

One-way MANOVA Model

- Data consist of g independent random samples $\{\mathbf{y}_{ij}\}_{1 \leq i \leq n_j}$ of sizes n_1, \dots, n_g from g groups or populations
- The additive linear model is $\mathbf{y}_{ij} = (\boldsymbol{\mu} + \boldsymbol{\alpha}_j) + \{\boldsymbol{\epsilon}_{ij}\}$, $j = 1, \dots, g$
 \mathbf{y}_{ij} , $\boldsymbol{\mu}$, $\boldsymbol{\alpha}_j$ and $\boldsymbol{\epsilon}_{ij}$ all $p \times 1$ and $E[\boldsymbol{\epsilon}_{ij}] = \mathbf{0}$.

The other assumptions are:

- Equal variance matrices
 $\boldsymbol{\Sigma}_1 = \boldsymbol{\Sigma}_2 = \dots = \boldsymbol{\Sigma}_g = \boldsymbol{\Sigma} = [\sigma_{\ell m}]$
 with $\boldsymbol{\Sigma}_j = [\sigma_{\ell m}^{(j)}] = V[\boldsymbol{\epsilon}]$ for group j .

Equality of $\boldsymbol{\Sigma}$'s is strong condition:

1. Equal variances among groups
 $\sigma_{\ell \ell}^{(1)} = \sigma_{\ell \ell}^{(2)} = \dots = \sigma_{\ell \ell}^{(g)} = \sigma_{\ell \ell}$, $\ell = 1, \dots, p$
 2. Equal correlations among groups
 $\rho_{\ell m}^{(1)} = \rho_{\ell m}^{(2)} = \dots = \rho_{\ell m}^{(g)} = \rho_{\ell m}$, $1 \leq \ell \neq m \leq p$
- Exact small sample inference requires that $\boldsymbol{\epsilon}$ is $N_p(\mathbf{0}, \boldsymbol{\Sigma})$.

Displays for Statistics 5401/8401

Lecture 17

October 14, 2005

Christopher Bingham, Instructor

612-625-1024, kb@umn.edu

372 Ford Hall

Class Web Page

<http://www.stat.umn.edu/~kb/classes/5401>

© 2005 by Christopher Bingham

You can also parametrize the one-way MANOVA model in terms of group mean vectors

$$\mu_1 = \mu + \alpha_1, \dots, \mu_g = \mu + \alpha_g$$

instead of a grand mean μ and effects α_j :

$$y_{ij} = \mu_j + \epsilon_{ij}$$

$$y_{ij}, \mu_j, \epsilon_{ij} \text{ all } p \times 1.$$

MANACOVA - Multivariate ANACOVA

$$y_{ij} = \mu + Z_{ij,1} \beta_1 + Z_{ij,2} \beta_2 + \dots + Z_{ij,k} \beta_k + \alpha_j + \epsilon_{ij}$$

- The Z's are covariates
- The β 's don't differ among groups.
- $\Sigma = V[\epsilon]$ is constant and doesn't depend on group or any of the Z_j 's.

The standard approach to multivariate linear models assumes the same model for every variable.

Regression:

$$y_i = \beta_0 + \beta_1 Z_{i1} + \dots + \beta_k Z_{ik} + \epsilon_i$$

is equivalent to p univariate regressions

$$y_{i\ell} = \beta_{0\ell} + \beta_{1\ell} Z_{i1} + \dots + \beta_{k\ell} Z_{ik} + \epsilon_{i\ell}$$

$$\ell = 1, \dots, p$$

all with the same predictors.

2 factor MANOVA

$$y_{ij} = \mu + \alpha_i + \beta_j + (\alpha\beta)_{ij} + \epsilon_{ij}$$

is equivalent to p univariate ANOVA models

$$y_{ij\ell} = \mu + \alpha_{i\ell} + \beta_{j\ell} + (\alpha\beta)_{ij\ell} + \epsilon_{ij\ell}$$

all with the both main effects and interaction.

The situation when different variables have different models is called Seemingly Unrelated Regression or **SUR**. The best estimates are *not* least squares.

