

## intro 6 — Comparing groups

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## Description

`sem` and `gsem` have a unique feature that allows you to easily compare groups—compare males with females, compare age group 1 with age group 2 with age group 3, and so on—with respect to any SEM. Said more technically, any model fit by `sem` or `gsem` can be simultaneously estimated for different groups with some parameters constrained to be equal across groups and others allowed to vary, and those estimates can be used to perform statistical tests for comparing the groups.

`sem` and `gsem` have similar syntax for multiple group models, but there are important differences in default assumptions and in how you specify constraints. So we discuss how to compare groups using each command separately, first with `sem` and then with `gsem`.

## Remarks and examples

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Remarks are presented under the following headings:

*Comparing groups with sem**The generic SEM model**sem: Fitting the model for different groups of the data**sem: Which parameters vary by default, and which do not**sem: Specifying which parameters are allowed to vary in broad, sweeping terms**sem: Adding constraints for path coefficients across groups**sem: Adding constraints for means, variances, or covariances across groups**sem: Adding constraints for some groups but not others**sem: Adding paths for some groups but not others**sem: Relaxing constraints**Comparing groups with gsem**gsem: Fitting the model for different groups of the data**gsem: Which parameters vary by default, and which do not**gsem: Specifying which parameters are allowed to vary in broad, sweeping terms**gsem: Adding constraints for path coefficients across groups**gsem: Adding constraints for means, variances, or covariances across groups**gsem: Adding constraints for some groups but not others**gsem: Adding paths for some groups but not others**gsem: Relaxing constraints*

## Comparing groups with sem

`sem` and `gsem` have similar syntax for multiple group models. There are, however, important differences in default assumptions and in how you specify constraints, so we discuss how to compare groups using each command separately. We begin with `sem`'s syntax for group comparisons.



where `agegrp` is a variable in our dataset, perhaps taking on values 1, 2, 3, . . . . We can specify the model by using the command language or by drawing the model in the Builder and then choosing and filling in the `group()` option.

After estimation, you can use `estat ginvariant` (see [SEM] [estat ginvariant](#)) to obtain Wald tests of whether constraints should be added and score tests of whether constraints should be relaxed.

### sem: Which parameters vary by default, and which do not

When we specify `group(groupvar)`, the measurement parts of the model (parts 1 and 3) are constrained by default to be the same across the groups, whereas the middle part (part 2) will have separate parameters for each group. More specifically, parts 1 and 3 are constrained to be equal across groups except that the variances of the errors will be estimated separately for each group.

If there is no measurement component to the model—if there are no latent variables—then by default all parameters are estimated separately for each group.

### sem: Specifying which parameters are allowed to vary in broad, sweeping terms

You can control which parameters are constrained to be equal across groups by specifying the `ginvariant()` option:

```
(Y1->...) (Y2->...)          /// part 3
(...)          ///
(...)          /// part 2
(...)          ///
(X1->...) (X2->...),          /// part 1
                    group(agegrp) ginvariant(pclasses)
```

The parameter classes (*pclasses*) for `sem` are as follows:

Parameter class description	Parameter class name
1. structural coefficients	<code>scoef</code>
2. structural intercepts	<code>scons</code>
3. measurement coefficients	<code>mcoef</code>
4. measurement intercepts	<code>mcons</code>
5. covariances of structural errors	<code>serrvar</code>
6. covariances of measurement errors	<code>merrvar</code>
7. covariances between structural and measurement errors	<code>smerrcov</code>
8. means of exogenous variables	<code>meanex</code> (*)
9. covariances of exogenous variables	<code>covex</code> (*)
10. all the above	<code>all</code> (*)
11. none of the above	<code>none</code>

(\*) Be aware that 8, 9, and 10 (`meanex`, `covex`, and `all`) exclude the observed exogenous variables—that is, they include only the latent exogenous variables—unless you specify the `noxconditional` option or the `noxconditional` option is otherwise implied; see [SEM] [sem option noxconditional](#). This is what you would desire in most cases.

