ECON 490ML Final Project Report

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Introduction

For our Final Project, we will be attempting to predict the yield of wild blueberries. These are known as "lowbush berries", which grow on low-level bushes and are typically pea-sized. Our dataset can be found at https://www.kaggle.com/datasets/saurabhshahane/wild-blueberry-yield-prediction. This data was generated by a simulation model called the Wild Blueberry Pollination Model, which has been validated by experimental data collected in the State of Main over the last 30 years.

Generally, in the field of agriculture, an increasing quantity of research has gone underway to understand the determinants of crop yield. Machine learning has enabled scientists and farmers to investigate which factors have the greatest impact on crop yield. In this specific case, the crop of interest is the wild blueberry. The paper for which this data was created is called "Simulation-based modeling of wild blueberry pollination". It was published in the January 2018 version of *Computers and Electronics in Agriculture*. The paper can be found at: https://www.sciencedirect.com/science/article/pii/S0168169916310274?via%3Dihub

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Handling Data

Below the dataset will be loaded from a .csv file. The tidyverse library has been loaded so that the data will automatically be loaded as a tibble, which provides some advantages over a standard data frame. One advantage being that the read_csv function will automatically assign types to each column in the tibble.

```
library(tidyverse)
data = read_csv("blueberryData.csv")
colnames(data)
```

```
[1] "Row#"
                                 "clonesize"
                                                         "honeybee"
                                                         "osmia"
    [4] "bumbles"
                                 "andrena"
##
    [7] "MaxOfUpperTRange"
                                 "MinOfUpperTRange"
                                                         "AverageOfUpperTRange"
  [10] "MaxOfLowerTRange"
                                 "MinOfLowerTRange"
                                                         "AverageOfLowerTRange"
   [13] "RainingDays"
                                 "AverageRainingDays"
                                                         "fruitset"
## [16] "fruitmass"
                                 "seeds"
                                                         "yield"
dim(data)
```

[1] 777 18

The dataset contains 18 columns and 777 observations. Along with downloaded the dataset from Kaggle, we downloaded the descriptions of the predictor variables in the dataset.

Table 1. Features and their description

Features	Unit	Description
Clonesize	m ²	The average blueberry clone size in the field
Honeybee	bees/m²/min	Honeybee density in the field
Bumbles	bees/m²/min	Bumblebee density in the field
Andrena	bees/m²/min	Andrena bee density in the field
Osmia	bees/m²/min	Osmia bee density in the field
MaxOfUpperTRange	°C	The highest record of the upper band daily air temperature during the
		bloom season
MinOfUpperTRange	°C	The lowest record of the upper band daily air temperature
AverageOfUpperTRange	°C	The average of the upper band daily air temperature
MaxOfLowerTRange	°C	The highest record of the lower band daily air temperature
MinOfLowerTRange	°C	The lowest record of the lower band daily air temperature
AverageOfLowerTRange	°C	The average of the lower band daily air temperature
RainingDays	Day	The total number of days during the bloom season, each of which has
		precipitation larger than zero
AverageRainingDays	Day	The average of raining days of the entire bloom season

There are 13 predictor variables in the dataset. Their information includes clone size (size of bush), bee density by species, temperature, and precipitation. fruitset, fruitmass, seeds, and yield are all response variables. All predictor and response variables are quantitative. There are no qualitative variables. Row# can be deleted because it serves no purpose.

```
data = select(data, -'Row#')
dim(data)
```

```
## [1] 777 17
```

The dataset does not have any NA values. We also will not need to convert any columns into factors, since there is no categorical data.

```
sum(is.na(data))
```

[1] 0

Exploration

A look at the density plots of 9 of our predictor variables shows that none of the distributions are approximately normal. Although this will not be detrimental to our algorithms, this was a surprise considering this is a simulation of natural factors (temperature, bees, precipitation).

```
library(gridExtra)

plot1 = ggplot(data, aes(clonesize)) + geom_density()

plot2 = ggplot(data, aes(honeybee)) + geom_density() + xlim(0, 1)

plot3 = ggplot(data, aes(bumbles)) + geom_density()

plot4 = ggplot(data, aes(andrena)) + geom_density()

plot5 = ggplot(data, aes(osmia)) + geom_density()

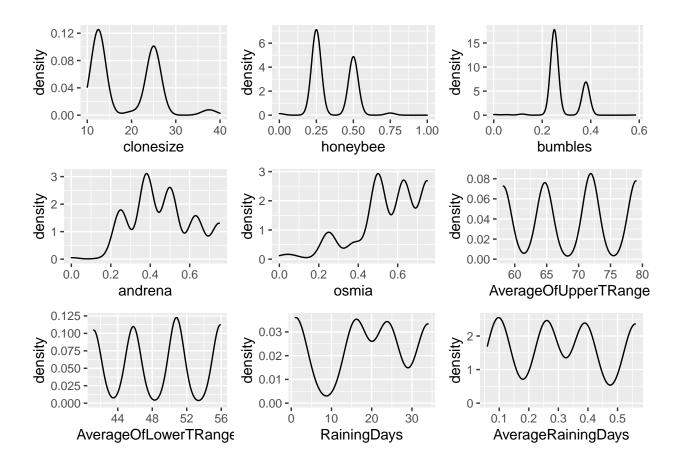
plot6 = ggplot(data, aes(AverageOfUpperTRange)) + geom_density()

plot7 = ggplot(data, aes(AverageOfLowerTRange)) + geom_density()

plot8 = ggplot(data, aes(RainingDays)) + geom_density()

plot9 = ggplot(data, aes(AverageRainingDays)) + geom_density()

grid.arrange(plot1, plot2, plot3, plot4, plot5, plot6, plot7, plot8, plot9, ncol = 3)
```



