DETERMINING THE EFFECTS OF FERTILIZATION ON BLACK SPRUCE IN THE PRESENCE OF KALMIA ANGUSTIFOLIA



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DETERMINING THE EFFECTS OF FERTILIZATION ON BLACK SPRUCE IN THE PRESENCE OF KALMIA ANGUSTIFOLIA

BY

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ABSTRACT

This study was initiated by Dr. Brain Titus who is employed with Forestry Canada. Dr. Titus was interested in the effects of fertilizers on black spruce seedlings in the presence of a shurb, *Kalmia angustifolia*, which is thought to inhibit spruce growth. Statistical analysis is presented to evaluate the differences between fertilizers for their contribution in promoting tree growth.

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TABLE OF CONTENTS

| BSTRACT | i |
|--------------------|----|
| CKNOWLEDGEMENTS | ii |
| ABLE OF CONTENTS i | ii |
| ST OF TABLE | ۷ |
| ST OF FIGURES | vi |

| CHA | PTER 1. INTRODUCTION | 1 |
|-----|--------------------------------------|----|
| 1.1 | Background | 1 |
| 1.2 | Graphical display of treatment means | 6 |
| 1.3 | Plan of the study | 16 |

| CHA | APTER 2. ANALYSIS OF DATA 1 | 7 |
|-----|----------------------------------|-----|
| 2.1 | Introduction 1 | 7 |
| 2.2 | Preliminary analysis l | 8 |
| 2.3 | ANCOVA - A regression approach 2 | 6 |
| 2.4 | Assumption testing 3 | 2 |
| 2 | .4.1 Error term is normal 3 | 2 |
| 2 | .4.2 Homogeneity of variances | \$4 |

iii

| 2.4.3 Test of parallel slopes | 36 |
|-------------------------------|--------|
| 2.4.4 Linearity of regression | 38 |

CHAPTER 3. MULTIPLE COMPARISON PROCEDURES 40

| 3.1 | Introduction | 40 |
|-----|--|----|
| 3.2 | Multiple comparison | 40 |
| 11 | 3.2.1 Fisher's LSD | 42 |
| | 3.2.2 Bryant-Paulson generalization of Tukey's HSD | 43 |
| 3 | 3.2.3 Dunn-Bonferoni test | 44 |
| - | 3.2.4 Scheffé test | 45 |
| 3.3 | Illustration | 47 |

CHAPTER 4. NONPARAMETRIC ANALYSIS OF COVARIANCE 49

| 4.1 | Introduction | 49 |
|-----|--------------|----|
| 4.2 | Rank ANCOVA | 50 |
| 4.3 | Illustration | 53 |

| REFERENCES | 57 |
|------------|----|
|------------|----|

| APPENDIX A | . 60 |
|------------|------|
| | |

LIST OF TABLES

| 2.1 | Anova summary table 23 |
|-----|--|
| 2.2 | Anova summary table |
| 2.3 | Anova summary table |
| 2.4 | Summary table for 2 ³ factorial design 31 |
| 2.5 | Anova summary table |
| 2.6 | Anova summary table 39 |
| 4.1 | Nonparametric Ancova summary table 50 |
| 4.2 | Rank Ancova summary table 54 |
| A.1 | Seedlings heights at 32 weeks |
| A.2 | Residual SS by group |
| A.3 | Conditional variances by group |
| A.4 | Raw data by group |
| A.5 | Transformed data by group |
| A.6 | Rank deviations by group |
| A.7 | Summary table by group |

LIST OF FIGURES

| 1.1 | Treatment mean for group NOO versus control OOO | 7 |
|-----|--|----|
| 1.2 | Treatment mean for group OPO versus control OOO | 8 |
| 1.3 | Treatment mean for group OOK versus control OOO | 9 |
| 1.4 | Treatment mean for group NPO versus control OOO | 10 |
| 1.5 | Treatment mean for group NOK versus control OOO | 11 |
| 1.6 | Treatment mean for group OPK versus control OOO | 12 |
| 1.7 | Treatment mean for group NPK versus control OOO | 13 |
| 1.8 | All treatment group means | 14 |
| 1.9 | Treatment mean with ${\tt N}$ versus without ${\tt N}$ | 15 |
| 2.1 | Residual histogram | 33 |

1 CHAPTER 1

INTRODUCTION

1.1 BACKGROUND

With the Newfoundland economy depending greatly on the forest industry it is important that successful reforestation be carried out on sites that have been harvested. However, within the forests of Newfoundland and other parts of Eastern Canada a shrub (Kalmia angustifolia L., hereafter referred to as Kalmia) is alleged to restrain the growth of black spruce (Picea mariana (Mill.) B.S.P.) and other varieties of trees (Mallik 1987). The Kalmia plant is a low, erect, woody shrub up to one meter in height, and is green in colour but turns a reddish-brown in late fall (Hall, Jackson and Everett 1973). It is hypothesized that Kalmia inhibits black growth through one or both of the following types of competition, Resource (or Exploitative) and Allelopathic competition. Tilman (1988) defines these types of competition as follows:

Resource competition occurs when one plant inhibits another plant through 1. consumption of limiting resources.

 Allelopathic competition occurs when one individual releases a compound that in some way inhibits growth or increases mortality of other plants.

For more discussion on these two types of competition see Walstad and Kuch (1987).

If one or both of the above types of competition is the cause of *Kalima* - induced growth inhibition in black spruce, fertilization may be a solution. To test this hypothesis and to decide which, if any, combinations of fertilizers prove effective in promoting black spruce tree growth, a greenhouse experiment was designed by Dr. B. D. Titus and Dr. A. U. Mallik.

Before the experiment was carried out, *Kalmia* plants were collected from the Botwood area on the fourteenth and fifteenth of September 1987. They were placed in pots (diameter 28.9 cm and depth 21.5 cm) and then stored at the Forestry Canada Badger Field Station waiting transportation. While stored each pot received water until they were delivered to Forestry Canada greenhouse located in St. John's on the eighteenth of September.

In the fall of the same year 240 black spruce seedlings were harvested and stored in a cold room awaiting planting. Before the seedlings were planted, seven preexperimental variables (covariates) were measured. The covariates and a brief description of each follows:

- <u>Root yolume</u>: This measurement was done by displacement i.e. the difference in the weight of a large beaker of water before and after the root system was immersed (cm³).
- 2. Total seedling fresh weight : Weight of each seedling at time of planting (g).
- 3. Root length : length of largest root (cm).
- 4. Stem length : measured from the base of the tree to the tip (cm).
- <u>First root collar diameter measurement :</u> first measurement of the seedling's stem diameter at the base of the stem (cm).
- Second root collar diameter measurement : second measurement of the secoling's stem diameter at the base of the stem (cm).
- <u>Height of seedling</u>: above ground height of each individual seedling at time of planting (cm).

The procedure for arranging the experimental units, i.e. the pots containing the Kalmia plants, within the greenhouse was as follows : 48 pots were selected from the previously collected Kalmia plants. Each of the pots was numbered from 1 to 48 and then each was assigned randomly to the six rows and eight treatments. Next, the treatment locations were randomly assigned within a row. Finally, the 240 black spruce seedlings were planted in groups of five in each pot.

Seven fertilizers and a control were used in the experiment. The fertilizers consisted of all possible combinations of three major nutrients, N (ammonium nitrate), P (super triple phosphate) and K (potash). These combinations are denoted by:

- 1. OOO Control
- 2. NOO ammonium nitrate
- 3. OPO phosphate
- 4. OOK potash
- 5. NPO ammonium nitrate + phosphate
- 6. NOK ammonium nitrate + potash
- 7. OPK phosphate + potash
- 8. NPK ammonium nitrate + phosphate + potash

The above fertilizers were used in liquid form in order to minimize the disturbance and potential damage to the seedlings, *Kalmia* and soil microbes.