The default when `ginvariant()` is not specified is `ginvariant(mcoef mcons)`:

```
(Y1->...) (Y2->...)          /// part 3, measurement
(...)          ///
(...)          /// part 2, structural
(...)          ///
(X1->...) (X2->...),          /// part 1, measurement
                    group(agegrp) ginvariant(mcoef mcons)
```

If you also wanted covariances of errors associated with measurement to be constrained across groups, you could type

```
(Y1->...) (Y2->...)          /// part 3, measurement
(...)          ///
(...)          /// part 2, structural
(...)          ///
(X1->...) (X2->...),          /// part 1, measurement
                    group(agegrp) ginvariant(mcoef mcons merrvar)
```

### sem: Adding constraints for path coefficients across groups

The `ginvariant()` option allows you to state in sweeping terms which parameters vary and which are invariant across groups. You can also constrain individual parameters to be equal across groups.

Pretend that in the substantive part of the generic model, we have  $Y1 < -Y2$ . Assume that we fit the model and allow the structural part to vary across groups:

```
(Y1->...) (Y2->...)          /// part 3, measurement
(...)          ///
(Y1<-Y2)          /// part 2, structural
(...)          ///
(X1->...) (X2->...),          /// part 1, measurement
                    group(agegrp)
```

In this model, the  $Y1 < -Y2$  path coefficient is allowed to vary across groups by default. We could constrain the coefficient to be equal across groups by typing

```
(Y1->...) (Y2->...)          /// part 3, measurement
(...)          ///
(Y1<-Y2@b)          /// part 2, structural
(...)          ///
(X1->...) (X2->...),          /// part 1, measurement
                    group(agegrp)
```

Note the `@b` in  $(Y1 < -Y2@b)$  where we previously typed  $Y1 < -Y2$ .

Constraining a coefficient to equal a symbolic name such as `b` is how we usually constrain equality, but in the usual case, the symbolic name appears at least twice in our model. For instance, we might have  $(Y1 < -Y2@b)$  and  $(Y1 < -Y3@b)$  and thus constrain path coefficients to be equal.

In the case above, however, `@b` appears only once. Because we specified `group(agegrp)`, results are as if we specified this model separately for each age group, and in each group, we are specifying `@b`. Thus we are constraining the path coefficient to be equal across all groups.

**sem: Adding constraints for means, variances, or covariances across groups**

You use the same technique for adding constraints to means, variances, and covariances as you would for adding constraints to path coefficients. Remember that means are specified by the `means()` option, variances by the `variance()` option, and covariances by the `covariance()` option. The `variance()` and `covariance()` options are abbreviated `var()` and `cov()`, respectively.

You can specify, for instance,

```
(Y1->...) (Y2->...)          /// part 3, measurement
(...)                          ///
(Y1<-Y2)                       /// part 2, structural
(...)                          ///
(X1->...) (X2->...),           /// part 1, measurement
      group(agegrp)          ///
      means(X1@M)
```

to constrain the mean of X1 to be the same across groups. The means would have been different across groups by default.

You can specify

```
(Y1->...) (Y2->...)          /// part 3, measurement
(...)                          ///
(Y1<-Y2)                       /// part 2, structural
(...)                          ///
(X1->...) (X2->...),           /// part 1, measurement
      group(agegrp)          ///
      var(e.Y1@V)
```

to constrain the variance of the error of Y1 to be the same across groups.

If we wanted to allow the errors of Y1 and Y2 to be correlated (by default, errors are uncorrelated), we could add the `cov()` option:

```
(Y1->...) (Y2->...)          /// part 3, measurement
(...)                          ///
(...)                          /// part 2, structural
(...)                          ///
(X1->...) (X2->...),           /// part 1, measurement
      group(agegrp)          ///
      var(e.Y1@V)            ///
      cov(e.Y1*e.Y2)
```

