6. <u>AN</u>alysis <u>Of VA</u>riance (ANOVA)

Setup and Notation

$$y = X\beta + \epsilon, \ \epsilon \sim N(0, \ \sigma^2 I)$$

Let $X_1 = 1$, $X_m = X$, and $X_{m+1} = I$.

Suppose X_2, \ldots, X_m are matrices satisfying

$$C(X_1) \subset C(X_2) \subset \cdots \subset C(X_{m-1}) \subset C(X_m).$$

Let
$$P_j = P_{X_j}$$
 and $r_j = \operatorname{rank}(X_j) \ \forall \ j = 1, \dots, m+1$.

The Total Sum of Squares

The total sum of squares (also known as the corrected total sum of squares) is

$$\sum_{i=1}^{n} (y_{i} - \bar{y}_{.})^{2} = \begin{bmatrix} y_{1} - \bar{y}_{.} \\ \vdots \\ y_{n} - \bar{y}_{.} \end{bmatrix}' \begin{bmatrix} y_{1} - \bar{y}_{.} \\ \vdots \\ y_{n} - \bar{y}_{.} \end{bmatrix} = [\mathbf{y} - \bar{y}_{.}\mathbf{1}]'[\mathbf{y} - \bar{y}_{.}\mathbf{1}]$$

$$= [\mathbf{y} - \mathbf{P}_{1}\mathbf{y}]'[\mathbf{y} - \mathbf{P}_{1}\mathbf{y}] = [\mathbf{I}\mathbf{y} - \mathbf{P}_{1}\mathbf{y}]'[\mathbf{I}\mathbf{y} - \mathbf{P}_{1}\mathbf{y}]$$

$$= [(\mathbf{I} - \mathbf{P}_{1})\mathbf{y}]'[(\mathbf{I} - \mathbf{P}_{1})\mathbf{y}] = \mathbf{y}'(\mathbf{I} - \mathbf{P}_{1})'(\mathbf{I} - \mathbf{P}_{1})\mathbf{y}$$

$$= \mathbf{y}'(\mathbf{I} - \mathbf{P}_{1})(\mathbf{I} - \mathbf{P}_{1})\mathbf{y} = \mathbf{y}'(\mathbf{I} - \mathbf{P}_{1})\mathbf{y}.$$

Partitioning the Total Sum of Squares

$$\sum_{i=1}^{n} (y_{i} - \bar{y}_{.})^{2} = y'(I - P_{1})y = y'(P_{m+1} - P_{1})y$$

$$= y'\left(\sum_{j=2}^{m+1} P_{j} - \sum_{j=1}^{m} P_{j}\right)y$$

$$= y'(P_{m+1} - P_{m} + P_{m} - P_{m-1} + \cdots + P_{2} - P_{1})y$$

$$= y'(P_{m+1} - P_{m})y + \ldots + y'(P_{2} - P_{1})y$$

$$= \sum_{j=1}^{m} y'(P_{j+1} - P_{j})y.$$

The sums of squares in the equation

$$\mathbf{y}'(\mathbf{I} - \mathbf{P}_1)\mathbf{y} = \sum_{j=1}^m \mathbf{y}'(\mathbf{P}_{j+1} - \mathbf{P}_j)\mathbf{y}$$

are often arranged in an ANOVA table.

Some Additional Sum of Squares Notation

Sum of Squares	Sum of Squares
$\mathbf{y}'(\mathbf{P}_2 - \mathbf{P}_1)\mathbf{y}$	$SS(2 \mid 1)$
$y'(P_3-P_2)y$	SS(3 2)
:	:
$\mathbf{y}'(\mathbf{P}_m - \mathbf{P}_{m-1})\mathbf{y}$	$SS(m \mid m-1)$
$y'(P_{m+1}-P_m)y$	$SSE = y'(I - P_X)y$
$y'(I-P_1)y$	$SSTo = \sum_{i=1}^{n} (y_i - \bar{y}_{\cdot})^2$

Note that

$$SS(j+1 \mid j) = y'(P_{j+1} - P_j)y$$

$$= y'(P_{j+1} - P_j + I - I)y$$

$$= y'(I - P_j - I + P_{j+1})y$$

$$= y'(I - P_j)y - y'(I - P_{j+1})y$$

$$= SSE_j - SSE_{j+1}$$

Thus, $SS(j+1 \mid j)$ is the amount the error sum of square decreases when y is projected onto $C(X_{j+1})$ instead of $C(X_j)$.

