Description

_xtologit_ fits random-effects ordered logistic models. The actual values taken on by the dependent variable are irrelevant, although larger values are assumed to correspond to “higher” outcomes. The conditional distribution of the dependent variable given the random effects is assumed to be multinomial with success probability determined by the logistic cumulative distribution function.

Quick start

Random-effects ordered logistic model of y as a function of x using _xtset_ data
   xtologit y x

Add indicators for levels of categorical variable a
   xtologit y x i.a

As above, but report odds ratios
   xtologit y x i.a, or

With cluster–robust standard errors for panels nested within cvar
   xtologit y x i.a, vce(cluster cvar)

Menu

Statistics > Longitudinal/panel data > Ordinal outcomes > Logistic regression (RE)
Syntax

\[ \text{xtologit} \ \text{depvar} \ [ \text{indepvars} ] \ [ \text{if} ] \ [ \text{in} ] \ [ \text{weight} ] \ [ , \ \text{options} ] \]

**options**

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A panel variable must be specified; see [XT] xtset.

\text{indepvars} may contain factor variables; see [U] 11.4.3 Factor variables.

\text{depvar} and \text{indepvars} may contain time-series operators; see [U] 11.4.4 Time-series varlists.

by, \text{fp}, and \text{statsby} are allowed; see [U] 11.1.10 Prefix commands.

\text{fweight}, \text{iweight}, and \text{pweight} are allowed; see [U] 11.1.6 weight.

\text{startgrid()}, \text{nodisplay}, and \text{coeflegend} do not appear in the dialog box.

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

### Model

offset(varname), constraints(constraints); 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 [XT] vce_options.

Specifying vce(robust) is equivalent to specifying vce(cluster panelvar); see xtologit and the robust VCE estimator in Methods and formulas.

### Reporting

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

or reports the estimated coefficients transformed to odds ratios, that is, \( e^b \) rather than \( b \). Standard errors and confidence intervals are similarly transformed. This option affects how results are displayed, not how they are estimated. or may be specified at estimation or when replaying previously estimated results.

lrmodel, nocnsreport; see [R] Estimation options.

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

### Integration

intmethod(intmethod), intpoints(#); see [R] Estimation options.

### Maximization

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

The following options are available with xtologit but are not shown in the dialog box:

startgrid(numlist) performs a grid search to improve the starting value of the random-intercept parameter. No grid search is performed by default unless the starting value is found to not be feasible; in this case, xtologit runs startgrid(0.1 1 10) and chooses the value that works best. You may already be using a default form of startgrid() without knowing it. If you see xtologit displaying Grid node 1, Grid node 2, ... following Grid node 0 in the iteration log, that is xtologit doing a default search because the original starting value was not feasible.

nodisplay is for programmers. It suppresses the display of the header and the coefficients.

collinear, coeflegend; see [R] Estimation options.
xtologit — Random-effects ordered logistic models

Remarks and examples

Remarks are presented under the following headings:

Overview
Video example

Overview

xtologit fits random-effects ordered logistic models. Ordered logistic models are used to estimate relationships between an ordinal dependent variable and a set of independent variables. An ordinal variable is a variable that is categorical and ordered, for instance, “poor”, “good”, and “excellent”, which might indicate a person’s current health status or the repair record of a car. If there are only two outcomes, see [XT] xtlogit, [XT] xtprobit, and [XT] xtctloglog. This entry is concerned only with more than two outcomes.

Example 1

We use the data from the “Television, School, and Family Smoking Prevention and Cessation Project” (Flay et al. 1988; Rabe-Hesketh and Skrondal 2012, chap. 11), where schools were randomly assigned into one of four groups defined by two treatment variables. Students within each school are nested in classes, and classes are nested in schools. In this example, we ignore the variability of classes within schools; see example 2 of [ME] meologit for a model that incorporates the additional class-level variance component. The dependent variable is the tobacco and health knowledge score (thk) collapsed into four ordered categories. We regress the outcome on the treatment variables and their interaction and control for the pretreatment score.
. use https://www.stata-press.com/data/r16/tvsfpors
. xtset school
    panel variable: school (unbalanced)
. xtologit thk prethk cc##tv

Fitting comparison model:
Iteration 0:  log likelihood =  -2212.775
Iteration 1:  log likelihood =  -2125.509
Iteration 2:  log likelihood =  -2125.103
Iteration 3:  log likelihood =  -2125.103

Refining starting values:
Grid node 0:  log likelihood =  -2136.243

Fitting full model:
Iteration 0:  log likelihood =  -2136.243  (not concave)
Iteration 1:  log likelihood =  -2120.258
Iteration 2:  log likelihood =  -2119.757
Iteration 3:  log likelihood =  -2119.743
Iteration 4:  log likelihood =  -2119.743