For all these models, the ϵ_i 's or ϵ_{ij} 's are assumed to have these properties in decreasing order of importance (most important first)

- 1 $E[\epsilon] = \mathbf{0}$
- 2 Independent cases (data matrix rows)
- 3 $V[\epsilon] = \Sigma$ (constant variance)
- 4 $\epsilon \sim N_p(\mathbf{0}, \Sigma)$ Needed for "exact" small sample inference .

Most tests and confidence procedures related to elements of \mathbf{B} are resistant to non-normality - they "work as advertised" adequately even with non-normal ϵ 's.

The assumption that $E[\epsilon] = \mathbf{0}$ is really just a statement that the fixed part of the model is correct. That's why I list it as the most important assumption.

You can put any multivariate linear model (regression, MANOVA, MANACOVA) in the form of a multivariate linear regression (involving "dummy" variables for MANOVA and MANACOVA).

This means you can express *all* the models in the form

$$Y = (ZB) + \{\epsilon\}, N \text{ by } p$$

- $Y = [Y_1, Y_2, \dots, Y_p]$, N by p matrix of response (dependent) variables
- $Z = [Z_0, Z_1, \dots, Z_k]$ is a n by k+1 matrix of predictor (independent) variables, possibly including dummy variables

- $B = [\beta_{j\ell}] = \begin{bmatrix} \beta_0' \\ \beta_1' \\ \dots \\ \beta_k' \end{bmatrix} = [b_1 \ b_2 \ \dots \ b_p]$

is a k+1 by p matrix of coefficients.

Each row β_ℓ' of \mathbf{B} goes with a predictor Z_ℓ . Each column \mathbf{b}_m of \mathbf{B} goes with a response variable Y_m .

One-way MANOVA $\mathbf{B} = \begin{bmatrix} \mu' \\ \alpha_1' \\ \alpha_2' \\ \dots \\ \alpha_{g-1}' \\ \alpha_g' \end{bmatrix}$

Linking with the general notation, $k = g$

$$\beta_0 = \mu, \beta_1 = \alpha_1, \dots, \beta_g = \alpha_g$$

$$\mathbf{b}_\ell = \begin{bmatrix} \mu_\ell \\ \alpha_{1\ell} \\ \dots \\ \alpha_{g\ell} \end{bmatrix}, \ell = 1, \dots, p$$

Caution: The \mathbf{Z} matrix for this parameter matrix is not full rank. It is, if \mathbf{B} omits the last row (α_g').

Estimation

For normal errors, it turns out that the best way (maximum likelihood) to estimate \mathbf{B} is by univariate ordinary least squares (OLS) for each column of \mathbf{B}

$\mathbf{b}_\ell = [\beta_{0\ell}, \beta_{1\ell}, \beta_{2\ell}, \dots, \beta_{k\ell}]', \ell = 1, \dots, p$.
separately.

The matrix formula for the univariate OLS estimates is

$$\hat{\mathbf{b}}_\ell \equiv [\hat{\beta}_{0\ell}, \hat{\beta}_{1\ell}, \dots, \hat{\beta}_{k\ell}]' = (\mathbf{Z}'\mathbf{Z})^{-1}\mathbf{Z}'\mathbf{Y}_\ell, \ell = 1, \dots, p$$

This assumes \mathbf{Z} is of full rank so $\mathbf{Z}'\mathbf{Z}$ is invertible and the coefficients are all estimable.

You can combine these into one matrix equation:

$$\hat{\mathbf{B}} = [\hat{\mathbf{b}}_1, \hat{\mathbf{b}}_2, \dots, \hat{\mathbf{b}}_p] = (\mathbf{Z}'\mathbf{Z})^{-1}\mathbf{Z}'\mathbf{Y}, k+1 \text{ by } p$$

- $\hat{\mathbf{B}} = (\mathbf{Z}'\mathbf{Z})^{-1}\mathbf{Z}'\mathbf{Y}$ is a "clone" of the univariate formula, that is, it has the *same algebraic form*.

- $\hat{\mathbf{B}}$ maximizes the normal likelihood.