If we then wanted to constrain the covariance to be the same across groups, we would type

```
(Y1->...) (Y2->...)          /// part 3, measurement
(...)                          ///
(...)                          /// part 2, structural
(...)                          ///
(X1->...) (X2->...),           /// part 1, measurement
      group(agegrp)          ///
      var(e.Y1@V)            ///
      cov(e.Y1*e.Y2@C)
```

**sem: Adding constraints for some groups but not others**

Consider the following model:

```
... (Y1<-Y2) ..., group(agegrp)
```

Above we saw how to constrain the Y1<-Y2 path coefficient to be the same across groups:

```
... (Y1<-Y2@b) ..., group(agegrp)
```

To constrain the path coefficients Y1<-Y2 to be equal for groups 1 and 2 but leave the Y1<-Y2 path coefficients unconstrained for the remaining groups, we would type

```
... (Y1<-Y2) (1: Y1<-Y2@b) (2: Y1<-Y2@b) ..., group(agegrp)
```

Think of this as follows:

1. (Y1<-Y2): We set a path for all the groups.
2. (1: Y1<-Y2@b): We modify the path for `agegrp = 1`.
3. (2: Y1<-Y2@b): We modify the path for `agegrp = 2`.
4. We do not modify the path for any other `agegrp` value.

The result is that we constrain age groups 1 and 2 to have the same value of the path, and we do not constrain the path for the other age groups.

You can constrain variance and covariance estimates to be the same across some groups but not others in the same way. You can specify, for instance,

```
..., group(agegrp) var(1: e.Y1@V) var(2: e.Y1@V)
```

or

```
..., group(agegrp) cov(e.Y1*e.Y2) cov(1: e.Y1*e.Y2@C) ///  
                                cov(2: e.Y1*e.Y2@C)
```

Similarly, you can constrain means for some groups but not others, although this is rarely done:

```
..., group(agegrp) means(1: X1@b) means(2: X1@b)
```

**sem: Adding paths for some groups but not others**

In the same way that you can constrain coefficients for some groups but not others, you can add paths for some groups but not others. Consider the following model:

```
... (Y1<-Y2) ..., group(agegrp)
```

You can add the path Y1<-Y3 for groups 1 and 2 by typing

```
... (Y1<-Y2) (1: Y1<-Y3) (2: Y1<-Y3) ..., group(agegrp)
```

You can add covariances for some groups but not others in the same way. For instance, to allow the errors of Y1 and Y2 to be correlated in groups 1 and 2 only, you can specify

```
..., group(agegrp) cov(1: e.Y1*e.Y2) cov(2: e.Y1*e.Y2)
```

**sem: Relaxing constraints**

Just as you can specify

```
..., group(agegrp) ginvariant(pclasses)
```

and then add constraints, you can also specify

```
..., group(agegrp) ginvariant(pclasses)
```

and then relax constraints that the parameter classes impose.

For instance, if we specified `ginvariant(scoef)`, then we would be constraining ( $Y1 <- Y2$ ) to be invariant across groups. We could then relax that constraint by typing

```
... (Y1<-Y2) (1: Y1<-Y2@b1) (2: Y1<-Y2@b2) ..., ///
                                group(agegrp) ginvariant(scoef)
```

The path coefficients would be free in groups 1 and 2 and constrained in the remaining groups, if there are any. The path coefficient is free in group 1 because we specified symbolic name `b1`, and `b1` appears nowhere else in the model. The path coefficient is free in group 2 because symbolic name `b2` appears nowhere else in the model. If there are remaining groups and we want to relax the constraint on them, too, we would need to add `(3: Y1<-Y2@b3)`, and so on.

The same technique can be used to relax constraints on means, variances, and covariances:

```
..., group(agegrp) ginvariant(... meanex ...) ///
                                means(1: X1@b1) means(2: X1@b2)
..., group(agegrp) ginvariant(... serrvar ...) ///
                                var(1: e.Y1@V1) var(2: e.Y1@V2)
..., group(agegrp) ginvariant(... serrvar ...) ///
                                cov(1: e.Y1*e.Y2@C1) cov(2: e.Y1*e.Y2@C2)
```

**Comparing groups with gsem**

With `gsem`, we fit multiple-group models in a similar way by using the `group()` and `ginvariant()` options. However, the allowed parameter classes and the default constraints differ between with `sem` and `gsem`. Here, we will discuss how compare groups and how to set and release constraints across groups for models fit with `gsem`.

