

sem and gsem path notation — Command syntax for path diagrams
[Description](#)[Syntax](#)[Options](#)[Remarks and examples](#)[Also see](#)

Description

The command syntax for describing your SEM is fully specified by *paths*, `covariance()`, `variance()`, and `means()`. How this works is described below.

If you are using `sem`, also see [\[SEM\] sem path notation extensions](#) for documentation of the `group()` option for comparing different groups in the data. The syntax of the elements described below is modified when `group()` is specified.

If you are using `gsem`, also see [\[SEM\] gsem path notation extensions](#) for documentation on specification of family-and-link for generalized (nonlinear) response variables, specification of multilevel latent variables, specification of categorical latent variables, and specification of multiple-group models. The syntax of the elements described below is modified when the `group()` option for comparing different groups or the `lclass()` option for categorical latent variables is specified.

Either way, read this section first.

Syntax

```
sem paths ... [ , covariance() variance() means() ]
gsem paths ... [ , covariance() variance() means() ]
```

paths specifies the direct paths between the variables of your model.

The model to be fit is fully described by *paths*, `covariance()`, `variance()`, and `means()`.

Options

`covariance()` is used to

1. specify that a particular covariance path of your model that usually is assumed to be 0 be estimated,
2. specify that a particular covariance path that usually is assumed to be nonzero is not to be estimated (to be constrained to be 0),
3. constrain a covariance path to a fixed value, such as 0, 0.5, 1, etc., and
4. constrain two or more covariance paths to be equal.

`variance()` does the same as `covariance()` except it does it with variances.

`means()` does the same as `covariance()` except it does it with means.

Remarks and examples

[stata.com](#)

Path notation is used by the `sem` and `gsem` commands to specify the model to be fit, for example,

```
. sem (x1 x2 x3 x4 <- X)
. gsem (L1 -> x1 x2 x3 x4 x5, logit) (L2 -> x6 x7 x8 x9 x10)
```

In the path notation,

1. Latent variables are indicated by a *name* in which at least the first letter is capitalized.
2. Observed variables are indicated by a *name* in which at least the first letter is lowercased. Observed variables correspond to variable names in the dataset.
3. Error variables, while mathematically a special case of latent variables, are considered in a class by themselves. For `sem`, every endogenous variable (whether observed or latent) automatically has an error variable associated with it. For `gsem`, the same is true of Gaussian endogenous variables (and latent variables, which are Gaussian). The error variable associated with endogenous variable *name* is `e.name`.

4. Paths between variables are written as

`(name1 <- name2)`

or

`(name2 -> name1)`

There is no significance to which coding is used.

5. Paths between the same variables can be combined: The paths

`(name1 <- name2) (name1 <- name3)`

can be combined as

`(name1 <- name2 name3)`

or as

`(name2 name3 -> name1)`

The paths

`(name1 <- name3) (name2 <- name3)`

can be combined as

`(name1 name2 <- name3)`

or as

`(name3 -> name1 name2)`

The paths

`(name1 <- name2 name3)`

`(name4 <- name2 name3)`

may be written as

`(name1 name4 <- name2 name3)`

or as

`(name2 name3 -> name1 name4)`

6. Variances and covariances (curved paths) between variables are indicated by options. Variances are indicated by

```
..., ... var(name1)
```

Covariances are indicated by

```
..., ... cov(name1*name2)
```

```
..., ... cov(name2*name1)
```

There is no significance to the order of the names.

The actual names of the options are `variance()` and `covariance()`, but they are invariably abbreviated as `var()` and `cov()`, respectively.

The `var()` and `cov()` options are the same option, so a variance can be typed as

```
..., ... cov(name1)
```

and a covariance can be typed as

```
..., ... var(name1*name2)
```

7. Variances may be combined, covariances may be combined, and variances and covariances may be combined.

If you have

```
..., ... var(name1) var(name2)
```

you may code this as

```
..., ... var(name1 name2)
```

If you have

```
..., ... cov(name1*name2) cov(name2*name3)
```

you may code this as

```
..., ... cov(name1*name2 name2*name3)
```

All the above combined can be coded as

```
..., ... var(name1 name2 name1*name2 name2*name3)
```

or as

```
..., ... cov(name1 name2 name1*name2 name2*name3)
```

8. All variables except endogenous variables are assumed to have a variance; it is only necessary to code the `var()` option if you wish to place a constraint on the variance or specify an initial value. See items 11, 12, 13, and 16 below. (In `gsem`, the variance and covariances of observed endogenous variables are not estimated and thus `var()` cannot be used with them.)

Endogenous variables have a variance, of course, but that is the variance implied by the model. If *name* is an endogenous variable, then `var(name)` is invalid. The error variance of the endogenous variable is `var(e.name)`.

9. Variables mostly default to being correlated:

- All exogenous variables are assumed to be correlated with each other, whether observed or latent.
- Endogenous variables are never directly correlated, although their associated error variables can be.
- All error variables are assumed to be uncorrelated with each other.

You can override these defaults on a variable-by-variable basis with the `cov()` option.

To assert that two variables are uncorrelated that otherwise would be assumed to be correlated, constrain the covariance to be 0:

```
..., ... cov(name1*name2@0)
```

To allow two variables to be correlated that otherwise would be assumed to be uncorrelated, simply specify the existence of the covariance:

```
..., ... cov(name1*name2)
```

This latter is especially commonly done with errors:

```
..., ... cov(e.name1*e.name2)
```

(In `gsem`, you may not use the `cov()` option with observed exogenous variables. You also may not use `cov()` with error terms associated with family Gaussian, link log.)