If you do the math you find that the MLE $\hat{\mathbf{B}}$ minimizes the *determinant* of the residual cross product (RCP) matrix

$$\det((\mathbf{Y}-\mathbf{ZB})'(\mathbf{Y}-\mathbf{ZB})) = \det(\text{RCP}).$$

The matrix $\mathbf{Y} - \mathbf{ZB}$ consists of residuals from the regression

Math shows that $\hat{\mathbf{B}}$ also minimizes all the diagonal elements of RCP, the residual sums of squares..

In the **SUR** situation (different models for different variables), although the maximum likelihood estimates minimize $\det(\text{RCP})$, the solution isn't the same as the univariate least squares estimates.

Sampling distribution of $\hat{\mathbf{B}} = (\mathbf{Z}'\mathbf{Z})^{-1}\mathbf{Z}'\mathbf{Y}$

If you know the univariate facts you know a lot.

- Univariate LS estimates are unbiased ($E[\hat{\mathbf{b}}] = \mathbf{b}$) \Rightarrow $\hat{\mathbf{B}}$ is *unbiased* ($E[\hat{\mathbf{B}}] = \mathbf{B}$).

- The variance matrix of a *column* $\hat{\mathbf{b}}_\ell$ of $\hat{\mathbf{B}}$ is (from the univariate result):

$$V[\hat{\mathbf{b}}_\ell] = \sigma_{\ell\ell}(\mathbf{Z}'\mathbf{Z})^{-1} = \sigma_{\ell\ell}\mathbf{C} = \sigma_{\ell\ell}[c_{ij}],$$

where $\mathbf{C} = [c_{ij}] = (\mathbf{Z}'\mathbf{Z})^{-1}$, and

$$\sigma_{\ell\ell} = V[\varepsilon_\ell], \ell = 1, \dots, p.$$

- The $(k+1) \times (k+1)$ matrix of covariances between elements in different *columns* of $\hat{\mathbf{B}}$ (coefficients for different variables) is

$$\text{Cov}[\hat{\mathbf{b}}_\ell, \hat{\mathbf{b}}_m] = E[(\hat{\mathbf{b}}_\ell - \mathbf{b}_m)(\hat{\mathbf{b}}_m - \mathbf{b}_m)']$$

$$p \times p \quad = \sigma_{\ell m}(\mathbf{Z}'\mathbf{Z})^{-1} = \sigma_{\ell m}\mathbf{C},$$

where $\sigma_{\ell m} = \text{Cov}[\varepsilon_\ell, \varepsilon_m], \ell \neq m$

- Each element $\hat{\beta}_{j\ell}$ in column ℓ of $\hat{\mathbf{B}}$ is a linear combination of the elements of \mathbf{Y}_ℓ .
- Each *column* $\hat{\mathbf{b}}_\ell$ (estimated coefficients for y_ℓ) is $N_{k+1}(\mathbf{b}_\ell, \sigma_{\ell\ell}(\mathbf{Z}'\mathbf{Z})^{-1})$
- Each *row* $\hat{\boldsymbol{\beta}}_j$ (estimated coefficients of Z_j for all y_ℓ 's) is $N_p(\boldsymbol{\beta}_j, c_{jj}\boldsymbol{\Sigma})$.
- All the $p(k+1)$ elements $\hat{\beta}_{j\ell}$ together are multivariate normal $N_{p(k+1)}$.

What is the variance matrix of all $p(k+1)$ estimated coefficients $\hat{\beta}_{j\ell}$?

There a neat mathematical notation you can use to describe the variance matrix of all $p \times (k+1)$ elements $\hat{\beta}_{j\ell}$:

Let

$$\mathbf{b} \equiv \text{vec}(\mathbf{B}) = \begin{bmatrix} \mathbf{b}_1 \\ \mathbf{b}_2 \\ \dots \\ \mathbf{b}_p \end{bmatrix} = [\mathbf{b}_1', \mathbf{b}_2', \dots, \mathbf{b}_p']'$$

be the length $p(k+1)$ vector obtained by stringing the columns \mathbf{b}_ℓ of \mathbf{B} one after the other. Similarly, let

$$\hat{\mathbf{b}} \equiv \text{vec}(\hat{\mathbf{B}}) = [\hat{\mathbf{b}}_1' \ \hat{\mathbf{b}}_2' \ \dots \ \hat{\mathbf{b}}_p']'$$

Then

- $\hat{\mathbf{b}}$ is $N_{p(k+1)}(\mathbf{b}, \boldsymbol{\Sigma} \otimes (\mathbf{Z}'\mathbf{Z})^{-1})$, where the $p(k+1)$ by $p(k+1)$ matrix $V[\hat{\mathbf{b}}] = \boldsymbol{\Sigma} \otimes (\mathbf{Z}'\mathbf{Z})^{-1}$ is the *Kronecker product* of $\boldsymbol{\Sigma}$ and $(\mathbf{Z}'\mathbf{Z})^{-1}$.

