Interpreting Models for Categorical and Count Outcomes

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Interpreting Models for Categorical and Count Outcomes Handout page: 1



- Learn how to fit models that include categorical variables and/or interactions using factor variable syntax
- Get an overview of tools available for investigating models
- Learn a bit about how Stata partitions model fitting and model testing tasks



Factor Variables

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A Logistic Regression Model

- We'll use data from the National Health and Nutrition Examination Survey (NHANES) for our examples
 - . webuse nhanes2
- We'll start with a model for high blood pressure (highbp) using age, body mass index (bmi) and sex (female)
- Before we fit the model, let's investigate the variables
 - . codebook highbp age bmi female
- Now we can fit the model
 - . logit highbp age bmi female

Factor Variables

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Working with Categorical Variables

- Now we would like to include region in the model, let's take a look at this variable
 - . codebook region
- region cannot simply be added to the list of covariates because it has 4 categories
- To include a categorical variable, put an i. in front of its name—this declares the variable to be a categorical variable, or in Stataese, a *factor variable*
- For example
 - . logit highbp age bmi i.female i.region

Factor Variables

Niceities

- Starting in Stata 13, value labels associated with factor variables are displayed in the regression table
- We can tell Stata to show the base categories for our factor variables
 - . set showbaselevels on
 - This means the base category will always be clearly documented in the output

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Factor Variables

Factor Notation as Operators

- The i. operator can be applied to many variables at once:
 - . logit highbp age bmi i.(female region)
- In other words, it understands the distributive property
 - This is useful when using variable ranges, for example
- For the curious, factor variable notation works with wildcards
 - If there were many variables starting with u, then i.u* would include them all as factor variables

Factor Variables

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Using Different Base Categories

- By default, the smallest-valued category is the base category
- This can be overridden within commands
 - b#. specifies the value # as the base
 - b(##). specifies the #'th largest value as the base
 - b(first). specifies the smallest value as the base
 - b(last). specifies the largest value as the base
 - b(freq). specifies the most prevalent value as the base
 - bn. specifies there should be no base
- The base can also be permanently changed using fvset; see help fvset for more information

Factor Variables

Playing with the Base

- We can use region=3 as the base class on the fly:
 - . logit highbp age bmi i.female b3.region
- We can use the most prevalent category as the base
 - . logit highbp age bmi i.female b(freq).region
- Factor variables can be distributed across many variables
 - . logit highbp age bmi b(freq).(female region)
- The base category can be omitted (with some care here)
 - . logit highbp age bmi i.female bn.region, noconstant

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- We can also include a term for region=4 only
 - . logit highbp age bmi i.female 4.region

Factor Variables

Specifying Interactions

- Factor variables are also used for specifying interactions
 - This is where they really shine
- To include both main effects and interaction terms in a model, put ## between the variables
- To include only the interaction terms, put # between the terms
- Variables involved in interactions are treated as categorical by default
 - Prefix a variable with c. to specify that a variable is continuous
- Here is our model with an interaction between age and female
 - . logit highbp bmi c.age##female i.region

Factor Variables

Some Factor Variable Notes

- If you plan to look at marginal effects of any kind, it is best to
 - Explicitly mark all categorical variables with i.
 - Specify all interactions using # or ##
 - Specify powers of a variable as interactions of the variable with itself
- There can be up to 8 categorical and 8 continuous interactions in one expression
 - Have fun with the interpretation

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Introduction to Postestimation

- In Stata jargon, postestimation commands are commands that can be run after a model is fit, for example
 - Predictions
 - Additional hypothesis tests
 - Checks of assumptions
- We'll explore postestimation tools that can be used to help interpret model results
 - The main example here is after logit models, but these tools can be used with most estimation commands
- The usefulness of specific tools will depend on the types of hypotheses you wish to examine

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Finding the Coefficient Names

- Some postestimation commands require that you know the names used to store the coefficients
- To see these names we can replay the model showing the *coefficient legend*
 - . logit, coeflegend
- From here, we can see the full specification of the factor levels:
 - _b[2.region] corresponds to region=2 which is "MW" or midwest
 - _b[3.region] corresponds to region=3 which is "S" or south

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• The coefficient for the female by age interaction is stored as _b[1.female#c.age]