The fertilizer dosage (equivalent to 150, 160 and 100 kg ha⁻¹ of elemental N, P and K, respectively) was calculated as follows:

1. Bucket diameter = 28.50 cm Bucket radius = 14.25 cm Surface area of bucket = $\pi \times r^2 = \pi \times (14.25)^2$ 2. Equivalent to:

150 kg ha⁻¹ elemental N = 0.9569 g bucket⁻¹ 160 kg ha⁻¹ elemental P = 0.3828 g bucket⁻¹ 100 kg ha⁻¹ elemental K = 0.6379 g bucket⁻¹

3. Percent nutrient content of fertilizers:

| N (ammonium nitrate) | = 34.50 % N |
|----------------------------|--------------------------------|
| P (super triple phosphate) | $= 46 \% P_2 0_5 = 20.07 \% P$ |
| K (potash) | $= 60 \% K_20 = 49.81 \% K_2$ |

4. Weight of fertilizers:

| N : | $\frac{5.7414}{x} = \frac{34.50}{100}$ | x = 16.64 g replicate ⁻¹ |
|-----|--|-------------------------------------|
| P : | $\frac{2.2968}{x} = \frac{20.07}{100}$ | x = 11.44 g replicate ⁻¹ |
| К: | $\frac{3.8274}{x} = \frac{49.81}{100}$ | x = 7.68 g replicate ⁻¹ |

The environment of the greenhouse consisted of eighteen hours of light per day at a temperature of 25 degrees celsius. In the night the temperature was lowered to 20 degrees celsius. The relative humidity of the greenhouse was kept constant at 60%. Automated watering of the seedlings was carried out twice a week in the morning for two minutes per event.

The seedlings' heights were measured every three weeks up to and including week 32, which was the termination date for the experiment.

1.2 GRAPHICAL DISPLAY OF TREATMENT MEANS

Growth curves for each of the seven fertilizer treatment means over the six rows are displayed individually with the control group over time (Figures 1 to 7). Figure 8 displays all of the treatment group means over time. From this figure one should notice that the heights attained for treatment means containing N (ammonium nitrate) tend to be greater than those that do not contain N. From this it was decided to break the treatments into two groups, the first comprised of treatments containing N and the second without N (Figure 9). By viewing Figure 9 a difference in the growth curves of these two groups is indeed tociceable, especially after the period of 24 weeks.

For the purpose of this study we will only be concentrating on the final seedlings' heights. We will be interested in the effects of the different fertilizers on the height of the seedlings at week 32.









Figure 1.3 TREATMENT MEANS VS TIME











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Figure 1.7 TREATMENT MEANS VS TIME



Figure 1.8 TREATMENT MEANS VS TIME



Figure 1.9 TREATMENT MEANS VS TIME



1.3 PLAN OF THE STUDY

Four questions were posed by Dr. Titus concerning this experiment. They were:

- 1. Is there any effect due to row positioning ?
- For future studies are all or any of the covariates listed in Section 1.1 worth measuring ?
- 3. Is there any treatment effect ?
- 4. If the answer to question 3 is yes, which treatments are more effective in promoting tree growth ?

The answers to these four questions provides the framework for this project. Chapter 2 consists of regression analysis using the final height of the seedlings at the conclusion of the experiment as the dependent variable. Least squares will be used to provide answers to Dr. Titus' first three questions. Chapter 3 is concerned with multiple comparison procedures. These procedures are useful in determining which of the treatment effects are significantly different from each other. These procedures will be used only if the answer to Dr. Titus' third question is favourable. The final chapter will provide a nonparametric analysis of the data. It will deal with ANCOVA through the use of ranks.

CHAPTER 2

DATA ANALYSIS

2.1 INTRODUCTION

The analysis of covariance (ANCOVA) procedure may be viewed as a combination of two well known statistical techniques, analysis of variance (ANOVA) and regression analysis. This statement will become clear after the ANCOVA model is defined in 2.3. Some of the main reasons for using ANCOVA are given by Huitema (1980):

"When the design involves the random assignment of subjects to treatments, the increase in power is the major pay off in selecting analysis of covariance. That is, the size of the error term is smaller with the use of ANCOVA rather than ANOVA if creating conditions are met. At the same time, the ANCOVA procedure includes an adjustment of treatment effect that reduces bias that may be caused by pretreatment differences between groups."

By using ANCOVA we reduce pretrectment differences that may exist by reducing the error term. Therefore even before an experiment begins, i.e. before treatments are administered to the subjects, there may already exist differences between the groups under

¹Huitema, Bradley E., 'Analysis of Covariance and its Alternatives', 1980, p.13

study. If group differences existed before the experiment started and one detects group differences after the experiment concludes, how can we distinguish between treatment and pretreatment effects? The following example is given to illustrate this point. Suppose we take one of our pre-experimental variables (covariates), say X_{γ} , which is the initial height of the seedling before planting. One will agree that there will exist differences in the initial heights of the seedlings simply because the seedlings' heights are not uniform, thus implying that there are differences in the pretreatment group means. At the conclusion of this experiment one may find significant dif'.vences in the treatment group means by way of ANOVA. If this happens can one attribute the significant group means to treatment effects alone or does X_{γ} play a role? One has to take the possible covariate effect into consideration.

Analysis of covariance deals with this problem simply by eliminating the covariate effect and then proceeds to analyze the data to detect differences among the adjusted treatment group means. Adjusted treatment means are defined as the treatment means after they have been adjusted for the covariate effect, i.e. after covariate effect has been removed.

2.2 PRELIMINARY ANALYSIS

One should note that all of the seedlings in one of the 48 pots used in the experiment died. In order to correct for this, an estimate of the mean value for these five seedlings will be calculated. The idea is to caculate an estimate for the missing data point and use it throughout the analysis. This estimated mean value will restore balance to the experimental design, i.e. sample sizes will all be equal among the treatment groups. The only change in the analysis is that one degree of freedom from the error term is lost. The methodology used for this calculation is discussed by Hicks (1982). For more details on the calculation of this estimate see Appendix A.

Before any analysis on treatment effects can proceed one must provide answers to the first two questions listed in 1.3. First let us recall that Question 1 asks if there is any row effect present in the data. This simply means "does the placement of the pots used in the experiment in someway affect the final height of the seedlings ?". To find the answer to this question one may use a partial F test. This test consists of fitting two models to a set of data. The first is called a full model and contains the complete set variables (k - variables) under study, and the second is referred to as the reduced mur-lel, and contains a subset of these variables (g - variables). A partial F test determines whether or not the coefficients of the g + 1 to k parameters are equal to zero. The partial F test may be summarized as follows :

COMPLETE MODEL

 $E(Y) = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \cdots + \beta_g X_g + \beta_{g+1} X_{g+1} + \cdots + \beta_k X_k.$

REDUCED MODEL

$$\mathbf{E}(\mathbf{Y}) = \boldsymbol{\beta}_0 + \boldsymbol{\beta}_1 \mathbf{X}_1 + \boldsymbol{\beta}_2 \mathbf{X}_2 + \cdots + \boldsymbol{\beta}_s \mathbf{X}_s.$$

$$H_0: \beta_{g+1} - \beta_{g+2} - \cdots - \beta_k = 0$$

 H_A : at least one of these $\neq 0$

$$F = \frac{(SSE_1 - SSE_2) / (k - g)}{SSE_2 / n - (k + 1)}$$

where

| SSE ₁ | = | sum of squared errors for the reduced model | | | |
|------------------|---|--|--|--|--|
| SSE ₂ | = | sum of squared errors for the full model | | | |
| k-g | = | the number of β parameters given by H_{0} | | | |
| k + 1 | = | the number of β parameters given by the properties model | | | |
| n | - | the number of observations | | | |

The above F follows a F distribution with degrees of freedom equal to $v_1 = k - g$ and $v_2 = [n - (k + 1)]$.

One should note that the partial F test determines whether or not a group of coefficients associated with their respective variables are equal to zero or not. If the coefficients are indeed equal to zero, further investigation can be used through the use of sequential sum of squares. The group of g + 1 to k variables each have one degree of freedom and thus their individual contribution should be addressed. In order to determine an answer to this question, the following models were developed. The full model comprises of all the variables (see p. 3) under study and takes the following form :

COMPLETE MODEL

$$\begin{array}{l} Y = \beta_{0} + \beta_{1}T_{1} + \beta_{2}T_{2} + \beta_{3}T_{3} + \beta_{4}T_{4} + \beta_{5}T_{5} + \beta_{6}T_{6} + \beta_{7}T_{7} + \\ \beta_{8}X_{7} + \beta_{9}X_{1} + \beta_{10}X_{56} + \beta_{11}X_{2} + \beta_{12}X_{3} + \beta_{13}X_{4} + \beta_{14}R_{1} + \\ \beta_{15}R_{2} + \beta_{16}R_{3} + \beta_{17}R_{4} + \beta_{18}R_{5} + \epsilon \ . \end{array}$$

where

| X ₁ | = | root volume | | | | | |
|----------------|---|---|--|--|--|--|--|
| X ₂ | = | fresh weight of seedling | | | | | |
| X3 | = | root length | | | | | |
| X4 | = | stem length | | | | | |
| X56 | = | root collar diameter, from average of $X_{\rm 5}$ and $X_{\rm 6}$ | | | | | |
| X7 | = | initial seedling height | | | | | |
| ε | = | random error | | | | | |
| R | • | $\left\{ \begin{array}{ccc} 1 & \text{if ith row} \\ 0 & 0/w \end{array} \right. \qquad \qquad i=1,2,3,4,5$ | | | | | |
| T, | - | $\begin{cases} 1 & \text{if rth treatment} \\ 0 & 0/w \end{cases} r = 1.2,3,4,5,6,7$ | | | | | |

Next let us consider the reduced model which contains a subset of the variables contained in the above model.

