# Week 7: Mixed Models MATH-516 Applied Statistics

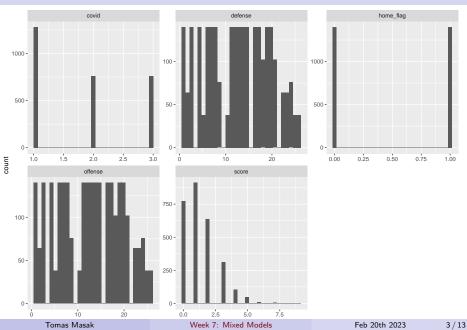
Tomas Masak

Feb 20th 2023

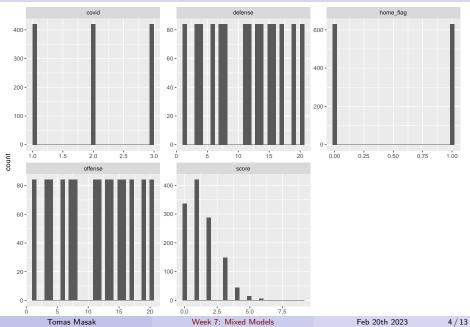
#### Section 1

Project 3: Premier League

## Data Visualized



## Balanced Data



# Models (Balanced Data)

## Model 2: score ~ home\_flag + defense + offense
## Resid. Df Resid. Dev Df Deviance Pr(>Chi)

1230 1357.5 -4 -4.7254 0.3167

1352.8

1226

## 2

```
m <- glm(score-covid*home_flag+defense+offense, data=subDat, family="poisson")
msub <- glm(score-home_flag+defense+offense, data=subDat, family="poisson")
anova(m,msub,test="LRT")

## Analysis of Deviance Table
##
## Model 1: score ~ covid * home flag + defense + offense</pre>
```

## covid:home\_flag 5.77 2 0.055895 . ## covid:defense 60.04 35 0.005292 \*\*

## covid:offense ## --- 36.35 35 0.405718

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

```
m <- glm(score~covid*(home flag+defense+offense), data=Data, family="poisson")
msub <- glm(score~home flag+defense+offense, data=Data, family="poisson")
anova(m.msub.test="LRT")
## Analysis of Deviance Table
##
## Model 1: score ~ covid * (home flag + defense + offense)
## Model 2: score ~ home flag + defense + offense
     Resid. Df Resid. Dev Df Deviance Pr(>Chi)
         2674
                 2952.6
## 1
         2748 3057.7 -74 -105.14 0.0101 *
## 2
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
library(car)
Anova (m, type=2)
## Analysis of Deviance Table (Type II tests)
##
## Response: score
##
                  LR Chisq Df Pr(>Chisq)
                    0.91 2 0.633917
## covid
                   16.99 1 3.761e-05 ***
## home_flag
## defense
                  184.35 25 < 2.2e-16 ***
## offense
                345.44 25 < 2.2e-16 ***
```

# Models (Full Data)

## [1] 30

```
minter <- glm(score~covid*(home flag+defense)+offense, data=Data, family="poisson")
msub <- glm(score~home_flag+covid*defense+offense, data=Data, family="poisson")
anova(m.msub.test="LRT")
## Analysis of Deviance Table
##
## Model 1: score ~ covid * (home flag + defense + offense)
## Model 2: score ~ home_flag + covid * defense + offense
    Resid. Df Resid. Dev Df Deviance Pr(>Chi)
          2674
                  2952 6
         2711
                  2994.7 -37 -42.102 0.2596
anova(minter.msub.test="LRT")
## Analysis of Deviance Table
##
## Model 1: score ~ covid * (home flag + defense) + offense
## Model 2: score ~ home_flag + covid * defense + offense
    Resid. Df Resid. Dev Df Deviance Pr(>Chi)
## 1
         2709
                  2988 9
## 2
          2711
                  2994.7 -2 -5.7544 0.05629 .
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
sum(is.na(coefficients(m)))
```

 we are on the edge of significance with a model that has too few observations to rely on asymptotics and to estimate all the parameters

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```
library(lme4)
m <- glmer(score ~ covid*home_flag+(1|defense/covid)+(1|offense)
msub <- glmer(score ~ covid*home_flag+(1|defense)+(1|offense)</pre>
```

- is m a good model?
- can it be simplified to msub?
- in both m and msub, the covid:home\_flag interaction looks significant
  - home advantage reduced during covid and bounced back after covid (not to the original level, but the difference not significant)

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## Common Feedback to Reports

- describe data you use, not data you were given
  - (nobody cares whether you got 4 csv files or a single one, and which variables were available but never used because they have absolutely nothing to do with anything, like the betting odds)
- there are some reserved words in statistics such as "significant" or "robust" that are better paraphrased when not used in the reserved meaning (statistical testing, robustness against outliers)
- multiple models vs. a single model
- not taking into account which teams are playing leads to dependence between data
- not including the intercept (i.e. manually discarding the intercept) is problematic since more parameters become inconsistent
  - Poisson vs. multinomial likelihoods
  - it is never a good idea to discard the intercept!
- doesn't vs. does not; let's vs. let us
- description of pre-processing your data (e.g. every match coded twice)

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#### Section 2

Common Feedback to Code by Charles

#### General comments

Overall good code, just a few remarks:

### Major:

- variable names should be descriptive data, data2 (use raw\_data, clean\_data)
- Code style (see next slides)
- Colors (see next slides)

#### Minor:

• commit messages: should describe work done by the commit (ahhhhh might reflect how you feel at the time of commit, but it's not very informative)

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## Code style

Code spacing and linting are important<sup>1</sup>

```
# Good
do something very complicated(
  something = "that",
  requires = many,
  arguments = "some of which may be long"
# Bad
do_something_very_complicated("that", requires, many, argument
                              "some of which may be long"
```

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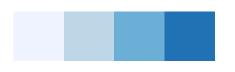
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<sup>&</sup>lt;sup>1</sup>check [https://style.tidyverse.org/]([https://style.tidyverse.org/) for a reference

## Color palette

- understand color scheme selection<sup>2</sup>
- use preset palettes:

```
library(RColorBrewer)
par(mfrow=c(1,2))
display.brewer.pal(name = "Blues", n = 4)
display.brewer.pal(name = "Set1", n = 4)
```





Blues (sequential)

Set1 (qualitative)

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 $<sup>^2</sup> see \ [https://www.gastonsanchez.com/] (https://www.gastonsanchez.com/) for more details$