Vocabulary: When \mathbf{A} is a M by N matrix and \mathbf{B} is a m by n matrix, their Kronecker product is the $M \times m$ by $N \times n$ matrix

$$\mathbf{A} \otimes \mathbf{B} \equiv \begin{bmatrix} a_{11} \mathbf{B} & a_{12} \mathbf{B} & \dots & a_{1N} \mathbf{B} \\ a_{21} \mathbf{B} & a_{22} \mathbf{B} & \dots & a_{2N} \mathbf{B} \\ \dots & \dots & \dots & \dots \\ a_{M1} \mathbf{B} & a_{M2} \mathbf{B} & \dots & a_{MN} \mathbf{B} \end{bmatrix},$$

MacAnova example using `kronecker()`:

```
Cmd> a <- matrix(run(4),2); a # arbitrary 2 by 2 matrix
(1,1)      1      3      m = 2, n = 2
(2,1)      2      4

Cmd> b <- matrix(vector(1,1,1, 1,-1,0), 3); b # 3 by 2 matrix
(1,1)      1      1      m = 3, n = 2
(2,1)      1     -1
(3,1)      1      0

Cmd> kronecker(a,b) # macro distributed with MacAnova
WARNING: searching for unrecognized macro kronecker near
kronecker(
(1,1)      1      1      |      3      3
(2,1)      1     -1     |      3     -3
(3,1)      1 a[1,1]*b 0  |      3 a[1,2]*b 0
-----|-----
(4,1)      2      2      |      4      4
(5,1)      2     -2     |      4     -4
(6,1)      2 a[2,1]*b 0  |      4 a[2,2]*b 0

Cmd> dim(kronecker(a,b)) # 2*3 by 2*2 matrix
(1)      6      4
```

Facts

- $(\mathbf{A} \otimes \mathbf{B})^{-1} = \mathbf{A}^{-1} \otimes \mathbf{B}^{-1}$
- $\hat{\mathbf{b}}' \mathbf{V}[\hat{\mathbf{b}}]^{-1} \hat{\mathbf{b}} = \text{tr } \boldsymbol{\Sigma}^{-1} \hat{\mathbf{B}}' (\mathbf{Z}' \mathbf{Z})^{-1} \hat{\mathbf{B}}$
 = sum of diagonals of $\boldsymbol{\Sigma}^{-1} \hat{\mathbf{B}}' (\mathbf{Z}' \mathbf{Z})^{-1} \hat{\mathbf{B}}$

Application

Suppose $\mathbf{S} = \hat{\boldsymbol{\Sigma}}$ estimates $\boldsymbol{\Sigma}$. Then

$$T^2 = \hat{\mathbf{b}}' \hat{\mathbf{V}}[\hat{\mathbf{b}}]^{-1} \hat{\mathbf{b}} = \text{tr } \mathbf{S}^{-1} (\hat{\mathbf{B}}' (\mathbf{Z}' \mathbf{Z})^{-1} \hat{\mathbf{B}})$$

is a form of Hotelling's T^2 statistic that tests $H_0: \mathbf{B} = \mathbf{0}$, that is

$$H_0: \beta_{j\ell} = 0, j = 0, \dots, k, \ell = 1, \dots, p$$

Under wide conditions, in large samples, the null distribution of T^2 is $\chi_{p(k+1)}^2$.

There is no easy exact small sample distribution as there is for the two-sample and paired Hotelling T^2 statistics.