**gsem: Fitting the model for different groups of the data**

Let's begin by considering a measurement model that has binary measurements of a latent variable. If we model the measurements using probit regression, we can fit a model with the command syntax as

```
(L -> y1 y2 y3 y4), probit
```

We can fit this model for different groups such as age groups by specifying the `group(varname)` option:

```
(L -> y1 y2 y3 y4), probit group(agegrp)
```

where `agegrp` is a variable in our dataset, perhaps taking on values 1, 2, 3, . . . We can specify the model by using the command language or by drawing the model in the Builder and then choosing and filling in the `group()` option.

After estimation, you can use `test` to obtain Wald tests of whether constraints should be added. See [SEM] [example 49g](#) for an example.

### **gsem: Which parameters vary by default, and which do not**

When we specify `group(groupvar)` with `gsem`, the coefficients, intercepts, and cutpoints in the model are constrained by default to be the same across the groups. All other types of parameters will be estimated separately in each group. More specifically, all variances, covariances, means, and scale parameters will be estimated separately for each group.

### **gsem: Specifying which parameters are allowed to vary in broad, sweeping terms**

You can control which parameters are constrained to be equal across groups by specifying the `ginvariant()` option:

```
(L -> y1 y2 y3 y4), probit ///
      group(agegrp) ginvariant(pclasses)
```

The parameter classes (*pclasses*) for `gsem` are as follows:

Parameter class description	Parameter class name
1. intercepts and cutpoints	<code>cons</code>
2. fixed coefficients	<code>coef</code>
3. latent variable coefficients	<code>loading</code>
4. covariances of errors	<code>errvar</code>
5. scaling parameters	<code>scale</code>
6. means of exogenous variables	<code>means</code>
7. covariances of exogenous latent variables	<code>covex</code>
8. all the above	<code>all</code>
9. none of the above	<code>none</code>

The default when `ginvariant()` is not specified is `ginvariant(cons coef loading)`:

```
(L -> y1 y2 y3 y4), probit ///
      group(agegrp) ginvariant(cons coef loading)
```

If you also wanted the variance of the latent variable to be constrained across groups, you could type

```
(L -> y1 y2 y3 y4), probit ///
      group(agegrp) ginvariant(cons coef loading covex)
```

because `L` is an exogenous variable and `covex` refers to all covariances and variances of exogenous variables.



**gsem: Adding constraints for path coefficients across groups**

The `ginvariant()` option allows you to state in sweeping terms which parameters vary and which are invariant across groups. You can also constrain individual parameters to be equal across groups.

If we fit a model by typing

```
(L -> y1 y2 y3 y4), probit ///
      group(agegrp) ginvariant(cons)
```

In this model, only the intercepts are constrained to be equal across groups. The L->y1, L->y2, L->y3, and L->y4 path coefficients are allowed to vary across groups because we did not include `loading` in the `ginvariant()` option. We could constrain the L->y4 coefficient to be equal across groups by typing

```
(L -> y1 y2 y3 y4@b), probit ///
      group(agegrp) ginvariant(cons)
```

Note the `@b` attached to `y4`.

Constraining a coefficient to equal a symbolic name such as `b` is how we usually constrain equality, but in the usual case, the symbolic name appears at least twice in our model. For instance, we might have `(L->y2@b)` and `(L->y3@b)` and thus constrain path coefficients to be equal.

In the case above, however, `@b` appears only once. Because we specified `group(agegrp)`, results are as if we specified this model separately for each age group, and in each group, we are specifying `@b`. Thus we are constraining the path coefficient to be equal across all groups.

