STATISTICSII PROJECT REPORT

Submitted by:

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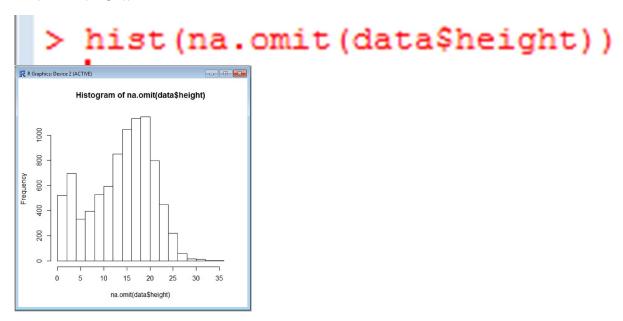
HNEE Matriculation No: 14209583

Contents

1) plot a histogram of *height* variable without *NA* records3
3) check if the average heights of *BRZ* (brzoza = birch) and *SO* (sosna = pine) species are the same4
4) check if the distribution of height for *SO* species follows the normal distribution5
5) using two-way ANOVA - check if there is a signifficant difference in stand heights between species (species_cd) and age classes (0-20, 21-40, 41-60, 61-80, 81-100, 101-120, above 120 years, based on species_age variable). Use data only for *SO*, *BRZ* and *DB* (dab = oak) species
Using R5
Using SPSS:6
6) using data for *SO* species (without *NA* records) find two variables with a non-linear relationship (e.g. age <> height). Build a nonlinear model for this relationship (choose the best out of at least 3 candidates from example from "CurveCatalog.pdf" publication of from other source). Perform a detailed regression analysis as well as justify the choice of the final model. Please do not limit yourself to the automatic procedure in SPSS, but perform a real process of choosing the proper candidate functions.
Using R9
Using SPSS:

1) plot a histogram of *height* variable without *NA* records

Hist(na.omit(height))



2) calculate average height of all stands (not counting *NAs*) as well as average heights of stands by species

```
> summary(na.omit(height))
Min. 1st Qu. Median Mean 3rd Qu. Max.
1.00 10.00 15.00 14.27 19.00 35.00
```

3) check if the average heights of *BRZ* (brzoza = birch) and *SO* (sosna = pine) species are the same

levels(species_cd)

- 1. "¦W"
- 2. "AK"
- 3. "BK"
- 4. "BRZ"
- 5. "DB"
- 6. "DB.B"
- 7. "DB.C"
- 8. "DB.S"
- 9. "DG"
- 10. "GB"
- 11. "JS"
- 12. "JW"
- 13. "LP"
- 14. "MD"
- 15. "OL"
- 16. "OL.S"
- 17. "OS"
- 18. "SO"
- 19. "TP"
- 20. "WZ"

So "BRZ" is the 4th in the list which shows mean as 14.710037 and "SO" is the 18th which gives 14.161620 as the average.

```
1 2 3 4
2.296875 15.661017 6.942857 14.710037 9.142857 2.750000 12.435897 3.133333 26.333333
17 18 19 20
6.000000 14.161620 27.600000 21.000000
```

4) check if the distribution of height for *SO* species follows the normal distribution

Is not a normal distribution.

[precise mean and standard variation are used so that the test is not done on a standard gaussian distribution.]

5) using two-way ANOVA - check if there is a signifficant difference in stand heights between species (species_cd) and age classes (0-20, 21-40, 41-60, 61-80, 81-100, 101-120, above 120 years, based on species_age variable). Use data only for *SO*, *BRZ* and *DB* (dab = oak) species.

Using R

Step1: Getting the right data without "NA" values

Note that we had unequal sample sizes, so I put a little more effort to get sample with equal size for each species and age class. But the species "BRZ" does not have the "above 120" age class. However, the data satisfies the assumption of homogeneity of variance through Levene's test (shown in SPSS result part).

Step 2: Creating categorical variable age_classes from the continuous variable "Species_age"

Step3:Performing the test.

Null hypothesis:

H0:Age has no significant effect on stand height

H0: Species type has no significant effect on stand height

H0: Age and species type interaction have no significant effect in stand height

```
| Names (af, noNA) | Names (af,
```

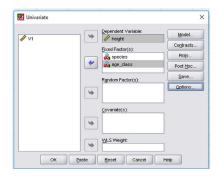
Discussion: Both age and species type have got impact on stand height. But the interaction between age and species have got no significant effect on stand height.

At the first age class, "BRZ" and "DB" have greater height than "SO". At the second age class(21-40) "BRZ" displays the greatest height and "SO" the least.

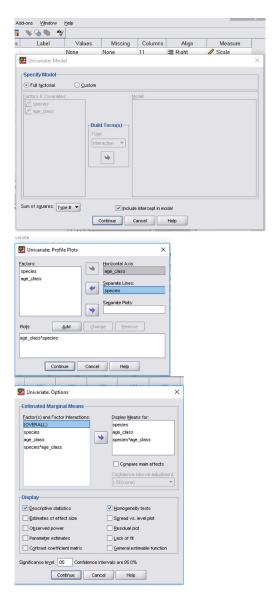
Using SPSS:.

