()
data(erikson)
table <- ftable(erikson)
mydata <- as.data.frame(table)
levels(mydata$country) <- c("England-Wales", "France", "Sweden")

# Save Levels variables. To be used later.
levels.origin <- levels(mydata$origin)
levels.destination <- levels(mydata$destination)
levels.country <- levels(mydata$country)
```

This is what the data looks like:

```
## # A tibble: 243 x 4
##   origin destination country      Freq
##   <fct>   <fct>      <fct>    <dbl>
## 1 I      I          England-Wales 311
## 2 II     I          England-Wales 161
## 3 III    I          England-Wales 128
## 4 IVa    I          England-Wales 88
## 5 IVb    I          England-Wales 36
## 6 IVc    I          England-Wales 43
## 7 V/VI   I          England-Wales 356
## 8 VIIa   I          England-Wales 150
```

```
## 9 VIIb I England-Wales 12
## 10 I II England-Wales 130
## # ... with 233 more rows
```

3. Next, I set the values to be used as reference categories.

```
# Set reference categories
mydata$origin <- relevel(mydata$origin, ref = "V/VI")
mydata$destination <- relevel(mydata$destination, ref = "V/VI")
mydata$country <- relevel(mydata$country, ref = "England-Wales")
mydata$Freq <- mydata$Freq + 1 # add small constant to avoid problems with empty cells.
```

4. Fit different model specifications. Some of these models are log-linear and other are log-multiplicative.

```
# Fit models

# independence
indep <- glm(Freq ~ (origin + destination)*country, family = poisson, data = mydata)

# quasi-perfect mobility
qpm <- glm(Freq ~ (origin + destination)*country + Diag(origin, destination)*country, family = poisson, data = mydata)

# row-column association 1
rc1 <- glm(Freq ~ (origin + destination)*country + Mult(origin, destination) + Diag(origin, destination)*country, family = poisson, data = mydata)

## Initialising
## Running start-up iterations..
## Running main iterations.....
.....
## .....
## Done

# quasi-symmetry
qsymm <- glm(Freq ~ (origin + destination)*country + Symm(origin, destination)*country, family = poisson, data = mydata)

# unidiff or log-multiplicative layers
unidiff <- glm(Freq ~ (origin + destination)*country + Mult(Exp(country), origin:destination), family = poisson, data = mydata)

## Initialising
## Running start-up iterations..
```

```
## Running main iterations.....
## Done

# saturated
sat <- glm(Freq ~ origin*destination*country, family = poisson, data = mydata)

# Compare models via goodness of fit statistics

models <- glmlist(indep,qpm,rc1,qsym,unidiff,sat)
LRstats(models)

## Likelihood summary table:
##           AIC      BIC LR Chisq Df Pr(>Chisq)
## indep    6498.4  6676.5   5152.6 192 < 2.2e-16 ***
## qpm      3130.9  3403.4   1731.1 165 < 2.2e-16 ***
## rc1      1973.4  2298.3    543.7 150 < 2.2e-16 ***
## qsym     1689.1  2244.5    127.3  84  0.001605 **
## unidiff  1649.0  2057.7    171.2 126  0.004580 **
## sat      1729.8  2578.6      0.0   0  1.000000
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

