Title

mlogit — Multinomial (polytomous) logistic regression

Syntax Menu Description Options
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Also see

Syntax

```
\begin{tabular}{ll} $\underline{\tt mlogit}$ $depvar $ [indepvars ] [if ] [in] [weight] [, options ] \end{tabular}
```

options	Description
Model	
<u>nocons</u> tant	suppress constant term
<pre><u>b</u>aseoutcome(#)</pre>	value of <i>depvar</i> that will be the base outcome
\underline{c} onstraints($clist$) \underline{col} linear	apply specified linear constraints; <i>clist</i> has the form $\#[-\#][,\#[-\#]]$ keep collinear variables
SE/Robust	
vce(vcetype)	vcetype may be oim, robust, cluster clustvar, bootstrap, or jackknife
Reporting	
<u>l</u> evel(#)	set confidence level; default is level(95)
<u>rr</u> r	report relative-risk ratios
<u>nocnsr</u> eport	do not display constraints
display_options	control column formats, row spacing, line width, display of omitted variables and base and empty cells, and factor-variable labeling
Maximization	
maximize_options	control the maximization process; seldom used
<u>coefl</u> egend	display legend instead of statistics

indepvars may contain factor variables; see [U] 11.4.3 Factor variables.

indepvars may contain time-series operators; see [U] 11.4.4 Time-series varlists.

bootstrap, by, fp, jackknife, mfp, mi estimate, rolling, statsby, and svy are allowed; see [U] 11.1.10 Prefix commands.

vce(bootstrap) and vce(jackknife) are not allowed with the mi estimate prefix; see [MI] mi estimate.

Weights are not allowed with the bootstrap prefix; see [R] bootstrap.

vce() and weights are not allowed with the svy prefix; see [SVY] svy.

fweights, iweights, and pweights are allowed; see [U] 11.1.6 weight.

coeflegend does not appear in the dialog box.

See [U] 20 Estimation and postestimation commands for more capabilities of estimation commands.

Menu

Statistics > Categorical outcomes > Multinomial logistic regression

Description

mlogit fits maximum-likelihood multinomial logit models, also known as polytomous logistic regression. You can define constraints to perform constrained estimation. Some people refer to conditional logistic regression as multinomial logit. If you are one of them, see [R] clogit.

See [R] logistic for a list of related estimation commands.

Options

_____ Model

noconstant; see [R] estimation options.

baseoutcome(#) specifies the value of *depvar* to be treated as the base outcome. The default is to choose the most frequent outcome.

constraints(clist), collinear; see [R] estimation options.

SE/Robust

vce(vcetype) specifies the type of standard error reported, which includes types that are derived from asymptotic theory (oim), that are robust to some kinds of misspecification (robust), that allow for intragroup correlation (cluster clustvar), and that use bootstrap or jackknife methods (bootstrap, jackknife); see [R] vce_option.

If specifying vce(bootstrap) or vce(jackknife), you must also specify baseoutcome().

Reporting

level(#); see [R] estimation options.

rrr reports the estimated coefficients transformed to relative-risk ratios, that is, e^b rather than b; see Description of the model below for an explanation of this concept. Standard errors and confidence intervals are similarly transformed. This option affects how results are displayed, not how they are estimated. rrr may be specified at estimation or when replaying previously estimated results.

nocnsreport; see [R] estimation options.

display_options: noomitted, vsquish, noemptycells, baselevels, allbaselevels, nofvlabel, fvwrap(#), fvwrapon(style), cformat(%fmt), pformat(%fmt), sformat(%fmt), and nolstretch; see [R] estimation options.

Maximization

maximize_options: difficult, technique(algorithm_spec), iterate(#), [no]log, trace,
 gradient, showstep, hessian, showtolerance, tolerance(#), ltolerance(#),
 nrtolerance(#), nonrtolerance, and from(init_specs); see [R] maximize. These options are
 seldom used.

The following option is available with mlogit but is not shown in the dialog box:

coeflegend; see [R] estimation options.

Remarks and examples

Remarks are presented under the following headings:

Description of the model Fitting unconstrained models Fitting constrained models

mlogit fits maximum likelihood models with discrete dependent (left-hand-side) variables when the dependent variable takes on more than two outcomes and the outcomes have no natural ordering. If the dependent variable takes on only two outcomes, estimates are identical to those produced by logistic or logit; see [R] logistic or [R] logit. If the outcomes are ordered, see [R] ologit.

Description of the model

For an introduction to multinomial logit models, see Greene (2012, 763–766), Hosmer, Lemeshow, and Sturdivant (2013, 269–289), Long (1997, chap. 6), Long and Freese (2014, chap. 8), and Treiman (2009, 336–341). For a description emphasizing the difference in assumptions and data requirements for conditional and multinomial logit, see Davidson and MacKinnon (1993).

Consider the outcomes $1, 2, 3, \ldots, m$ recorded in y, and the explanatory variables X. Assume that there are m=3 outcomes: "buy an American car", "buy a Japanese car", and "buy a European car". The values of y are then said to be "unordered". Even though the outcomes are coded 1, 2, and 3, the numerical values are arbitrary because 1 < 2 < 3 does not imply that outcome 1 (buy American) is less than outcome 2 (buy Japanese) is less than outcome 3 (buy European). This unordered categorical property of y distinguishes the use of mlogit from regress (which is appropriate for a continuous dependent variable), from ologit (which is appropriate for ordered categorical data), and from logit (which is appropriate for two outcomes, which can be thought of as ordered).

