GOV 2001 / 1002 / E-200 Section 5 Binary Dependent Variable Regression¹

Anton Strezhnev

Harvard University

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¹These section notes are heavily indebted to past Gov 2001 TFs for slides and R code.

LOGISTICS

Reading Assignment- 4 papers on binary dependent variable models - pay attention particularly to the applications and common pitfalls.

Problem Set 5- Due by 6pm, 3/9 on Canvas.

Assessment Question- Due by 6pm, 3/9 on on Canvas. You must work alone and only <u>one</u> attempt.

OVERVIEW

- ► In this section you will...
 - ► learn why the Fisher information is used to approximate standard errors of MLEs
 - learn about generalized linear models and how logit models fit into that framework
 - ▶ learn how to evaluate binary dependent variable models.

OUTLINE

MLES IN ASYMPTOPIA

- ► For the entirety of this course, we'll be working with the "large sample" properties of Maximum Likelihood Estimators.
- ► This is because most MLEs particularly for generalized linear models are biased in finite samples. But that bias goes away as the sample gets large.
- ► Also, small sample standard errors are hard to calculate analytically no easy formula like OLS.
- Warning: Be wary of published GLM results with small samples - they don't have the same small-sample unbiasedness properties as OLS

LARGE-SAMPLE PROPERTIES OF MLES

- ► Two big properties:
- ► Consistency: $\hat{\theta}_{MLE} \xrightarrow{p} \theta_0$. The MLE estimator converges in probability to the true value θ_0 as n gets large.
- ► **Asymptotic Normality:** $\hat{\theta}_{MLE} \sim \text{Normal}(\theta_0, \sigma_{MLE}^2)$ in large samples.
- ▶ But how do we calculate σ_{MLE}^2 !?

REVIEW: ESTIMATOR VARIANCE

- ▶ Why does the MLE have a variance?
- Because the data is random! We have a stochastic component in our model that describes how the observations are generated.
- ► If we drew another sample, or re-ran the experiment on another hypothetical group, we would get a different MLE. Imagining repeating this process over and over gives us the theoretical *sampling distribution*.
- ► More simply, likelihoods are sums of random variables (e.g. *Y*_i). Therefore functions of them are also random variables!

ASYMPTOTIC VARIANCE OF MLES

- ▶ When we did OLS, we could get σ_{MLE}^2 by just taking $Var(\hat{\theta})$ since there was a closed form solution
- $Var(\hat{\theta}_{OLS}) = Var\left((X'X)^{-1}X'Y\right) = \sigma^2(X'X)^{-1}$
- ▶ But we don't have closed form solutions for almost all MLEs $\hat{\theta}_{MLE}$. What can we use?

THE SCORE

- ▶ Remember that the likelihood of θ : $L(\theta|X) = p(X|\theta)$.
- ► The log-likelihood $\ell(\theta|X) = \ln p(X|\theta)$.
- ► The derivative of the log-likelihood is known as the "score" function.
 - $S(\theta) = \frac{\partial}{\partial \theta} \ln p(X|\theta) = \ell'(\theta|X)$
- ► At the MLE, by definition, the score is 0: $S(\hat{\theta}) = 0$.
- We can also show that the expectation of the score function is also 0

$$E[S(\theta)] = 0$$

THE INFORMATION

► Where does the "information matrix" we talk about come from? Well, it's the variance of the score.

$$I(\theta) = Var[S(\theta)]$$
$$= E[S(\theta)^{2}] - E[S(\theta)]^{2}$$

From before, the second term is 0. So

$$I(\theta) = E[S(\theta)^2]$$

It turns out that we can also show that it equals the expectation of the negative of the Hessian of the likelihood.

$$I(\theta) = E[S(\theta)^2] = E[-\ell''(\theta)]$$

POWER OF THE I.I.D. ASSUMPTION

- ► A lot of intuition for asymptotics comes from our i.i.d. assumption. Under this assumption, our log-likelihoods are sums of *n* separate log-likelihoods for each observation.
- ► So $\ell(\theta|X) = \sum_{i=1}^{n} \ell(\theta|X_i)$ under i.i.d. observations.
- ► Likewise, $S(\theta) = \ell'(\theta|X) = \sum_{i=1}^{n} \ell'(\theta|X_i)$
- ▶ Under the i.i.d. assumption, the information also grows with *n*, so we often denote for i.i.d. observations

$$I_n(\theta) = E\left[-\sum_{i=1}^n \ell''(\theta|X_i)\right] = -nE\left[\ell''(\theta|X_1)\right]$$

► **Key takeaway:** Under the i.i.d. assumption, likelihoods are sums of *i.i.d.* random variables. This lets us invoke the Law of Large Numbers and Central Limit Theorem.