 $SS(j+1 \mid j), \ j=1, \ \dots, \ m-1$ are called Sequential Sums of Squares.

SAS calls these Type I Sums of Squares.

Properties of the Matrices of the Quadratic Forms

The matrices of the quadratic forms in the ANOVA table have several useful properties:

- Symmetry
- Idempotency
- $rank(P_{j+1} P_j) = r_{j+1} r_j$
- Zero Cross-Products

Symmetry and Idempotency

Note that $\forall j = 1, \ldots, m$

$$(\mathbf{P}_{j+1} - \mathbf{P}_j)' = \mathbf{P}'_{j+1} - \mathbf{P}'_j = \mathbf{P}_{j+1} - \mathbf{P}_j$$

and

$$(P_{j+1} - P_j)(P_{j+1} - P_j) = P_{j+1}P_{j+1} - P_{j+1}P_j - P_jP_{j+1} + P_jP_j$$

 $= P_{j+1} - P_j - P_j + P_j$
 $= P_{j+1} - P_j.$

By idempotency and symmetry,

$$y'(P_{j+1} - P_j)y = y'(P_{j+1} - P_j)(P_{j+1} - P_j)y$$

$$= y'(P_{j+1} - P_j)'(P_{j+1} - P_j)y$$

$$= [(P_{j+1} - P_j)y]'[(P_{j+1} - P_j)y]$$

$$= ||(P_{j+1} - P_j)y||^2$$

$$= ||P_{j+1}y - P_jy||^2$$

$$\equiv ||\hat{y}^{(j+1)} - \hat{y}^{(j)}||^2$$

$$= \sum_{i=1}^{n} (\hat{y}_i^{(j+1)} - \hat{y}_i^{(j)})^2,$$

which is why we call $y'(P_{i+1} - P_i)y$ a "sum of squares."

$$\operatorname{rank}(P_{j+1} - P_j) = r_{j+1} - r_j$$

Because rank is equal to trace for idempotent matrices, we have

$$\operatorname{rank}(\boldsymbol{P}_{j+1} - \boldsymbol{P}_j) = \operatorname{tr}(\boldsymbol{P}_{j+1} - \boldsymbol{P}_j) = \operatorname{tr}(\boldsymbol{P}_{j+1}) - \operatorname{tr}(\boldsymbol{P}_j)$$

$$= \operatorname{rank}(\boldsymbol{P}_{j+1}) - \operatorname{rank}(\boldsymbol{P}_j)$$

$$= \operatorname{rank}(\boldsymbol{X}_{j+1}) - \operatorname{rank}(\boldsymbol{X}_j)$$

$$= r_{j+1} - r_j.$$

Zero Cross-Products

 $\forall j < \ell$

$$(P_{j+1} - P_j)(P_{\ell+1} - P_\ell) = P_{j+1}P_{\ell+1} - P_{j+1}P_\ell - P_jP_{\ell+1} + P_jP_\ell$$

 $= P_{j+1} - P_{j+1} - P_j + P_j$
 $= 0.$

Transposing both sides and using symmetry gives

$$(\mathbf{P}_{\ell+1} - \mathbf{P}_{\ell})(\mathbf{P}_{j+1} - \mathbf{P}_{j}) = \mathbf{0}.$$

Distribution of Scaled ANOVA Sums of Squares

Because

$$\left(\frac{\boldsymbol{P}_{j+1}-\boldsymbol{P}_{j}}{\sigma^{2}}\right)\left(\sigma^{2}\boldsymbol{I}\right)=\boldsymbol{P}_{j+1}-\boldsymbol{P}_{j}$$

is idempotent,

$$\frac{\mathbf{y}'(\mathbf{P}_{j+1} - \mathbf{P}_j)\mathbf{y}}{\sigma^2} \sim \chi^2_{r_{j+1} - r_j} \left(\frac{\boldsymbol{\beta}' \mathbf{X}'(\mathbf{P}_{j+1} - \mathbf{P}_j) \mathbf{X} \boldsymbol{\beta}}{2\sigma^2} \right)$$

for all $j = 1, \ldots, m$.