In the multinomial logit model, you estimate a set of coefficients, $\beta^{(1)}$, $\beta^{(2)}$, and $\beta^{(3)}$, corresponding to each outcome:

$$\Pr(y=1) = \frac{e^{X\beta^{(1)}}}{e^{X\beta^{(1)}} + e^{X\beta^{(2)}} + e^{X\beta^{(3)}}}$$

$$\Pr(y=2) = \frac{e^{X\beta^{(2)}}}{e^{X\beta^{(1)}} + e^{X\beta^{(2)}} + e^{X\beta^{(3)}}}$$

$$\Pr(y=3) = \frac{e^{X\beta^{(1)}}}{e^{X\beta^{(1)}} + e^{X\beta^{(2)}} + e^{X\beta^{(3)}}}$$

The model, however, is unidentified in the sense that there is more than one solution to $\beta^{(1)}$, $\beta^{(2)}$, and $\beta^{(3)}$ that leads to the same probabilities for y=1, y=2, and y=3. To identify the model, you arbitrarily set one of $\beta^{(1)}$, $\beta^{(2)}$, or $\beta^{(3)}$ to 0—it does not matter which. That is, if you arbitrarily set $\beta^{(1)}=0$, the remaining coefficients $\beta^{(2)}$ and $\beta^{(3)}$ will measure the change relative to the y=1 group. If you instead set $\beta^{(2)}=0$, the remaining coefficients $\beta^{(1)}$ and $\beta^{(3)}$ will measure the change relative to the y=2 group. The coefficients will differ because they have different interpretations, but the predicted probabilities for y=1, 2, and 3 will still be the same. Thus either parameterization will be a solution to the same underlying model.

Setting $\beta^{(1)} = 0$, the equations become

$$\Pr(y=1) = \frac{1}{1 + e^{X\beta^{(2)}} + e^{X\beta^{(3)}}}$$

$$\Pr(y=2) = \frac{e^{X\beta^{(2)}}}{1 + e^{X\beta^{(3)}} + e^{X\beta^{(3)}}}$$

$$\Pr(y=3) = \frac{e^{X\beta^{(3)}}}{1 + e^{X\beta^{(2)}} + e^{X\beta^{(3)}}}$$

The relative probability of y = 2 to the base outcome is

$$\frac{\Pr(y=2)}{\Pr(y=1)} = e^{X\beta^{(2)}}$$

Let's call this ratio the relative risk, and let's further assume that X and $\beta_k^{(2)}$ are vectors equal to (x_1, x_2, \ldots, x_k) and $(\beta_1^{(2)}, \beta_2^{(2)}, \ldots, \beta_k^{(2)})'$, respectively. The ratio of the relative risk for a one-unit change in x_i is then

$$\frac{e^{\beta_1^{(2)}x_1 + \dots + \beta_i^{(2)}(x_i + 1) + \dots + \beta_k^{(2)}x_k}}{e^{\beta_1^{(2)}x_1 + \dots + \beta_i^{(2)}x_i + \dots + \beta_k^{(2)}x_k}} = e^{\beta_i^{(2)}}$$

Thus the exponentiated value of a coefficient is the relative-risk ratio for a one-unit change in the corresponding variable (risk is measured as the risk of the outcome relative to the base outcome).

Fitting unconstrained models

Example 1: A first example

We have data on the type of health insurance available to 616 psychologically depressed subjects in the United States (Tarlov et al. 1989; Wells et al. 1989). The insurance is categorized as either an indemnity plan (that is, regular fee-for-service insurance, which may have a deductible or coinsurance rate) or a prepaid plan (a fixed up-front payment allowing subsequent unlimited use as provided, for instance, by an HMO). The third possibility is that the subject has no insurance whatsoever. We wish to explore the demographic factors associated with each subject's insurance choice. One of the demographic factors in our data is the race of the participant, coded as white or nonwhite:

- . use http://www.stata-press.com/data/r13/sysdsn1 (Health insurance data)
- . tabulate insure nonwhite, chi2 col

Key
frequency column percentage

	nonw	hite	
insure	0	1	Total
Indemnity	251	43	294
	50.71	35.54	47.73
Prepaid	208	69	277
	42.02	57.02	44.97
Uninsure	36	9	45
	7.27	7.44	7.31
Total	495	121	616
	100.00	100.00	100.00

Pearson chi2(2) = 9.5599Pr = 0.008

Although insure appears to take on the values Indemnity, Prepaid, and Uninsure, it actually takes on the values 1, 2, and 3. The words appear because we have associated a value label with the numeric variable insure; see [U] 12.6.3 Value labels.

When we fit a multinomial logit model, we can tell mlogit which outcome to use as the base outcome, or we can let mlogit choose. To fit a model of insure on nonwhite, letting mlogit choose the base outcome, we type

. mlogit insure nonwhite

Iteration 0: $log\ likelihood = -556.59502$ Iteration 1: $log\ likelihood = -551.78935$ Iteration 2: $log\ likelihood = -551.78348$ Iteration 3: $log\ likelihood = -551.78348$

Multinomial logistic regression

LR chi2(2) 9.62 Prob > chi2 0.0081 Pseudo R2 0.0086

616

Number of obs

Log likelihood = -551.78348

	,					
insure	Coef.	Std. Err.	z	P> z	[95% Conf.	Interval]
Indemnity	(base outco	ome)				
Prepaid						
nonwhite	.6608212	.2157321	3.06	0.002	.2379942	1.083648
_cons	1879149	.0937644	-2.00	0.045	3716896	0041401
Uninsure						
nonwhite	.3779586	.407589	0.93	0.354	4209011	1.176818
_cons	-1.941934	.1782185	-10.90	0.000	-2.291236	-1.592632

mlogit chose the indemnity outcome as the base outcome and presented coefficients for the outcomes prepaid and uninsured. According to the model, the probability of prepaid for whites (nonwhite = 0) is

$$\Pr(\texttt{insure} = \texttt{Prepaid}) = \frac{e^{-.188}}{1 + e^{-.188} + e^{-1.942}} = 0.420$$

Similarly, for nonwhites, the probability of prepaid is

$$\Pr(\texttt{insure} = \texttt{Prepaid}) = \frac{e^{-.188 + .661}}{1 + e^{-.188 + .661} + e^{-1.942 + .378}} = 0.570$$

These results agree with the column percentages presented by tabulate because the mlogit model is fully saturated. That is, there are enough terms in the model to fully explain the column percentage in each cell. The model chi-squared and the tabulate chi-squared are in almost perfect agreement; both test that the column percentages of insure are the same for both values of nonwhite.

