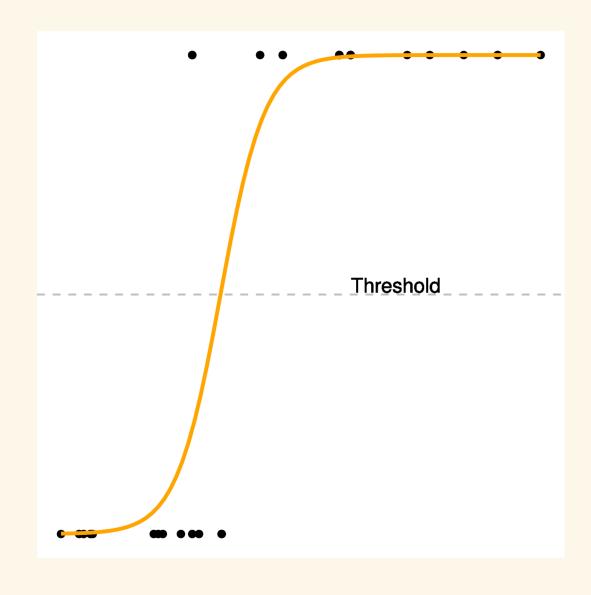
Logistic regression for binary responses: Diagnostics

Stat 230

May 16 2022

### Overview



### Today:

Checking log-odds linearity: log-odds plots

Residuals and Case influence stats

## Model Assumptions

$$Y_i \mid X_i \stackrel{indep.}{\sim} \mathrm{Bern}(\pi\left(X_i
ight)) \ \eta_i = \log\Bigl(rac{\pi(X_i)}{1-\pi(X_i)}\Bigr) = eta_0 + eta_1 x_{1,i} + \dots + eta_p x_{p,i}$$

#### **Independence**

- how are data collected?!
- Are the cases naturally clustered together in a way that isn't accounted for in the model?
- More on this in binomial logistic regression!

#### **Log-odds linearity**

• quantitative predictors are linearly related to the log odds of success

# EDA for logistic models: Empirical log odds plot

$$\eta_i = \logigg(rac{\pi\left(X_i
ight)}{1-\pi\left(X_i
ight)}igg) = eta_0 + eta_1 x_{1,i} + \dots + eta_p x_{p,i}$$

Plot the empirical (sample) log-odds against the predictor and look for linearity. Get empirical log odds for binned (grouped) data

- 1. Group cases into groups with similar predictor values using ntile
- 2. Compute (summarize) the proportion of successes within each group (group\_by)

$$ilde{\pi}_{emp} = rac{ ext{number of successes in the group}}{ ext{group size}}$$

3. Compute (summarize) the log odds of success in the group, within each group.

$$ext{logit}_{emp} = ext{ln}igg(rac{ ilde{\pi}_{emp}}{1- ilde{\pi}_{emp}}igg)$$

### Example: BWCA

#### 1999 windstorm in **northern MN**

what factors are associated with a tree blow down?

#### sample of 659 trees

- y = 1 means the tree died during the storm
- D: tree diameter (inches)

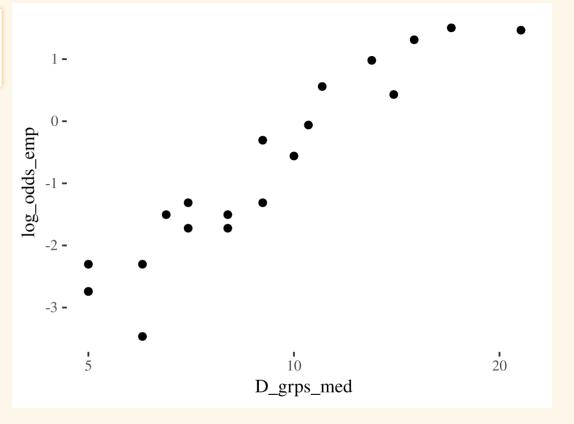
```
library(dplyr)
blowBF <- read.csv("https://raw.githubuserconte
mean(blowBF$y) # proportion died
[1] 0.353566</pre>
```

```
table(blowBF$status) # character response

died survived
233 426
```

### Linearity in empirical log odds vs. (median) diameter?

```
ggplot(blowBF_empLO, aes(x=D_grps_med, y=log_odds_emp)) +
  geom_point() +
scale_x_log10() # log is better
```