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Joint Tests

- The test command performs a Wald test of the specified null hypothesis
 - The default test is that the listed terms are equal to 0
- test takes a list of terms, which may be variable names, but can also be terms associated with factor variables
- To specify a joint test of the null hypothesis that the coefficients for the levels of region are all equal to 0
 - . test 2.region 3.region 4.region

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Testing Sets of Coefficients

- If you are testing a large number of terms, typing them all out can be laborious
- testparm also performs Wald tests, but it accepts lists of variables, rather than coefficients in the model
- For example, to test all coefficients associated with i.region . testparm i.region

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Likelihood Ratio Tests

- Likelihood ratio tests provide an alternative method of testing sets of coefficients
- To test the coefficients associated with region we need to store our model results. The name is arbitrary, we'll call them m1 . estimates store m1
- Now we can rerun our model without region
 - . logit highbp bmi c.age##female if e(sample)
- Adding if e(sample) makes sure the same sample, what Stata calls the *estimation sample*, is used for both models

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Likelihood Ratio Tests (Continued)

- Now we store the second set of estimates
 - . estimates store m2
- And use the lrtest command to perform the likelihood ratio test
 - . lrtest m1 m2
- We'll restore the results from m1 which includes region even though the terms are not collectively significant
 - . estimates restore m1
- Now it's as though we just ran the model stored as m1

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Tests of Differences

- test can also be used to the equality of coefficients
 - . test 3.region = 4.region
- A likelihood ratio test can also be used; see help constraint for information on setting the necessary constraints
- The lincom command calculates linear combinations of coefficients, along with standard errors, hypothesis tests, and confidence intervals
- For example, to obtain the difference in coefficients
 - . lincom 3.region 4.region

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What are margins?

- Stata defines margins as "statistics calculated from predictions of a previously fit model at fixed values of some covariates and averaging or otherwise integrating over the remaining covariates."
 - Also known as counterfactuals, or when we fix a categorical variable, potential outcomes
- What sorts of predictions does margins work with?
 - Predicted means, probabilities, and counts
 - Derivatives
 - Elasticities
- We'll also see contrasts and pairwise comparisons of the above

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Average Predictions

- Let's start with margins in its most basic form
 - . margins
- What happened here?
 - The predicted probability of highbp=1 was calculated for each case, using each case's observed values of bmi, age, female, and region

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- 2. The average of those predictions was calculated and displayed
- Unless we tell it to do otherwise, margins works with the estimation sample



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Predictions at the Average

- An alternative is to calculate the predicted probability fixing all the covariates at some value, often the mean
 - . margins, atmeans
- What happened here?
 - 1. The mean of each independent variable was calculated
 - The predicted probability of highbp=1 was calculated using the means from step 1

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Predictions at Each Level of a Factor Variable

- Adding a factor variable specifies that the predictions be repeated at each level of the variable, for example
 - . margins region
- What happened here?
 - The predicted probability is calculated treating all cases as if region=1 and using each case's observed values of bmi, age, and female

- 2. The mean of the predictions from step 1 is calculated
- 3. Repeat steps 1 and 2 for each value of region

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Multiple Factor Variables

- We can obtain margins for multiple variables
 - . margins region female
- Or combinations of values, for example each combination of region and female
 - . margins region#female
- We can graph the resulting predictions using the marginsplot command

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Graphing Predicted Probabilities

- For example to graph the last set of margins
 - . marginsplot



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Predictions at Specified Values of Covariates

- The at() option is used to specify values at which margins should be calculated
- To obtain the average predicted probability setting age=40 specify
 - . margins, at(age=40)
- at() accepts number lists, so we can obtain predictions setting age to 20, 30, ..., 70

. margins, at(age=(20(10)70)) vsquish

• The vsquish option reduces the amount of vertical space the header for margins takes up

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Graphing Across Values of Continuous Variables

. marginsplot



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Specifying Values of Multiple Variables

- We can specify values of multiple variables using at()
- If we set values of all the independent variables in our model, we can ask very specific questions
- For example, what is the predicted probability of high blood pressure for an male who is age 40, with a bmi of 25 and living in the midwest (region=2)? What is the predicted probability if the person is female?
 - . margins female, at(age=40 bmi=25 region=2)
- We can use the contrast operator r. to compare the predicted probabilities for males and females

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- . margins r.female, at(age=40 bmi=25 region=2)
- We'll see more on contrasts below

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Specifying Ranges of Multiple Variables

