# 5.1: Logistic Regression

Dr. Bean - Stat 5100

## 1 Why Logistic Regression?

Recall the linear regression model

$$Y = \beta_0 + \beta_1 X_1 + \dots + \beta_{p-1} X_{p-1} + \epsilon \qquad (\epsilon \sim N(0, \sigma^2)).$$

(Individual) What are some properties of the variable Y that are required for  $\epsilon$  to be normally distributed.

- Y must be linearly related to  $X_1, \ldots X_{p-1}$ .
- Y must be a continuous, quantitative variable

### 1.1 Why not regression on categorical data?

Consider fitting a regression model where we use age to try and predict whether or not a person has a disease (a 0-1 variable).

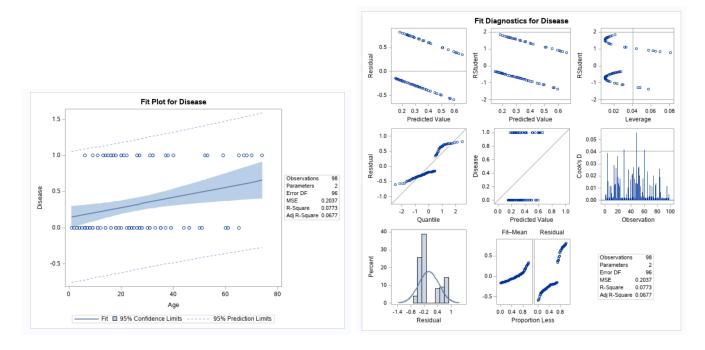


Figure 1: Fit plot and residual diagnostics for regression model that uses age to predict the presence/absence of a disease.

It is for this reason that instead of trying to predict the **value** of a categorical predictor, we should rather try to predict the **probability** of occurrence  $\pi_i$ ,

$$\pi_i = \beta_0 + \beta_1 X_{i,1} + \dots + \beta_{p-1} X_{i,p-1} + \epsilon_i \qquad (\epsilon \sim N(0, \sigma^2)).$$
 (1)

(Individual) However, based on the previous example, what are some of the issues with trying to predict the probability using (1)?

- We don't actually know  $\pi_i$ .
- Model can predict negative probabilities or probabilities above 1.
- Residual assumptions never satisfied (impossible for residuals to be normally distributed).

## 2 Transforming Probabilities

Because regression works best with **unconstrained** variables (i.e. variables that can theoretically take on any value). We need to find a transformation that maps  $\pi \in [0, 1]$  to  $f(\pi) \in (-\infty, \infty)$ .

Solution: log-odds ratio.

- $\pi \to [0, 1]$
- $\frac{\pi}{1-\pi} \to [0,\infty)$
- $L = \log\left(\frac{\pi}{1-\pi}\right) \to (-\infty, \infty)$

The **probit** function is another common transformation that achieves similar results.

• Probit:  $Q_i = Z_{\pi_i} \to \mathbb{Z}$  score (of a standard normal distribution) associated with the percentile  $\pi_i$ .

Other "S" shape curves exist, which tend to reach similar conclusions.

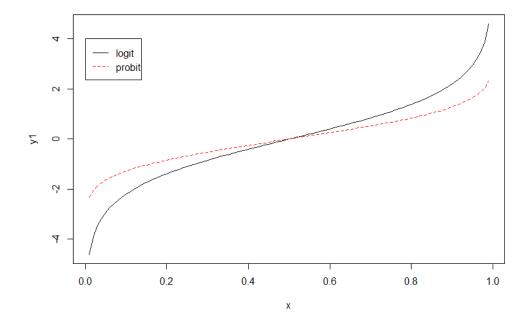


Figure 2: Visualization of logit and probit function for various probabilities.

## 3 Logistic Regression

$$L_i = \beta_0 + \beta_1 X_{i,1} + \dots + \beta_{p-1} X_{i,p-1} + \epsilon_i$$

- $b_k$  estimates obtained from MLE-based iterative procedure (Newton-Raphson, Fisher)
- Transform estimates  $\hat{L}_i = b_0 + b_1 X_{i,1} + \dots + b_{p-1} X_{i,p-1}$  back to probability scale.

$$\hat{\pi}_i = \frac{1}{1 + e^{-\hat{L}_i}} \qquad O\hat{d}ds_i = e^{\hat{L}_i}$$

#### 3.1 Interpretation of Estimates

- $X_{i,1} = \cdots = X_{i,p-1} = 0 \implies \hat{L}_i = b_0 \implies O\hat{d}ds_i = e^{b_0}$
- Hold  $X_{i,2} = \cdots = X_{i,p-1} = 0$ , increase  $X_{i,1}$  from 0 to 1

$$\implies \hat{L}_i = b_0 + b_1 \implies O\hat{d}ds_i = e^{b_0 + b_1} = e^{b_0}e^{b_1}$$

- Thus, an increase in one unit in  $X_j$  multiplies the odds (in favor of Y=1) by a factor of  $e^{b_j}$ .
  - Note that it is the *odds* that are multiplied, **not** the probability.
- Alternative Interpretation: the odds of Y = 1 change by  $100(e^{b_j} 1)\%$  per unit increase in  $X_i$  while holding other predictors constant.
  - Example (Handout 5.1.1):  $b_i$  for sector is 1.57  $\implies e^{1.57} = 4.83$ .
  - "Holding all other predictors constant, the odds of having disease are 100(4.83 1) = 383% greater in Sector 2 than in Sector 1.