ANOVA Table with Degrees of Freedom

Sum of Squares	Degrees of Freedom	DF
$y'(P_2-P_1)y$	$\operatorname{rank}(\boldsymbol{X}_2) - \operatorname{rank}(\boldsymbol{X}_1)$	$r_2 - 1$
$y'(P_3-P_2)y$	$\operatorname{rank}(\boldsymbol{X}_3) - \operatorname{rank}(\boldsymbol{X}_2)$	$r_3 - r_2$
:	:	÷
$y'(P_m-P_{m-1})y$	$\operatorname{rank}(\boldsymbol{X}_m) - \operatorname{rank}(\boldsymbol{X}_{m-1})$	$r-r_{m-1}$
$y'(P_{m+1}-P_m)y$	$\operatorname{rank}(\boldsymbol{X}_{m+1}) - \operatorname{rank}(\boldsymbol{X}_m)$	n-r
$y'(I-P_1)y$	$\operatorname{rank}(\boldsymbol{X}_{m+1}) - \operatorname{rank}(\boldsymbol{X}_1)$	n-1

Mean Squares

For $j = 1, \ldots, m-1$, define

$$MS(j+1 \mid j) = \frac{SS(j+1 \mid j)}{r_{j+1} - r_j} = \frac{\mathbf{y}'(\mathbf{P}_{j+1} - \mathbf{P}_j)\mathbf{y}}{r_{j+1} - r_j}.$$

These sums of squares divided by their degrees of freedom are known as *mean squares*.

ANOVA Table with Mean Squares

Sum of Squares	Degrees of Freedom	Mean Square
SS(2 1)	$r_2 - 1$	<i>MS</i> (2 1)
<i>SS</i> (3 2)	$r_3 - r_2$	MS(3 2)
÷	:	:
$SS(m \mid m-1)$	$r-r_{m-1}$	MS(m m-1)
SSE	n-r	MSE
SSTo	n-1	

Independence of ANOVA Sums of Squares

Because

$$(\boldsymbol{P}_{j+1} - \boldsymbol{P}_j) (\sigma^2 \boldsymbol{I}) (\boldsymbol{P}_{\ell+1} - \boldsymbol{P}_{\ell}) = \boldsymbol{0}$$

for all $j \neq \ell$, any two ANOVA sums of squares (not including SSTo) are independent.

It is also true that the ANOVA sums of squares (not including *SSTo*) are mutually independent by Cochran's Theorem, but that stronger result is not usually needed.

ANOVA F Statistics

For $j = 1, \ldots, m-1$ we have

$$F_j = \frac{MS(j+1 | j)}{MSE} = \frac{y'(P_{j+1} - P_j)y/(r_{j+1} - r_j)}{y'(I - P_X)y/(n-r)}$$

$$\sim F_{r_{j+1}-r_j,n-r}\left(rac{oldsymbol{eta}'X'(oldsymbol{P}_{j+1}-oldsymbol{P}_j)Xoldsymbol{eta}}{2\sigma^2}
ight).$$

ANOVA Table with F Statistics

Sum of Squares	Degrees of Freedom	Mean Square	F Stat
SS(2 1)	$r_2 - 1$	<i>MS</i> (2 1)	$\overline{F_1}$
<i>SS</i> (3 2)	$r_3 - r_2$	MS(3 2)	F_2
÷	i i	:	÷
$SS(m \mid m-1)$	$r-r_{m-1}$	MS(m m-1)	F_{m-1}
SSE	n-r	MSE	
SSTo	n-1		

Relationship with Reduced vs. Full Model F Statistic

The ANOVA F_i statistic:

$$F_{j} = \frac{\mathbf{y}'(\mathbf{P}_{j+1} - \mathbf{P}_{j})\mathbf{y}/(r_{j+1} - r_{j})}{\mathbf{y}'(\mathbf{I} - \mathbf{P}_{X})\mathbf{y}/(n - r)} = \frac{MS(j+1 \mid j)}{MSE}$$

The reduced vs. full model *F* statistic:

$$F = \frac{y'(P_X - P_{X_0})y/(r - r_0)}{y'(I - P_X)y/(n - r)}$$

In general, an *F* statistic is used to test

 H_0 : "The non-centrality parameter of the F statistic is zero."