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Example 2: Specifying the base outcome

By specifying the baseoutcome() option, we can control which outcome of the dependent variable is treated as the base. Left to its own, mlogit chose to make outcome 1, indemnity, the base outcome. To make outcome 2, prepaid, the base, we would type

. mlogit insure nonwhite, base(2)

Iteration 0: log likelihood = -556.59502
Iteration 1: log likelihood = -551.78935
Iteration 2: log likelihood = -551.78348
Iteration 3: log likelihood = -551.78348

Multinomial logistic regression

Number of obs = 616 LR chi2(2) = 9.62 Prob > chi2 = 0.0081 Pseudo R2 = 0.0086

Log likelihood = -551.78348

insure	Coef.	Std. Err.	z	P> z	[95% Conf.	Interval]
Indemnity nonwhite _cons	6608212 .1879149	.2157321 .0937644	-3.06 2.00	0.002 0.045	-1.083648 .0041401	2379942 .3716896
Prepaid	(base outco	ome)				
Uninsure nonwhite _cons	2828627 -1.754019	.3977302 .1805145	-0.71 -9.72	0.477	-1.0624 -2.107821	.4966742 -1.400217

The baseoutcome() option requires that we specify the numeric value of the outcome, so we could not type base(Prepaid).

Although the coefficients now appear to be different, the summary statistics reported at the top are identical. With this parameterization, the probability of prepaid insurance for whites is

$$\Pr(\texttt{insure} = \texttt{Prepaid}) = \frac{1}{1 + e^{.188} + e^{-1.754}} = 0.420$$

This is the same answer we obtained previously.

Example 3: Displaying relative-risk ratios

By specifying rrr, which we can do at estimation time or when we redisplay results, we see the model in terms of relative-risk ratios:

	DDD	G + 3	F		Ds.II	F0E%	a c	Tt
Log likelihood	= -551.78348				Pseudo	R2	=	0.0086
					Prob >	chi2	=	0.0081
					LR chi	2(2)	=	9.62
Multinomial log	gistic regres:	sion			Number	of obs	3 =	616
. mlogit, rrr								

insure	RRR	Std. Err.	z	P> z	[95% Conf.	<pre>Interval]</pre>
Indemnity	540405					
nonwhite	.516427	.1114099	-3.06	0.002	.3383588	.7882073
_cons	1.206731	.1131483	2.00	0.045	1.004149	1.450183
Prepaid	(base outco	ome)				
Uninsure						
nonwhite	.7536233	.2997387	-0.71	0.477	.3456255	1.643247
_cons	. 1730769	.0312429	-9.72	0.000	.1215024	. 2465434

Looked at this way, the relative risk of choosing an indemnity over a prepaid plan is 0.516 for nonwhites relative to whites.

To illustrate, from the output and discussions of examples 1 and 2 we find that

$$\Pr\left(\texttt{insure} = \texttt{Indemnity} \mid \texttt{white}\right) = \frac{1}{1 + e^{-.188} + e^{-1.942}} = 0.507$$

and thus the relative risk of choosing indemnity over prepaid (for whites) is

$$\frac{\Pr\left(\texttt{insure} = \texttt{Indemnity} \mid \texttt{white}\right)}{\Pr\left(\texttt{insure} = \texttt{Prepaid} \mid \texttt{white}\right)} = \frac{0.507}{0.420} = 1.207$$

For nonwhites,

$$\Pr\left(\text{insure} = \text{Indemnity} \mid \text{not white}\right) = \frac{1}{1 + e^{-.188 + .661} + e^{-1.942 + .378}} = 0.355$$

and thus the relative risk of choosing indemnity over prepaid (for nonwhites) is

$$\frac{\Pr\left(\texttt{insure} = \texttt{Indemnity} \mid \texttt{not white}\right)}{\Pr\left(\texttt{insure} = \texttt{Prepaid} \mid \texttt{not white}\right)} = \frac{0.355}{0.570} = 0.623$$

The ratio of these two relative risks, hence the name "relative-risk ratio", is 0.623/1.207 = 0.516, as given in the output under the heading "RRR".

□ Technical note

In models where only two categories are considered, the mlogit model reduces to standard logit. Consequently the exponentiated regression coefficients, labeled as RRR within mlogit, are equal to the odds ratios as given when the or option is specified under logit; see [R] logit.

As such, always referring to mlogit's exponentiated coefficients as odds ratios may be tempting. However, the discussion in example 3 demonstrates that doing so would be incorrect. In general mlogit models, the exponentiated coefficients are ratios of relative risks, not ratios of odds.

Example 4: Model with continuous and multiple categorical variables

One of the advantages of mlogit over tabulate is that we can include continuous variables and multiple categorical variables in the model. In examining the data on insurance choice, we decide that we want to control for age, gender, and site of study (the study was conducted in three sites):

```
. mlogit insure age male nonwhite i.site
```

Iteration 0: log likelihood = -555.85446
Iteration 1: log likelihood = -534.67443
Iteration 2: log likelihood = -534.36284
Iteration 3: log likelihood = -534.36165
Iteration 4: log likelihood = -534.36165

 ${\tt Multinomial\ logistic\ regression}$

Number of obs = 615 LR chi2(10) = 42.99 Prob > chi2 = 0.0000 Pseudo R2 = 0.0387

Log likelihood = -534.36165

	T					
insure	Coef.	Std. Err.	z	P> z	[95% Conf.	Interval]
Indemnity	(base outco	ome)				
Prepaid						
age	011745	.0061946	-1.90	0.058	0238862	.0003962
male	.5616934	.2027465	2.77	0.006	.1643175	.9590693
nonwhite	.9747768	.2363213	4.12	0.000	.5115955	1.437958
site						
2	.1130359	.2101903	0.54	0.591	2989296	.5250013
3	5879879	.2279351	-2.58	0.010	-1.034733	1412433
_cons	.2697127	.3284422	0.82	0.412	3740222	.9134476
Uninsure						
age	0077961	.0114418	-0.68	0.496	0302217	.0146294
male	.4518496	.3674867	1.23	0.219	268411	1.17211
nonwhite	.2170589	.4256361	0.51	0.610	6171725	1.05129
site						
2	-1.211563	.4705127	-2.57	0.010	-2.133751	2893747
3	2078123	.3662926	-0.57	0.570	9257327	.510108
_cons	-1.286943	.5923219	-2.17	0.030	-2.447872	1260134

These results suggest that the inclination of nonwhites to choose prepaid care is even stronger than it was without controlling. We also see that subjects in site 2 are less likely to be uninsured.