When $k = 0$, $T^2 \approx (p f_e / (f_e - p + 1)) F_{p, f_e - p + 1}$

When $p = 1$, $T^2 \approx F_{k+1, f_e}$

Unbiased estimate of Σ

Define the p by p **error matrix**

$$\mathbf{E} = \sum_{1 \leq i \leq N} (\mathbf{y}_i - \hat{\mathbf{y}}_i)(\mathbf{y}_i - \hat{\mathbf{y}}_i)' = (\mathbf{Y} - \mathbf{Z}\hat{\mathbf{B}})'(\mathbf{Y} - \mathbf{Z}\hat{\mathbf{B}})$$

where $\hat{\mathbf{y}}_i = \hat{\mathbf{B}}' \mathbf{z}_i = (\mathbf{z}_i' \hat{\mathbf{B}})'$ is the predicted value based on \mathbf{z}_i' , (row i of \mathbf{Z}).

- $\mathbf{Y} - \mathbf{Z}\hat{\mathbf{B}}$ is the matrix of least squares residuals.
- \mathbf{E} is the multivariate analogue of SS_e in univariate ANOVA and regression. To get a formula for \mathbf{E} , replace $(\dots)^2$ in a formula for SS_e by $(\dots)(\dots)'$.
- $e_{\ell\ell} = \sum_{1 \leq i \leq N} (y_{i\ell} - \hat{y}_{i\ell})^2 = SS_e^{(\ell)}$ (ANOVA residual sum of squares for y_ℓ)
- $e_{\ell m} = e_{m\ell} = \sum_{1 \leq i \leq N} (y_{i\ell} - \hat{y}_{i\ell})(y_{im} - \hat{y}_{im})$ (residual sum of products for y_ℓ and y_m)

Johnson and Wichern use \mathbf{W} (for **W**ithin) instead of \mathbf{E} in some contexts.

The minimum number of linearly independent parameter *vectors*, each of length p , required in the model is $r = \text{rank}(\mathbf{Z})$. If \mathbf{Z} is of full rank, $r = k+1$. Thus at least $r \times p$ parameters are required in all.

Define

- $\mathbf{S} \equiv (1/f_e)\mathbf{E} = (1/f_e)\sum_{1 \leq i \leq N} (\mathbf{y}_i - \hat{\mathbf{y}}_i)(\mathbf{y}_i - \hat{\mathbf{y}}_i)'$ where
- $f_e = N - r$ ($f_e = N - k - 1$ for full rank \mathbf{Z})

Facts:

- $E[\mathbf{S}] = \Sigma \Rightarrow \mathbf{S}$ is an unbiased estimate of Σ .
- When \mathbf{y} is MVN with $V[\mathbf{y}] = \Sigma$, \mathbf{E} is $W_p(f_e, \Sigma)$ ($\sigma^2 \chi_{f_e}^2$ when $p = 1$)

\mathbf{S} is multivariate analog of the univariate

$$s^2 = (1/f_e)\sum_{1 \leq i \leq N} (y_i - \hat{y}_i)^2$$

MacAnova MANOVA Example

```
Cmd> irisdata <- read("", "t11_05", quiet:T)
Read from file "TP1:Stat5401:Data:JWData5.txt"
Cmd> varieties <- factor(irisdata[,1])
```

Using `factor()` is essential to mark varieties as a categorical variable rather than a quantitative variable.

```
Cmd> y <- irisdata[, -1] # strip off variety numbers
Cmd> list(varieties, y)
varieties      REAL    150    1      FACTOR with 3 levels
y              REAL    150    4      N = 150, p = 4
```

```
Cmd> manova("y=varieties") # like anova()
Model used is y=varieties
WARNING: summaries are sequential
```

SS and SP Matrices				
	DF			
CONSTANT	1			
SepLen	5121.7	SepWid	PetLen	PetWid
SepWid	2679.8	1402.1	1723.4	550.01
PetLen	3293.9	1723.4	2118.4	676.06
PetWid	1051.2	550.01	676.06	215.76
varieties	2			
SepLen	63.212	SepWid	PetLen	PetWid
SepWid	-19.953	11.345	-57.24	-22.933
PetLen	165.25	-57.24	437.1	186.77
PetWid	71.279	-22.933	186.77	80.413
ERROR1	147			
SepLen	38.956	SepWid	PetLen	PetWid
SepWid	13.63	16.962	8.1208	4.8084
PetLen	24.625	8.1208	27.223	6.2718
PetWid	5.645	4.8084	6.2718	6.1566

This is default `manova()` output when $p \leq 5$.

