# **Exercises Day 4**

### **PSY8003**

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# **Exercise 1: Logistic regression**

- as expected, survival probability drops for cheaper tickets and lower Ticket class
- when including Pclass the Fare effect is no longer significant (within each ticket class the tickets seem to have not varied much in price) but including both predictors is still better in terms of model fit
- Pclass and Fare add to the overall model-fit
- looking at the ORs, the drop is dramatic (94% reduced survival probability for 3rd vs. 1st class!); probably the cheap cabins were located deep within the ship?

```
library(tidyverse)
```

```
-- Attaching packages ----- tidyverse 1.3.1 --
v ggplot2 3.3.5
                v purrr
                        0.3.4
v tibble 3.1.6
               v dplyr
                        1.0.7
       1.1.4
                v stringr 1.4.0
v tidyr
v readr
       2.1.1
                v forcats 0.5.1
-- Conflicts ----- tidyverse_conflicts() --
x dplyr::filter() masks stats::filter()
x dplyr::lag()
            masks stats::lag()
  library(lmtest)
```

```
Loading required package: zoo
Attaching package: 'zoo'
The following objects are masked from 'package:base':
   as.Date, as.Date.numeric
  titanic <- haven::read_dta("../data/titanic.dta") |> na.omit()
  mod1 <- glm(Survived ~ Fare, data=titanic, family=binomial)</pre>
  summary(mod1)
Call:
glm(formula = Survived ~ Fare, family = binomial, data = titanic)
Deviance Residuals:
               Median
   Min
            1Q
                             3Q
                                    Max
-2.5870 -0.9073 -0.8714 1.3331 1.5741
Coefficients:
           Estimate Std. Error z value Pr(>|z|)
Fare
           Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
   Null deviance: 964.52 on 713 degrees of freedom
Residual deviance: 901.25 on 712 degrees of freedom
AIC: 905.25
Number of Fisher Scoring iterations: 5
  mod2 <- glm(Survived ~ factor(Pclass), data=titanic, family=binomial)</pre>
  summary(mod2)
```

```
Call:
glm(formula = Survived ~ factor(Pclass), family = binomial, data = titanic)
Deviance Residuals:
   Min
             1Q
                Median
                              3Q
                                     Max
-1.4607 -0.7399 -0.7399 0.9184
                                   1.6908
Coefficients:
              Estimate Std. Error z value Pr(>|z|)
             (Intercept)
factor(Pclass)2 -0.7261
                          0.2168 -3.350 0.000808 ***
factor(Pclass)3 -1.8009
                          0.1982 -9.086 < 2e-16 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
   Null deviance: 964.52 on 713 degrees of freedom
Residual deviance: 869.81 on 711 degrees of freedom
AIC: 875.81
Number of Fisher Scoring iterations: 4
  mod3 <- glm(Survived ~ factor(Pclass) + Fare, data=titanic, family=binomial)</pre>
  summary(mod3)
Call:
glm(formula = Survived ~ factor(Pclass) + Fare, family = binomial,
   data = titanic)
Deviance Residuals:
             1Q Median
                              3Q
                                     Max
-2.0170 -0.7789 -0.7275 1.0612
                                   1.7320
Coefficients:
               Estimate Std. Error z value Pr(>|z|)
(Intercept)
              0.098422 0.252624 0.390 0.6968
factor(Pclass)2 -0.326010 0.259653 -1.256
                                            0.2093
factor(Pclass)3 -1.345719  0.257121 -5.234 1.66e-07 ***
```

```
Fare
                ___
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
   Null deviance: 964.52 on 713 degrees of freedom
Residual deviance: 861.50 on 710 degrees of freedom
AIC: 869.5
Number of Fisher Scoring iterations: 4
  lrtest(mod1,mod2,mod3)
Likelihood ratio test
Model 1: Survived ~ Fare
Model 2: Survived ~ factor(Pclass)
Model 3: Survived ~ factor(Pclass) + Fare
 #Df LogLik Df Chisq Pr(>Chisq)
  2 -450.63
  3 -434.91 1 31.4428 2.054e-08 ***
  4 -430.75 1 8.3077 0.003948 **
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
  mod4 <- glm(Survived ~ Age*Sex + factor(Pclass) + Fare, data=titanic, family=binomial)</pre>
  summary(mod4)
Call:
glm(formula = Survived ~ Age * Sex + factor(Pclass) + Fare, family = binomial,
   data = titanic)
Deviance Residuals:
             1Q
                 Median
                              ЗQ
                                      Max
-2.4957 -0.6416 -0.3714 0.6598
                                   2.5934
```