5. Goodness of fit statistics suggest that the unidiff models is the one that better fits the data. At this point I compute prediction from this model and from them, log odd ratios.

```
# Create a synthetic dataset with all possible combinations of values

dummy.model <- lm(Freq ~ origin + destination + country, data=mydata)
new_x <- mydata %>% data_grid(origin,destination,country,.model=dummy.model)

# Compute predictions from different models. In this case: unidiff, quasi-symmetry and saturated model.

for ( m in c("unidiff","qsym","sat")) {

  chosen_model <- eval(parse(text = m ))

  # Predicted counts
  predictions <- cbind(mydata%>% data_grid(origin,destination,country,.model=dummy.model), pred = predict(chosen_model, newdata=new_x)) %>%
    as_tibble()

  # Intercept
  intercept <- predictions %>% filter(origin=="V/VI", destination=="V/VI", country=="England-Wales") %>% dplyr::summarise(pred) %>% as.numeric()
```

```

# Log odd ratios for marginal distributions
predictions <- predictions %>% mutate(pred = pred - intercept) # remove intercept

predictions_country <- predictions %>% filter(origin=="V/VI", destination=="V/VI") %>% rename(margin_country=pred) %>% select(country,margin_country)
predictions_origin <- predictions %>% filter(country=="England-Wales", destination=="V/VI") %>% rename(margin_origin=pred) %>% select(origin,margin_origin)
predictions_destination <- predictions %>% filter(country=="England-Wales", origin=="V/VI") %>% rename(margin_destination=pred) %>% select(destination,margin_destination)

# match
predictions <- predictions %>% left_join(predictions_country, by="country")
predictions <- predictions %>% left_join(predictions_origin, by="origin")
predictions <- predictions %>% left_join(predictions_destination, by="destination")

# Log odd ratios for marginal distributions origin and destination by country
predictions_country_origin <- predictions %>% filter(origin!="V/VI",country!="England-Wales",destination=="V/VI") %>%
  rename(margin_country_origin=pred) %>% mutate(margin_country_origin = margin_country_origin - (margin_country + margin_origin )) %>%
  select(country,origin,margin_country_origin)

predictions_country_destination <- predictions %>% filter(origin=="V/VI",country!="England-Wales",destination!="V/VI") %>%
  rename(margin_country_destination=pred) %>% mutate(margin_country_destination = margin_country_destination - (margin_country + margin_destination )) %>%
  select(country,destination,margin_country_destination)

predictions <- predictions %>% left_join(predictions_country_origin, by=c("country","origin")) %>% replace_na(list(margin_country_origin = 0))
predictions <- predictions %>% left_join(predictions_country_destination, by=c("country","destination")) %>% replace_na(list(margin_country_destination = 0))

# Margin-free Log-odd ratios (LORs)
predictions <- predictions %>%

```

```

    mutate(`log-OR` = pred - (margin_country + margin_origin + margin_destination + margin_country_origin + margin_country_destination) )

# Save predictions
assign(paste0("predictions_",m),predictions)

}

```

6. Finally, for each model I visualize the estimated log odd ratios capturing margin-free association between origin and destination across countries . Of course, other quantities can also be visualized in the same way.

```

# Combine models

predictions_unidiff <- predictions_unidiff %>% mutate(model = "Unidiff")
predictions_qsymm <- predictions_qsymm %>% mutate(model = "Quasi-symmetry")
predictions_sat <- predictions_sat %>% mutate(model = "Saturated")

predictions <- bind_rows(predictions_unidiff,predictions_qsymm,predictions_sat) %>%
  mutate(model = factor(model, levels=c("Unidiff","Quasi-symmetry","Saturated")))

# Plot

plot <- predictions %>%
  ggplot(aes(y=factor(origin, levels = rev(levels.origin)),
             x=factor(destination, levels = levels.destination))) + facet_grid(
    model ~ country) + geom_raster(aes(fill= `log-OR`)) +
  scale_fill_gradientn(limits=c(-6,6), colours=c("red","white","blue")) +
  labs(y="Father's occupation", x= "Children's occupation", colour="") +
  theme_bw() + theme(axis.text.x = element_text(size=9, angle=45, vjust=-1, hjust=0),
                    axis.text.y = element_text(size=9, angle=0),
                    plot.title= element_text(size=11)) +
  scale_x_discrete(position="top")

# Add Labels
plot <- plot %>% add_sub(., "I+II: Service class, III: Routine non-manual employees, IVa+b: Petty bourgeoisie, IVc: Farmers, V/VI: Skilled working class, VI Ia: Semi and unskilled working class, VIIb: Agricultural workers", size= 9) %>%
  ggdraw()

print(plot)

```