VS.

 H_A : "The non-centrality parameter of the F statistic is not zero."

The ANOVA F statistic

$$F_{j} = \frac{\mathbf{y}'(\mathbf{P}_{j+1} - \mathbf{P}_{j})\mathbf{y}/(r_{j+1} - r_{j})}{\mathbf{y}'(\mathbf{I} - \mathbf{P}_{X})\mathbf{y}/(n - r)} = \frac{MS(j+1 \mid j)}{MSE}$$

has non-centrality parameter

$$\frac{\boldsymbol{\beta}'\boldsymbol{X}'(\boldsymbol{P}_{j+1}-\boldsymbol{P}_{j})\boldsymbol{X}\boldsymbol{\beta}}{2\sigma^{2}}.$$

Thus, F_i can be used to test

$$H_{0j}:rac{oldsymbol{eta}'X'(oldsymbol{P}_{j+1}-oldsymbol{P}_{j})Xoldsymbol{eta}}{2\sigma^{2}}=0 ext{ vs. } H_{Aj}:rac{oldsymbol{eta}'X'(oldsymbol{P}_{j+1}-oldsymbol{P}_{j})Xoldsymbol{eta}}{2\sigma^{2}}
eq 0.$$

 H_{0i}

The following are equivalent ways to write the null and alternative hypotheses tested by F_j .

 $egin{aligned} eta'X'(P_{j+1}-P_j)Xeta&=0 &eta'X'(P_{j+1}-P_j)Xeta&
otag&=0 &(P_{j+1}-P_j)Xeta&\neq 0 \ P_jE(y)&=P_{j+1}E(y) &P_jE(y)&
otag&=P_{j+1}E(y) &P_{j+1}E(y) &P_{j+1}E(y) &
otag&=P_{j+1}E(y)&
otag&=P_{j+1}E(y)&$

 H_{Ai}

$$H_{0j}:(extbf{ extit{P}}_{j+1}- extbf{ extit{P}}_{j}) extbf{ extit{X}}oldsymbol{eta}= extbf{0} \quad ext{vs.} \quad H_{Aj}:(extbf{ extit{P}}_{j+1}- extbf{ extit{P}}_{j}) extbf{ extit{X}}oldsymbol{eta}
eq extbf{0}$$
 is of the form $H_{0j}: extbf{ extit{C}}_{j}^{st}oldsymbol{eta}= extbf{0} \quad ext{vs.} \quad H_{Aj}: extbf{ extit{C}}_{j}^{st}oldsymbol{eta}
eq extbf{0},$

where $C_{i}^{*} = (P_{i+1} - P_{i})X$.

As written, H_{0j} is not a testable hypothesis because C_j^* has n rows but rank $r_{j+1} - r_j < n$ (homework problem).

We can rewrite H_{0j} as a testable hypothesis by replacing C_j^* with any matrix C_j whose $q = r_{j+1} - r_j$ rows form a basis for the row space of C_i^* .

Example: Multiple Regression

$$egin{array}{lcl} m{X}_1 &=& m{1} \ m{X}_2 &=& [m{1}, \, m{x}_1] \ m{X}_3 &=& [m{1}, \, m{x}_1, \, m{x}_2] \ & dots \ m{X}_m &=& [m{1}, \, m{x}_1, \, \dots, \, m{x}_{m-1}] \end{array}$$

 $SS(j+1 \mid j)$ is the decrease in SSE that results when the explanatory variable x_j is added to a model containing an intercept and explanatory variables x_1, \ldots, x_{j-1} .

Example: Polynomial Regression

$$egin{array}{lll} X_1 &=& 1 \ X_2 &=& [1, \, x] \ X_3 &=& [1, \, x, \, x^2] \ &dots \ X_m &=& [1, \, x, \, x^2, \, \dots, \, x^{m-1}] \end{array}$$

 $SS(j+1 \mid j)$ is the decrease in SSE that results when the explanatory variable x^j is added to a model containing an intercept and explanatory variables x, x^2, \dots, x^{j-1} .