Coefficients:

```
Estimate Std. Error z value Pr(>|z|)
(Intercept)
                3.1432058   0.4960243   6.337   2.35e-10 ***
               -0.0072077 0.0116843 -0.617 0.537320
Age
Sexmale
               -1.1585842 0.4385724 -2.642 0.008249 **
factor(Pclass)2 -1.5322436  0.3407047  -4.497  6.88e-06 ***
factor(Pclass)3 -2.7279214  0.3462059  -7.879  3.29e-15 ***
               -0.0003952 0.0023346 -0.169 0.865561
Age:Sexmale
               Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
    Null deviance: 964.52 on 713 degrees of freedom
Residual deviance: 635.36 on 707 degrees of freedom
AIC: 649.36
Number of Fisher Scoring iterations: 5
  exp(coef(mod4))
    (Intercept)
                                      Sexmale factor(Pclass)2 factor(Pclass)3
                          Age
     23.1780517
                     0.9928182
                                    0.3139303
                                                   0.2160504
                                                                   0.0653550
                   Age:Sexmale
          Fare
     0.9996048
                     0.9510885
  lrtest(mod3,mod4)
Likelihood ratio test
Model 1: Survived ~ factor(Pclass) + Fare
Model 2: Survived ~ Age * Sex + factor(Pclass) + Fare
  #Df LogLik Df Chisq Pr(>Chisq)
   4 -430.75
   7 -317.68 3 226.15 < 2.2e-16 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

# **Exercise 2: Logistic regression for classification**

- all variables except make are significant
- ORs are the multiplicative effect on the probability that an email is spam
- the dollar and noon variables have huge ORs
- the confusion matrix show perfect categorization (all emails are correctly categorized as spam or not)
- this is due to overfitting on the training dataset; a better way to test this is cross-validation (hold-out datasets)

```
spam <- haven::read_dta("../data/spam.dta")</pre>
  mod <- glm(isspam ~ crltot + dollar + bang + money + n000 + make, data=spam, family=binomi
Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
  summary(mod)
Call:
glm(formula = isspam ~ crltot + dollar + bang + money + n000 +
    make, family = binomial(), data = spam)
Deviance Residuals:
    Min
              1Q
                 Median
                               3Q
                                       Max
-8.4904 -0.6153 -0.5816 0.4439
                                    1.9323
Coefficients:
             Estimate Std. Error z value Pr(>|z|)
(Intercept) -1.700e+00 5.361e-02 -31.717 < 2e-16 ***
            6.917e-04 9.745e-05 7.098 1.27e-12 ***
crltot
            8.013e+00 6.175e-01 12.976 < 2e-16 ***
dollar
            1.572e+00 1.115e-01 14.096 < 2e-16 ***
bang
            2.142e+00 2.418e-01 8.859 < 2e-16 ***
money
            4.149e+00 4.371e-01 9.492 < 2e-16 ***
n000
make
            1.698e-02 1.434e-01 0.118
                                            0.906
___
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
(Dispersion parameter for binomial family taken to be 1)
```

```
Null deviance: 6170.2 on 4600 degrees of freedom Residual deviance: 4058.8 on 4594 degrees of freedom
```

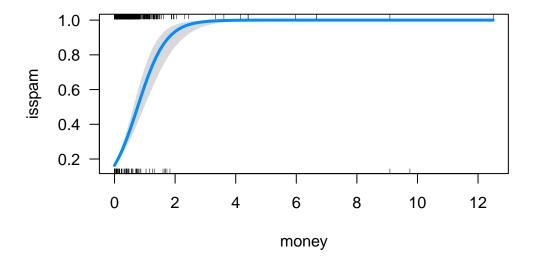
AIC: 4072.8

Number of Fisher Scoring iterations: 16

```
exp(coef(mod))
```

```
(Intercept) crltot dollar bang money n000 0.1826354 1.0006919 3018.4867001 4.8156296 8.5141268 63.3512064 make 1.0171234
```

```
library(visreg)
library(modelr)
visreg(mod, scale = "response", xvar = "money" )
```



Confusion Matrix and Statistics

confusionMatrix(predicted, observed)

Reference
Prediction no spam spam
no spam 3273 0
spam 0 1328

Accuracy : 1

95% CI: (0.9992, 1)

No Information Rate : 0.7114 P-Value [Acc > NIR] : < 2.2e-16

Kappa : 1

Mcnemar's Test P-Value : NA

Sensitivity: 1.0000 Specificity: 1.0000 Pos Pred Value: 1.0000 Neg Pred Value: 1.0000 Prevalence: 0.7114 Detection Rate: 0.7114 Detection Prevalence: 0.7114
Balanced Accuracy: 1.0000

'Positive' Class : no spam

## **Exercise 3: Poisson regression**

```
affairs <- haven::read_dta("../data/affairs.dta")</pre>
  affairs |>
   mutate(female=as.integer(gender==2)) -> affairs
  # a few example models...
  mod <- glm(affairs ~ female*age, data=affairs, family = poisson)</pre>
  summary(mod)
Call:
glm(formula = affairs ~ female * age, family = poisson, data = affairs)
Deviance Residuals:
  Min
          1Q Median
                        3Q
                               Max
-2.346 -1.758 -1.580 -1.394
                             6.186
Coefficients:
           Estimate Std. Error z value Pr(>|z|)
female
          0.281204 0.245582 1.145 0.25219
           age
female:age -0.008972 0.006845 -1.311 0.18993
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for poisson family taken to be 1)
   Null deviance: 2925.5 on 600 degrees of freedom
Residual deviance: 2885.2 on 597 degrees of freedom
AIC: 3387.2
```

