# IDS 702: Module 2.7

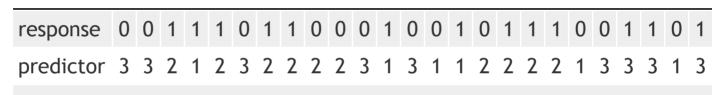
AGGREGATED OUTCOMES; PROBIT REGRESSION

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#### AGGREGATED BINARY OUTCOMES

- In the datasets we have seen so far under logistic regression, we observe the binary outcomes for each observation, that is, each  $y_i \in \{0,1\}$ .
- This is not always the case. Sometimes, we get an aggregated version, with the outcome summed up by combinations of other variables.
- For example, for individual-level data, suppose we had



where predictor is a factor with 3 levels: 1,2,3.

The aggregated version of the same data could look like

predictor	n	successes
1	31	17
2	35	16
3	34	14

#### AGGREGATED BINARY OUTCOMES

- Recall that if  $Y \sim \operatorname{Bin}(n,p)$  (that is, Y is a random variable that follows a binomial distribution with parameters n and p), then Y follows a  $\operatorname{Bernoulli}(p)$  distribution when n=1.
- lacksquare Alternatively, we also have that if  $Z_1,\ldots,Z_n\sim \mathrm{Bernoulli}(p)$ , then  $Y=\sum_i^n Z_i\sim \mathrm{Bin}(n,p).$
- That is, the sum of n "iid"  $\operatorname{Bernoulli}(p)$  random variables gives a random variable with the  $\operatorname{Bin}(n,p)$  distribution.
- The logistic regression model can be used either for Bernoulli data (as we have done so far) or for data summarized as binomial counts (that is, aggregated counts).
- In the aggregated form, the model is

$$y_i|x_i\sim ext{Bin}(n_i,\pi_i); \ \ \log\left(rac{\pi_i}{1-\pi_i}
ight)=eta_0+eta_1x_{i1}+eta_2x_{i2}+\ldots+eta_px_{ip},$$

### BERNOULLI VERSUS BINOMIAL OUTCOMES

#### Normally, for individual-level data, we would have

response predictor

```
## 1
## 2
           1
                     1
## 5
## 6
                     3
M1 <- glm(response~predictor,data=Data,family=binomial)
summary(M1)
##
## Call:
## glm(formula = response ~ predictor, family = binomial, data = Data)
## Deviance Residuals:
              10 Median
     Min
                              30
                                     Max
## -1.261 -1.105 -1.030 1.251
                                  1.332
##
## Coefficients:
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) 0.1942
                           0.3609
                                   0.538
                                             0.591
## predictor2 -0.3660
                           0.4954 - 0.739
                                             0.460
## predictor3 -0.5508
                           0.5017 -1.098
                                             0.272
##
## (Dispersion parameter for binomial family taken to be 1)
##
      Null deviance: 138.27 on 99 degrees of freedom
## Residual deviance: 137.02 on 97 degrees of freedom
## AIC: 143.02
## Number of Fisher Scoring iterations: 4
```

### BERNOULLI VERSUS BINOMIAL OUTCOMES

But we could also do the following with the aggregate level data instead

```
M2 <- glm(cbind(successes, n-successes)~predictor, data=Data agg, family=binomial)
summary (M2)
##
## Call:
## glm(formula = cbind(successes, n - successes) ~ predictor, family = binomial,
      data = Data agg)
##
## Deviance Residuals:
## [1] 0 0 0
##
## Coefficients:
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) 0.1942
                           0.3609
                                   0.538
                                              0.591
## predictor2 -0.3660
                         0.4954 -0.739
                                              0.460
## predictor3 -0.5508
                           0.5017 - 1.098
                                              0.272
## (Dispersion parameter for binomial family taken to be 1)
##
      Null deviance: 1.2524e+00 on 2 degrees of freedom
## Residual deviance: 1.3323e-14 on 0 degrees of freedom
## AIC: 17.868
## Number of Fisher Scoring iterations: 2
```

Same results overall! Deviance and AIC are different because of the different likelihood functions.

Note that some glm functions use n in the formular instead of n-successes.

# PROBIT REGRESSION



#### PROBIT REGRESSION

Recall the "Bernoulli" logistic regression model:

$$y_i|x_i \sim \mathrm{Bernoulli}(\pi_i); \ \ \log\left(rac{\pi_i}{1-\pi_i}
ight) = eta_0 + eta_1 x_{i1} + eta_2 x_{i2} + \ldots + eta_p x_{ip},$$

for 
$$i = 1, \ldots, n$$
.