An Example in R

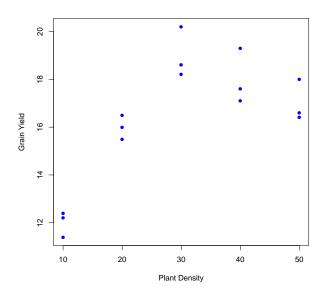
```
> #An example from "Design of Experiments: Statistical
> #Principles of Research Design and Analysis"
> #2nd Edition by Robert O. Kuehl
>
> d=read.delim("https://.../S510/PlantDensity.txt")
```

The Data

> d		
]	PlantDensity	GrainYield
1	10	12.2
2	10	11.4
3	10	12.4
4	20	16.0
5	20	15.5
6	20	16.5
7	30	18.6
8	30	20.2
9	30	18.2
10	40	17.6
11	40	19.3
12	40	17.1
13	50	18.0
14	50	16.4
15	50	16.6

Renaming the Variables and Plotting the Data

```
> names (d) = c ("x", "v")
> head(d)
  X y
1 10 12.2
2 10 11.4
3 10 12.4
4 20 16.0
5 20 15.5
6 20 16.5
>
> plot(d[,1],d[,2],col=4,pch=16,xlab="Plant Density",
 ylab="Grain Yield")
```



Matrices with Nested Column Spaces

	1		1	10		1	10	100]
	1		1	10		1	10	100	
	1		1	10		1	10	100	İ
	1		1	20		1	20	400	
	1		1	20		1	20	400	
	1		1	20		1	20	400	
	1		1	30		1	30	900	
$X_1 =$	1	$, X_2 =$	1	30	$, X_3 =$	1	30	900	١,
	1		1	30		1	30	900	
	1		1	40		1	40	1600	
	1		1	40		1	40	1600	
	1		1	40		1	40	1600	
	1		1	50		1	50	2500	
	1		1	50		1	50	2500	
	1		1	50		1	50	2500	

Matrices with Nested Column Spaces

	_			_		_				_	
	1	10	100	1000		1	10	100	1000	10000	ĺ
$X_4 =$	1	10	100	1000		1	10	100	1000	10000	
	1	10	100	1000		1	10	100	1000	10000	ı
	1	20	400	8000		1	20	400	8000	160000	ĺ
	1	20	400	8000		1	20	400	8000	160000	ĺ
	1	20	400	8000		1	20	400	8000	160000	ĺ
	1	30	900	27000		1	30	900	27000	810000	ĺ
	1	30	900	27000	$, X_{5} =$	1	30	900	27000	810000	ı
	1	30	900	27000		1	30	900	27000	810000	ĺ
	1	40	1600	64000		1	40	1600	64000	2560000	ĺ
	1	40	1600	64000		1	40	1600	64000	2560000	ĺ
	1	40	1600	64000		1	40	1600	64000	2560000	ĺ
	1	50	2500	125000		1	50	2500	125000	6250000	ĺ
	1	50	2500	125000		1	50	2500	125000	6250000	
	1	50	2500	125000		1	50	2500	125000	6250000	

Centering and Standardizing for Numerical Stability

It is typically best for numerical stability to center and scale a quantitative explanatory variable prior to computing higher order terms.

In the plant density example, we could replace x by (x-30)/10 and work with the matrices on the next two slides.

Because these matrices have the same column spaces as the original matrices, the ANOVA table entries are mathematically identical for either set of matrices.

Matrices with Centered and Scaled x

Matrices with Centered and Scaled x

$$X_4 = \begin{bmatrix}
1 & -2 & 4 & -8 \\
1 & -2 & 4 & -8 \\
1 & -1 & 1 & -1 \\
1 & -1 & 1 & -1 \\
1 & -1 & 1 & -1 \\
1 & 0 & 0 & 0 \\
1 & 0 & 0 & 0 \\
1 & 1 & 1 & 1 \\
1 & 1 & 1 & 1 \\
1 & 2 & 4 & 8 \\
1 & 2 & 4 & 8 \\
1 & 2 & 4 & 8 \\
1 & 2 & 4 & 8
\end{bmatrix}, X_5 = \begin{bmatrix}
1 & -2 & 4 & -8 & 16 \\
1 & -2 & 4 & -8 & 16 \\
1 & -2 & 4 & -8 & 16 \\
1 & -1 & 1 & -1 & 1 \\
1 & -1 & 1 & -1 & 1 \\
1 & -1 & 1 & -1 & 1 \\
1 & 0 & 0 & 0 & 0 \\
1 & 0 & 0 & 0 & 0 \\
1 & 1 & 1 & 1 & 1 \\
1 & 1 & 1 & 1 & 1 \\
1 & 1 & 1 & 1 & 1 \\
1 & 2 & 4 & 8 & 16 \\
1 & 2 & 4 & 8 & 16 \\
1 & 2 & 4 & 8 & 16 \\
1 & 2 & 4 & 8 & 16
\end{bmatrix}$$