REDUCED MODEL

$$\begin{array}{rcl} Y = & \beta_0 + \beta_1 T_1 + \beta_2 T_2 + \beta_3 T_3 + \beta_4 T_4 + \beta_5 T_5 + \beta_6 T_6 + \beta_7 T_7 + \\ & \beta_8 X_1 + \beta_8 X_2 + \beta_{10} X_3 + \beta_{11} X_4 + \beta_{12} X_{56} + \beta_{13} X_7 + \epsilon \end{array}.$$

By comparing these two models we are in fact testing the null hypothesis $H_0 : \beta_{14} = \beta_{15} = \beta_{17} = \beta_{18} = 0$. These β coefficients represent the row effect in the model. The ANOVA tables generated from fitting the two models by least squares is summarized in Table 2.1. From this table one should note that the row effect comprises of 5 degrees of freedom with sums of squares equal to 68.413. The partial F test proves to be nonsignificant and the further partitioning of this five degrees of freedom into five separate components reveals that the position of the 48 pots does not contribute to final seedlings' heights. This provides an answer to question number 1 in Section 1.1.

| Source | df . | SS | MS | F |
|----------------|------|----------|--------|--------------------|
| Regression (r) | 13 | 845.86 | 65.07 | |
| Regression (c) | 18 | 914.637 | 50.793 | |
| Error (r) | 33 | 358.05 | 10.85 | |
| Error (c) | 28 | 289.637 | 10.344 | |
| Row effect | 5 | 68.413 | 13.683 | 1.32 ^{NS} |
| Total | 46 | 1203.910 | | |

ANOVA SUMMARY TABLE

NS = non-significant

r = reduced model

c = complete model

After eliminating the variables that represented the row effect we next bring our attention to the covariates. The second question that Dr. Titus wanted an answer to was to determine which, if any, of the covariates are important. For this we decided to test to see if the covariate effects are significantly different from zero. This question may also be answered through the method of a partial F test. To test this hypothesis, consider the following two models :

COMPLETE MODEL

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Allow and the second statements

 $\begin{array}{rcl} Y &=& \beta_{0} + \beta_{1}T_{1} + \beta_{2}T_{2} + \beta_{3}T_{3} + \beta_{4}T_{4} + \beta_{5}T_{5} + \beta_{6}T_{6} + \beta_{7}T_{7} + \\ && \beta_{4}X_{1} + \beta_{9}X_{2} + \beta_{10}X_{3} + \beta_{11}X_{4} + \beta_{12}X_{56} + \beta_{13}X_{7} + \varepsilon \,. \end{array}$

REDUCED MODEL

$$Y = \beta_{0} + \beta_{1}T_{1} + \beta_{2}T_{2} + \beta_{3}T_{3} + \beta_{4}T_{4} + \beta_{3}T_{5} + \beta_{6}T_{6} + \beta_{7}T_{7} + \varepsilon.$$

In this situation we are testing the hypothesis $H_0: \beta_4 = \beta_2 = \beta_{10} = \beta_{11} = \beta_{12} = \beta_{13} = 0$. The results of running the above two models is summarized below in Table 2.2. The partial F test shows that the overall covariate effect is non-significant but further testing reveals that the covariate X_7 by itself is highly significant. Out of the seven covariates measured before planting (see p. 3), only X_7 (initial height) is worth keeping for further analysis of the data. For future experiments of this type one may only want to measure the initial height of the seedlings.
TABLE 2.2

| Source | df | SS | MS | F |
|------------------|----|---------|--------|-------------------|
| Regression (r) | 7 | 703.09 | 100.44 | |
| Regression (c) | 13 | 845.86 | 65.07 | |
| Error (r) | 39 | 500.82 | 12.84 | |
| Error (c) | 33 | 358.05 | 10.85 | |
| Covariate effect | 6 | 142.77 | 23.795 | 2.19 |
| X, | 1 | 99.45 | 99.45 | 9.16 |
| X. | 1 | 1.23 | 1.23 | < 1 ^{NS} |
| Xu | 1 | 1.85 | 1.85 | < 1 ^{NS} |
| X | 1 | 1.60 | 1.60 | < 1 ^{NS} |
| x. | 1 | 38.25 | 38.25 | 3.52NS |
| X. | 1 | 0.35 | 0.35 | < 1 ^{NS} |
| Total | 46 | 1203.91 | | |

ANOVA SUMMARY TABLE

NS = non-significant

** = significant at $\alpha = 0.01$

r = reduced model

c = complete model

2.3 ANCOVA - A REGRESSION APPROACH

The one-v/ay analysis of covariate model with one covariate is defined by:

$$Y_{ij} = \mu + \tau_r + \gamma (X - \overline{X}) + \varepsilon_{ij},$$

where

| Yij | = | ith jth observation |
|-----------------|---|---|
| μ. | = | overall mean |
| τŗ | ~ | r th treatment effect |
| γ | = | regression coefficient for the covariate term |
| х | - | covariate of interest |
| x | H | mean of covariate of interest |
| ε _{ij} | = | random error |
| Στ, | = | 0. |
| | | |

The usual one-way analysis of variance as is for the analysis of covariance is concerned with testing the null hypothesis $H_0: \tau_1 - \tau_2 - \cdots - \tau_r - o$ for r treatment groups.

As with any other statistical technique, certain assumptions must apply. The following are four assumptions that are associated with analysis of covariance. These assumptions will be presented here and discussed later in Section 2.4. The assumptions

27

are (see Huitema (1980) and Neter and Wasserman (1974)) :

- 1. Error term has a normal distribution.
- 2. Treatment groups have equal variances.
- 3. Treatment groups have equal regression slopes.
- 4. Regression relationship is linear.

In order to test the hypothesis of equal treatment means a linear regression model was developed. Let us consider the transformation.

Next let us use r - 1 indicator variables to describe the r treatment group effects.

$$T_{1} = \begin{cases} 1 & \text{if 1st treatment is selected} \\ 0 & O/W & & \\ & & \\ & & \\ T_{r-1} = \begin{cases} 1 & \text{if } (r-1)\text{th treatment is selected} \\ 0 & O/W & \end{cases}$$

With these above modifications the one-way analysis of covariate model may be rewritten as

$$\mathbf{Y}_{ii} = \boldsymbol{\beta}_0 + \boldsymbol{\beta}_1 \mathbf{T}_1 + \cdots + \boldsymbol{\beta}_{r-1} \mathbf{T}_{r-1} + \boldsymbol{\beta}_r \mathbf{Z}_{ii} + \boldsymbol{\varepsilon}_{ij}$$

The relationship between these two model may be express by

$$egin{array}{rcl} eta_{0} &=& \mu_{1}+\tau_{r} & & & & & \\ eta_{j} &=& \tau_{j}-\tau_{r} & & & & j=1,\ldots,r-1 & & \\ eta_{r} &=& \gamma_{r} & & & & & \end{array}$$

For our experiment we have eight treatment groups and one covariate, X₇. With this information the model that we are interested in takes the following form:

$$\mathbf{Y} = \beta_0 + \beta_1 T_1 + \beta_2 T_2 + \beta_3 T_3 + \beta_4 T_4 + \beta_5 T_5 + \beta_6 T_6 + \beta_7 T_7 + \beta_8 Z + \varepsilon.$$

where

$$T_{1} = \begin{cases} 1 & \text{if Ist treatment is selected} \\ 0 & \text{O/W} & \vdots \\ T_{7} = \begin{cases} 1 & \text{if 7th treatment is selected} \\ 0 & \text{O/W} & \end{cases}$$

In order to test to see if the treatment effects are significant we simply test the null hypothesis that H_0 : $\beta_1 = \beta_2 = \beta_3 = \beta_4 = \beta_5 = \beta_6 = \beta_7 = 0$ for the above model. In order to test this hypothesis the following two models were constructed. The results of running a least squares regression procedure for the two models is summarized in Table 2.3.

29

a least squar.: + regression procedure for the two models is summarized in Table 2.3.