Fitting constrained models

mlogit can fit models with subsets of coefficients constrained to be zero, with subsets of coefficients constrained to be equal both within and across equations, and with subsets of coefficients arbitrarily constrained to equal linear combinations of other estimated coefficients.

Before fitting a constrained model, you define the constraints with the constraint command; see [R] constraint. Once the constraints are defined, you estimate using mlogit, specifying the constraint() option. Typing constraint(4) would use the constraint you previously saved as 4. Typing constraint (1,4,6) would use the previously stored constraints 1, 4, and 6. Typing constraint (1-4,6) would use the previously stored constraints 1, 2, 3, 4, and 6.

Sometimes you will not be able to specify the constraints without knowing the omitted outcome. In such cases, assume that the omitted outcome is whatever outcome is convenient for you, and include the baseoutcome() option when you specify the mlogit command.

Example 5: Specifying constraints to test hypotheses

We can use constraints to test hypotheses, among other things. In our insurance-choice model, let's test the hypothesis that there is no distinction between having indemnity insurance and being uninsured. Indemnity-style insurance was the omitted outcome, so we type

```
. test [Uninsure]
(1)
      [Uninsure]age = 0
(2)
      [Uninsure]male = 0
(3)
     [Uninsure] nonwhite = 0
(4) [Uninsure]1b.site = 0
(5) [Uninsure] 2.site = 0
( 6) [Uninsure] 3. site = 0
      Constraint 4 dropped
          chi2(5) =
                         9.31
        Prob > chi2 =
                         0.0973
```

If indemnity had not been the omitted outcome, we would have typed test [Uninsure=Indemnity].

The results produced by test are an approximation based on the estimated covariance matrix of the coefficients. Because the probability of being uninsured is low, the log likelihood may be nonlinear for the uninsured. Conventional statistical wisdom is not to trust the asymptotic answer under these circumstances but to perform a likelihood-ratio test instead.

To use Stata's 1rtest (likelihood-ratio test) command, we must fit both the unconstrained and constrained models. The unconstrained model is the one we have previously fit. Following the instruction in [R] Irtest, we first store the unconstrained model results:

```
. estimates store unconstrained
```

To fit the constrained model, we must refit our model with all the coefficients except the constant set to 0 in the Uninsure equation. We define the constraint and then refit:

```
. constraint 1 [Uninsure]
```

. mlogit insure age male nonwhite i.site, constraints(1)

Iteration 0: $log\ likelihood = -555.85446$ Iteration 1: log likelihood = -539.80523Iteration 2: $log\ likelihood = -539.75644$ Iteration 3: log likelihood = -539.75643

Multinomial logistic regression Number of obs 615 Wald chi2(5) 29.70 Prob > chi2 0.0000

Log likelihood = -539.75643

- (1) [Uninsure]o.age = 0
- (2) [Uninsure]o.male = 0
- (3) [Uninsure]o.nonwhite = 0 (4) [Uninsure]2o.site = 0
- (5) [Uninsure]3o.site = 0

insure	Coef.	Std. Err.	z	P> z	[95% Conf.	Interval]
Indemnity	(base outc	ome)				
Prepaid						
age	0107025	.0060039	-1.78	0.075	0224699	.0010649
male	.4963616	. 1939683	2.56	0.010	.1161907	.8765324
nonwhite	.9421369	.2252094	4.18	0.000	.5007346	1.383539
site						
2	.2530912	.2029465	1.25	0.212	1446767	.6508591
3	5521773	.2187237	-2.52	0.012	9808678	1234869
_cons	.1792752	.3171372	0.57	0.572	4423023	.8008527
Uninsure						
age	0	(omitted)				
male	0	(omitted)				
nonwhite	0	(omitted)				
site						
2	0	(omitted)				
3	0	(omitted)				
_cons	-1.87351	.1601099	-11.70	0.000	-2.18732	-1.5597

We can now perform the likelihood-ratio test:

```
. lrtest unconstrained .
Likelihood-ratio test
                                                       LR chi2(5) =
                                                                         10.79
(Assumption: . nested in unconstrained)
                                                       Prob > chi2 =
                                                                        0.0557
```

The likelihood-ratio chi-squared is 10.79 with 5 degrees of freedom—just slightly greater than the magic p = 0.05 level—so we should not call this difference significant. 1

□ Technical note

In certain circumstances, you should fit a multinomial logit model with conditional logit; see [R] clogit. With substantial data manipulation, clogit can handle the same class of models with some interesting additions. For example, if we had available the price and deductible of the most competitive insurance plan of each type, mlogit could not use this information, but clogit could.

Stored results

mlogit stores the following in e():

```
Scalars
    e(N)
                               number of observations
                               number of completely determined observations
    e(N_cd)
                               number of outcomes
    e(k_out)
    e(k)
                               number of parameters
    e(k_eq)
                               number of equations in e(b)
    e(k_eq_model)
                               number of equations in overall model test
                               number of dependent variables
    e(k_dv)
                               model degrees of freedom
    e(df_m)
    e(r2_p)
                                pseudo-R-squared
    e(11)
                               log likelihood
    e(11_0)
                               log likelihood, constant-only model
                               number of clusters
    e(N_clust)
                               \chi^2
    e(chi2)
    e(p)
                               significance
    e(k_eq_base)
                               equation number of the base outcome
                               the value of depvar to be treated as the base outcome
    e(baseout)
                               index of the base outcome
    e(ibaseout)
                               rank of e(V)
    e(rank)
                               number of iterations
    e(ic)
    e(rc)
                               return code
    e(converged)
                                1 if converged, 0 otherwise
Macros
    e(cmd)
                               mlogit
    e(cmdline)
                               command as typed
                               name of dependent variable
    e(depvar)
    e(wtype)
                               weight type
    e(wexp)
                               weight expression
    e(title)
                               title in estimation output
    e(clustvar)
                               name of cluster variable
                               Wald or LR; type of model \chi^2 test
    e(chi2type)
    e(vce)
                               vcetype specified in vce()
                               title used to label Std. Err.
    e(vcetype)
                               names of equations
    e(eqnames)
    e(baselab)
                               value label corresponding to base outcome
    e(opt)
                               type of optimization
    e(which)
                               max or min; whether optimizer is to perform maximization or minimization
                               type of ml method
    e(ml_method)
                               name of likelihood-evaluator program
    e(user)
    e(technique)
                               maximization technique
    e(properties)
    e(predict)
                               program used to implement predict
                               predictions disallowed by margins
    e(marginsnotok)
    e(asbalanced)
                               factor variables fyset as asbalanced
    e(asobserved)
                                factor variables fyset as asobserved
```

```
Matrices
    e(b)
                                 coefficient vector
    e(out)
                                 outcome values
    e(Cns)
                                 constraints matrix
    e(ilog)
                                 iteration log (up to 20 iterations)
    e(gradient)
                                 gradient vector
                                 variance-covariance matrix of the estimators
    e(V)
    e(V_modelbased)
                                 model-based variance
Functions
    e(sample)
                                 marks estimation sample
```

Methods and formulas

The multinomial logit model is described in Greene (2012, 763–766).