- We can also specify ranges of values for multiple variables, for example multiple values of age and bmi
 - . margins, at(age=(20(10)70) bmi=(20(10)40))
- We can also combine the use of factor and continuous variables, for example
 - . margins female, at(age=(20(10)70)) vsquish

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More Plots

. marginsplot, legend(order(3 "Males" 4 "Females"))



• The standard errors are drawn before the lines for the predictions, so we want the legend to show the third and fourth plots

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More Predictions

- We can use at() with the generate() suboption to answer different sorts of questions
- For example, what would the averaged predicted probability be if everyone aged 5 years, while their values female and region remained the same?
- The generate(age+5) requests margins calculated at each observations value of age plus 5
 - margins, at(age=generate(age+5))
- We can specify at() multiple times, to obtain predictions under different scenarios

. margins, at(age=generate(age)) ///
 at(age=generate(age+5)) at(age=generate(age+10))

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Predictions Over Groups

- The over() option produces predictions averaging within groups defined by the factor variable, for example, female
 - . margins, over(female)
- What happened here?
 - 1. The predicted probability for each case is calculated, using the case's observed values on all variables
 - The average predicted probability is calculated using only cases where female=0

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3. Repeat step 2 using only cases where female=1

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Pairwise Comparisons of Predictions

- Earlier we obtained average predicted probabilities at each level of region using
 - . margins region
- For pairwise comparisons of these margins we can add the pwcompare option
 - . margins region, pwcompare
- Adding the groups option will allow us to see which levels are statistically distinguishable
 - . margins region, pwcompare(groups)
- The pwcompare() option can be used to specify other suboptions; see help margins pwcompare for more information

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Contrasts of Predictions

- The margins command allows *contrast operators* which are used to request comparisons of the margins
 - In this case the margins are predicted probabilities
- For example, to compare average predicted probabilities setting female=0 versus female=1 add the r. prefix
 - . margins r.female
- We can use the @ operator to contrast female at each level of region
 - . margins r.female@region
- This reports the differences in predicted probabilities when female=1 versus female=0 at each level of region

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Contrasts of Predictions (Continued)

- To perform contrasts at different values of a continuous variable use the at() option
 - . margins r.female, at(age=(20(10)70)) vsquish
- The output gives tests of the differences in predicted probabilities for female=1 versus female=0 at each of the specified values of age
 - The joint test is statistically significant
 - The differences get smaller in absolute value as age increases

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Predictions Postestimation

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Plotting Contrasts

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Contrast Operators

- A few common contrast operators are
 - r. differences from the base (a.k.a. reference) level
 - a. differences from the next (adjacent) level
 - ar. differences from the previous level (reverse adjacent)
 - ${\ensuremath{\,\circ\,}}$ g . differences from the balanced grand mean
 - gw. differences from the observeration-weighted grand mean
 - There are also operators for Helmert contrats and contrasts using orthogonal polynomials for balanced and unbalanced cases

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contrast suboptions

- So far we've obtained contrasts using *contrast operators*, but margins also allows a contrast() option
- The contrast() option is particularly useful for specifying options to contrast
- For example, to obtain contrasts for continuous variables the atcontrast() suboption is used
 - The effects suboption requests a table showing the contrasts along with confidence intervals and p-values
 - In atcontrast(a) the a contrast operator requests comparisons of adjacent categories
 - . margins, at(age=(20(10)70)) contrast(atcontrast(a) effects)

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Contrasts with generate()

- Earlier we used the generate() suboption to obtain predicted probabilities modifying the observed values
- Specifically, we obtained predicted probabilities using each case's observed value of age and each case's observed value +5 years
 - . margins, at(age=generate(age)) at(age=generate(age+5))
- Using the contrast option, we can compare the two



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Contrasts of Differences

- We can also request contrasts of contrasts by combining contrast operators
- For example, to compare the differences between males and females across levels of region use
 - . margins r.female#r.region

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Adjusting for Multiple Comparisons

- Use of contrast and pwcompare can result in a large number of hypothesis tests
- The mcompare() option can be used to adjust p-values and confidence intervals for multiple comparisons within factor variable terms
- The available methods are
 - noadjust
 - bonferroni
 - sidak
 - scheffe

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Using mcompare()

- To apply Bonferroni's adjustment to an earlier contrast
 - . margins r.female@region, mcompare(bonferroni)
- Specifying adjusted p-values with the pwcompare option . margins region, mcompare(sidak) pwcompare