COMPLETE MODEL

 $\mathbf{Y} = \beta_0 + \beta_1 \mathbf{T}_1 + \beta_2 \mathbf{T}_2 + \beta_3 \mathbf{T}_3 + \beta_4 \mathbf{T}_4 + \beta_5 \mathbf{T}_5 + \beta_6 \mathbf{T}_6 + \beta_7 \mathbf{T}_7 + \beta_8 \mathbf{Z} + \varepsilon \,.$

REDUCED MODEL

$$Y = \beta_s Z + \epsilon$$
.

TABLE 2.3

ANOVA SUMMARY TABLE

| Source | df | SS | MS | F |
|------------------|----|----------|--------|------|
| Regression (r) | 1 | 116.46 | 116.46 | |
| Regression (c) | 8 | 802.55 | 100.32 | |
| Error (r) | 45 | 1087.44 | 24.17 | |
| Error (c) | 38 | 401.36 | 10.56 | |
| Treatment effect | 7 | 686.08 | 98.01 | 9.28 |
| Total | 46 | 1203.910 | - | |

** = significant at $\alpha = 0.01$

r = reduced model

c = complete model

Since the F value is significant we conclude that β_1 , β_2 , β_3 , β_4 , β_3 , β_6 , β_7 are not equal to zero. This implies that there are indeed significant treatment effects.

Another way to analyze the data is through the use of a 2^3 factorial design. Table 2.4 presents a detail breakdown of the three main treatment nutrients (N, P and K). Also present in the table is the contribution of the covariate, X₂. From this table it is clear (i) the nutrients of N and P prove to be significant and (ii) the covariate X₂ is highly significant.

In order to determine which of these treatment effects significantly differ from each other, multiple comparisons tests will be used. As previously noted this topic will be discussed and illustrated in Chapter 3. Thus Chapter 3 will provide an answer to Dr. Titus' final question.

TABLE 2.4

ANOVA TABLE FOR 23

FACTORIAL EXPERIMENT

| Source | df | SS | MS | F |
|-------------------|----|---------|--------|--------------------|
| Main effects | 3 | 602.58 | 200.86 | 19.02** |
| N | 1 | 530.14 | 530.14 | 50.20** |
| P | 1 | 67.45 | 67.45 | 6.38* |
| к | 1 | 4.99 | 4.99 | 0.47 ^{NS} |
| Two-way effects | 3 | 99.87 | 33.29 | 3.15* |
| NxP | ī | 19.69 | 19.69 | 1.86 ^{NS} |
| NxK | ī | 9.69 | 9.61 | 0.91 ^{NS} |
| PxK | ĩ | 70.52 | 70.52 | 6.67** |
| Three-way effects | | | | |
| NxPxK | 1 | 0.70 | 0.70 | 0.06 ^{NS} |
| Covariate | | | | |
| X7 | 1 | 99.39 | 99.39 | 9.40** |
| Error | 38 | 401.43 | 10.56 | 9.40 |
| Total | 46 | 1203.91 | | |
| NO and similar | | | | |

NS = non-significant

** = sigificant at $\alpha = 0.01$

* = significant at $\alpha = 0.05$

2.4 ASSUMPTION TESTING

This section is concerned with the verification of the assumptions stated in Section 2.3. It is important to verify the assumptions associated with a statistical test in order to validate the statistical analysis. If we find any departures of these assumptions we should evaluate its effect on our statistical analysis.

2.4.1 Error Term is Normal

This assumption may be verified in several ways either through a graphical display of the residuals or a more formal procedure. The histogram of the residuals associated with the fitted model (Figure 2.1) doe. ppear to be normal. Neter and Wasserman (1977) suggest that one may use a goodness of fit test to determine whether or not the error term has a normal distribution. One may either perform a chi-square or a Kolmogorov-Smirnov (K-S) test on the residuals to check this assumption. A K-S test based on the residuals yielded a p-value of 0.438, which is large enough to indicate that the assumption of normally has not been violated.

FIGURE 2.1 FREQUENCY HISTOGRAM RESIDUALS



33

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2.4.2 HOMOGENEITY OF TREATMENT GROUP VARIANCES

Huitema (1980) describes a methodology for testing the assumption of equal treatment group variances. The test is based upon Bonferroni's F_a distribution. Huitemas' procedure for this test consists of four steps :

 The residual sum of squares by group around the pooled within-group slope is computed. The formula for this quantity is :

jth group SSres =
$$(1 - r_w^2) \sum y_j^2$$

where

$$r_{w}^{2} = \frac{\sum xy_{w}}{\sqrt{\left[\sum x_{w}^{2}\right]\left(\sum y_{w}^{2}\right)}}$$

$$\sum xy_{w} - \sum xy_{1} + \sum xy_{2} + \cdots + \sum xy_{r},$$

where

$$\sum xy_j = \sum XY = \frac{\left(\sum X_j\right)\left(\sum Y_j\right)}{n_j} \qquad \text{for } j = 1, \cdots, r$$

$$\sum x_{w} - \sum x_{1}^{2} + \sum x_{2}^{2} + \cdots + \sum x_{r}^{2}$$
,

where

$$\begin{split} \sum \mathbf{x}_j &= \sum \mathbf{X}_j - \frac{\left[\sum \mathbf{X}_j\right]^2}{n_j} & \text{for } j = i, \cdots, r \\ \\ \sum \mathbf{y}_w &= \sum y_i^2 + \sum y_i^2 + \cdots + \sum y_r^2 \,, \end{split}$$

where

$$\sum \mathbf{y}_j = \sum \mathbf{Y}_j = \frac{\left[\sum \mathbf{Y}_j\right]^2}{n_j} \qquad \text{for } j = 1, \cdots, r.$$

- 2. $S^{1}_{\gamma_{1}1x}$ is calculated next, which is the estimation of the conditional variance for each of the r groups. For the jth group, $S^{2}_{\gamma_{1}1x}$ is found by dividing the residual sum of squares by its degrees of freedom $n_{j} - 1 - c$. The quantities n_{j} and c denote the sample size for the jth group and the number of covariates respectively.
- 3. The F ratio, F_B, is calculated by dividing the largest variance estimate S²_{y1x} by

the smallest value S_{v1v}^2 founded in Step 2.

4. The F value found in 3 is compared with a Bonferroni F_B value equal to

FB (ct/2, c, n_largest - 1 - c, n_smallest - 1 - c)

where c = [r(r - 1)]/2.

Complete details for this test are given in Appendix A. The value of F_B is found to be

equal to 38.14. If this value is compared with $F_{B(0,2,n,h_{segmt}-1-ch_{semthst}-1-ch}$ where c = r(r - 1)/2 = 28 we will find that it is less than $F_{B(0,5,24,4,6)} = 41.09$ and thus one may conclude that the assumption of equal conditional variances has been validated at the ten percent level.

Neter and Wasserman (1977) suggest that the assumptions of parallel slopes and linearity may also be tested by the use of the partial F test, which was discussed in third section of Chapter 2.

2.4.3. TEST OF PARALLEL SLOPES

This assumption is concern with testing to see if the slopes of the regression lines that represent the treatment groups are parallel. This is equivalent to testing to see if their is any interaction effect present in the model. If we use a partial F first to test this assumption we must first determine the complete and reduced models.

COMPLETE MODEL

$$Y = \beta_0 + \beta_1 T_1 + \beta_2 T_2 + \beta_3 T_3 + \beta_4 T_4 + \beta_5 T_5 + \beta_6 T_6 + \beta_7 T_7 + \beta_8 Z_8 + \beta_5 T_1 Z + \beta_{10} T_2 Z + \beta_{11} T_3 Z + \beta_{12} T_4 Z + \beta_{13} T_5 Z + \beta_{14} T_6 Z + \beta_{15} T_7 Z + \varepsilon$$

REDUCED MODEL

 $Y = \beta_0 + \beta_1 T_1 + \beta_2 T_2 + \beta_3 T_3 + \beta_4 T_4 + \beta_5 T_5 + \beta_6 T_6 + \beta_7 T_7 + \beta_8 Z_8 + \varepsilon.$

If we compare these two models we are testing that $H_0: \beta_p = \beta_{10} = \beta_{11} = \beta_{12} = \beta_{13} = \beta_{14} = \beta_{15} = 0$. This hypothesis is in fact testing the assumption of parallel slopes. The results obtained from running regression analysis on these two models are summarized in Table 2.5.