Regardless of whether we center and scale x, the column space of X_5 is the same as the column space of the cell means model matrix

$$\boldsymbol{X} = \left[\begin{array}{cccccc} 1 & 0 & 0 & 0 & 0 \\ 1 & 0 & 0 & 0 & 0 \\ 1 & 0 & 0 & 0 & 0 \\ 1 & 0 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 & 0 \\ 0 & 0 & 1 & 0 & 0 \\ 0 & 0 & 1 & 0 & 0 \\ 0 & 0 & 1 & 0 & 0 \\ 0 & 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 0 & 1 \\ 0 & 0 & 0 & 0 & 1 \\ 0 & 0 & 0 & 0 & 1 \end{array} \right] .$$

ANOVA Table for the Plant Density Data

Source	Sum of Squares	DF
x 1	$y'(P_2-P_1)y$	2 - 1 = 1
$x^{2} 1,x$	$\mathbf{y}'(\mathbf{P}_3 - \mathbf{P}_2)\mathbf{y}$	3 - 2 = 1
$x^3 1, x, x^2$	$oldsymbol{y}'(oldsymbol{P}_4-oldsymbol{P}_3)oldsymbol{y}$	4 - 3 = 1
$x^4 1, x, x^2, x^3$	$\mathbf{y}'(\mathbf{P}_5 - \mathbf{P}_4)\mathbf{y}$	5 - 4 = 1
Error	$\mathbf{y}'(\mathbf{I}-\mathbf{P}_5)\mathbf{y}$	15 - 5 = 10
C. Total	$\mathbf{y}'(\mathbf{I} - \mathbf{P}_1)\mathbf{y}$	15 - 1 = 14

Creating the Matrices in R

```
> y=d$y
> x = (d$x-mean(d$x))/10
> x
 [1] -2 -2 -2 -1 -1 -1 0 0 0 1 1 1 2 2 2
>
> n=nrow(d)
>
> x1=matrix(1,nrow=n,ncol=1)
> x2 = cbind(x1,x)
> x3 = cbind(x2, x^2)
> x4 = cbind(x3, x^3)
> x5=matrix(model.matrix(~0+factor(x)),nrow=n)
> I=diag(rep(1,n))
```

Creating the Projection Matrices in R

Computing the Sums of Squares in R

```
> t(y) % * % (p2-p1) % * % y
   [,1]
[1,1 43,2
> t(y) % * % (p3-p2) % * % y
   [,1]
[1,] 42
> t(y) %*% (p4-p3) %*%y
   [,1]
[1,] 0.3
> t(v)%*%(p5-p4)%*%v
  [,1]
[1,] 2.1
> t(v)%*%(I-p5)%*%v
  [,1]
[1,] 7.48
> t(v)% *% (I-p1)% *%v
  [,1]
[1,] 95.08
```

The ANOVA Table in R

```
> anova(o)
Analysis of Variance Table
Response: y
        Df Sum Sq Mean Sq F value Pr(>F)
x 1 43.20 43.200 57.7540 1.841e-05 ***
I(x^2) 1 42.00 42.000 56.1497 2.079e-05 ***
I(x^3) 1 0.30 0.300 0.4011 0.5407
I(x^4) 1 2.10 2.100 2.8075 0.1248
Residuals 10 7.48 0.748
Signif. codes: 0 *** 0.001 ** 0.01 * 0.05 . 0.1 1
```

 $> o=lm(y^x+I(x^2)+I(x^3)+I(x^4), data=d)$

What do these ANOVA *F* statistics test?

1st line: Does a linear mean function fit the data significantly better than a constant mean function?