Random-effects ordered logistic regression  Number of obs =  1,600
Group variable: school Number of groups =  28
Random effects u_i ~ Gaussian Obs per group:
            min =  18
            avg =  57.1
            max =  137
Integration method: mvaghermite Integration pts. =  12
Wald chi2(4) =  128.06
Log likelihood =  -2119.743  Prob > chi2 =  0.0000

|          | Coef.  | Std. Err. |    z  | P>|z|   | [95% Conf. Interval] |
|----------|--------|-----------|------|------|---------------------|
| thk      |        |           |      |      |                     |
| prethk   | .4032892 | .03886    | 10.38| 0.000| .327125 .4794534   |
| 1.cc     | .9237904 | .204074   | 4.53 | 0.000| .5238127 1.323768  |
| 1.tv     | .2749937 | .1977424  | 1.39 | 0.164| -.1125744 .6625618 |

| cc##tv   |        |           |      |      |                     |
| 1 1      | -.4659256 | .2845963  | -1.64| 0.102| -1.023724 .0918728 |

|        |        |           |      |      |                     |
| /cut1  | -.0884493 | .1641062  | -.410916| 0.233193 |
| /cut2  | 1.153364  | .165616   | .8287625| 1.477965  |
| /cut3  | 2.33195   | .1734199  | 1.992053| 2.671846  |

| /sigma2_u| .0735112 | .0383106 | .0264695| .2041551 |

LR test vs. ologit model:  chibar2(01) = 10.72  Prob >= chibar2 = 0.0005

The estimation table reports the parameter estimates, the estimated cutpoints (κ₁, κ₂, κ₃), and the estimated panel-level variance component labeled sigma2_u. The parameter estimates can be interpreted just as the output from a standard ordered logistic regression would be interpreted; see [R] ologit. For example, we find that students with higher preintervention scores tend to have higher postintervention scores.

Underneath the parameter estimates and the cutpoints, the table shows the estimated variance component. The estimate of σ_u² is 0.074 with standard error 0.038. The reported likelihood-ratio test shows that there is enough variability between schools to favor a random-effects ordered logistic regression over a standard ordered logistic regression.
Technical note

The random-effects model is calculated using quadrature, which is an approximation whose accuracy depends partially on the number of integration points used. We can use the `quadchk` command to see if changing the number of integration points affects the results. If the results change, the quadrature approximation is not accurate given the number of integration points. Try increasing the number of integration points using the `intpoints()` option and run `quadchk` again. Do not attempt to interpret the results of estimates when the coefficients reported by `quadchk` differ substantially. See `[XT] quadchk` for details and `[XT] xtprobit` for an example.

Because the `xtologit` likelihood function is calculated by Gauss–Hermite quadrature, on large problems the computations can be slow. Computation time is roughly proportional to the number of points used for the quadrature.

Video example

Ordered logistic and probit for panel data

Stored results

`xtologit` stores the following in `e()`:

Scalars

- `e(N)` number of observations
- `e(N_g)` number of groups
- `e(k)` number of parameters
- `e(k_aux)` number of auxiliary parameters
- `e(k_eq)` number of equations in `e(b)`
- `e(k_eq_model)` number of equations in overall model test
- `e(k_dv)` number of dependent variables
- `e(k_cat)` number of categories
- `e(df_m)` model degrees of freedom
- `e(ll)` log likelihood
- `e(ll_0)` log likelihood, constant-only model
- `e(ll_c)` log likelihood, comparison model
- `e(chi2)` \( \chi^2 \)
- `e(chi2_c)` \( \chi^2 \) for comparison test
- `e(N_clust)` number of clusters
- `e(sigma_u)` panel-level standard deviation
- `e(n_quad)` number of quadrature points
- `e(g_min)` smallest group size
- `e(g_avg)` average group size
- `e(g_max)` largest group size
- `e(p)` \( p \)-value for model test
- `e(rank)` rank of `e(V)`
- `e(rank0)` rank of `e(V)` for constant-only model
- `e(ic)` number of iterations
- `e(rc)` return code
- `e(converged)` 1 if converged, 0 otherwise

Macros

- `e(cmd)` `meglm`
- `e(cmd2)` `xtologit`
- `e(cmdline)` command as typed
- `e(covariates)` list of covariates
- `e(covariates)` list of covariates
- `e(covariates)` list of covariates
- `e(ivar)` variable denoting groups
**xtologit** fits via maximum likelihood the random-effects model

\[
\Pr(y_{it} > k | \kappa, x_{it}, \nu_i) = H(x_{it}\beta + \nu_i - \kappa_k)
\]

for \(i = 1, \ldots, n\) panels, where \(t = 1, \ldots, n_i\), \(\nu_i\) are independent and identically distributed \(N(0, \sigma^2_{\nu_i})\), and \(\kappa\) is a set of cutpoints \(\kappa_1, \kappa_2, \ldots, \kappa_{K-1}\), where \(K\) is the number of possible outcomes; and \(H(\cdot)\) is the logistic cumulative distribution function.