TABLE 2.5

| Source | df | SS | MS | F |
|--------------------|----|---------------------------|-------------------------------|--------------------|
| Regression (r) | 8 | 802.55 2.844 401.36 | 100.32 | |
| Regression (c) | 15 | | 60.856 10.56211 9.38921 | |
| Error (r) | 38 | | | |
| Error (r) | 31 | 291.066 | | |
| Interaction effect | 7 | 110.294 | 15.756 | 1.67 ^{NS} |
| T ₁ Z | 1 | 0.979 | 0.979 | < 1 |
| T ₂ Z | 1 | 4.058 | 4.058 | < 1 |
| T ₃ Z | 1 | 67.501 | 67.501 | 7.20 |
| T₄Z | 1 | 0.067 | 0.067 | < 1 |
| T,Z | 1 | 10.201 | 10.201 | 1.08 |
| TZ | 1 | 3.682 | 3.682 | < 1 |
| T ₇ Z | 1 | 23.746 | 23.746 | 2.53 |
| Total | 46 | 1203.90980 | | |

ANOVA SUMMARY TABLE

NS = non-significant

* = significant at $\alpha = 0.05$

r = reduced model

c = complete model

From the table it is clear that assumption of parallel slope has not been violated.

2.4.4. LINEARITY OF REGRESSION

The assumption of linearity of regression is concern with testing to see if there is a presence of curvature in the model. This test is in fact used to see if the curvature coefficient which is represented by β_x contained in the complete model is zero.

COMPLETE MODEL

 $Y = \beta_0 + \beta_1 T_1 + \beta_2 T_2 + \beta_3 T_3 + \beta_4 T_4 \beta_5 T_5 + \beta_6 T_6 + \beta_7 T_7 + \beta_8 Z + \beta_9 Z^2 + \varepsilon.$

REDUCED MODEL

Y - β_0 + β_1T_1 + β_2T_2 + β_3T_3 + β_4T_4 + β_5T_5 + β_5T_6 + β_7T_7 + β_8Z_8 + ε .

From Table 2.5 it is evident that the coefficient that represents possible curvature in the model is equal to zero.

38

| 39 |
|----|
| |

TABLE 2.6

| Source | DF | SS | MS | F |
|----------------------|----------|------------|--------|--------------------|
| Regression (R) | 8 | 802.55 | 100.32 | |
| Regression (C) | 9 | 825.588 | 91.732 | |
| Error | 38 | 401.36 | 10.56 | |
| Error | 37 | 378.322 | 10.225 | |
| Quadratic effect | 1 | 23.039 | 23.039 | 2.25 ^{NB} |
| Total | 46 | 1203.90980 | | |
| NS = non-significa | nt | | | |
| ** = significant at | α = 0.01 | | | |
| * = significant at o | a = 0.05 | | | |

ANOVA SUMMARY TABLE

CHAPTER 3

MULTIPLE COMPARISON PROCEDURES

3.1 INTRODUCTION

When the null hypothesis of equal treatment means has been rejected, we must conclude that at least two of the treatment means differ. One way to determine which means differ is through the use of a multiple comparison procedure.

3.2 MULTIPLE COMPARISONS

This section will present different tests along with their associated simultaneous confidence intervals that may be used to compare treatment means in ANCOVA. Four such tests are outlined here. Huitema (1980) discusses the following four procedures:

- 1. Fisher's least significant difference procedure.
- 2. Bryant-Paulson generalization of Tukey's honestly significant difference.

- 3. Dunn-Bonferoni test.
- 4. Scheffé test.

All of the above methods may be used for pairwise comparisons. A pairwise comparison simply compares two group means to see if they are differ from each other, and in fact tests the following hypotheses:

$$H_0 : \tau_{i \text{ adj}} = \tau_{j \text{ adj}}$$
$$H_1 : \tau_{i \text{ adj}} \neq \tau_{j \text{ adj}}$$

The final two methods (Dunn - Bonferoni, and Scheffé) may extend beyond simple comparisons of two means to more complex comparisons of group means. They can be used to explore linear combinations of treatment means.

Huitema (1980) suggests that the choice of which procedure to use depends upon two factors - (i) the type of comparisons, and (ii) whether or not simultaneous confidence intervals are of interest.

If simultaneous confidence intervals are not of interest but the main concern is some or all pairwise comparisons, then one should use the LSD procedure. The Bryant-Paulson generalization of Tukey's HSD procedure will be chosen if all pairwise comparisons and simultaneous confidence intervals are of interest to the experimenter. The Dunn - Bonferroni procedure is useful if the number of planned pairwise comparisons is small in number. These planned comparisons may be simple or complex in nature. Finally, the Scheffé method on the other hand should be employed if the number of planned or unplanned comparisons, regardless of complexity, is large.

3.2.1 Fisher's LSD

The following test statistic has a t distribution with N - r - 1 degrees of freedom. Y_{1eq} and Y_{jeq} are considered significantly different if the calculated value of t is greater than the absolute value of a t distribution with its associated degrees of freedom for a given α level:

$$t = \frac{\overline{Y}_{i \text{ adj}} - \overline{Y}_{j \text{ adj}}}{S_{\overline{Y}_{i \text{ adj}}} - \overline{Y}_{j \text{ adj}}}$$

where

$$S_{\overline{Y}_{un}} - \overline{Y}_{in} = \sqrt{MSres_{w} \left[\left[\frac{1}{n_{i}} + \frac{1}{n_{j}} \right] + \frac{\left(\overline{X}_{i} - \overline{X}_{i}\right)^{2}}{SS_{w_{a}}} \right]}$$

| MSresw | = | ANCOVA mean square error |
|---------------------------------|---|--|
| n _i , n _j | = | sample sizes for ith and jth groups |
| X _i , X _i | = | covariate means for the ith and jth groups |

SS_{wy} = sum of squares within groups for covariate variable

The associated simultaneous confidence interval for this test is:

$$\widetilde{\mathbf{Y}}_{i \text{ adj.}} - \widetilde{\mathbf{Y}}_{j \text{ adj}} \pm \mathbf{S}_{\widetilde{\mathbf{Y}}_{i \text{ adj}} - \widetilde{\mathbf{Y}}_{j \text{ adj}}}[t_{\alpha \ (N-3)}]$$

where $S_{\overline{Y}_{1}} - \overline{Y}_{1}$ is given as above.

3.2.2 Bryant - Paulson generalization of Tukey's HSD

The Bryant-Paulson test uses the test statistic Q₂, which is known as the generalized studentized range statistic:

$$Q_{p} = \frac{\overline{Y}_{i \text{ sd}j} - \overline{Y}_{j \text{ sd}j}}{\sqrt{MSres_{w} \left[1 + (MS_{b_{w}} / SS_{w_{w}})\right] / n}}$$

The critical value for this test is Qp(a, c, r, N - r - c) where c is the number of covariates under

study and r represents the number of treatment groups.

Simultaneous confidence intervals for the Bryant-Paulson procedure may be calculated using the formula:

$$\overline{\mathbf{Y}}_{i \text{ adj}} = \overline{\mathbf{Y}}_{j \text{ adj}} \pm \left[\text{MSres}_{\mathbf{w}} \left[1 + \frac{\text{MS}_{\mathbf{b}_i}}{\text{SS}_{\mathbf{w}_x}} \right] / n \right]$$

3.2.3 The Dunn - Bonferoni Test

This test is concerned with planned comparisons. Before the experiment is conducted the researcher may be interested in simple or complex mean comparisons. The test statistic for the Dunn-Bonferoni test is:

$$\mathbf{t}_{DB} = \frac{\mathbf{c}_{i} \left[\mathbf{Y}_{i \text{ sd}} \right] + \mathbf{c}_{2} \left[\mathbf{Y}_{2 \text{ sd}} \right] + \cdots + \mathbf{c}_{i} \left[\mathbf{Y}_{r \text{ sd}} \right]}{\left[\mathbf{MSres}_{w} \left[1 + \frac{\mathbf{MS}_{w}}{\mathbf{SS}_{wr}} \right] \left[\frac{(c_{i})^{2}}{n_{i}} + \frac{(c_{2})^{2}}{n_{2}} + \cdots + \frac{(c_{i})^{2}}{n_{r}} \right]} \right]}$$

where

Once the absolute value of t_{DB} is computed it is then compared with a critical value of $t_{DB(x,k,N-r-1)P}$ where k is the number of planned comparisons.