► Let's start with the quadratic Taylor approximation of the likelihood around the true value θ_0 .

$$\ell(\theta) \approx \ell(\theta_0) + \ell'(\theta_0)(\theta - \theta_0) + \frac{1}{2}\ell''(\theta_0)(\theta - \theta_0)^2$$

► Take the derivative to get the score

$$\ell'(\theta) = \ell'(\theta_0) + \ell''(\theta_0)(\theta - \theta_0)$$

▶ At the MLE, $\hat{\theta}$, the score is 0, so we can write

$$0 = \ell'(\theta_0) + \ell''(\theta_0)(\hat{\theta} - \theta_0)$$

► Re-arranging terms

$$(\hat{\theta} - \theta_0) = \frac{\ell'(\theta_0)}{-\ell''(\theta_0)}$$

► Multiply both sides by \sqrt{n}

$$\sqrt{n}(\hat{\theta} - \theta_0) = \frac{\frac{1}{\sqrt{n}}\ell'(\theta_0)}{-\frac{1}{n}\ell''(\theta_0)}$$

► Now we make use of two convergence rules. Recall that both the score and the information are sums of i.i.d. random variables

► First, by the Law of Large Numbers, the denominator converges to the information for a single observation

$$-\frac{1}{n}\ell''(\theta_0) = -\frac{1}{n}\sum_{i=1}^n \ell''(\theta_0|X_i) \xrightarrow{p} -E[\ell''(\theta_0|X_i)] = I(\theta)$$

► Second, by the Central Limit Theorem, the numerator converges in distribution to a normal distribution with mean $E[S(\theta|X_i)] = 0$ and variance $Var(S(\theta|X_i)] = I(\theta)$

$$\sqrt{n}\frac{1}{n}\sum_{i=1}^{n}\ell'(\theta|X_i) \xrightarrow{d} \text{Normal}(0,I(\theta))$$

▶ By Slutsky's theorem, therefore, we have

$$\sqrt{n}(\hat{\theta} - \theta_0) \xrightarrow{d} \text{Normal}\left(0, \frac{1}{I(\theta)}\right)$$

► And by extension

$$\hat{\theta} \xrightarrow{d} \text{Normal} \left(\theta_0, \frac{1}{I_n(\theta)} \right)$$

► Finally, $I_n(\theta)$ is an expectation that's hard to calculate. Instead, we plug in a consistent estimate – the observed information $I_n(\hat{\theta})$. This is the negative Hessian evaluated at the MLE.

REGULARITY CONDITIONS IN PRACTICE

- We've been handwaving a lot about what makes a likelihood "nice" for the asymptotics to apply. But what does this mean in practice?
- ► Three big "regularity" conditions for convergence:
- ▶ **Identifiability** If $L(\hat{\theta}) = L(\theta)$, then $\hat{\theta} = \theta$. That is, there is a single value of θ that maximizes the likelihood. One place where this doesn't hold is when there are more parameters than data points.
- ▶ **I.i.d. observations** The likelihood can be factored into n identical and independent densities that is, $L(\theta|X) = \prod_{i=1}^{n} L(\theta|X_i)$
- ► Parameter space fixed relative to *n* As sample size increases, the number of parameters being estimated *doesn't* grow with it.

OUTLINE

GENERALIZED LINEAR MODELS

- ► Most models that we work with in this course are part of a class of models called *generalized linear models (GLM)*.
- ► Takes the "linear" component $X\beta$ from OLS and allows it to model outcomes with different types of distributions.
- ► Three components:
 - ► A distribution for *Y* (stochastic component)
 - A linear predictor for $X\beta$ (systematic component)
 - ► A link function that connects the linear predictor to parameters of the distribution on *Y*

1. PICK A DISTRIBUTION FOR Y

- We start by assuming our data comes from some distribution.
- ► Examples:
 - ► Continuous and Unbounded: **Normal**(μ , σ^2)
 - ▶ Binary: **Bernoulli**(π)
 - ► Event Count: **Poisson**(λ), **Negative Binomial**(r, p)
 - ▶ Duration: **Exponential**(λ), **Weibull**(λ ,k)
 - ▶ Unordered Categories: **Multinomial**(π)
- ► Sometimes, instead of directly putting a distribution on *Y*, we can put a distribution on an unobserved "latent" variable *Y** and treat *Y* as a function of *Y** e.g. for ordered categorical data, *Y** is unbounded, but *Y* is a piece-wise function of *Y**.

2. Specify a linear predictor $\eta = X\beta$

- ▶ Our covariates X enter into a GLM in a very specific way. As in OLS, the linear predictor is a linear combination of the parameters β .
- ▶ If we have *k* covariates, our linear predictor $\eta = X\beta$ is:

$$\eta = X\beta = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \ldots + x_k \beta_k$$

► Just like OLS, we can have (non-linear) functions of *X* as covariates (e.g. *X*²), but our *parameters* are a linear combination.