# Residuals in logistic models

• **Response:** response minus estimated mean

$$r_i = y_i - \hat{\pi}\left(X_i
ight)$$

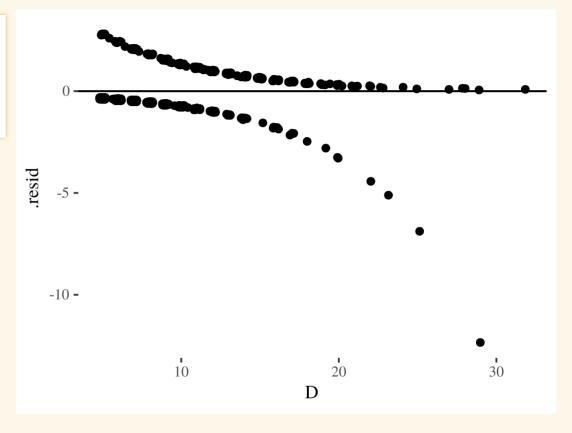
• **Pearson:** response residuals standardized based on the binomial SD:

$$pr_{i}=rac{y_{i}-\hat{\pi}\left(X_{i}
ight)}{\sqrt{\hat{\pi}\left(X_{i}
ight)\left(1-\hat{\pi}\left(X_{i}
ight)
ight)}}$$

• **Deviance:** each case's contribution to the residual deviance

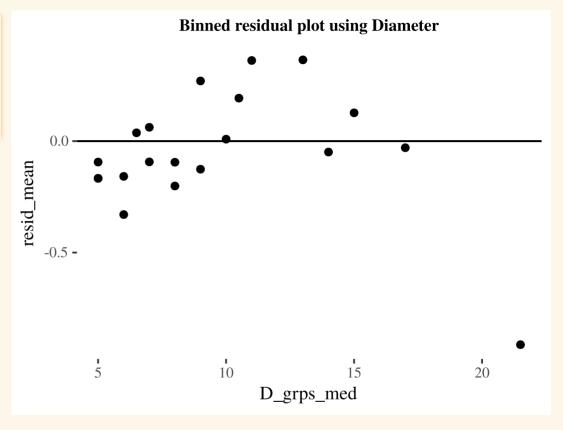
$$ext{Dres}_i = ext{sign}(y_i - \hat{\pi}\left(X_i
ight)) \sqrt{2\left[y_i \ln\!\left(rac{y_i}{\hat{\pi}\left(X_i
ight)}
ight) + (1-y_i) \ln\!\left(rac{1-y_i}{1-\hat{\pi}\left(X_i
ight)}
ight)
ight]}$$

#### Interpretation of binary model's residual plots



### Binned response residuals

```
ggplot(blowBF_resid1, aes(x=D_grps_med, y=resid_mean)) +
geom_point() +
geom_hline(yintercept = 0) +
labs(title="Binned residual plot using Diameter")+
    theme(plot.title = element_text(hjust=0.5, size=9, face='bol
```



Low diameter cases overestimated and higher diameters underestimated

## Case influence in logistic models

leverage and Cook's distance are used with logistic models, just as they are with regular linear models.

#### Use the usual R commands:

- plot(my\_glm, which = 5) (or type 4)
- ggResidpanel::resid\_panel(my\_glm, plots =c("cookd", "lev"))

- The odds of death as a function of diameter?
- Effect of doubling tree diameter?

• The odds of death as a function of diameter:

$$\widehat{odds}(D) = e^{-7.89 + 3.26 \ln(D)} = e^{-7.89} D^{3.26}$$

Doubling diameter is associated with a 9.58-fold increase in the odds of death (95% CI 6.68 to 14.12).

$$m^{\hat{eta}_1} = 2^{3.26} = 9.58$$

```
2^3.26 # estimate
[1] 9.57983
2^2.74 # lower bound
[1] 6.680703
2^3.82 # upper bound
[1] 14.12325
```

```
anova(fir_glm2)
```

```
Analysis of Deviance Table

Model: binomial, link: logit

Response: y

Terms added sequentially (first to last)

Df Deviance Resid. Df Resid. Dev

NULL
658 856.21
log(D) 1 200.97 657 655.24
```

# Example: BWCA with status response

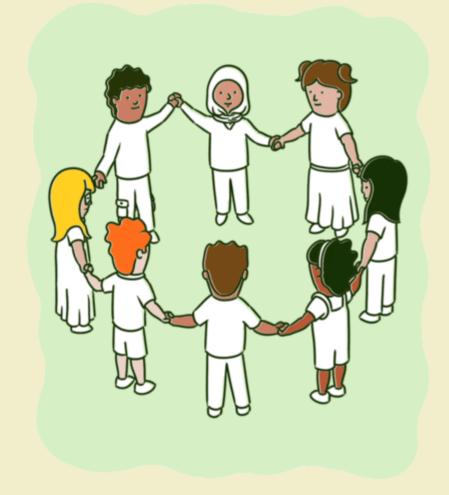
- status's second level is survived
- a model with status as the response will model the probability of survival!
  - glm will want this as.factor in order to fit the model!

```
table(blowBF$status)

died survived
233 426
```



05:00



- Go over to the in class activity file
- Go over the class activity in your group