2nd line: Does a quadratic mean function fit the data significantly better than a linear mean function?

3rd line: Does a cubic mean function fit the data significantly better than a quadratic mean function?

4th line: Does a quartic mean function fit the data significantly better than a cubic mean function?

To answer each question, the error variance σ^2 is estimated from the fit of the full model with one mean for each plant density.

What do these ANOVA *F* statistics test?

In general, we have

$$H_{0j}:(\pmb{P}_{j+1}-\pmb{P}_j)\pmb{X}\pmb{\beta}=\pmb{0}$$
 vs. $H_{Aj}:(\pmb{P}_{j+1}-\pmb{P}_j)\pmb{X}\pmb{\beta}\neq\pmb{0}$

which, in testable form, is

$$H_{0j}: \boldsymbol{C}_{j}\boldsymbol{\beta} = \boldsymbol{0}$$
 vs. $H_{Aj}: \boldsymbol{C}_{j}\boldsymbol{\beta} \neq \boldsymbol{0},$

where C_j is any matrix whose $q = r_{j+1} - r_j$ rows form a basis for the row space of $(P_{j+1} - P_j)X$.

First Line of the ANOVA Table as Test of H_0 : $C\beta = 0$

```
> X=x5
> (p2-p1)%*%X
     [,1] [,2] [,3] [,4] [,5]
[1,] 0.4 0.2 0 -0.2 -0.4
[2,] 0.4 0.2 0 -0.2 -0.4
 [3,] 0.4 0.2 0 -0.2 -0.4
[4,] 0.2 0.1 0 -0.1 -0.2
[5,] 0.2 0.1 0 -0.1 -0.2
 [6,] 0.2 0.1 0 -0.1 -0.2
 [7,] 0.0 0.0 0.0 0.0
[8,] 0.0 0.0 0.0 0.0
[9,] 0.0 0.0 0 0.0 0.0
[10,] -0.2 -0.1 0 0.1 0.2
[11,] -0.2 -0.1 0 0.1 0.2
[12,] -0.2 -0.1 0 0.1 0.2
[13,] -0.4 -0.2 0 0.2 0.4
[14,] -0.4 -0.2 0 0.2 0.4
[15,] -0.4 -0.2 0 0.2 0.4
```

First Line of the ANOVA Table as Test of H_0 : $C\beta = 0$

Because

 $\operatorname{rank}[(\boldsymbol{P}_2-\boldsymbol{P}_1)\boldsymbol{X}]=\operatorname{rank}(\boldsymbol{P}_2-\boldsymbol{P}_1)=\operatorname{rank}(\boldsymbol{X}_2)-\operatorname{rank}(\boldsymbol{X}_1)=2-1=1,$ any nonzero constant times any one nonzero row of $(\boldsymbol{P}_2-\boldsymbol{P}_1)\boldsymbol{X}$ forms a basis for the row space of $(\boldsymbol{P}_2-\boldsymbol{P}_1)\boldsymbol{X}$.

For example, we could choose C to be the following one-row matrix:

Some text books would describe these as "the coefficients of a contrast to test for linear trend." (Note this is different than a test for "lack of linear fit.")

We can add consecutive lines in an ANOVA table.

Source	Sum of Squares	DF
x 1	$\mathbf{y}'(\mathbf{P}_2 - \mathbf{P}_1)\mathbf{y}$	2 - 1 = 1
$x^{2} 1,x$	$\mathbf{y}'(\mathbf{P}_3 - \mathbf{P}_2)\mathbf{y}$	3 - 2 = 1
$x^{3} 1,x,x^{2}$	$\mathbf{y}'(\mathbf{P}_4 - \mathbf{P}_3)\mathbf{y}$	4 - 3 = 1
$x^4 1, x, x^2, x^3$	$y'(P_5-P_4)y$	5 - 4 = 1
Error	$y'(I-P_5)y$	15 - 5 = 10
C. Total	$\mathbf{y}'(\mathbf{I} - \mathbf{P}_1)\mathbf{y}$	15 - 1 = 14

We can add consecutive lines in an ANOVA table.