Suppose that there are k categorical outcomes and—without loss of generality—let the base outcome be 1. The probability that the response for the jth observation is equal to the ith outcome is

$$p_{ij} = \Pr(y_j = i) = \begin{cases} \frac{1}{1 + \sum\limits_{m=2}^{k} \exp(\mathbf{x}_j \boldsymbol{\beta}_m)}, & \text{if} \quad i = 1\\ \frac{1}{1 + \sum\limits_{m=2}^{k} \exp(\mathbf{x}_j \boldsymbol{\beta}_i)}, & \text{if} \quad i > 1\\ \frac{1}{1 + \sum\limits_{m=2}^{k} \exp(\mathbf{x}_j \boldsymbol{\beta}_m)}, & \text{if} \quad i > 1 \end{cases}$$

where \mathbf{x}_j is the row vector of observed values of the independent variables for the jth observation and β_m is the coefficient vector for outcome m. The log pseudolikelihood is

$$\ln L = \sum_{i} w_{j} \sum_{i=1}^{k} I_{i}(y_{j}) \ln p_{ik}$$

where w_i is an optional weight and

$$I_i(y_j) = \begin{cases} 1, & \text{if } y_j = i \\ 0, & \text{otherwise} \end{cases}$$

Newton-Raphson maximum likelihood is used; see [R] maximize.

For constrained equations, the set of constraints is orthogonalized, and a subset of maximizable parameters is selected. For example, a parameter that is constrained to zero is not a maximizable parameter. If two parameters are constrained to be equal to each other, only one is a maximizable parameter.

Let \mathbf{r} be the vector of maximizable parameters. \mathbf{r} is physically a subset of the solution parameters, \mathbf{b} . A matrix, \mathbf{T} , and a vector, \mathbf{m} , are defined as

$$b = Tr + m$$

so that

$$\frac{\partial f}{\partial \mathbf{b}} = \frac{\partial f}{\partial \mathbf{r}} \mathbf{T}'$$
$$\frac{\partial^2 f}{\partial \mathbf{b}^2} = \mathbf{T} \frac{\partial^2 f}{\partial \mathbf{r}^2} \mathbf{T}'$$

T consists of a block form in which one part is a permutation of the identity matrix and the other part describes how to calculate the constrained parameters from the maximizable parameters.

This command supports the Huber/White/sandwich estimator of the variance and its clustered version using vce(robust) and vce(cluster *clustvar*), respectively. See [P] _robust, particularly Maximum likelihood estimators and Methods and formulas.

mlogit also supports estimation with survey data. For details on VCEs with survey data, see [SVY] variance estimation.

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Also see

- [R] mlogit postestimation Postestimation tools for mlogit
- [R] **clogit** Conditional (fixed-effects) logistic regression
- [R] logistic Logistic regression, reporting odds ratios
- [R] **logit** Logistic regression, reporting coefficients
- [R] **mprobit** Multinomial probit regression
- [R] **nlogit** Nested logit regression
- [R] **ologit** Ordered logistic regression
- [R] rologit Rank-ordered logistic regression
- [R] **slogit** Stereotype logistic regression
- [MI] estimation Estimation commands for use with mi estimate
- [SVY] svy estimation Estimation commands for survey data
- [U] 20 Estimation and postestimation commands

Title

mlogit postestimation — Postestimation tools for mlo

Description	Syntax for predict	Menu for predict	Options for predict
Remarks and examples	Reference	Also see	

Description

The following postestimation commands are available after mlogit:

Command	Description
contrast	contrasts and ANOVA-style joint tests of estimates
estat ic	Akaike's and Schwarz's Bayesian information criteria (AIC and BIC)
estat summarize	summary statistics for the estimation sample
estat vce	variance-covariance matrix of the estimators (VCE)
estat (svy)	postestimation statistics for survey data
estimates	cataloging estimation results
${ t forecast}^1$	dynamic forecasts and simulations
hausman	Hausman's specification test
lincom	point estimates, standard errors, testing, and inference for linear combinations of coefficients
${\sf lrtest}^2$	likelihood-ratio test
margins	marginal means, predictive margins, marginal effects, and average marginal effects
marginsplot	graph the results from margins (profile plots, interaction plots, etc.)
nlcom	point estimates, standard errors, testing, and inference for nonlinear combinations of coefficients
predict	predictions, residuals, influence statistics, and other diagnostic measures
predictnl	point estimates, standard errors, testing, and inference for generalized predictions
pwcompare	pairwise comparisons of estimates
suest	seemingly unrelated estimation
test	Wald tests of simple and composite linear hypotheses
testnl	Wald tests of nonlinear hypotheses

 $^{^{1}\,}$ forecast is not appropriate with mi or svy estimation results.

 $^{^{2}}$ lrtest is not appropriate with svy estimation results.

Syntax for predict

```
predict [type] { stub* | newvar | newvarlist } [if] [in] [, statistic outcome(outcome)]

predict [type] { stub* | newvarlist } [if] [in], scores

statistic Description

Main

pr probability of a positive outcome; the default linear prediction
```

If you do not specify outcome(), pr (with one new variable specified), xb, and stdp assume outcome(#1). You must specify outcome() with the stddp option.

standard error of the difference in two linear predictions

You specify one or k new variables with pr, where k is the number of outcomes.

standard error of the linear prediction

You specify one new variable with xb, stdp, and stddp.

These statistics are available both in and out of sample; type predict ... if e(sample) ... if wanted only for the estimation sample.

