# Module recap and Q&A PSM 2

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Probability, Statistics & Modeling II

# Today

Module recap and Q&A

- Model comparison
- The loglinear model
- Interpretation

Model comparison

#### When do we need it?

- you can model an outcome variable in many ways
  - income ~ age + gender
  - income ~ age + gender + education
  - $income \sim ethnicity + family status$
- Which model explains the data (i.e. the outcome) better?

How to do that comparison?

One model is nested in another if you can always obtain the first model by constraining some of the parameters of the second model.

Nice explanation in this SO answer

Model 1:  $Y \sim x1 + x2 + x1 : x2 + x3$ 

Model 2:  $Y \sim x1 + x2$ 

Can we constrain the parameters of Model 1 to obtain Model 2?

#### Model parameters

Model 1: 
$$Y \sim \beta_1 x 1 + \beta_2 x 2 + \beta_3 (x 1 : x 2) + \beta_4 x 3$$
  
Model 2:  $Y \sim \beta_1 x 1 + \beta_2 x 2$ 

Can we constrain the parameters of Model 1 to obtain Model 2?

-> Yes: set  $\beta_3 = \beta_4 = 0$  so that Model1 = Model2

Model 1:  $Y \sim x1 + x2$ 

Model 2:  $Y \sim x1$ 

Model 1:  $Y \sim x1 + x2$ 

Model 2:  $Y \sim x1 + x3$ 

Model 1:  $Y \sim x1 + x2 + x3 + x4$ 

Model 2:  $Y \sim x5$ 

#### Nested models

 $income \sim age + gender$ 

 $income \sim age + gender + education$ 

#### Nested models

M1:  $income \sim age + gender$ 

 $M2:income \sim age + gender + education$ 

M3:  $income \sim ethnicity + family status$ 

#### Nested models

 $income \sim age + gender$ 

 $income \sim age + gender + education$ 

In essence: do we really need the additional predictor education?

Nested structure allows for formal statistical tests!

# Formal model comparison logic

- if nested, we can test whether a simpler model is significantly worse than a more complex model
- if the model comparison is sign., then choose the more complex model
- if the test is not sign., choose the simpler one (Ockham's razor principle)

#### Non-nested models

 $income \sim age + gender + education$  vs.

 $income \sim ethnicity + familystatus$ No formal test possible?

#### Non-nested models

- for non-nested models, compare goodness of fit indices
- e.g. sum of squared residuals, mean absolute error, ...
- other fit indices: AIC, Log-likelihood, BIC

In essence: you have to make a judgment without formal statistical test.

The loglinear model

# A step back:

	No anti-virus software	Anti-virus software	Sum
Hacked	300	250	550
Not hacked	200	250	450
Sum	500	500	1000

# For r-by-c tables

Idea of the Chi-square test:

- Observed values O
- ullet Expected values (if there were no association) E
- rows: *i*
- columns: *j*

$$E_{i,j} = \frac{(total_i * total_j)}{total}$$

# Chi-square test for r-by-c

$$\chi^2 = \sum \frac{(O - E^2)}{E}$$

- Null-hypothesis: there is no association between the two factors
- Alt. hypothesis: there is a significant association

Thus: if sign. -> there is a sign. association between r and c

#### More dimensions?

# The Log-Linear Model

- GLM with link function for count data
- count data aptly modelled as a Poisson distrubuted variable

# Stepwise

- 1. we build the "independence" model
  - no relationships between variables
- 2. we assess the  $H_0$  of model adequacy
  - if significant: model not adequate for the data
  - if non-sign.: model is considered adequate
- 3. we build more complex models
  - e.g. with dependencies (i.e. interactions) between variables

#### Stepwise

- remember, we're modelling counts that come about due to a combination of factors
- thus: the saturated (= full) model will explain the data perfectly

```
## UK yes no
## gender
## male 40 80
## female 70 30
```

```
(exampledata = as.data.frame(as.table(example)))
```

```
## gender UK Freq
## 1 male yes 40
## 2 female yes 70
## 3 male no 80
## 4 female no 30
```

#### Next:

- look at  $H_0$  of model adequacy
- look at predicted values

#### Model adequacy hyp.

```
summary(indep model)
```

```
##
## Call:
## glm(formula = Freq ~ gender + UK, family = poisson, data = exampledata
##
## Deviance Residuals:
##
## -2.750 2.666 2.455 -3.058
## Coefficients:
##
                Estimate Std. Error z value Pr(>|z|)
## (Intercept) 4.094e+00 1.135e-01 36.078 <2e-16 ***
## genderfemale -1.823e-01 1.354e-01 -1.347 0.178
## UKno
           7.856e-12 1.348e-01 0.000 1.000
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 '' 1
##
## (Dispersion parameter for poisson family taken to be 1)
```

# Model adequacy hyp.

```
pchisq(30.048, 1, lower.tail = F)
```

## [1] 4.21483e-08

#### Predicted values

gender	UK	Freq	round(fitted(indep_model), 2)
male	yes	40	60
female	yes	70	50
male	no	80	60
female	no	30	50

Next step: full model

gender	UK	Freq	round(fitted(full_model), 2)
male	yes	40	40
female	yes	70	70
male	no	80	80
female	no	30	30

#### Conclusion

```
## UK yes no
## gender
## male 40 80
## female 70 30
```

There is an association between gender and "UK".

# Making sense of the coefficients

```
coefficients(full model)
##
          (Intercept)
                            genderfemale
                                                        UKno genderfemale: UKn
##
                               0.5596158
            3.6888795
                                                  0.6931472
                                                                     -1.54044
exp(coefficients(full model))
##
                            genderfemale
                                                        UKno genderfemale: UKn
          (Intercept)
           40.000000
                               1.7500000
                                                  2.0000000
                                                                      0.21428
```

Remeber: we're modelling the log (hence log-linear model)

- UK\_no: 2.00
  - The odds of a person being from the UK are 1:2.00, regardless of their gender.
- gender\_female: 1.75
  - The odds of a person being female are 1.75:1, regardless of their UK status.
- gender\_female:UK\_no: 0.21
  - People that are female have estimated odds of not being from the UK is 0.21 times the odds for males of not being from the UK.

# Log-linear model

- extends Chisquare idea to mutliple dimensions
- brings in the modelling aspect
- aim: find a model that is simpler than the full model
- core: simplest model to explain the data

Interpretation

#### Interpretation of results

#### General strategy:

- there's always a hypothesis
- make the hypothesis explicit
- every RQ must come down to one or multiple hypotheses

# Hypotheses

- difference in means 2 groups (t-test, rank sum test)
- difference in means 2+ groups (ANOVA, Kruskal-Wallis test)
- predictor combinations to explain an outcome (model comparison tests)
  - linear models
  - logistic regression models
  - log linear models

# Interpretation strategy

When you ran your test/model:

- ask yourself: what did I test?
- which hypothesis was behind the test?
- what does the hyp. testing result reveal?
- how does this feed back to my RQ?

#### **Pitfalls**

- forgetting to re-transform coefficients in logistic regression or loglinear models
- forgetting the unit of interpretation of coefficients
- forgetting the direction of effects
- attributing causality to correlational data

Your interpretation becomes very difficult if you do not know the question you want to answer.

#### Easiest trick

Always start with the question!

Open Q&A session

# Next week CLASS TEST

- Tuesday, 19 March 2019
- 10am-12pm
- 60 min
- 10 questions (5 MC, 5 open)

# **END**