# (Groups) How would you interpret the coefficient associated with Age in the Handout 5.1.1 logistic model?

"Holding all other predictors constant, the odds of having disease are  $100(e^{0.297}-1)=3.01\%$  greater for each year increase in age."

• The "Odds Ratio" for  $X_j$  (odds of Y=1 when  $X_j+1$  vs odds of Y=1 when  $X_j$ )

$$\frac{e^{b_0+b_1X_1+\cdots+\mathbf{b_j}(\mathbf{X_j}+1)+\cdots+b_{p-1}X_{p-1}}}{e^{b_0+b_1X_1+\cdots+\mathbf{b_j}(\mathbf{X_j})+\cdots+b_{p-1}X_{p-1}}}=e^{b_j}$$

#### 3.2 Inference with Estimates

- Single Variable Test:
  - $-H_0: \beta_i = 0 \ (X_i \text{ has no effect on } P(Y=1)).$
  - Test statistic:  $t = \frac{b_j}{SE\{b_i\}}$  (standard normal for "large" N).
  - $-\implies t^2 \sim \chi_1^2$  (obtain confidence intervals from here)
    - $\ast\,$  This approach is called the "Wald Test"
- Subset variables test:

$$- H_0: \beta_{p-H} = \dots = \beta_{p-1} = 0$$

- \* reorder the X variables so that the subset we are checking for comes last
- Let  $L_{full}$  be the likelihood associated with the full model
- Test statistics:  $\chi^2 = -2 \log \frac{L_{red}}{L_{full}}$
- Under  $H_0: \chi^2 \sim \chi_H^2$
- Overall model test:

$$Model\chi^2 = -2 \log L_{intercept} + 2 \log L_{int&covariates}$$

- Often called the **deviance**, DEV or  $DEV(X_0, X_1, X_{n-1})$
- $\bullet$  Conditional Effect plot: predicted  $\hat{pi}$  vs one predictor  $X_j$ 
  - While holding all other predictors at some constant level. The default level in SAS is the mean (average) of each variable.

### 4 Goodness of Fit Measures:

- Pseudo R-square:  $\frac{\chi^2}{\chi^2+n}$  ( $\chi^2$  from model test)
- Hosmer-Lemeshow Goodness of Fit Test
  - $-H_0$ : logistic regression response function is appropriate
  - Based on sorted  $\hat{\pi}$  values, group observations into 5-10 roughly equal sized groups.
  - Within each group, look at the total observed numbers of Y = 1 and Y = 0
  - Based on the model fit, calculate the total expected numbers of Y = 1 and Y = 0.
  - Test statistic  $\chi^2$  is sum (across groups) of  $\frac{(observed-expected)^2}{expected}$
- $\bullet$  "Concordance" look at all pairs of observations with different Y
  - Let  $n_c$  be the # of "concordant" pairs (observed Y = 1 has larger  $\hat{\pi}$ )
  - Let  $n_d$  be the # of "discordant" pairs (observed Y = 1 has smaller  $\hat{\pi}$ )
  - Let  $n_t$  be the # of "tied" paired (observed Y = 1 and Y = 0 have same  $\hat{\pi}$  (likely due to identical X-profiles)
  - Define rank correlation indices (larger is better):

Somers' 
$$D = \frac{n_c - n_d}{n_c + n_d + n_t}$$

$$\gamma = \frac{n_c - n_d}{n_c + n_d}$$

Tau-a = 
$$\frac{n_c - n_d}{0.5(n-1)n}$$

$$AUC = \frac{n_c + 0.5n_t}{n_c + n_d + n_t}$$

- ROC (Receiver Operating Characteristic) Curve
  - Sort all observations from the smallest to biggest  $\hat{pi}$ .
  - At each position in the list:
    - \* Use  $\hat{\pi}$  as threshold for  $\hat{Y} = 1$ , moving cutoff from the standard 0.5 threshold.
    - \* Calculate sensitivity: (proportion  $Y_i = 1$  values with  $\hat{Y}_i = 1$ ).
    - \* Calculate specificity: (proportion Y = 0 values with  $\hat{Y} = 0$ ).
      - · Sensitivity and Specificity think smoke alarms and pregnancy tests.
    - \* False positive rate (prop Y = 0 values with  $\hat{Y} = 1$ ) = 1 specificity
    - \* Plot false positives against true positive rates (sensitivity)
    - \* Calculate the area under the curve.