Menu for predict

stdp

stddp

Statistics > Postestimation > Predictions, residuals, etc.

Options for predict

Main

pr, the default, calculates the probability of each of the categories of the dependent variable or the probability of the level specified in outcome(outcome). If you specify the outcome(outcome) option, you need to specify only one new variable; otherwise, you must specify a new variable for each category of the dependent variable.

xb calculates the linear prediction. You must also specify the outcome(outcome) option.

stdp calculates the standard error of the linear prediction. You must also specify the outcome(outcome) option.

stddp calculates the standard error of the difference in two linear predictions. You must specify the outcome(outcome) option, and here you specify the two particular outcomes of interest inside the parentheses, for example, predict sed, stddp outcome(1,3).

outcome(outcome) specifies the outcome for which the statistic is to be calculated. equation() is a synonym for outcome(): it does not matter which you use. outcome() or equation() can be specified using

#1, #2, ..., where #1 means the first category of the dependent variable, #2 means the second category, etc.;

the values of the dependent variable; or

the value labels of the dependent variable if they exist.

scores calculates equation-level score variables. The number of score variables created will be one less than the number of outcomes in the model. If the number of outcomes in the model were k, then

```
the first new variable will contain \partial \ln L/\partial(\mathbf{x}_i\boldsymbol{\beta}_1);
the second new variable will contain \partial \ln L/\partial(\mathbf{x}_i\beta_2);
the (k-1)th new variable will contain \partial \ln L/\partial (\mathbf{x}_i \boldsymbol{\beta}_{k-1}).
```

Remarks and examples

Remarks are presented under the following headings:

Obtaining predicted values Calculating marginal effects Testing hypotheses about coefficients

Obtaining predicted values

Example 1: Obtaining predicted probabilities

After estimation, we can use predict to obtain predicted probabilities, index values, and standard errors of the index, or differences in the index. For instance, in example 4 of [R] mlogit, we fit a model of insurance choice on various characteristics. We can obtain the predicted probabilities for outcome 1 by typing

```
. use http://www.stata-press.com/data/r13/sysdsn1
(Health insurance data)
. mlogit insure age i.male i.nonwhite i.site
 (output omitted)
. predict p1 if e(sample), outcome(1)
(option pr assumed; predicted probability)
(29 missing values generated)
. summarize p1
    Variable
                     Obs
                                 Mean
                                         Std. Dev.
                                                          Min
                     615
                             .4764228
                                         .1032279
                                                     .1698142
```

We added the i. prefix to the male, nonwhite, and site variables to explicitly identify them as factor variables. That makes no difference in the estimated results, but we will take advantage of it in later examples. We also included if e(sample) to restrict the calculation to the estimation sample. In example 4 of [R] mlogit, the multinomial logit model was fit on 615 observations, so there must be missing values in our dataset.

Max

.71939

Although we typed outcome(1), specifying 1 for the indemnity outcome, we could have typed outcome (Indemnity). For instance, to obtain the probabilities for prepaid, we could type

```
. predict p2 if e(sample), outcome(Prepaid)
(option pr assumed; predicted probability)
(29 missing values generated)
```

. summarize p2

Max	Min	Std. Dev.	Mean	Obs	Variable
.7885724	.1964103	.1125962	.4504065	615	p2

We must specify the label exactly as it appears in the underlying value label (or how it appears in the mlogit output), including capitalization.

Here we have used predict to obtain probabilities for the same sample on which we estimated. That is not necessary. We could use another dataset that had the independent variables defined (in our example, age, male, nonwhite, and site) and use predict to obtain predicted probabilities; here, we would not specify if e(sample).

4

Example 2: Obtaining index values

predict can also be used to obtain the index values—the $\sum x_i \widehat{eta}_i^{(k)}$ —as well as the probabilities:

- . predict idx1, outcome(Indemnity) xb
 (1 missing value generated)
- . summarize idx1

Variable	Obs	Mean	Std. Dev.	Min	Max
idx1	643	0	0	0	0

The indemnity outcome was our base outcome—the outcome for which all the coefficients were set to 0—so the index is always 0. For the prepaid and uninsured outcomes, we type

- . predict idx2, outcome(Prepaid) xb
 (1 missing value generated)
 . predict idx3, outcome(Uninsure) xb
- (1 missing value generated)
- . summarize idx2 idx3

Variable	Obs	Mean	Std. Dev.	Min	Max
idx2 idx3	643 643	0566113 -1.980747		-1.298198 -3.112741	

We can obtain the standard error of the index by specifying the stdp option:

- . predict se2, outcome(Prepaid) stdp
 (1 missing value generated)
- . list p2 idx2 se2 in 1/5

	p2	idx2	se2
1. 2. 3. 4.	.3709022 .4977667 .4113073 .5424927	4831167 .055111 1712106 .3788345 0925817	.2437772 .1694686 .1793498 .2513701 .1452616

We obtained the probability, p2, in the previous example.

Finally, predict can calculate the standard error of the difference in the index values between two outcomes with the stddp option:

```
. predict se_2_3, outcome(Prepaid,Uninsure) stddp
(1 missing value generated)
```

. list $idx2 idx3 se_2_3 in 1/5$

	idx2	idx3	se_2_3
1.	4831167	-3.073253	.5469354
2.	.055111	-2.715986	.4331918
3.	1712106	-1.579621	.3053815
4.	.3788345	-1.462007	.4492552
5.	0925817	-2.814022	.4024784

In the first observation, the difference in the indexes is -0.483 - (-3.073) = 2.59. The standard error of that difference is 0.547. 4

Example 3: Interpreting results using predictive margins

It is more difficult to interpret the results from mlogit than those from clogit or logit because there are multiple equations. For example, suppose that one of the independent variables in our model takes on the values 0 and 1, and we are attempting to understand the effect of this variable. Assume that the coefficient on this variable for the second outcome, $\beta^{(2)}$, is positive. We might then be tempted to reason that the probability of the second outcome is higher if the variable is 1 rather than 0. Most of the time, that will be true, but occasionally we will be surprised. The probability of some other outcome could increase even more (say, $\beta^{(3)} > \beta^{(2)}$), and thus the probability of outcome 2 would actually fall relative to that outcome. We can use predict to help interpret such results.

