Multivariate Analysis of Variance

If you fit several dependent variables to the same effects, you might want to make joint tests involving parameters of several dependent variables. Suppose you have *P*dependent variables, *k*parameters for each dependent variable, and *n*observations. The models can be collected into one equation:

$$\mathbf{Y} = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\varepsilon}$$

where Y is $n \times p$, X is $n \times k$, β is $k \times p$, and ϵ is $n \times p$. Each of the *P* models can be estimated and tested separately. However, you might also want to consider the joint distribution and test the *P* models simultaneously.

For multivariate tests, you need to make some assumptions about the errors. With *P*dependent variables, there are $n \times p$ errors that are independent across observations but not across dependent variables. Assume

$$\operatorname{vec}(\boldsymbol{\varepsilon}) \sim N(\boldsymbol{0}, \mathbf{I}_n \otimes \boldsymbol{\Sigma})$$

where $vec^{(\varepsilon)}$ strings ε out by rows, \otimes denotes Kronecker product multiplication, and Σ is $p \times p$. Σ can be estimated by

$$\mathbf{S} = \frac{\mathbf{e}'\mathbf{e}}{n-r} = \frac{(\mathbf{Y} - \mathbf{X}\mathbf{b})'(\mathbf{Y} - \mathbf{X}\mathbf{b})}{n-r}$$

where $\mathbf{b} = (\mathbf{X}'\mathbf{X})^{-}\mathbf{X}'\mathbf{Y}$, *r* is the rank of the **X** matrix, and **e** is the matrix of residuals.

If Sis scaled to unit diagonals, the values in Sare called *partial correlations of the Ys adjusting for the Xs*. This matrix can be displayed by PROC GLM if <u>PRINTE</u> is specified as a <u>MANOVA</u> option.

The multivariate general linear hypothesis is written

$L\beta M = 0$

You can form hypotheses for linear combinations across columns, as well as across rows of β .

The <u>MANOVA</u> statement of the GLM procedure tests special cases where L corresponds to Type I, Type II, Type II, or Type IV tests, and **M** is the $P \times P$ identity matrix. These tests are joint tests that the given type of hypothesis holds for all dependent variables in the model, and they are often sufficient to test all hypotheses of interest.

Finally, when these special cases are not appropriate, you can specify your own Land Mmatrices by using the <u>CONTRAST</u> statement before the <u>MANOVA</u> statement and the <u>M=</u> specification in the <u>MANOVA</u> statement, respectively. Another alternative is to use a <u>REPEATED</u> statement, which automatically generates a variety of ^Mmatrices useful in repeated measures analysis of variance. See the section <u>REPEATED</u> statement and the section <u>Repeated Measures Analysis of Variance</u> for more information.

One useful way to think of a MANOVA analysis with an **M** matrix other than the identity is as an analysis of a set of transformed variables defined by the columns of the **M** matrix. You should note, however, that PROC GLM always displays the **M** matrix in such a way that the transformed variables are defined by the rows, not the columns, of the displayed **M** matrix.

All multivariate tests carried out by the GLM procedure first construct the matrices **H**and **E**corresponding to the numerator and denominator, respectively, of a univariate *F* test:

$$H = M'(Lb)'(L(X'X)^{-}L')^{-1}(Lb)M$$

$$\mathbf{E} = \mathbf{M}'(\mathbf{Y}'\mathbf{Y} - \mathbf{b}'(\mathbf{X}'\mathbf{X})\mathbf{b})\mathbf{M}$$

The diagonal elements of **H**and **E**correspond to the hypothesis and error SS for univariate tests. When the **M**matrix is the identity matrix (the default), these tests are for the original dependent variables on the left side of the <u>MODEL</u> statement. When an **M**matrix other than the identity is specified, the tests are for transformed variables defined by the columns of the **M**matrix. These tests can be studied by requesting the SUMMARY option, which produces univariate analyses for each original or transformed variable.

Four multivariate test statistics, all functions of the eigenvalues of $\mathbf{E}^{-1}\mathbf{H}$ (or $(\mathbf{E}+\mathbf{H})^{-1}\mathbf{H}$), are constructed:

Wilks' lambda = det(\mathbf{E})/det($\mathbf{H} + \mathbf{E}$) Pillai's trace = trace($\mathbf{H}(\mathbf{H} + \mathbf{E})^{-1}$) Hotelling-Lawley trace = trace($\mathbf{E}^{-1}\mathbf{H}$) Roy's greatest root = λ , largest eigenvalue of $\mathbf{E}^{-1}\mathbf{H}$

By default, all four are reported with *p*-values based on *F* approximations, as discussed in the "Multivariate Tests" section in Chapter 4, <u>Introduction to Regression Procedures</u>. Alternatively, if you specify MSTAT=EXACT in the associated <u>MANOVA</u> or <u>REPEATED</u> statement, *p*-values for three of the four tests are computed exactly (Wilks' lambda, the Hotelling-Lawley trace, and Roy's greatest root), and the *p*-values for the fourth (Pillai's trace) are based on an *F* approximation that is more accurate than the default. See the "Multivariate Tests" section in Chapter 4, <u>Introduction to Regression Procedures</u>, for more details on the exact calculations.

MANOVA Statement

MANOVA <test-options> <detail-options> ;

If the <u>MODEL</u> statement includes more than one dependent variable, you can perform multivariate analysis of variance with the MANOVA statement. The *test-options* define which effects to test, while the *detail-options* specify how to execute the tests and what results to display.

When a MANOVA statement appears before the first RUN statement, PROC GLM enters a multivariate mode with respect to the handling of missing values; in addition to observations with missing independent variables, observations with *any* missing dependent variables are excluded from the analysis. If you want to use this mode of handling missing values and do not need any multivariate analyses, specify the <u>MANOVA</u> option in the <u>PROC GLM</u> statement.

If you use both the <u>CONTRAST</u> and MANOVA statements, the MANOVA statement must appear after the <u>CONTRAST</u> statement.

Test Options

The following options can be specified in the MANOVA statement as *test-options* in order to define which multivariate tests to perform.

H=effects | INTERCEPT | _ALL_

specifies effects in the preceding model to use as hypothesis matrices. For each **H**matrix (the SSCP matrix associated with an effect), the H= specification displays the characteristic roots and vectors of $\mathbf{E}^{-1}\mathbf{H}$ (where **E** is the matrix associated with the error effect), along with the Hotelling-Lawley trace, Pillai's trace, Wilks' lambda, and Roy's greatest root. By default, these statistics are tested with approximations based on the *F* distribution. To test them with exact (but computationally intensive) calculations, use the <u>MSTAT=</u>EXACT option.

Use the keyword INTERCEPT to produce tests for the intercept. To produce tests for all effects listed in the <u>MODEL</u> statement, use the keyword _ALL_ in place of a list of effects.

For background and further details, see the section Multivariate Analysis of Variance.

E=effect

specifies the error effect. If you omit the E= specification, the GLM procedure uses the error SSCP (residual) matrix from the analysis.