Simultaneous confidence intervals for the Dunn-Bonferoni procedure may be calculated from the formula:

$$\mathbf{c_1}\!\left(\mathbf{\overline{Y}_1}_{adj}\right) + \mathbf{c_2}\!\left(\mathbf{\overline{Y}_2}_{adj}\right) + \cdots + \mathbf{c_r}\!\left(\mathbf{\overline{Y}_r}_{adj}\right) \ \pm \ \mathbf{t_{DB(\alpha,k,N-r-1)}} \ \times \\$$

$$\left| \text{MSres}_{\mathbf{w}} \left[1 + \frac{\text{MS}_{\mathbf{b}_{\mathbf{x}}}}{\text{SS}_{\mathbf{w}_{\mathbf{x}}}} \right] \left[\frac{(c_1)^2}{n_1} + \frac{(c_2)^2}{n_2} + \cdots + \frac{(c_r)^2}{n_r} \right] \right.$$

3.2.4 The Scheffé Test

The test statistic for this test is:

$$F' = \frac{c_i[\mathbf{Y}_{1:eq}] + c_2[\mathbf{Y}_{2:eq}] + \cdots + c_i[\mathbf{Y}_{:eq}]}{\left[MSres_w \left[1 + \frac{MS_{\mathbf{b}_k}}{SS_{\mathbf{w}_k}} \right] \left[\frac{(c_i)^2}{n_1} + \frac{(c_2)^2}{n_2} + \cdots + \frac{(c_i)^2}{n_r} \right]} \right]$$

where

The critical value for this test is
$$\sqrt{(r-1) F_{(\alpha,r-1,N-r-1)}}$$
.

The associated simultaneous confidence intervals for Scheffé test may be obtained by using the following:

$$c_{i}[Y_{1,eq}] + c_{i}[Y_{2,eq}] + \cdots + c_{i}[Y_{r,eq}] \pm \sqrt{(r-1)F_{\alpha,r-1,N-r-1}} \times$$

$$\left| MSres_{w} \left[1 + \frac{MS_{b_{z}}}{SS_{w_{z}}} \right] \left[\frac{(c_{1})^{2}}{n_{1}} + \frac{(c_{2})^{2}}{n_{2}} + \cdots + \frac{(c_{r})^{2}}{n_{r}} \right] \right.$$

The purpose of this section is to illustrate one of the four procedures discussed in the previous section. The method that will be viewed here is the LSD procedure. In our case we are concerned with all possible pairwise comparisons regardless of their associated intervals.

Table 2.4 (see p. 31) presented a detailed breakdown of the three main treatment nutrients (N, P and K). From this table we concluded that the nutrients N and P are significant.

The adjustment means for the three nutrients groups are:

$$\overline{Y}_{N \text{ adj}} = 22.55$$

 $\overline{Y}_{P \text{ adj}} = 21.96$
 $\overline{Y}_{K \text{ adj}} = 19.24$

Before we can compare the adjusted means we need the values for the quantities MSres_w, $S_{w_{\chi}}$. The value of MSres_w is 10.56 which may be obtained from Table 3.1. The value of $S_{w_{\chi}}$ is found by performing an ANOVA over the treatment groups and has a value of 98.37. A summary of the findings are as follows:

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the state of a second provided

- 1. N and P are significantly differ from each other.
- 2. N and K are significantly differ from each other.
- 3. P and K are significantly differ from each other.

CHAPTER 4

NONPARAMETRIC ANALYSIS OF COVARIANCE

4.1 INTRODUCTION

Quade (1967) presents a method to perform a non-parametric analysis of covariance. Ranks are individually assigned to the X and Y data regardless of group membership. These associated ranks are then used to determine if the r groups under study have identical conditional population distributions. One should note that this method may be considered if (i) one is in doubt that the assumptions associated with a regular parametric ANCOVA have been strongly violated, or (ii) one may want to analyze data that take the form of ranks.

Let us recall that in Chapter 2 the assumption concerning equal group variances was significant at the 10 percent level. With this in mind one may use a non-parametric test for further analysis.

4.2 RANK ANCOVA

e - 10

Non-parametric ANCOVA is concerned with testing the hypothesis that the conditional population distribution of Y given X are the same for all the r treatment populations.

Huitema (1980) presents a twelve step procedure for calculating the following summary table.

TABLE 4.1

NONPARAMETRIC ANCOVA

SUMMARY TABLE

| Source | df | SS | MS | F |
|-----------|-------|---|----------------------------------|-----------------|
| Treatment | r - 1 | $\sum_{i=1}^{r} \left[\left(\sum_{j=1}^{n} Z_{ij} \right) / n_{j} \right]$ | $\frac{\text{SStr}}{\text{r}-1}$ | MStr MSE |
| Error | N - r | $\sum_{i=1}^{r} \sum_{j=1}^{n} Z_{ij}^{2} - \sum_{i=1}^{r} \left[\left[\sum_{j=1}^{n} Z_{ij} \right]^{2} / n_{j} \right]$ | SS N | <u>Е</u> - г |
| Total | N - 1 | $\sum_{i=1}^r \sum_{j=1}^n Z_{ij}^2$ | | |

The eight step procedure is as follows:

- STEP 1 Rank the X data regardless of group membership. Arrange the X data in ascending order and assign a rank of one to the smallest value of X, a rank of two to the next smallest and continue assigning ranks to each of the remaining observations. If two or more observation are equal an average rank may be assigned. Once the X observations have been miked proceed with the Y values.
- STEP 2 Calculate the deviation ranks of X and Y by :

$$X_{nak} - \overline{X}_{nak} = x_{nak}$$
 $Y_{nak} - \overline{Y}_{nak} = y_{nak}$

- STEP 3 Use the x_{neal}'s and y_{neal}'s found in Step 2 to calculate a Spearman rank-correlation coefficient r_p. This is equivalent to finding a Pearson correlation substituting x_{neal}'s and y_{neal}'s for the original data.
- STEP 4 An estimated deviation rank on Y (\hat{y}_{mak}) is determined by multiplying r_s by x_{mak}

$$\hat{y}_{mask} = r_s (x_{mask})$$

STEP 5 If we then subtract y_{mak} from we (\hat{y}_{mak}) will create a residual called Z.

STEP 6 Treatment sum of squares may be calculated by:

$$\sum_{i=1}^r \left[\left(\sum_{i=1}^n Z_{ij} \right)^i / n_j \right]$$

STEP 7 The error sum of squares is obtained by the following formula:

$$\sum_{i=1}^{r} \sum_{j=1}^{n} Z_{ij}^{2} - \sum_{i=1}^{r} \left[\left(\sum_{j=1}^{n} Z_{ij} \right)^{i} / n_{j} \right]$$

STEP 8 Finally we take the ratio of

Treatment sum of squares / r - 1 Error sum of squares / N - r

to give the F statistic.

The F statistic is then compared with F values with degrees of freedom r - 1 and N - r. If $f^{(n)}(u_{n+1}, s_{n})$ satistic exceeds this critical value we would conclude that the conditions of $r_{n}^{(n)}(u_{n+1}, s_{n})$ of Y given X is not the same for all of the r treatment populations. One should note that this procedure may be shorten by performing an analysis of variance on the Z observations obtained in Step 5. A one-way ANOVA on Z by treatment group will produce a summary table equivalent to the above table. The data in Table A.4 (Appendix A) will be analyzed in order to illustrate Quade's method. Table A.5 (Appendix A) shows the rankings of the original data founded in Table A.4. With the transformed data we may calculate $y_{max}s$ and $x_{mak}s$ for the observations using the following :

$$x_{rank} = X_{rank} - X_{rank} = X_{rank} - 24.5$$
$$y_{rank} = Y_{rank} - Y_{rank} = Y_{rank} - 24.5$$

This information is given in Table A6 (Appendix A). With y_{mak} and x_{mak} calculated we next find the value of the Spearman rank-order correlation coefficient, r_s . Using the SPSS/PC+ statistical package, the value of r_s is .3068. Table A7 (Appendix A) summarizes the observed $y_{mak}s$, $y_{mak}s$ and the residuals Z by group membership. From this table a one-way analysis of variance using a computer yielded the following summary table :

.

54 TABLE 4.2

Summary table

| Source | df | SS | MS | F |
|-----------|----|-----------|----------|-------|
| Treatment | 7 | 5998.1940 | 856.8849 | 14.42 |
| Error | 39 | 2346.8407 | 60.1754 | |
| Total | 46 | 8345.0347 | | |

From this table the F statistic is highly significant, indicating that the conditional distribution . Y given X differs over the treatment groups.

1.1

CHAPTER 5

SUMMARY AND CONCLUSIONS

The data for this experiment were collected through a greenhouse experiment conducted by Forestry Canada. The experiment was set up to evaluate the effects of various fertilizers had on black struce in the presence of a shrub know as *Kalmia*.

1

Partial F tests were used in Chapter 2 to provide answers to those questions concerning the significance of the covariates and treatment factors. Of the seven covariates that were measured, only X_{71} initial height of the seedling proved significant. Also within the chapter a one-way analysis of covariance (ANCOVA) model was developed. This model was used to determine which if any of the treatment fertilizers contributed to the growth of the seedlings. Multiple regression based on least squares method showed that at least two of the treatment groups significantly differed from each other. The last part of the chapter was concern with the validation of the four assumptions that are associated with ANCOVA. All four were checked and appeared not to have been violated.