Hypothesis matrix

$$H = B = \sum_{1 \leq j \leq g} n_j (\bar{y}_{.j} - \bar{y}_{..}) (\bar{y}_{.j} - \bar{y}_{..})'$$

This generalizes the univariate formula

$$SS_h = SSB = \sum_{1 \leq j \leq g} n_j (\bar{y}_{.j} - \bar{y}_{..})^2$$

Error matrix is multiple of pooled variance matrix estimate

$$E = W = \sum_{1 \leq j \leq g} (n_j - 1) S_j$$

$$S = S_{pooled} = (N - g)^{-1} \sum_{1 \leq j \leq g} (n_j - 1) S_j$$

This generalizes the univariate formula

$$s_{pooled}^2 = (N - g)^{-1} \sum_{1 \leq j \leq g} (n_j - 1) s_j^2$$

MacAnova computes variables `DF`, `RESIDUALS` and `SS` just as `anova()` and `regress()` do.

```
Cmd> list(DF, RESIDUALS, SS)
DF          REAL    3      (labels)
RESIDUALS   REAL   150    4      (labels)
SS          REAL    3      4      4      (labels)

Cmd> DF # computed by manova(); same as anova() DF
CONSTANT varieties ERROR1
      1         2         147
```

```

Cmd> SS # 3 by 4 by 4 array; also computed by manova()
          SepLen      SepWid      PetLen      PetWid
CONSTANT SepLen      5121.7      2679.8      3293.9      1051.2
          SepWid      2679.8      1402.1      1723.4      550.01
          PetLen      3293.9      1723.4      2118.4      676.06
          PetWid      1051.2      550.01      676.06      215.76
varieties SepLen      63.212     -19.953     165.25      71.279
          SepWid      -19.953     11.345     -57.24      -22.933
          PetLen      165.25      -57.24      437.1       186.77
          PetWid      71.279      -22.933     186.77      80.413
ERROR1    SepLen      38.956      13.63      24.625      5.645
          SepWid      13.63      16.962     8.1208      4.8084
          PetLen      24.625      8.1208     27.223      6.2718
          PetWid      5.645      4.8084     6.2718      6.1566
    
```

ss is a 3 dimensional array, with the first subscript indexing matrices.

```

Cmd> list(SS) # SS is a three dimension array
SS          REAL      3      4      4      (labels)
Cmd> e <- SS[3,,]; e # third matrix E; SS[2,,] is H
          SepLen      SepWid      PetLen      PetWid
ERROR1    SepLen      38.956      13.63      24.625      5.645
          SepWid      13.63      16.962     8.1208      4.8084
          PetLen      24.625      8.1208     27.223      6.2718
          PetWid      5.645      4.8084     6.2718      6.1566
    
```

The diagonal elements of `ss[j,,]` are the univariate SS:

```

Cmd> ss <- SS # save it
Cmd> anova("{y[,3]} = varieties") # univariate ANOVA
Model used is {y[,3]} = varieties
          DF      SS      MS
CONSTANT      1      2118.4      2118.4
varieties      2      437.1      218.55
ERROR1        147      27.223      0.18519
Cmd> ss[,3,3] # 3rd diagonal element of matrices
          PetLen
CONSTANT PetLen      2118.4
varieties PetLen      437.1
ERROR1    PetLen      27.223
    
```

MacAnova computes MANOVA as multi-variate regression with dummy variables with values 0, 1 and -1. You can see what they are using through `modelinfo()`. Here is an example with "toy" data, $g = 3$, $p = 3$, $N = 10$.

```

Cmd> a <- factor(1,1,1,2,2,2,3,3,3,3) # n_1=3, n_2=3, n_3=4
Cmd> Y <- matrix(rnorm(30),10) # N = 10, p = 3
Cmd> manova("Y = a", silent:T)
Cmd> xvariables() # gets the actual Z matrix used
(1,1)      1      1      0
(2,1)      1      1      0
(3,1)      1      1      0
(4,1)      1      0      1
(5,1)      1      0      1
(6,1)      1      0      1
(7,1)      1     -1     -1
(8,1)      1     -1     -1
(9,1)      1     -1     -1
(10,1)     1     -1     -1
    
```

