

gsem — Generalized structural equation model estimation command

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Description

`gsem` fits generalized SEMs. When you use the Builder in `gsem` mode, you are using the `gsem` command.

Menu

Statistics > SEM (structural equation modeling) > Model building and estimation

Syntax

`gsem paths [if] [in] [weight] [, options]`

where *paths* are the paths of the model in command-language path notation; see [SEM] **sem and gsem path notation**.

| <i>options</i> | Description |
|----------------------------------|-------------------------------------------------------------|
| <i>model_description_options</i> | fully define, along with <i>paths</i> , the model to be fit |
| <i>estimation_options</i> | method used to obtain estimation results |
| <i>reporting_options</i> | reporting of estimation results |
| <i>syntax_options</i> | controlling interpretation of syntax |

Factor variables and time-series operators are allowed.

`bootstrap`, `by`, `jackknife`, `permute`, `statsby`, and `svy` are allowed; see [U] **11.1.10 Prefix commands**.

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**.

Also see [SEM] **gsem postestimation** for features available after estimation.

Options

model_description_options describe the model to be fit. The model to be fit is fully specified by *paths*—which appear immediately after `gsem`—and the options `covariance()`, `variance()`, and `means()`. See [SEM] **gsem model description options** and [SEM] **sem and gsem path notation**.

estimation_options control how the estimation results are obtained. These options control how the standard errors (VCE) are obtained and control technical issues such as choice of estimation method. See [SEM] **gsem estimation options**.

reporting_options control how the results of estimation are displayed. See [SEM] **gsem reporting options**.

syntax_options control how the syntax that you type is interpreted. See [SEM] **sem and gsem syntax options**.

Remarks and examples

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gsem provides important features not provided by **sem** and correspondingly omits useful features provided by **sem**. The differences in capabilities are the following:

1. **gsem** allows generalized linear response functions as well as the linear response functions allowed by **sem**.
2. **gsem** allows for multilevel models, something **sem** does not.
3. **gsem** allows Stata's factor-variable notation to be used in specifying models, something **sem** does not.
4. **gsem**'s method ML is sometimes able to use more observations in the presence of missing values than can **sem**'s method ML. Meanwhile, **gsem** does not provide the MLMV method provided by **sem** for explicitly handling missing values.
5. **gsem** lacks the group-comparison capabilities that **sem** provides.
6. **gsem** cannot produce standardized coefficients.
7. **gsem** cannot use summary statistic datasets (SSDs); **sem** can.

gsem has nearly identical syntax to **sem**. Differences in syntax arise because of differences in capabilities. The resulting differences in syntax are the following:

1. **gsem** adds new syntax to *paths* to handle latent variables associated with multilevel modeling.
2. **gsem** adds new options to handle the family and link of generalized linear responses.
3. **gsem** deletes options related to features it does not have, such as group-comparison capabilities and SSDs.
4. **gsem** adds technical options for controlling features not provided by **sem**, such as numerical integration (quadrature choices), number of integration points, and a number of options dealing with starting values, which are a more difficult proposition in the generalized SEM framework.

For a readable explanation of what **gsem** can do and how to use it, see the intro sections. You might start with [SEM] **intro 1**.

For examples of **gsem** in action, see the example sections. You might start with [SEM] **example 1**.

For detailed syntax and descriptions, see the references below.

Remarks on three advanced topics are presented under the following headings:

- Default normalization constraints*
- Default covariance assumptions*
- How to solve convergence problems*

Default normalization constraints

`gsem` applies the same rules as `sem` to identify models; see [SEM] `sem` and see [SEM] `intro 4`. Everything said there about latent variables applies to multilevel latent variables such as M1[school] and M2[school>teacher].

Default covariance assumptions

`gsem` assumes the same covariance structures as does `sem`; see [SEM] `sem` and see [SEM] `intro 4`. `gsem`, however, treats covariances between observed exogenous variables as given. Actually, so does `sem` unless you specify an override. The override cannot be specified with `gsem`.

How to solve convergence problems

See [SEM] `intro 12`.