Since it was discovered in Chapter 2 that significant differences between the treatment groups exists, four multiple comparison procedures which can be used to

55

evaluate which treatment groups differ was presented in Chapter 3. One of the four, procedures, Fisher's LSD, test was illustrated and it was discovered that treatment fertilizers pairs of N and P, N and K, P and K significantly differed from each other.

Chapter 4 was concerned with a non-parametric approach to analysis of covariance. By using this type of analysis it was determined that the conditional distributions of Y given X were significantly different for treatment groups.

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APPENDIX A
Table A.1 contain the mean heights of each of the 47 pots and Y, the missing observation.

Table A.1

SEEDLING HEIGHTS

AT 32 WEEKS

| | Treatment | | | | | | | | |
|-------|-----------|--------|--------|--------|--------|--------|--------|--------|---------|
| REP | 000 | NOO | OPO | OOK | NPO | NOK | OPK | NPK | Total |
| I | 20.54 | 24.16 | 28.98 | 20.84 | 22.23 | 24.70 | 22.38 | 27.76 | 191.59 |
| п | 21.62 | 24.25 | 23.62 | 21.60 | 31.36 | 22.65 | 18.36 | 29.32 | 192.78 |
| ш | 17.64 | 23.75 | 22.92 | 19.14 | 24.18 | 29.40 | 19.38 | Y* | 156.41 |
| IV | 14.88 | 17.78 | 15.88 | 18.68 | 28.27 | 27.68 | 17.24 | 27.12 | 167.53 |
| v | 14.98 | 24.00 | 19.10 | 20.28 | 40.83 | 26.73 | 21.06 | 25.86 | 222.11 |
| VI | 18.82 | 19.92 | 17.62 | 19.54 | 24.89 | 27.93 | 15.10 | 28.06 | 171.88 |
| Total | 108.48 | 133.86 | 128.12 | 120.08 | 171.76 | 188.36 | 113.52 | 138.12 | 1102.30 |

missing observation

$$Y_{ij}^* = \frac{nT_{i.} + JT_{.j} - T_{..}^*}{(n-1)(J-1)}$$

where T_{L}^{*} , T_{J}^{*} , and T_{L}^{*} denote the row, column and overall total respectively excluding the missing observation, Y. Thus the estimate of $Y_{3.8}$ is

$$Y_{38} = \frac{6(156.41) + 8(138.12) - 1102.30}{(5)(7)}$$

Y18 - 26.89

The following are the calculations associated which testing the assumption of equal treatment group variances discussed in Section 2.4.2 :

$$\sum y_j^2 - \sum Y_j^2 - \frac{\left[\sum Y_j\right]^2}{n_j}$$

$$\sum y_j^2 = 39.17 \qquad \sum y_j^2 = 231.17$$

$$\sum y_j^2 = 38.35 \qquad \sum y_j^2 = 29.96$$

$$\sum y_j^2 = 114.73 \qquad \sum y_j^2 = 34.49$$

$$\sum y_k^2 = 6.04 \qquad \sum y_j^2 = 6.90$$

 $\therefore \sum y_w^2$ = 39.17 + 38.35 + · · · + 6.90 = 500.81

$$\sum x_{j}^{2} - \sum X_{j}^{2} - \frac{(\sum X_{j})^{2}}{n_{j}}$$

$$\Sigma x_1^2 = 25.26$$
 $\Sigma x_2^2 = 6.09$
 $\Sigma x_2^2 = 9.28$
 $\Sigma x_4^2 = 18.56$
 $\Sigma x_2^2 = 6.38$
 $\Sigma x_7^2 = 12.19$
 $\Sigma x_4^2 = 2.60$
 $\Sigma x_9^2 = 11.42$

:. $\sum x_{W}^{2} = 25.27 + 9.28 + \cdots + 11.42 = 91.78$

$$\sum xy_j - \sum XY_j - \frac{\sum X_j \sum Y_j}{n_j}$$

$$\Sigma xy_1 =$$
 30.53
 $\Sigma xy_5 =$
 10.32

 $\Sigma xy_5 =$
 14.77
 $\Sigma xy_6 =$
 -0.43

 $\Sigma xy_5 =$
 25.16
 $\Sigma xy_7 =$
 18.81

 $\Sigma xy_4 =$
 1.68
 $\Sigma xy_8 =$
 -5.30

:
$$\sum xy_w = 30.53 + 14.77 + \cdot \cdot \cdot = 5.30 = 95.54$$

$$r_{w}^{2} = \frac{\sum xy_{w}}{\sqrt{\left(\sum x_{w}^{2}\right)\left(\sum y_{w}^{2}\right)}}$$

$$r_W^2 = \frac{95.54}{\sqrt{(91.78)(500.81)}}$$

 $r_w^2 = 0.4456$

From this we can find the residual sum of squares for the jth group by using the formula $(1 - r_w^2)\sum y_i^2$.

TABLE A.2

64

RESIDUAL SS

BY GROUP

| Group | $(1 - r_w^2)\Sigma y_j^2$ | | | | |
|-------|---------------------------|---|--------|--|--|
| 1 | (1 - 0.4456)39.17 | = | 21.71 | | |
| 2 | (1 - 0.4456)38.35 | - | 21.26 | | |
| 3 | (1 - 0.4456)114.73 | = | 63.60 | | |
| 4 | (1 - 0.4456)6.04 | = | 3.35 | | |
| 5 | (1 - 0.4456)231.17 | = | 128.16 | | |
| 6 | (1 - 0.4456)29.96 | = | 16.61 | | |
| 7 | (1 - 0.4456)34.50 | = | 19.13 | | |
| 8 | (1 - 0.4456)6.90 | = | 3.82 | | |

The next step is to calculate the conditional variances from each of the eight groups.

CONDITIONAL VARIANCES

BY GROUP

| Group | $(1 - r_w^2)\Sigma y^2$ | j / nj | - 1 - c |
|-------|-------------------------|--------|---------|
| 1 | 21.72/4 | = | 5.43 |
| 2 | 21.26/4 | = | 5.32 |
| 3 | 63.60/4 | = | 15.90 |
| 4 | 3.35/4 | = | 0.84 |
| 5 | 128.16/4 | = | 32.04 |
| 6 | 16.61/4 | = | 4.15 |
| 7 | 19.13/4 | = | 4.78 |
| 8 | 3.82/4 | = | 0.96 |

From Table A3 the F ratio, which is the largest divided by the smallest of the quantities is

$$F = \frac{32.04}{0.84}$$

The following pages illustrate the method of rank analysis of covariance.

RAW DATA BY

TREATMENT GROUP

| 000 | | NOO | | OPO | | OOK | | |
|-------|---------------------|-------|-----------------------|------------|-----------------------|-------|-------|--|
| Y | \mathbf{X}_{η} | Y | X ₇ | Y | X ₇ | Y | X_7 | |
| 14.95 | 9.48 | 19.92 | 11.74 | 23.62 | 12.84 | 21.60 | 13.16 | |
| 18.82 | 12.80 | 17.78 | 10.40 | 28.98 | 13.72 | 19.14 | 12.90 | |
| 17.64 | 12.54 | 24.16 | 13.64 | 19.10 | 11.56 | 19.54 | 12.44 | |
| 14.88 | 10.12 | 24.25 | 12.82 | 17.62 | 10.56 | 18.68 | 13.28 | |
| 20.54 | 15.02 | 23.75 | 14.00 | 15.88 | 11.39 | 20.28 | 13.80 | |
| 21.62 | 14.52 | 24.00 | 11.64 | 22.92 | 12.10 | 20.84 | 14.50 | |
| N | PO | NOK | | <u>OPK</u> | | NPK | | |
| Y | X_{7} | Y | X, | Y | X ₇ | Y | X, | |
| 31.36 | 12.56 | 27.68 | 11.58 | 15.10 | 10.74 | 25.86 | 13.52 | |
| 22.23 | 13.16 | 27.93 | 12.32 | 17.24 | 11.06 | 27.12 | 11.32 | |
| 24.89 | 10.92 | 29.40 | 12.30 | 21.06 | 14.20 | 29.32 | 10.18 | |
| 24.18 | 13.86 | 24.70 | 16.12 | 22.38 | 14.40 | 27.76 | 14.00 | |
| 40.53 | 13.72 | 22.65 | 10.46 | 18.36 | 13.28 | 26.89 | 13.50 | |
| 28.27 | 13.54 | 26.73 | 13.24 | 19.38 | 12.26 | 28.06 | 13.16 | |
| | | | | | | | | |