3. Specify a link function $g(\mu) = X\beta$

- ▶ Finally, we need to connect the linear predictor to the mean μ of the distribution on Y. Often this will be a parameter of that distribution.
- ► Lots of choices we need the domain of the link to match the range of the mean.
- We pick a $g(\dot{})$ and set $g(\mu) = X\beta$
- ► Then solve back to get the inverse link $\mu = g^{-1}(X\beta)$.

LINK FUNCTION EXAMPLE: LOGIT

- ▶ If $Y_i \sim \text{Bernoulli}(\pi_i)$, then $E[Y_i] = \pi_i$.
- ▶ $\pi_i \in (0,1)$, so we need a function that maps from (0,1) to $(-\infty,\infty)$.
- ► One function is the "logit" logit(p) = ln $\left(\frac{p}{1-p}\right)$.
- ▶ So we set $\ln \left(\frac{\pi_i}{1-\pi_i} \right) = X_i \beta$. And take the inverse to solve for π_i .

$$\ln\left(\frac{1-\pi_i}{\pi_i}\right) = -X_i\beta$$

$$\frac{1-\pi_i}{\pi_i} = \exp(-X_i\beta)$$

$$\frac{1}{\pi_i} - 1 = \exp(-X_i\beta)$$

$$\frac{1}{\pi_i} = 1 + \exp(-X_i\beta)$$

$$\pi_i = \frac{1}{1 + \exp(-X_i\beta)}$$

VISUALIZING THE INVERSE LOGIT

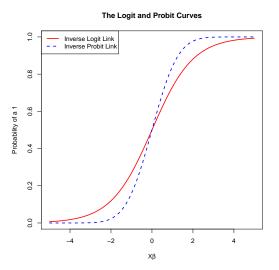


Figure: Comparison of inverse-logit and inverse-probit links

OUTLINE

FORECASTING CONGRESSIONAL ELECTIONS



Suppose we want to forecast whether or not the incumbent party will win the U.S. House general election.

FORECASTING - DEFINE THE MODEL

- ► First, let's define a model. We observe n observations each with outcome Y_i and covariates X_i .
- ► Our distribution on the data is simple:

$$Y_i \sim \text{Bernoulli}(\pi_i)$$

▶ Our linear predictor is a function of covariates. In this case, we observe three: whether the seat is open: *open*; whether the incumbent is a freshman: *fresh*; and the vote share of the incumbent party in the district in the last presidential election: *incshare*.

FORECASTING - DEFINE THE MODEL

▶ So our linear predictor η_i is

$$\eta_i = X_i \beta = \beta_0 + \beta_1 X_{i,open} + \beta_2 X_{i,fresh} + \beta_3 X_{i,incshare}$$

► Finally, we pick a link function. For simplicity, we'll pick the logit link, which yields

$$\pi_i = \frac{1}{1 + \exp(-X_i \beta)}$$

FORECASTING - ESTIMATE THE MODEL

► In class, we derived the log-likelihood of this model. Which we can maximize numerically in R.

```
## Load election results from 04-08
votes <- read.dta("votes0408.dta")
## Our log-likelihood function, logit.ll takes three arguments:
## par: the parameters
## outcome: the Y variable
## covariates: the X matrix (including an intercept column
## Create the X matrix
design.matrix <- as.matrix(cbind(1,votes[,c("open","freshman","incpres")]))</pre>
## Estimate the MLE:
opt <- optim(par = rep(0, ncol(votes[,2:4]) + 1),
             fn = logit.ll,
             covariates = design.matrix,
             outcome = votes$incwin.
             control = list(fnscale = -1),
             hessian = T.
             method = "BFGS")
```

Forecasting - Estimates for 2004-2008

► Our coefficient estimates are

```
coefs <- opt$par # Beta values for the intercept and 3 coefficients
names(coefs) <- c("Intercept", "open", "freshman", "incpres")
coefs
Intercept open freshman incpres
-2.9064379 -2.1266744 -0.3568115 0.1112137
```

And our estimates of the standard errors are

FORECASTING - QUANTITIES OF INTEREST

- ▶ By themselves, the coefficients are hard to interpret (log-odds ratios). We want to obtain more informative quantities. One intuitive quantity is a predicted probability $\hat{\pi}_i$ for some set of covariates X_i .
- ► By MLE invariance,

$$\hat{\pi_i} = \frac{1}{1 + \exp(-X_i \hat{\beta})}$$

▶ So applying the inverse-logit to our linear predictor us an MLE estimate of $\hat{\pi_i}$! This is how we get "fitted values" for a logit model. It's also how we make predictions for *new* or hypothetical observations of X_i .