10. Means of variables are indicated by the following option:

```
..., ... means(name)
```

Variables mostly default to having nonzero means:

- All observed exogenous variables are assumed to have nonzero means. In `sem`, the means can be constrained using the `means()` option, but only if you are performing `noxconditional` estimation; [SEM] **sem option noxconditional**.
- Latent exogenous variables are assumed to have mean 0. Means of latent variables are not estimated by default. If you specify enough normalization constraints to identify the mean of a latent exogenous variable, you can specify `means(name)` to indicate that the mean should be estimated in either.
- Endogenous variables have no separate mean. Their means are those implied by the model. The `means()` option may not be used with endogenous variables.
- Error variables have mean 0 and this cannot be modified. The `means()` option may not be used with error variables.

To constrain the mean to a fixed value, such as 57, code

```
..., ... means(name@57)
```

Separate `means()` options may be combined:

```
..., ... means(name1@57 name2@100)
```

11. Fixed-value constraints may be specified for a path, variance, covariance, or mean by using `@` (the “at” symbol). For example,

```
(name1 <- name2@1)
```

```
(name1 <- name2@1 name3@1)
```

```
..., ... var(name@100)
```

```
..., ... cov(name1*name2@223)
```

```
..., ... cov(name1@1 name2@1 name1*name2@.8)
```

```
..., ... means(name@57)
```

12. Symbolic constraints may be specified for a path, variance, covariance, or mean by using @ (the “at” symbol). For example,

```
(name1 <- name2@c1) (name3 <- name4@c1)
... , ... var(name1@c1 name2@c1)
... , ... cov(name1@1 name2@1 name3@1 name1*name2@c1 name1*name3@c1)
... , ... means(name1@c1 name2@c1)
(name1 <- name2@c1) ... , var(name3@c1) means(name4@c1)
```

Symbolic names are just names from 1 to 32 characters in length. Symbolic constraints constrain equality. For simplicity, all constraints below will have names `c1`, `c2`, ...

13. Linear combinations of symbolic constraints may be specified for a path, variance, covariance, or mean by using @ (the “at” symbol). For example,

```
(name1 <- name2@c1) (name3 <- name4@(2*c1))
... , ... var(name1@c1 name2@(c1/2))
... , ... cov(name1@1 name2@1 name3@1 name1*name2@c1 name1*name2@(c1/2))
... , ... means(name1@c1 name2@(3*c1+10))
(name1 <- name2@(c1/2)) ... , var(name3@c1) means(name4@(2*c1))
```

14. All equations in the model are assumed to have an intercept (to include observed exogenous variable `_cons`) unless the `noconstant` option (abbreviation `nocons`) is specified, and then all equations are assumed not to have an intercept (not to include `_cons`). (There are some exceptions to this in `gsem` because some generalized linear models have no intercept or even the concept of an intercept.)

Regardless of whether `noconstant` is specified, you may explicitly refer to observed exogenous variable `_cons`.

The following path specifications are ways of writing the same model:

```
(name1 <- name2) (name1 <- name3)
(name1 <- name2) (name1 <- name3) (name1 <- _cons)
(name1 <- name2 name3)
(name1 <- name2 name3 _cons)
```

There is no reason to explicitly specify `_cons` unless you have also specified the `noconstant` option and want to include `_cons` in some equations but not others, or regardless of whether you specified the `noconstant` option, you want to place a constraint on its path coefficient. For example,

```
(name1 <- name2 name3 _cons@c1) (name4 <- name5 _cons@c1)
```

15. The `noconstant` option may be specified globally or within a path specification. That is,

```
(name1 <- name2 name3) (name4 <- name5) , nocons
```

suppresses the intercepts in both equations. Alternatively,

```
(name1 <- name2 name3, nocons) (name4 <- name5)
```

suppresses the intercept in the first equation but not the second, whereas

```
(name1 <- name2 name3) (name4 <- name5, nocons)
```

suppresses the intercept in the second equation but not the first.

In addition, consider the equation

```
(name1 <- name2 name3, nocons)
```

This can be written equivalently as

```
(name1 <- name2, nocons) (name1 <- name3, nocons)
```

16. Initial values (starting values) may be specified for a path, variance, covariance, or mean by using the `init(#)` suboption:

```
(name1 <- (name2, init(0)))  
(name1 <- (name2, init(0)) name3)  
(name1 <- (name2, init(0)) (name3, init(5)))  
..., ... var((name3, init(1)))  
..., ... cov((name4*name5, init(.5)))  
..., ... means((name5, init(0)))
```

The initial values may be combined with symbolic constraints:

```
(name1 <- (name2@c1, init(0)))  
(name1 <- (name2@c1, init(0)) name3)  
(name1 <- (name2@c1, init(0)) (name3@c2, init(5)))  
..., ... var((name3@c1, init(1)))  
..., ... cov((name4*name5@c1, init(.5)))  
..., ... means((name5@c1, init(0)))
```

Also see

[SEM] **sem** — Structural equation model estimation command

[SEM] **gsem** — Generalized structural equation model estimation command

[SEM] **sem path notation extensions** — Command syntax for path diagrams

[SEM] **gsem path notation extensions** — Command syntax for path diagrams

[SEM] **intro 2** — Learning the language: Path diagrams and command language

[SEM] **intro 6** — Comparing groups