Basic confidence interval for one coefficient

A multivariate linear model can always be put in the form

$$Y = ZB + \epsilon, E[\epsilon] = 0, V[\epsilon] = \Sigma$$

Y and ϵ n by p , Z N by $k+1$,

$$B = [b_1, \dots, b_p] = [\beta_0, \beta_1, \dots, \beta_k]' \text{ } k+1 \text{ by } p$$

Let $C = [c_{ij}] = (Z'Z)^{-1}$. Then

$$V[\hat{b}_\ell] = \sigma_{\ell\ell} C, \ell = 1, \dots, p$$

In particular

$$V[\hat{\beta}_{j\ell}] = c_{jj} \sigma_{\ell\ell}, j = 0, \dots, k, \ell = 1, \dots, p$$

The estimated standard error of $\hat{\beta}_{j\ell}$ is

$$SE[\hat{\beta}_{j\ell}] = \sqrt{\{c_{jj} \hat{\sigma}_{\ell\ell}\}}$$

where $\hat{\sigma}_{\ell\ell}$, is the MSE for y_ℓ , and is a diagonal element of $\hat{\Sigma} = S = (1/f_e)E$.

MacAnova You can use `secoefs()` to retrieve all $\hat{\beta}_{j\ell}$'s and all $SE[\hat{\beta}_{j\ell}]$.

There are several different $100(1 - \alpha)\%$ confidence intervals for a coefficient $\beta_{j\ell}$, both "ordinary" (non-simultaneous) and simultaneous.

All have the form

$$\beta_{\ell j} = \hat{\beta}_{\ell j} \pm K_\alpha \sqrt{\{c_{jj} \hat{\sigma}_{\ell\ell}\}}, \text{ with constant } K_\alpha \text{ which depends on the type of interval}$$

- *Single non-simultaneous large sample* confidence interval has $K_\alpha = z(\alpha/2)$
- *Single non-simultaneous* confidence interval with normal or near normal errors has $K_\alpha = t_{f_e}(\alpha/2)$.
- *Simultaneous* intervals for *all* $M = (k+1)p$ coefficients by Bonferonizing Student's t by M :
 $K_\alpha = z((\alpha/M)/2)$ or $K_\alpha = t_{f_e}((\alpha/M)/2)$.

Example

```
Cmd> manova("y=varieties",silent:T)
```

```
Cmd> coefs()#describes most recent regress(), anova(), manova()
```

```
component: CONSTANT      Least squares estimates of  $\mu$ 
```

	SepLen	SepWid	PetLen	PetWid	$\hat{\mu}'$
(1)	5.8433	3.0573	3.758	1.1993	

component: varieties **Least squares of variety effects**

	SepLen	SepWid	PetLen	PetWid	$\hat{\alpha}'_1$
(1)	-0.83733	0.37067	-2.296	-0.95333	
(2)	0.092667	-0.28733	0.502	0.12667	$\hat{\alpha}'_2$
(3)	0.74467	-0.083333	1.794	0.82667	$\hat{\alpha}'_3$

```
Cmd> stats <- secoefs(); stats
```

```
component: CONSTANT      Estimates and their standard errors
```

```
component: coefs
```

	SepLen	SepWid	PetLen	PetWid
(1)	5.8433	3.0573	3.758	1.1993

```
component: se
```

	SepLen	SepWid	PetLen	PetWid
(1)	0.042032	0.027735	0.035137	0.01671

component: varieties

```
component: coefs      Alphahats
```

	SepLen	SepWid	PetLen	PetWid
(1)	-0.83733	0.37067	-2.296	-0.95333
(2)	0.092667	-0.28733	0.502	0.12667
(3)	0.74467	-0.083333	1.794	0.82667

```
component: se      Standard errors of alphahats
```

	SepLen	SepWid	PetLen	PetWid
(1)	0.059443	0.039224	0.049691	0.023631
(2)	0.059443	0.039224	0.049691	0.023631
(3)	0.059443	0.039224	0.049691	0.023631

(to be continued)