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Marginal Effects

 $\bullet\,$ In a straightforward linear model, the marginal effect of a variable is the coefficient $b\,$

$$y = b_0 + b_1 x_1 + b_2 x_2 + e$$

- In more complex models, this is no longer true
 - models with interactions
 - models with polynomial terms
 - generalized linear models when the margin is not on the linear scale
- For example, in a logistic regression model, the marginal effect of covariates is not constant on the probability scale
- margins can be used to estimate the margins of the derivative of a response

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A Closer Look at Slopes

- Here is a graph of predicted probabilities across values of bmi
 margins, at(bmi=(12(5)62))
 - . marginsplot



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Average Marginal Effects

- The slope of bmi is not constant, but we might want to know what it is on average
- We can obtain the average marginal effect of bmi
 - . margins, dydx(bmi)
- What happened here?
 - 1. Calculate the derivative of the predicted probability with respect to bmi for each observaton

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- 2. Calculate the average of derivatives from step 1 $% \left({{{\left({{{\left({{{\left({{{c}}} \right)}} \right)}_{c}}} \right)}_{c}}} \right)$
- We can do the same for all variables in our model
 - . margins, dydx(*)

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Marginal Effects Over the Response Surface

- $\bullet\,$ It can also be informative to estimate the marginal effect of x at different values of x
- For example, we can obtain the derviative with respect to age at age=20, 30, ..., 70
 - . margins, dydx(age) at(age=(20(10)70)) vsquish
- Here we do something similar, setting female=0 and then female=1
 - . margins female, dydx(age) at(age=(20(10)70)) vsquish

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Plots of Marginal Effects

- We can, of course, plot these marginal effects, to see how they change with different values of female and age
 - . marginsplot



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margins with Other Estimation Commands

- margins works after most estimation commands
- The default prediction for margins is the same as the default prediction for predict after a given command
- See help *command* postestimation for information on postestimation commands and their defaults after a given command
- You can specify different predictions from margins using the predict() option

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Modeling Household Size

- For the next set of examples we will model the number of individuals in a household (houssiz) using a Poisson model
- Our model will include covariates age, age², region, rural, and a region by rural interaction
- We've been working with age and region but we'll take a look at the new variables
 - . codebook houssiz rural
- Now we can fit our model
 - . poisson houssiz i.region##i.rural age c.age#c.age

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margins after poisson

- predict's default after poisson is the predicted count
- To obtain the average predicted count, using the observed values of all covarites use
 - . margins
- As before, we can request predicted counts at specified values of factor variables
 - . margins region#rural
- And continuous variables
 - . margins, at(age=(20(10)70)) vsquish

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Plotting Predicted Counts

. marginsplot



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Other Margins

• After poisson, margins can be used to predict the following

- n number of events; the default
- ir incidence rate, exp(xb), n when the exposure variable = 1

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- pr(n) probability that y=n
- pr(a,b) probability that $a \le y \le b$
- xb the linear predcition
- Predicted probability that houssiz=1
 - . margins rural, predict(pr(1))
- \bullet Predicted probability that $3 \leq \texttt{houssiz} \leq 5$
 - . margins region#rural, predict(pr(3,5))

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Multiple Responses

- Starting in Stata 14, margins can compute margins for multiple responses at the same time
 - After, for example, ologit, mlogit, mvreg
- To demonstrate this, we'll model self-rated health in a different version of the NHANES dataset

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- . webuse nhanes2f
- . codebook health
- Our model is
 - . ologit health i.female age c.age#c.age

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Specifying the Response

- By default margins will produce the average predicted probability of each value of health
 - . margins
- To request a single outcome we can use predict(outcome(#))
 . margins, predict(outcome(2))
- For multiple responses from a single command, repeat the predict() option
 - . margins, predict(outcome(1)) predict(outcome(2))
- To obtain predictions across values of age
 - . margins, at(age=(20(10)70)) pr(out(1)) pr(out(2)) vsquish

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Postestimation

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Plots with Multiple Responses

. marginsplot, legend(order(3 "Poor" 4 "Fair"))



Conclusion

Conclusion

- We've seen how to obtain a variety of predictions and marginal effects after regression models
- We now know how to perform contrasts of predictions and marginal effects
- We've also seen how to graph these results