Continuing with our previously fit insurance-choice model, we wish to describe the model's predictions by race. For this purpose, we can use the method of predictive margins (also known as recycled predictions), in which we vary characteristics of interest across the whole dataset and average the predictions. That is, we have data on both whites and nonwhites, and our individuals have other characteristics as well. We will first pretend that all the people in our data are white but hold their other characteristics constant. We then calculate the probabilities of each outcome. Next we will pretend that all the people in our data are nonwhite, still holding their other characteristics constant. Again we calculate the probabilities of each outcome. The difference in those two sets of calculated probabilities, then, is the difference due to race, holding other characteristics constant.

```
. gen byte nonwhold=nonwhite
                                          // save real race
. replace nonwhite=0
                                          // make everyone white
(126 real changes made)
. predict wpind, outcome(Indemnity)
                                          // predict probabilities
(option pr assumed; predicted probability)
(1 missing value generated)
. predict wpp, outcome(Prepaid)
(option pr assumed; predicted probability)
(1 missing value generated)
. predict wpnoi, outcome(Uninsure)
(option pr assumed; predicted probability)
(1 missing value generated)
. replace nonwhite=1
                                          // make everyone nonwhite
(644 real changes made)
```

643

```
. predict nwpind, outcome(Indemnity)
(option pr assumed; predicted probability)
(1 missing value generated)
. predict nwpp, outcome(Prepaid)
(option pr assumed; predicted probability)
(1 missing value generated)
. predict nwpnoi, outcome(Uninsure)
(option pr assumed; predicted probability)
(1 missing value generated)
 replace nonwhite=nonwhold
                                            // restore real race
(518 real changes made)
. summarize wp* nwp*, sep(3)
    Variable
                      Obs
                                 Mean
                                          Std. Dev.
                                                           Min
                                                                      Max
       wpind
                      643
                             .5141673
                                          .0872679
                                                      .3092903
                                                                   .71939
                      643
                              .4082052
                                          .0993286
                                                      .1964103
                                                                  .6502247
         qqw
                      643
                             .0776275
                                          .0360283
                                                      .0273596
                                                                 .1302816
       wpnoi
      nwpind
                      643
                             .3112809
                                          .0817693
                                                      .1511329
                                                                  .535021
                              .630078
                                          .0979976
                      643
                                                      .3871782
                                                                  .8278881
        nwpp
```

.0586411

In example 1 of [R] **mlogit**, we presented a cross-tabulation of insurance type and race. Those values were unadjusted. The means reported above are the values adjusted for age, sex, and site. Combining the results gives

.0287185

.0209648

.0933874

	Una	djusted	Ad	justed
	white	nonwhite	white	nonwhite
Indemnity	0.51	0.36	0.51	0.31
Prepaid	0.42	0.57	0.41	0.63
Uninsured	0.07	0.07	0.08	0.06

We find, for instance, after adjusting for age, sex, and site, that although 57% of nonwhites in our data had prepaid plans, 63% of nonwhites chose prepaid plans.

Computing predictive margins by hand was instructive, but we can compute these values more easily using the margins command (see [R] margins). The two margins for the indemnity outcome can be estimated by typing

. margins nonwhite, predict(outcome(Indemnity)) noesample

Predictive margins Number of obs = 643

Model VCE : OIM

nwpnoi

Expression : Pr(insure==Indemnity), predict(outcome(Indemnity))

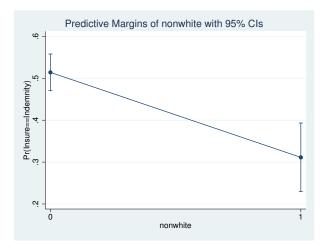
	Margin	Delta-method Std. Err.	z	P> z	[95% Conf.	Interval]
nonwhite						
0	.5141673	.0223485	23.01	0.000	.470365	.5579695
1	.3112809	.0418049	7.45	0.000	.2293448	.393217

margins also estimates the standard errors and confidence intervals of the margins. By default, margins uses only the estimation sample. We added the noesample option so that margins would use the entire sample and produce results comparable to our earlier analysis.

4

We can use marginsplot to graph the results from margins:

. marginsplot Variables that uniquely identify margins: nonwhite



The margins for the other two outcomes can be computed by typing

- . margins nonwhite, predict(outcome(Prepaid)) noesample (output omitted)
- . margins nonwhite, predict(outcome(Uninsure)) noesample (output omitted)

□ Technical note

You can use predict to classify predicted values and compare them with the observed outcomes to interpret a multinomial logit model. This is a variation on the notions of sensitivity and specificity for logistic regression. Here we will classify indemnity and prepaid as definitely predicting indemnity, definitely predicting prepaid, and ambiguous.

```
. predict indem, outcome(Indemnity) index
                                                         // obtain indexes
(1 missing value generated)
. predict prepaid, outcome(Prepaid) index
(1 missing value generated)
. gen diff = prepaid-indem
                                                         // obtain difference
(1 missing value generated)
. predict sediff, outcome(Indemnity, Prepaid) stddp
                                                         // & its standard error
(1 missing value generated)
. gen type = 1 if diff/sediff < -1.96
                                                         // definitely indemnity
(504 missing values generated)
. replace type = 3 if diff/sediff > 1.96
                                                         // definitely prepaid
(100 real changes made)
. replace type = 2 if type>=. & diff/sediff < .</pre>
                                                        // ambiguous
(404 real changes made)
. label def type 1 "Def Ind" 2 "Ambiguous" 3 "Def Prep"
. label values type type
                                                         // label results
```

. tabulate insure type

		type		
insure	Def Ind	Ambiguous	Def Prep	Total
Indemnity	78	183	33	294
Prepaid	44	177	56	277
Uninsure	12	28	5	45
Total	134	388	94	616

We can see that the predictive power of this model is modest. There are many misclassifications in both directions, though there are more correctly classified observations than misclassified observations.

Also the uninsured look overwhelmingly as though they might have come from the indemnity system rather than from the prepaid system.

Calculating marginal effects

Example 4

We have already noted that the coefficients from multinomial logit can be difficult to interpret because they are relative to the base outcome. Another way to evaluate the effect of covariates is to examine the marginal effect of changing their values on the probability of observing an outcome.