# Given the three ROC curves in Figure 3, which model has the best predictive power and why?

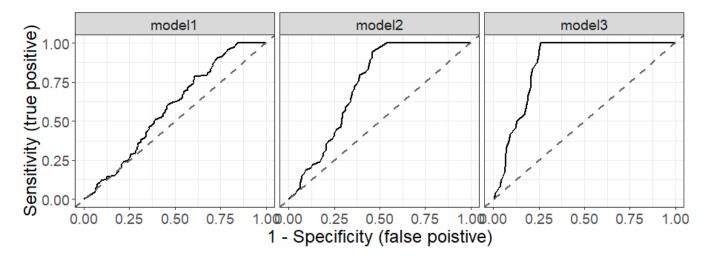


Figure 3: Comparison of three ROC curves.

Model 3 is the most accurate. The model sensitivity increases much faster than the false positive rate.

## 5 Multicollinearity

Recall that multicollinearity occurs when X variables are highly correlated with each other. It has **nothing** to do with the response variable Y.

As in OLS, multicollinearity inflates the variance of the  $b_k$  estimates, making them hard to interpret/test for significance.

As in OLS, stepwise selection and all possible regression methods exist to "score" each combination of explanatory variables and select a best model.

## 6 Outliers in Logistic Regression

(Individual) If Y can only take on two values (0 or 1), how are outlier values possible?

An outlier is a point for which the observation strongly disagrees with the predicted probability.

• Define "deviance residual" as

$$dev_i = \operatorname{sign}(Y_i - \hat{\pi}_i) \sqrt{-2 \left( Y_i \log \hat{\pi}_i + (1 - Y_i) \log(1 - \hat{p}i_i) \right)}$$

- The more certain we are (probability near 0 or 1), the more potential we have to be very wrong.
- $DEV(X_0, \cdots X_{p-1}) = \sum_i dev_i^2$
- "Outliers" are values not well represented by the model
- "Half-normal probability plot observed  $|dev_i|$  vs expected value under normality
  - **However**, since the residuals are not normally distributed, we asses differences from our expectation using simulations based on  $\hat{\pi}_i$ .
    - \* Create 19 simulations by generating a "new" response variable where the values of  $Y_{new,i} \sim Bernoulli(\hat{\pi}_i)$
  - Simulated envelop (SEE 5.1.1 MACRO ON CANVAS) plots the minimum, maximum, and mean of the 19 simulations
    - \* Why 19 simulations? Since our observed deviances represent the 20th observation, the probability that our deviances will fall outside the envelope is less than 5% IF the fitted model is appropriate.
    - \* Points falling outside in the envelop in the upper right corner of the plot are evidence of outliers/bad fits.

#### 7 Influential Observations

Influential observations have the same effect on model coefficients as they did in OLS.

Diagnostics (similar to Leverage and DFBETAS):

- $\delta D_i : DEV DEV_{(i)}$ 
  - Measures decrease in "misfit" when obs. i is ignored. (essentially measures the "poorness of fit for observation i).
  - "large"  $\delta D_i \implies \text{obs. } i \text{ overly influences model fit}$
  - SAS: DIFDEV on step difference in deviance
- $\delta B_i$ 
  - Similar to Cook's distance, measures influence of obs. i on the estimates  $b_j$

- SAS: C confidence interval displacement C
- $\delta \chi_i^2$ 
  - Similar to  $\delta D_i$ : "poorness of fit" for obs i
  - SAS: DIFCHISQ one step difference in Pearson  $\chi^2$

Unlike in OLS, there is no consistent numerical rule of thumbs to determine thresholds for the  $\delta$  measures.

Instead, we will simply rely on graphical diagnostics.

- $\delta D_i, \delta B_i, \delta X_i^2$  vs Observation Number look for extreme values
- $\delta D_i$  vs  $\hat{\pi}_i$  (or  $\delta X_i^2$  vs  $\hat{\pi}_i$ )
  - Look for points with low  $\hat{\pi}$  but  $Y_i = 1$  (upper left corner) OR high  $\hat{\pi}$  but Y = 0 (upper right corner) which are much different than the overall pattern
  - (Optional) plot different size points where point size is determined by  $\delta B_i$

#### 8 Remedial Measures

Similar to OLS:

- Look for typos in the data
- Consider transformations of the X variables
- Consider dropping problematic points (only if you have a good argument for removing them).

## 9 Final Thought

If you have a lot of explanatory variables, you should strongly consider classification trees and random forest for classification.