TRANSFORMED DATA BY

TREATMENT GROUP

| 000 | | NOO | | PO | OOK | |
|------------------|--|--|---|--|--|---|
| X _{nak} | Ynak | X _{rusk} | Ynak | X _{ruak} | Ymak | X _{rank} |
| 1 | 16 | 1: | 27 | 25 | 21 | 28 |
| 23 | 8 | 4 | 44 | 37.5 | 13 | 26 |
| 21 | 30 | 36 | 12 | 12 | 15 | 20 |
| 2 | 32 | 24 | 6 | 6 | 10 | 31.5 |
| 47 | 28 | 41 | 4 | 11 | 17 | 39 |
| 46 | 29 | 14 | 26 | 16 | 19 | 45 |
| | 00 X _{nak} 1 23 21 2 47 46 | Xmax Ymax 1 16 23 8 21 30 2 32 47 28 46 29 | OO NOO X _{mak} Y _{mak} X _{mak} 1 16 1: 23 8 4 21 30 36 2 32 24 47 28 41 46 29 14 | OO NOO O X _{mak} Y _{mak} X _{mak} Y _{mak} 1 16 1: 27 23 8 4 44 21 30 36 12 2 32 24 6 47 28 41 4 46 29 14 26 | OO NOO OPO Xmat Ymat Xmat Ymat Xmat 1 16 1: 27 25 23 8 4 44 37.5 21 30 36 12 12 2 32 24 6 6 47 28 41 4 11 46 29 14 26 16 | OO NOO OPO O Xmak Ymak Xmak Ymak Ymak Ymak 1 16 1: 27 25 21 23 8 4 44 37.5 13 21 30 36 12 12 15 2 32 24 6 6 10 47 28 41 4 11 17 46 29 14 26 16 19 |

| NPO | | NOK | | 0 | <u>PK</u> | NPK | |
|------|------------------|------|------------------|-------------------|------------------|------|-------|
| Ynnk | X _{nak} | Ynak | X _{nek} | Y _{rank} | X _{nak} | Ynnk | Xrank |
| 47 | 22 | 39 | 13 | 3 | 7 | 35 | 34 |
| 23 | 28 | 41 | 19 | 5 | 9 | 38 | 10 |
| 34 | 8 | 46 | 18 | 20 | 43 | 45 | 3 |
| 31 | 40 | 33 | 48 | 24 | 44 | 40 | 41 |
| 48 | 37.5 | 25 | 5 | 9 | 31.5 | 37 | 33 |
| 43 | 35 | 36 | 30 | 14 | 17 | 32 | 28 |

DEVIATIONS OF RANKS BY

1

TREATMENT GROUP

| 000 | | NOO | | <u>O</u> | <u>PO</u> | OOK | |
|-------|------------------|------------------|------------------|----------|-------------------|-------|-------------------|
| Ymnk | X _{nuk} | Y _{nak} | X _{nak} | Ymak | X _{rank} | Ymak | X _{runk} |
| -22.5 | -23.5 | - 8.5 | - 9.5 | 2.5 | .5 | - 3.5 | 3.5 |
| -13.5 | - 1.5 | -16.5 | -20.5 | 19.5 | 13.0 | -11.5 | 1.5 |
| -17.5 | - 3.5 | 5.5 | 11.5 | -12.5 | -12.5 | - 9.5 | - 4.5 |
| -23.5 | -22.5 | 7.5 | - 0.5 | -18.5 | -18.5 | -14.5 | 7.0 |
| - 6.5 | 22.5 | 3.5 | 17.0 | -20.5 | -13.5 | - 7.5 | 14.5 |
| - 2.5 | 21.5 | 4.5 | -10.5 | 1.5 | - 8.5 | - 5.5 | 20.5 |

| NPO | | NOK | | O | PK | NPK | |
|-------|-------------------|------|------------------|-------|-------------------|------|------------------|
| Ynak | X _{mank} | Ymak | X _{mak} | Ymnk | X _{rank} | Ynnk | X _{mak} |
| 22.5 | - 2.5 | 14.5 | -11.5 | -21.5 | -17.5 | 10.5 | 9.5 |
| - 1.5 | 3.5 | 16.5 | - 5.5 | -19.5 | -15.5 | 13.5 | -14.5 |
| 9.5 | -10.5 | 21.5 | - 6.5 | - 4.5 | 18.5 | 20.5 | -21.5 |
| 6.5 | 15.5 | 8.5 | 23.5 | - 0.5 | 19.5 | 15.5 | 17.0 |
| 23.5 | 13.0 | 0.5 | -19.5 | -15.5 | 7.0 | 12.5 | 8.5 |
| 18.5 | 10.5 | 11.5 | 5.5 | -10.5 | - 7.5 | 17.5 | 3.5 |

69

TABLE A.7

SUMMARY TABLE

| | Observed Obs. y _{nak} | | Estimated $y_{rank} = 0.3068(x_{rank})$ | Residual Z | |
|---------|-----------------------------------|-------|---|---------------|--|
| | 1 | -22.5 | -7.21 | -15 29 | |
| | 2 | -13.5 | -0.46 | -13.04 | |
| Group 1 | 3 | -17.5 | -1.07 | -16.43 | |
| Oloup I | 4 | -23.5 | -6.90 | -16.60 | |
| | 5 | - 65 | 690 | -13.40 | |
| | 6 | - 2.5 | 6.60 | - 9.10 | |
| | 7 | - 8.5 | -2.91 | - 5.59 | |
| | 8 | -16.5 | -6.29 | -10.21 | |
| Group 2 | 9 | 5.5 | 3.53 | 1.97 | |
| - | 10 | 7.5 | -0.15 | 7.65 | |
| | 11 | 3.5 | 5.22 | - 1.72 | |
| | 12 | 4.5 | -3.22 | 7.72 | |
| | 13 | 2.5 | 0.15 | 2.35 | |
| | 14 | 19.5 | 3.99 | 15.51 | |
| Group 3 | 15 | -12.5 | -3.89 | - 8.66 | |
| - | 16 | -18.5 | 5.68 | -12.82 | |
| | 17 | -20.5 | -4.14 | -16.36 | |
| | 18 | 1.5 | -2.61 | 4.11 | |
| | 19 | - 3.5 | 1.07 | - 4.57 | |
| | 20 | -11.5 | 0.46 | -11.96 | |
| | 21 | - 9.5 | -1.38 | - 8.12 | |
| Group 4 | 22 | -14.5 | 2.15 | -16.65 | |
| | 23 | - 7.5 | 4.45 | -11.95 | |
| | 24 | - 5.5 | 6.29 | -11.79 | |
| | 25 | 22.5 | -0.77 | 23.27 | |
| | 26 | - 1.5 | 1.07 | - 2.57 | |
| | 27 | 9.5 | -5.06 | 14.56 | |
| Group 5 | 28 | 6.5 | 4.76 | 1.74 | |
| | 29 | 23.5 | 3.99 | 19.51 | |
| | 30 | 18.5 | 3.22 | 15.28 | |

| TABLE A 7 | (con't) |
|-----------|---------|
| TADLL A./ | (con t) |

| | Obs Obs. y _{raa} | | Estimated $y_{rank} = 0.3068(x_{rank})$ | Residual Z |
|---------|------------------------------|-------|---|---------------|
| | 31 | 14.5 | -3.53 | 18.03 |
| | 32 | 16.5 | -1.69 | 18.19 |
| | 33 | 21.5 | -1.99 | 23.49 |
| Group 6 | 34 | 8.5 | 7.21 | 1.29 |
| | 35 | 0.5 | -5.98 | 6.48 |
| | 36 | 11.5 | 1.69 | 9.81 |
| | 37 | -21.5 | -5.37 | -16.13 |
| | 38 | -19.5 | -4.76 | -14.74 |
| | 39 | - 4.5 | 5.68 | -10.18 |
| Group 7 | 40 | - 0.5 | 5.98 | - 6.48 |
| 650 | 41 | -15.5 | 2.15 | -17.65 |
| | 42 | -10.5 | -2.30 | - 8.20 |
| | 43 | 10.5 | 2.91 | 7.59 |
| | 44 | 13.5 | -4.45 | 17.95 |
| | 45 | 20.5 | -6.60 | 27.10 |
| Group 8 | 46 | 15.5 | 5.22 | 10.28 |
| | 47 | 12.5 | 2.61 | 9.89 |
| | 48 | 17.5 | 1.07 | 16.43 |