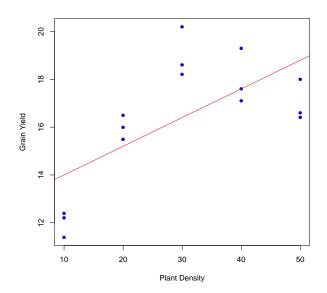
Source	Sum of Squares	DF
x 1	$y'(P_2-P_1)y$	2 - 1 = 1
$x^2, x^3, x^4, 1, x $	$\mathbf{y}'(\mathbf{P}_5 - \mathbf{P}_2)\mathbf{y}$	5 - 2 = 3
Error	$y'(I-P_5)y$	15 - 5 = 10
C. Total	$\mathbf{y}'(\mathbf{I} - \mathbf{P}_1)\mathbf{y}$	15 - 1 = 14

In this case, the combined rows test for lack of linear fit relative to a model with one unrestricted mean for each plant density.

Source	Sum of Squares	DF
x 1	$y'(P_2-P_1)y$	2 - 1 = 1
Lack of Linear Fit	$\mathbf{y}'(\mathbf{P}_5 - \mathbf{P}_2)\mathbf{y}$	5 - 2 = 3
Error	$y'(I - P_5)y$	15 - 5 = 10
C. Total	$\mathbf{y}'(\mathbf{I} - \mathbf{P}_1)\mathbf{y}$	15 - 1 = 14

```
> #Let's add the best fitting simple linear regression
> #line to our plot.
> 
> o=lm(y~x,data=d)
> 
> u=seq(0,60,by=.01) #overkill here but used later.
>
```

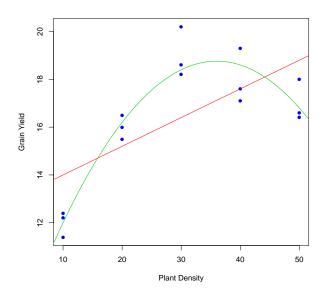
> lines(u, coef(o)[1]+coef(o)[2]*u, col=2)



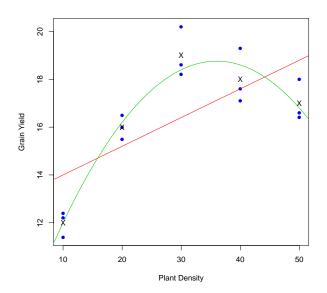
```
> #The linear fit doesn't look very good.
> #Let's formally test for lack of fit.
>
> o=lm(y~x+factor(x),data=d)
> anova(o)
Analysis of Variance Table
Response: y
         Df Sum Sq Mean Sq F value Pr(>F)
       1 43.20 43.200 57.754 1.841e-05 ***
X
factor(x) 3 44.40 14.800 19.786 0.0001582 ***
Residuals 10 7.48 0.748
Signif. codes: 0 *** 0.001 ** 0.01 * 0.05 . 0.1 1
```

```
> #It looks like a linear fit is inadequate.
> #Let's try a quadratic fit.
>
> o=lm(y^x+I(x^2)+factor(x),data=d)
> anova(o)
Analysis of Variance Table
Response: y
         Df Sum Sq Mean Sq F value Pr(>F)
x 1 43.20 43.200 57.7540 1.841e-05 ***
I(x^2) 1 42.00 42.000 56.1497 2.079e-05 ***
factor(x) 2 2.40 1.200 1.6043 0.2487
Residuals 10 7.48 0.748
Signif. codes: 0 *** 0.001 ** 0.01 * 0.05 . 0.1 1
```

```
> #It looks like a quadratic fit is adequate.
> #Let's estimate the coefficients for the best
> #quadratic fit.
>
> b=coef(lm(y~x+I(x^2),data=d))
>
> #Let's add the best fitting quadratic curve
> #to our plot.
> lines(u,b[1]+b[2]*u+b[3]*u^2,col=3)
```



```
> #Let's add the treatment group means to our plot.
>
> trt.means=tapply(d$y,d$x,mean)
>
> points(unique(d$x),trt.means,pch="X")
```



```
> #The quartic fit will pass through the treatment
> #means.
>
> b=coef(lm(y~x+I(x^2)+I(x^3)+I(x^4),data=d))
> lines(u,b[1]+b[2]*u+b[3]*u^2+b[4]*u^3+b[5]*u^4,col=1)
```