The margins command can be used for this too. We can estimate the marginal effect of each covariate on the probability of observing the first outcome—indemnity insurance—by typing

. margins, dydx(*) predict(outcome(Indemnity))

Average marginal effects Number of obs = 615

Model VCE : OIM

Expression : Pr(insure==Indemnity), predict(outcome(Indemnity))

dy/dx w.r.t. : age 1.male 1.nonwhite 2.site 3.site

	dy/dx	Delta-method Std. Err.	z	P> z	[95% Conf.	Interval]
age 1.male 1.nonwhite	.0026655 1295734 2032404	.001399 .0450945 .0482554	1.91 -2.87 -4.21	0.057 0.004 0.000	0000765 2179571 2978192	.0054074 0411898 1086616
site 2 3	.0070995 .1216165	.0479993 .0505833	0.15 2.40	0.882 0.016	0869775 .022475	.1011765 .220758

Note: dy/dx for factor levels is the discrete change from the base level.

By default, margins estimates the average marginal effect over the estimation sample, and that is what we see above. Being male decreases the average probability of having indemnity insurance by 0.130. We also see, from the note at the bottom of the table, that the marginal effect was computed as a discrete change in the probability of being male rather than female. That is why we made male a factor variable when fitting the model.

The dydx(*) option requested that margins estimate the marginal effect for each regressor, dydx(age) would have produced estimates only for the effect of age. margins has many options for controlling how the marginal effect is computed, including the ability to average over subgroups or to compute estimates for specified values of the regressors; see [R] margins.

We could evaluate the marginal effects on the other two outcomes by typing

```
. margins, dydx(*) predict(outcome(Prepaid))
 (output omitted)
. margins, dydx(*) predict(outcome(Uninsure))
 (output omitted)
```

4

Testing hypotheses about coefficients

Example 5

test tests hypotheses about the coefficients just as after any estimation command; see [R] test. Note, however, test's syntax for dealing with multiple-equation models. Because test bases its results on the estimated covariance matrix, we might prefer a likelihood-ratio test; see example 5 in [R] mlogit for an example of lrtest.

If we simply list variables after the test command, we are testing that the corresponding coefficients are zero across all equations:

```
. test 2.site 3.site
(1) [Indemnity]20.site = 0
(2) [Prepaid] 2.site = 0
( 3) [Uninsure] 2.site = 0
(4) [Indemnity]3o.site = 0
(5) [Prepaid]3.site = 0
( 6) [Uninsure]3.site = 0
      Constraint 1 dropped
      Constraint 4 dropped
          chi2(4) =
                        19.74
        Prob > chi2 =
                         0.0006
```

We can test that all the coefficients (except the constant) in an equation are zero by simply typing the outcome in square brackets:

```
. test [Uninsure]
(1)
      [Uninsure]age = 0
(2)
     [Uninsure]Ob.male = 0
(3) [Uninsure]1.male = 0
(4) [Uninsure]Ob.nonwhite = 0
(5) [Uninsure]1.nonwhite = 0
(6) [Uninsure]1b.site = 0
(7)
     [Uninsure]2.site = 0
(8)
     [Uninsure]3.site = 0
      Constraint 2 dropped
      Constraint 4 dropped
      Constraint 6 dropped
          chi2(5) =
                         9.31
                         0.0973
        Prob > chi2 =
```

We specify the outcome just as we do with predict; we can specify the label if the outcome variable is labeled, or we can specify the numeric value of the outcome. We would have obtained the same test as above if we had typed test [3] because 3 is the value of insure for the outcome uninsured.

We can combine the two syntaxes. To test that the coefficients on the site variables are 0 in the equation corresponding to the outcome prepaid, we can type

```
. test [Prepaid]: 2.site 3.site
 (1)
       [Prepaid] 2.site = 0
 (2)
       [Prepaid]3.site = 0
          chi2(2) =
                        10.78
        Prob > chi2 =
                       0.0046
```

We specified the outcome and then followed that with a colon and the variables we wanted to test.

We can also test that coefficients are equal across equations. To test that all coefficients except the constant are equal for the prepaid and uninsured outcomes, we can type

```
. test [Prepaid=Uninsure]
       [Prepaid]age - [Uninsure]age = 0
(2)
       [Prepaid] Ob.male - [Uninsure] Ob.male = 0
(3)
      [Prepaid]1.male - [Uninsure]1.male = 0
(4)
       [Prepaid] Ob.nonwhite - [Uninsure] Ob.nonwhite = 0
(5)
      [Prepaid] 1. nonwhite - [Uninsure] 1. nonwhite = 0
(6)
      [Prepaid]1b.site - [Uninsure]1b.site = 0
      [Prepaid] 2.site - [Uninsure] 2.site = 0
(7)
(8)
      [Prepaid]3.site - [Uninsure]3.site = 0
      Constraint 2 dropped
      Constraint 4 dropped
      Constraint 6 dropped
          chi2(5) =
                       13.80
        Prob > chi2 =
                       0.0169
```

To test that only the site variables are equal, we can type

```
. test [Prepaid=Uninsure]: 2.site 3.site
 (1)
       [Prepaid] 2.site - [Uninsure] 2.site = 0
 (2)
       [Prepaid]3.site - [Uninsure]3.site = 0
          chi2(2) =
                        12.68
        Prob > chi2 =
                         0.0018
```

Finally, we can test any arbitrary constraint by simply entering the equation and specifying the coefficients as described in [U] 13.5 Accessing coefficients and standard errors. The following hypothesis is senseless but illustrates the point:

```
. test ([Prepaid]age+[Uninsure]2.site)/2 = 2-[Uninsure]1.nonwhite
      .5*[Prepaid]age + [Uninsure]1.nonwhite + .5*[Uninsure]2.site = 2
(1)
          chi2( 1) =
                        22.45
        Prob > chi2 =
                         0.0000
```

See [R] test for more information about test. The information there about combining hypotheses across test commands (the accumulate option) also applies after mlogit.

1

Reference

Fagerland, M. W., and D. W. Hosmer, Jr. 2012. A generalized HosmerLemeshow goodness-of-fit test for multinomial logistic regression models. Stata Journal 12: 447-453.

Also see

- [R] **mlogit** Multinomial (polytomous) logistic regression
- [U] 20 Estimation and postestimation commands