**gsem: Adding constraints for means, variances, or covariances across groups**

You use the same technique for adding constraints to means, variances, and covariances as you would for adding constraints to path coefficients. Remember that means are specified by the `means()` option, variances by the `variance()` option, and covariances by the `covariance()` option. The `variance()` and `covariance()` options are abbreviated `var()` and `cov()`, respectively.

You can specify, for instance,

```
(L -> y1 y2 y3 y4), probit ///
      group(agegrp) ginvariant(cons) var(L@v)
```

to constrain the variance of `L` to be the same across groups.

**gsem: Adding constraints for some groups but not others**

Again, we consider the following model:

```
(L -> y1 y2 y3 y4), probit ///
      group(agegrp) ginvariant(cons)
```

Above we saw how to constrain the L->y4 path coefficient to be the same across groups:

```
(L -> y1 y2 y3 y4@b), probit ///
      group(agegrp) ginvariant(cons)
```

To constrain the path coefficients  $L \rightarrow y_4$  to be equal for groups 1 and 2 but leave the  $L \rightarrow y_4$  path coefficients unconstrained for the remaining groups, we would type

```
(L -> y1 y2 y3 y4) (1: L->y4@b) (2: L->y4@b), probit ///
      group(agegrp) ginvariant(cons)
```

Think of this as follows:

1. ( $L \rightarrow y_4$ ): We set a path for all the groups.
2. (1:  $L \rightarrow y_4@b$ ): We modify the path for  $\text{agegrp} = 1$ .
3. (2:  $L \rightarrow y_4@b$ ): We modify the path for  $\text{agegrp} = 2$ .
4. We do not modify the path for any other  $\text{agegrp}$  value.

The result is that we constrain age groups 1 and 2 to have the same value of the path, and we do not constrain the path for the other age groups.

You can constrain variance, covariance, and mean estimates to be the same across some groups but not others in the same way. You can specify, for instance,

```
(L -> y1 y2 y3 y4), probit ///
      group(agegrp) var(1: L@v) var(2: L@v)
```

### **gsem: Adding paths for some groups but not others**

In the same way that you can constrain coefficients for some groups but not others, you can add paths for some groups but not others. Consider the following model:

```
(L -> y1 y2 y3), probit group(agegrp)
```

You can add the path  $L \rightarrow y_4$  for groups 1 and 2 by typing

```
(L -> y1 y2 y3) (1: L->y4) (2: L->y4), probit group(agegrp)
```

### **gsem: Relaxing constraints**

Just as you can specify

```
..., group(agegrp) ginvariant(pclasses)
```

and then add constraints, you can also specify

```
..., group(agegrp) ginvariant(pclasses)
```

and then relax constraints that the parameter classes impose.

For instance, if we specified `ginvariant>Loading`, then we would be constraining ( $L \rightarrow y_4$ ) to be invariant across groups. We could then relax that constraint by typing

```
(L -> y1 y2 y3 y4) (1: L->y4@b1) (2: L->y4@b2), probit ///
      group(agegrp) ginvariant(... Loading)
```

The ( $L \rightarrow y_4$ ) path coefficients would be free in groups 1 and 2 and constrained in the remaining groups, if there are any. The path coefficient is free in group 1 because we specified symbolic name `b1`, and `b1` appears nowhere else in the model. The path coefficient is free in group 2 because symbolic name `b2` appears nowhere else in the model. If there are remaining groups and we want to relax the constraint on them, too, we would need to add (3:  $L \rightarrow y_4@b3$ ), and so on.

The same technique can be used to relax constraints on means, variances, and covariances. For instance,

```
(L -> y1 y2 y3 y4), probit ///  
    group(agegrp) ginvariant(... covex) var(1: L@V1) var(2: L@V2)
```

## Reference

Acock, A. C. 2013. *Discovering Structural Equation Modeling Using Stata*. Rev. ed. College Station, TX: Stata Press.

## Also see

[SEM] [intro 5](#) — Tour of models

[SEM] [intro 7](#) — Postestimation tests and predictions

[SEM] [sem group options](#) — Fitting models on different groups

[SEM] [sem and gsem option covstructure\(\)](#) — Specifying covariance restrictions