Step1

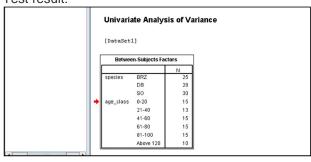
Stating the dependent and factor variables:

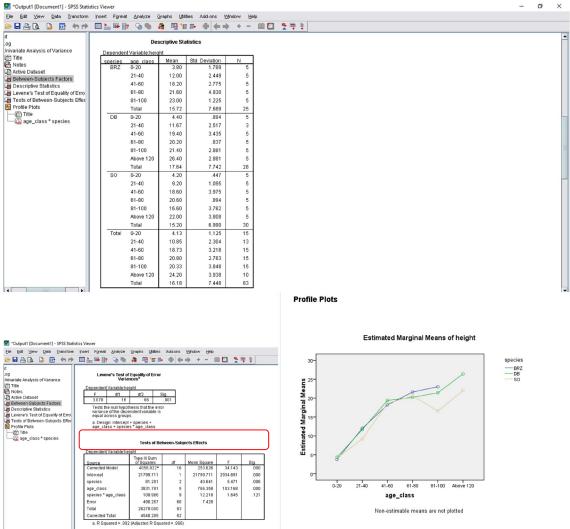


Step2:Specifying Model ,plots and other options



Test result:





Discussion: Both SPSS and R displays similar results that age and species have significant effect on stand height but not their interaction.

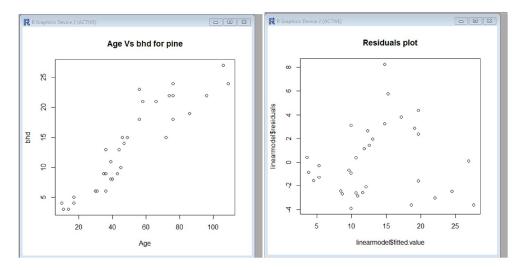
6) using data for *SO* species (without *NA* records) find two variables with a non-linear relationship (e.g. age <> height). Build a nonlinear model for this relationship (choose the best out of at least 3 candidates from example from "CurveCatalog.pdf" publication of from other source). Perform a detailed regression analysis as well as justify the choice of the final model. <u>Please do not limit yourself</u> to the automatic procedure in SPSS, but perform a real process of choosing the proper candidate functions.

Using R

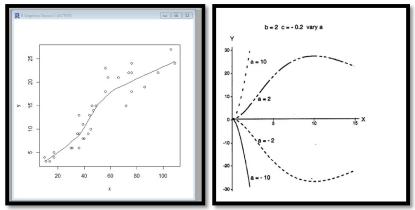
```
> raw_data=cbind(pine_data$species_age,pine_data$bhd)
> df=data.frame(raw_data)
> df_noNA = na.omit(df)
> mydata=df_nona
Error: object 'df_nona' not found
> raw_data=cbind(pine_data$species_age,pine_data$bhd)
> df=data.frame(raw_data)
> df=data.frame(raw_data)
> df=noNA = na.omit(df)
> mydata=df_noNA

> number=(1:200)
> mysample=sample(number,40,replace=T)
> x=X1[mysample]
Error: object 'X1' not found
> x=mydata$X1[mysample]
> y=mydata$X2[mysample]
```

```
> plot(y~x,main="Age Vs bhd for pine" ,xlab="Age",ylab="bhd")
> linearmodel = lm(y~x)
> plot(linearmodel$residuals~linearmodel$fitted.value , main="Residuals plot")
>
```



The residuals are not evenly dispersed which says that I need a nonlinear model for this relationship.



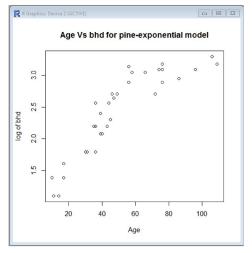
The scatter plot seems to be an exponential function graph . But to ensure the correct model , I try a power function and a polynomial function and finally choose one among the three.

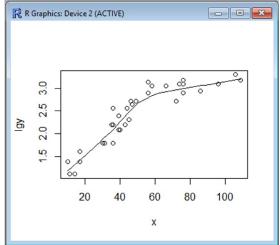
I. the exponential model:

A log transform is needed.

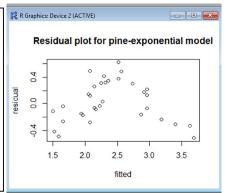
Linearized model and parameters: ln(Y) = b0 + b1Xa = eb0 b = b1

Description: The parameter $\bf a$ is the Y-intercept; the parameter $\bf b$ is the shape parameter of the curve.





```
Call:
lm(formula = lgy ~ x)
Residuals:
    Min
               10
                   Median
                                        Max
-0.51307 -0.24399 -0.05115 0.27050
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
                      0.105315
(Intercept) 1.274907
                                 12.11 1.31e-14
           0.022167
                      0.001853
                                11.96 1.86e-14 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.3254 on 38 degrees of freedom
Multiple R-squared: 0.7902,
                               Adjusted R-squared: 0.7847
F-statistic: 143.2 on 1 and 38 DF, p-value: 1.864e-14
```



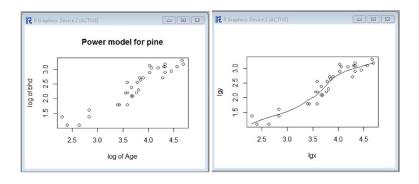
The p –values for intercept and slope are low, indicating that both of them are significantly different from zero.