- Here the link function is the logit function, which ensures that the probabilities lie between 0 and 1.
- We can also use the probit function  $\Phi^{-1}$ , which is the quantile function associated with the standard normal distribution N(0,1), as the link.

#### PROBIT REGRESSION

- lacktriangledown That is, suppose H follows a standard normal distribution, that is,  $H \sim N(0,1).$
- ullet Then  $\Phi$  is the CDF, that is,  $\Pr[H \leq h] = \Phi(h)$ .
- Formally, the probit regression model can be written as

$$y_i|x_i \sim \mathrm{Bernoulli}(\pi_i); \quad \Phi^{-1}\left(\pi_i
ight) = eta_0 + eta_1 x_{i1} + eta_2 x_{i2} + \ldots + eta_p x_{ip}.$$

It is then easy to see that

$$egin{align} \Pr[y_i=1|x_i] &= \pi_i = \Phi\left(eta_0 + eta_1 x_{i1} + eta_2 x_{i2} + \ldots + eta_p x_{ip}
ight) \ &= \Pr[H \leq eta_0 + eta_1 x_{i1} + eta_2 x_{i2} + \ldots + eta_p x_{ip}]. \end{split}$$

#### LATENT VARIABLE REPRESENTATION

It turns out that we can rewrite the probit regression model as

$$egin{aligned} y_i &= 1[z_i > 0]; \ z_i &= eta_0 + eta_1 x_{i1} + eta_2 x_{i2} + \ldots + eta_p x_{ip} + \epsilon_i; \ \ \epsilon_i \sim N(0,1) \end{aligned}$$

where  $y_i = \mathbb{1}[z_i > 0]$  means  $y_i = 1$  if  $z_i > 0$  and  $y_i = 0$  if  $z_i < 0$ .

To see that the two representations are equivalent, note that

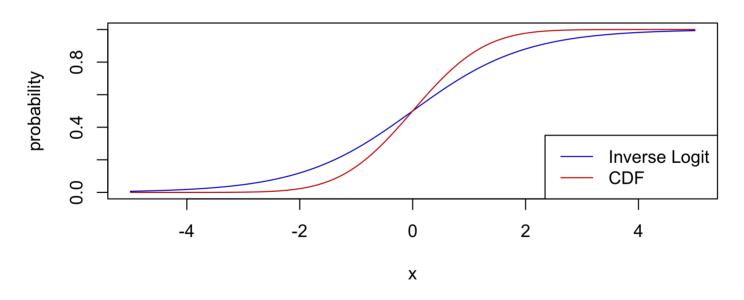
$$egin{aligned} \Pr[y_i = 1 | x_i] &= \Pr[z_i > 0] \ &= \Pr[eta_0 + eta_1 x_{i1} + eta_2 x_{i2} + \ldots + eta_p x_{ip} + \epsilon_i > 0] \ &= \Pr[\epsilon_i > -(eta_0 + eta_1 x_{i1} + eta_2 x_{i2} + \ldots + eta_p x_{ip})] \ &= \Pr[\epsilon_i < (eta_0 + eta_1 x_{i1} + eta_2 x_{i2} + \ldots + eta_p x_{ip})] \quad [\text{since} \quad \epsilon_i \sim N(0, 1)] \ &= \Phi\left(eta_0 + eta_1 x_{i1} + eta_2 x_{i2} + \ldots + eta_p x_{ip}\right) = \pi_i \end{aligned}$$

• Clearly, we do not observe  $Z=(z_1,z_2,\ldots,z_n)$  and it is thus referred to as an auxiliary variable.

## PROBIT VS LOGIT FUNCTIONS?

lacktriangle The plots below compares the inverse logit function  $\pi_i=rac{e^x}{1+e^x}$  and the CDF function (inverse probit)  $\pi_i=\Phi(x)$ .

#### **Probit vs logit functions**





 Notice that they are similar, but the CDF of the standard normal distribution has fatter tails (the inverse logit has thinner tails).

## PROBIT OR LOGISTIC REGRESSION?

- In practice, the decision to use one or the other is often based on preference: the overall conclusions from both are usually quite similar.
- The results based on logistic regression (using odds and odds ratio) can be more interpretable than those based on Probit regression.
- In some applications, interpreting the  $z_i$ 's may be meaningful but that is not always the case.
- For example, suppose  $y_i$  is a binary variable for whether or not person i chooses to buy the new iPhone, then  $z_i$  can be thought of as person i's "utility" in a way.
- Works in this example, but does not always work across different domains.
- In R, use the glm command but set the option family="binomial(link=probit) instead of family="binomial(link=logit).

# WHAT'S NEXT?

MOVE ON TO THE READINGS FOR THE NEXT MODULE!