```
Number of Fisher Scoring iterations: 6
```

```
mod <- glm(affairs ~ female*religiousness, data=affairs, family = poisson)</pre>
  summary(mod)
Call:
glm(formula = affairs ~ female * religiousness, family = poisson,
    data = affairs)
Deviance Residuals:
           1Q Median
                         3Q
                                  Max
-2.254 -1.936 -1.506 -1.227
                                6.630
Coefficients:
                    Estimate Std. Error z value Pr(>|z|)
(Intercept)
                    1.23588 0.12612 9.799 < 2e-16 ***
female
                    -0.06668
                                0.17446 -0.382
                                                   0.702
religiousness
                    -0.30395
                                0.04291 -7.083 1.41e-12 ***
                                0.05881
                                        0.731
                                                   0.465
female:religiousness 0.04299
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for poisson family taken to be 1)
    Null deviance: 2925.5 on 600 degrees of freedom
Residual deviance: 2830.8 on 597 degrees of freedom
AIC: 3332.8
Number of Fisher Scoring iterations: 7
  mod <- glm(affairs ~ female*yearsmarried , data=affairs, family = poisson)</pre>
  summary(mod)
Call:
glm(formula = affairs ~ female * yearsmarried, family = poisson,
   data = affairs)
```

```
Deviance Residuals:
```

Min 1Q Median 3Q Max -2.130 -1.764 -1.360 -1.162 6.729

#### Coefficients:

Estimate Std. Error z value Pr(>|z|)

(Intercept) -0.403473 0.105073 -3.840 0.000123 \*\*\*
female 0.091300 0.151603 0.602 0.547020
yearsmarried 0.081176 0.008938 9.082 < 2e-16 \*\*\*
female:yearsmarried -0.005737 0.012880 -0.445 0.656035

\_\_\_

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for poisson family taken to be 1)

Null deviance: 2925.5 on 600 degrees of freedom Residual deviance: 2766.4 on 597 degrees of freedom

AIC: 3268.4

Number of Fisher Scoring iterations: 7

exp(coef(mod))

(Intercept) female yearsmarried female:yearsmarried 0.6679960 1.0955976 1.0845621 0.9942798

library(AER)

Loading required package: car

Loading required package: carData

Attaching package: 'car'

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

recode

```
The following object is masked from 'package:purrr':
    some
Loading required package: sandwich
Loading required package: survival
Attaching package: 'survival'
The following object is masked from 'package:caret':
    cluster
  dispersiontest(mod)
    Overdispersion test
data: mod
z = 8.3884, p-value < 2.2e-16
alternative hypothesis: true dispersion is greater than 1
sample estimates:
dispersion
  7.097836
  library(MASS)
Attaching package: 'MASS'
The following object is masked from 'package:dplyr':
    select
```

```
mod <- glm.nb(affairs ~ female*yearsmarried , data=affairs)
summary(mod)</pre>
```

#### Call:

glm.nb(formula = affairs ~ female \* yearsmarried, data = affairs, init.theta = 0.1202267055, link = log)

### Deviance Residuals:

Min 1Q Median 3Q Max -0.8612 -0.7974 -0.7166 -0.6622 1.7407

#### Coefficients:

Estimate Std. Error z value Pr(>|z|) (Intercept) -0.455305 0.303679 -1.499 0.13380 female -0.027146 0.449822 -0.060 0.95188 yearsmarried 0.086814 0.030411 2.855 0.00431 \*\* female:yearsmarried 0.006638 0.044534 0.149 0.88150

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for Negative Binomial(0.1202) family taken to be 1)

Null deviance: 347.24 on 600 degrees of freedom Residual deviance: 332.98 on 597 degrees of freedom

AIC: 1498.5

Number of Fisher Scoring iterations: 1

Theta: 0.1202 Std. Err.: 0.0129

2 x log-likelihood: -1488.5420

### exp(coef(mod))

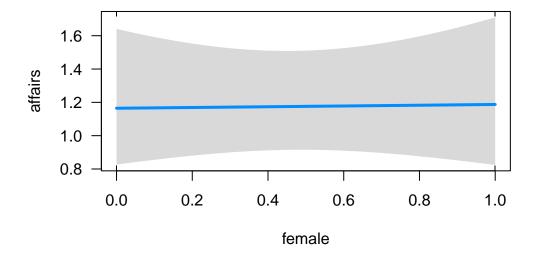
(Intercept) female yearsmarried female:yearsmarried 0.6342546 0.9732188 1.0906936 1.0066604

## visreg(mod, scale="response")

Warning: Note that you are attempting to plot a 'main effect' in a model that contains an interaction. This is potentially misleading; you may wish to consider using the 'by' argument.

Conditions used in construction of plot yearsmarried: 7

Warning: Note that you are attempting to plot a 'main effect' in a model that contains an interaction. This is potentially misleading; you may wish to consider using the 'by' argument.



Conditions used in construction of plot female: 0