From the above, we can derive the probability of observing outcome \(k\) for response \(y_{it}\) as

\[
p_{itk} \equiv \Pr(y_{it} = k | \kappa, x_{it}, \nu_i) = \Pr(\kappa_{k-1} < x_{it}\beta + \nu_i + \epsilon_{it} \leq \kappa_k)
\]

\[
= \Pr(\kappa_{k-1} - x_{it}\beta - \nu_i < \epsilon_{it} \leq \kappa_k - x_{it}\beta - \nu_i)
\]

\[
= H(\kappa_k - x_{it}\beta - \nu_i) - H(\kappa_{k-1} - x_{it}\beta - \nu_i)
\]

\[
= \frac{1}{1 + \exp(-\kappa_k + x_{it}\beta + \nu_i)} - \frac{1}{1 + \exp(-\kappa_{k-1} + x_{it}\beta + \nu_i)}
\]

where \(\kappa_0\) is taken as \(-\infty\) and \(\kappa_{K}\) is taken as \(+\infty\). Here \(x_{it}\) does not contain a constant term, because its effect is absorbed into the cutpoints.
We may also express this model in terms of a latent linear response, where observed ordinal responses $y_{it}$ are generated from the latent continuous responses, such that

$$y^*_it = \mathbf{x}_{it} \beta + \nu_i + \epsilon_{it}$$

and

$$y_{it} = \begin{cases} 
1 & \text{if } y^*_it \leq \kappa_1 \\
2 & \text{if } \kappa_1 < y^*_it \leq \kappa_2 \\
\vdots & \\
K & \text{if } \kappa_{K-1} < y^*_it \leq \kappa_K
\end{cases}$$

The errors $\epsilon_{it}$ are distributed as logistic with mean zero and variance $\pi^2/3$ and are independent of $\nu_i$.

Given a set of panel-level random effects $\nu_i$, we can define the conditional distribution for response $y_{it}$ as

$$f(y_{it}, \kappa, \mathbf{x}_{it} \beta + \nu_i) = \prod_{k=1}^{K} p_{itk} = \exp \left\{ \sum_{k=1}^{K} I_k(y_{it}) \log(p_{itk}) \right\}$$

where

$$I_k(y_{it}) = \begin{cases} 
1 & \text{if } y_{it} = k \\
0 & \text{otherwise}
\end{cases}$$

For panel $i$, $i = 1, \ldots, M$, the conditional distribution of $\mathbf{y}_i = (y_{i1}, \ldots, y_{in_i})'$ is

$$\prod_{t=1}^{n_i} f(y_{it}, \kappa, \mathbf{x}_{it} \beta + \nu_i)$$

and the panel-level likelihood $l_i$ is given by

$$l_i(\beta, \kappa, \sigma^2_{\nu}) = \int_{-\infty}^{\infty} e^{-\nu_i^2/2\sigma^2_{\nu}} \left\{ \prod_{t=1}^{n_i} f(y_{it}, \kappa, \mathbf{x}_{it} \beta + \nu_i) \right\} d\nu_i$$

$$\equiv \int_{-\infty}^{\infty} g(y_{it}, \kappa, \mathbf{x}_{it}, \nu_i) d\nu_i$$

This integral can be approximated with $M$-point Gauss–Hermite quadrature

$$\int_{-\infty}^{\infty} e^{-x^2} h(x) dx \approx \sum_{m=1}^{M} w_m^* h(a_m^*)$$

This is equivalent to

$$\int_{-\infty}^{\infty} f(x) dx \approx \sum_{m=1}^{M} w_m^* \exp \left\{ (a_m^*)^2 \right\} f(a_m^*)$$

where the $w_m^*$ denote the quadrature weights and the $a_m^*$ denote the quadrature abscissas. The log likelihood, $L$, is the sum of the logs of the panel-level likelihoods $l_i$. 
The default approximation of the log likelihood is by mean–variance adaptive Gauss–Hermite quadrature, which approximates the panel-level likelihood with

$$l_i \approx \sqrt{2} \tilde{\sigma}_i \sum_{m=1}^{M} w_m^* \exp \left\{ (a_m^*)^2 \right\} g(y_{it}, \kappa, x_{it}, \sqrt{2} \tilde{\sigma}_i a_m^* + \tilde{\mu}_i)$$