FORECASTING - VALIDATION

- How well does our model explain our data? Couple of ways of evaluating this in the binary D.V. context? The Greenhill et. al. reading for this week gives a general overview.
- ► People often look at accuracy given some "cut-off" probability, how many cases are correctly predicted by the model. Accuracy can be misleading!
- ► Suppose we have a very rare event e.g. only 1% of cases are 1s. Then a model that just always predicted 0 would have 99% accuracy!
- ► Instead, in binary classification, we often care about *sensitivity* vs. *specificity*

FORECASTING - SENSITIVITY AND SPECIFICITY

	Actual Outcome	
Predicted Outcome	Negative	Positive
Negative	True Negative	False Negative
Positive	False Positive	True Positive

Table: Confusion matrix for binary predictions

- ► Sensitivity = True Positive Rate = $\frac{\sum \text{True Positive}}{\sum \text{Actual Positives}}$
- ► Specificity = True Negative Rate = $\frac{\sum \text{True Negative}}{\sum \text{Actual Negatives}}$

FORECASTING - SENSITIVITY AND SPECIFICITY

- ► Given a model, there's a trade-off between Sensitivity and Specificity. In a naive model, we can always get 100% sensitivity by labeling everything as a positive. But this would yield 0% specificity.
- Sensitivity and specificity are going to depend on the "cutoff" $\hat{\pi}_0$ that we use to classify observations as either 0s or 1s. So we can control one, and see how well we do on the other.
- ► How do we quantify the trade-off for our particular model. Receiver Operating Characteristic (ROC) plots!
 - ► Basically, a plot of True Positive Rate on Y axis against False Positive Rate (1 True Negative Rate) on the X axis.

- ► How to create an ROC:
 - ▶ Pick a threshold $\pi_0 \in [0,1]$.
 - For your test data, generate predictions for each observation $\hat{\pi}_i$.
 - ▶ Predict $\hat{Y}_i = 0$ if $\pi_i < \pi_0$ and $\hat{Y}_i = 1$ otherwise.
 - Calculate sensitivity and specificity.
 - ▶ Repeat for values of π_0 from 0 to 1 and plot.

Here's what it looks like in R. First we get our predicted probabilities.

```
#### ROC curve
thresholds <- seq(0, 1, by=.001) ## Vector of thresholds to test
sensitivity <- rep(NA, length(thresholds))
specificity <- rep(NA, length(thresholds))

### Get predicted probabilities
pred.probs <- 1/(1 + exp(-design.matrix%*%coefs))</pre>
```

Then we calculate true positive rate and true negative rate for each threshold

```
### For each threshold
for(i in 1:length(thresholds)){
 ### Select the threshold
 thresh <- thresholds[i]
 ### Make a prediction
 v hat <- ifelse(pred.probs < thresh, 0, 1)
 ### Compare to true Y
 cross tab <- table(v hat, votes$incwin)
 ### R-hack - Make sure cross tab is a 2x2.
 if (nrow(cross tab) == 2 & ncol(cross tab) == 2) {
    ## True positive rate (1s correctly predicted/total 1s)
    tpr <- cross tab[2,2]/(cross tab[2,2] + cross tab[1,2])
    ## True negative rate
    tnr \leftarrow cross tab[1,1]/(cross tab[1,1] + cross tab[2,1])
  }else{
 ### If we only predicted one class
    if (max(y_hat) == 0){
      ### If we only predict zeroes, no false positives, but no true positives
      tpr <- 0
      tnr <- 1
    }else if (min(v hat) == 1){
      ### If we only predict 1s, no true negatives, but all true positives
      tpr <- 1
      tnr <- 0
  sensitivity[i] <- tpr
  specificity[i] <- tnr
```

Finally, we plot it!

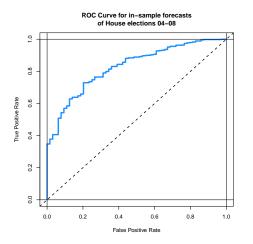


Figure: ROC Plot for House Election Logit Model

FORECASTING - IN-SAMPLE VS. OUT-OF-SAMPLE FORECASTING

- ► Careful when validating models to be wary of over-fitting. We can have a model fit perfectly to our sample that does terribly on other samples. This is because our model is *over-fit* the estimates are highly sensitive to arbitrary noise in the sample.
- ► Analogy Can always get a better *R*² by adding more junk to a linear model does that make a model with millions of covariates better?
- ► Solution is to fit a model to one part of the data and use it to forecast another part "cross-validation."
- ► Often see all of these prediction diagnostics used on a "held-out" set of data that the model does not "see" during estimation.

QUESTIONS

Questions?