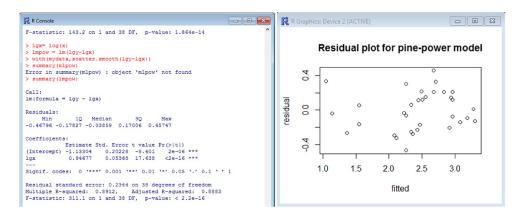
II. The power model

Functional form: $Y = aX^b$

Linearized model and parameters: $ln(Y) = b_0 + b_1 \ ln(X)$

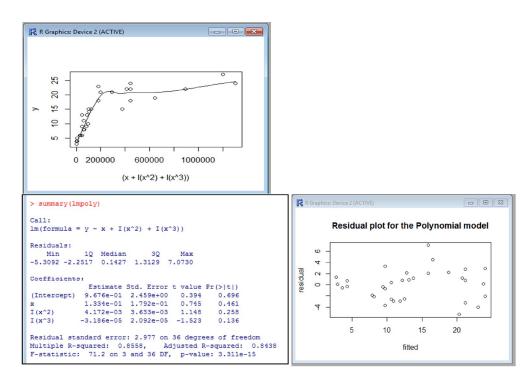
 $a=eb_0\ b=b_1$





III The polynomial model

The graph may have a cubic term. So I use a polynomial of degree 3.



This model fails as the p-values are high meaning that the estimates are not significantly different from zero.

Dignostics:

Model comparison

Model	Fitted plot	Adjusted R Square	Residual plot
Exponential model		0.7843	The best
Power model		0.8883	Better than the linear model
Polynomial model	Fails		

So I can select the power model. The equation for power model would be:

Functional form: $Y = aX^b$

Linearized model and parameters: $ln(Y) = b_0 + b_1 ln(X)$

 $a = e^{b0}$ $b = b_1$

From R, we have got

b0 = -1.3304

b1= 0.94677

ln(Y) = -1.3304 + 0.94677 * ln(X)

⇒ (Raised to the power with base e)

 \Rightarrow Y = 0.26 * X (0.94677)

 \Rightarrow BHD = 0.26 * (AGE) (0.94677)

Example 1:

Age = 44

BHD = $0.26 * (44)^{(0.94677)} = 9.35$

Example 2:

Age = 39

BHD = $0.26 * (39)^{(0.94677)} = 8.34$

Using SPSS:

Now I check my result from R with SPSS:

```
- - X
R Console
F-statistic: 143.2 on 1 and 38 DF, p-value: 1.864e-14
> lgx = log(x)
> lmpow = lm(lgy~lgx)
> with(mydata,scatter.smooth(lgy~lgx))
> summary(mlpow)
Error in summary(mlpow) : object 'mlpow' not found
> summary(lmpow)
Call:
lm(formula = lgy ~ lgx)
Residuals:
Min 1Q Median 3Q Max
-0.46796 -0.17827 -0.03859 0.17006 0.45747
Coefficients:

Estimate Std. Error t value Pr(>|t|)

20228 -5.601 2e-06
lgx
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.2344 on 38 degrees of freedom
Multiple R-squared: 0.8912, Adjusted R-squared: 0.8883 F-statistic: 311.1 on 1 and 38 DF, p-value: < 2.2e-16
```

Iteration History^b

	iteration mistory			
Iteration		Parameter		
Number	Residual Sum of			
a	Squares	Α	В	
1.0	253487.400	.260	.947	
1.1	95494.116	.339	.960	
2.0	95494.116	.339	.960	
2.1	95048.580	.340	.956	
3.0	95048.580	.340	.956	
3.1	95048.557	.340	.956	
4.0	95048.557	.340	.956	
4.1	95048.557	.340	.956	

Derivatives are calculated numerically.

Parameter Estimates				
Parame			95% Confidence Interval	
ter	Estimate	Std. Error	Lower Bound	Upper Bound
A	.340	.008	.325	.355
В	.956	.005	.946	.967

Correlations of Parameter

Estimates			
	A	В	
A	1.000	995	
В	995	1.000	

ANOVA*			
Source	Sum of Squares	.df.	Mean Squares
Regression	2251273.443	2	1125636.721
Residual	95048.557	8785	10.819
Uncorrected Total	2346322.000	8787	
Corrected Total	612739.425	8786	

Dependent variable: bhd

a. R squared = 1 - (Residual Sum of Squares) / (Corrected Sum of Squares) = .845.

SPSS performed 8 iterations and found the equation as :

Discussion: Values I have retrieved with R are used as initial values in SPSS which seems to further improve the equation.