where $\tilde{\sigma}_i$ and $\tilde{\mu}_i$ are the adaptive parameters for panel $i$. The method of calculating the posterior mean and variance and using those parameters for $\tilde{\mu}_i$ and $\tilde{\sigma}_i$ is described in detail in Naylor and Smith (1982) and Skrondal and Rabe-Hesketh (2004). We start with $\tilde{\sigma}_{i,0} = 1$ and $\tilde{\mu}_{i,0} = 0$, and the posterior means and variances are updated in the $j$th iteration. That is, at the $j$th iteration of the optimization for $l_i$, we use

$$l_{i,j} \approx \sum_{m=1}^{M} \sqrt{2} \tilde{\sigma}_{i,j-1} w_m^* \exp \left\{ (a_m^*)^2 \right\} g(y_{it}, \kappa, x_{it}, \sqrt{2} \tilde{\sigma}_{i,j-1} a_m^* + \tilde{\mu}_{i,j-1})$$

Letting

$$\tau_{i,m,j-1} = \sqrt{2} \tilde{\sigma}_{i,j-1} a_m^* + \tilde{\mu}_{i,j-1}$$

$$\tilde{\mu}_{i,j} = \sum_{m=1}^{M} \tau_{i,m,j-1} \sqrt{2} \tilde{\sigma}_{i,j-1} w_m^* \exp \left\{ (a_m^*)^2 \right\} g(y_{it}, \kappa, x_{it}, \tau_{i,m,j-1})$$

and

$$\tilde{\sigma}_{i,j} = \sum_{m=1}^{M} (\tau_{i,m,j-1})^2 \sqrt{2} \tilde{\sigma}_{i,j-1} w_m^* \exp \left\{ (a_m^*)^2 \right\} g(y_{it}, \kappa, x_{it}, \tau_{i,m,j-1}) \frac{l_{i,j}}{l_{i,j}} - (\tilde{\mu}_{i,j})^2$$

This is repeated until $\tilde{\mu}_{i,j}$ and $\tilde{\sigma}_{i,j}$ have converged for this iteration of the maximization algorithm. This adaptation is applied on every iteration.

The log likelihood can also be calculated by nonadaptive Gauss–Hermite quadrature with the option intmethod(ghermite), where $\rho = \sigma_i^2 / (\sigma_i^2 + 1)$:

$$L = \sum_{i=1}^{n} w_i \log \left\{ \Pr(y_{i1}, \ldots, y_{ini} | \kappa, x_{i1}, \ldots, x_{ini}) \right\} \approx \sum_{i=1}^{n} w_i \log \left[ \frac{1}{\sqrt{\pi}} \sum_{m=1}^{M} w_m^* \prod_{t=1}^{n_i} f \left\{ y_{it}, \kappa, x_{it} \beta + a_m^* \left( \frac{2\rho}{1-\rho} \right)^{1/2} \right\} \right]$$

Both quadrature formulas require that the integrated function be well approximated by a polynomial of degree equal to the number of quadrature points. The number of periods (panel size) can affect whether

$$\prod_{t=1}^{n_i} f(y_{it}, \kappa, x_{it} \beta + \nu_t)$$

is well approximated by a polynomial. As panel size and $\rho$ increase, the quadrature approximation can become less accurate. For large $\rho$, the random-effects model can also become unidentified. Adaptive quadrature gives better results for correlated data and large panels than nonadaptive quadrature; however, we recommend that you use the quadchk command (see [XT] quadchk) to verify the quadrature approximation used in this command, whichever approximation you choose.
xtologit and the robust VCE estimator

Specifying vce(robust) or vce(cluster clustvar) causes the Huber/White/sandwich VCE estimator to be calculated for the coefficients estimated in this regression. See [P] _robust, particularly Introduction and Methods and formulas. Wooldridge (2020) and Arellano (2003) discuss this application of the Huber/White/sandwich VCE estimator. As discussed by Wooldridge (2020), Stock and Watson (2008), and Arellano (2003), specifying vce(robust) is equivalent to specifying vce(cluster panelvar), where panelvar is the variable that identifies the panels.

Clustering on the panel variable produces a consistent VCE estimator when the disturbances are not identically distributed over the panels or there is serial correlation in $\epsilon_{it}$.

The cluster–robust VCE estimator requires that there are many clusters and the disturbances are uncorrelated across the clusters. The panel variable must be nested within the cluster variable because of the within-panel correlation that is generally induced by the random-effects transform when there is heteroskedasticity or within-panel serial correlation in the idiosyncratic errors.

References


Also see

[XT] xtologit postestimation — Postestimation tools for xtologit

[XT] quadchk — Check sensitivity of quadrature approximation

[XT] xtprobit — Random-effects ordered probit models

[XT] xtset — Declare data to be panel data

[ME] meologit — Multilevel mixed-effects ordered logistic regression

[R] logistic — Logistic regression, reporting odds ratios

[R] logit — Logistic regression, reporting coefficients

[U] 20 Estimation and postestimation commands