Stored results

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

Scalars

| | |
|---------------------------|----------------------------------------------------------|
| <code>e(N)</code> | number of observations |
| <code>e(N_clust)</code> | number of clusters |
| <code>e(k)</code> | number of parameters |
| <code>e(k_cat#)</code> | number of categories for the #th <i>depvar</i> , ordinal |
| <code>e(k_dv)</code> | number of dependent variables |
| <code>e(k_eq)</code> | number of equations in <code>e(b)</code> |
| <code>e(k_out#)</code> | number of outcomes for the #th <i>depvar</i> , mlogit |
| <code>e(k_rc)</code> | number of covariances |
| <code>e(k_rs)</code> | number of variances |
| <code>e(l1)</code> | log likelihood |
| <code>e(n_quad)</code> | number of integration points |
| <code>e(rank)</code> | rank of <code>e(V)</code> |
| <code>e(ic)</code> | number of iterations |
| <code>e(rc)</code> | return code |
| <code>e(converged)</code> | 1 if target model converged, 0 otherwise |

Macros

| | |
|---------------------------|-----------------------------------------------------------------------------------------------------|
| <code>e(cmd)</code> | <code>gsem</code> |
| <code>e(cmdline)</code> | command as typed |
| <code>e(depvar)</code> | names of dependent variables |
| <code>e(wtype)</code> | weight type |
| <code>e(wexp)</code> | weight expression |
| <code>e(fweightk)</code> | fweight variable for kth level, if specified |
| <code>e(pweightk)</code> | pweight variable for kth level, if specified |
| <code>e(iweightk)</code> | iweight variable for kth level, if specified |
| <code>e(title)</code> | title in estimation output |
| <code>e(clustvar)</code> | name of cluster variable |
| <code>e(family#)</code> | family for the #th <i>depvar</i> |
| <code>e(link#)</code> | link for the #th <i>depvar</i> |
| <code>e(offset#)</code> | offset for the #th <i>depvar</i> |
| <code>e(intmethod)</code> | integration method |
| <code>e(vce)</code> | <i>vcetype</i> specified in <code>vce()</code> |
| <code>e(vcetype)</code> | title used to label Std. Err. |
| <code>e(opt)</code> | type of optimization |
| <code>e(which)</code> | <code>max</code> or <code>min</code> ; whether optimizer is to perform maximization or minimization |
| <code>e(method)</code> | estimation method: <code>ml</code> |
| <code>e(ml_method)</code> | type of <code>ml</code> method |

| | |
|----------------------|-----------------------------------------------------------------------|
| e(user) | name of likelihood-evaluator program |
| e(technique) | maximization technique |
| e(datasignature) | the checksum |
| e(datasignaturevars) | variables used in calculation of checksum |
| e(properties) | b V |
| e(estat_cmd) | program used to implement <code>estat</code> |
| e(predict) | program used to implement <code>predict</code> |
| e(covariates) | list of covariates |
| e(footnote) | program used to implement the footnote display |
| e(asbalanced) | factor variables <code>fvset</code> as <code>asbalanced</code> |
| e(asobserved) | factor variables <code>fvset</code> as <code>asobserved</code> |
| e(marginsnotok) | predictions not allowed by <code>margins</code> |
| e(marginswtype) | weight type for <code>margins</code> |
| e(marginswexp) | weight expression for <code>margins</code> |
| e(marginsdefault) | default <code>predict()</code> specification for <code>margins</code> |
| Matrices | |
| e(b) | parameter vector |
| e(b_pclass) | parameter class |
| e(cat#) | categories for the #th <code>depvar</code> , ordinal |
| e(out#) | outcomes for the #th <code>depvar</code> , mlogit |
| e(Cns) | constraints matrix |
| e(iolog) | iteration log (up to 20 iterations) |
| e(gradient) | gradient vector |
| e(V) | covariance matrix of the estimators |
| e(V_modelbased) | model-based variance |
| Functions | |
| e(sample) | marks estimation sample |

Also see

- [SEM] **intro 1** — Introduction
- [SEM] **sem and gsem path notation** — Command syntax for path diagrams
- [SEM] **gsem path notation extensions** — Command syntax for path diagrams
- [SEM] **gsem model description options** — Model description options
- [SEM] **gsem estimation options** — Options affecting estimation
- [SEM] **gsem reporting options** — Options affecting reporting of results
- [SEM] **sem and gsem syntax options** — Options affecting interpretation of syntax
- [SEM] **gsem postestimation** — Postestimation tools for gsem
- [SEM] **methods and formulas for gsem** — Methods and formulas for gsem
- [SVY] **svy estimation** — Estimation commands for survey data