Prediction

We will attempt to predict yield with the given predictor variables in the dataset. We chose yield as opposed to the other response variables because it is measured in quantity of blueberries, which we felt to the most important measure of a harvest.

Before beginning any machine learning algorithms, we decided to only retain AverageOfLowerTRange and AverageOfUpperTRange amongst out six temperature variables. The researchers used historical weather data to simulate the ranges of temperatures. Because of this we are comfortable using these two variables as our high and low temperatures. Our decision to remove these variables will simplify the model, and minimize multicolinearity. This will result in 9 predictor variables for our modeling.

```
data = select(data, -c(MaxOfUpperTRange, MinOfUpperTRange, MaxOfLowerTRange, MinOfLowerTRange)) %>%
    select(-c(fruitset, fruitmass, seeds))
dim(data)
```

[1] 777 10

We will also create training and test sets from our data before starting our modeling. Out split will be 70% training, 30% testing.

```
set.seed(42)
train = sample(c(TRUE, FALSE), size = nrow(data), prob = c(0.7, 0.3), replace = TRUE)
traindata = data[train, ]
testdata = data[!train, ]
```

Multiple Linear Regression

Our first method will be to model the data using Multiple Linear Regression, with all 9 predictor variables. The result of the regression is the formula

```
yield = 79.28.955 - 98.207 clone size + 118.492 honeybee + 5980.501 bumbles + 520.207 adrena + 2448.218 osmia \\ + 236.815 Average Of Upper TRange - 369.804 Average Of Lower TRange + 51.718 Raining Days \\ - 8375.642 Average Raining Days
```

Surprisingly, AverageOfUpperTRange and AverageOfLowerTRange are not statistically significant in this model. Perhaps temperature does not have much an effect yield. It's too early to know for sure, but more evidence may be presented as we continue. The RMSE produced a value of 590.3259. Another insight this model has provided is that bumblebees may have the largest impact on blueberry yield of the four species. Lastly, it was not expected that the coefficient of clonesize would be negative. We assumed that larger bushes produce more berries on average.

```
set.seed(42)
MLR = lm(yield ~ ., data = traindata)
summary(MLR)
```

```
##
## Call:
## lm(formula = yield ~ ., data = traindata)
##
## Residuals:
```

```
##
       Min
                 1Q
                      Median
                                   3Q
                                           Max
## -1596.85 -367.76
                       28.41
                               411.26 1399.72
##
## Coefficients:
##
                        Estimate Std. Error t value Pr(>|t|)
                                    292.619 27.096 < 2e-16 ***
                        7928.955
## (Intercept)
## clonesize
                         -98.207
                                      3.778 -25.998 < 2e-16 ***
                                              4.973 8.91e-07 ***
## honeybee
                         118.492
                                     23.828
## bumbles
                        5980.501
                                    409.568 14.602 < 2e-16 ***
## andrena
                         520.207
                                    173.790
                                            2.993 0.00289 **
                                    180.627 13.554 < 2e-16 ***
## osmia
                        2448.218
## AverageOfUpperTRange
                                    306.793
                                            0.772 0.44051
                         236.815
## AverageOfLowerTRange -369.804
                                    434.823 -0.850 0.39544
                                             3.168 0.00162 **
## RainingDays
                          51.718
                                     16.323
## AverageRainingDays
                       -8375.642
                                   1159.401 -7.224 1.75e-12 ***
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## Residual standard error: 591.5 on 535 degrees of freedom
## Multiple R-squared: 0.8115, Adjusted R-squared: 0.8084
## F-statistic:
                 256 on 9 and 535 DF, p-value: < 2.2e-16
MLRtest = predict(MLR, newdata = testdata)
sqrt(mean((MLRtest - testdata$yield) ^ 2))
```

[1] 590.3259

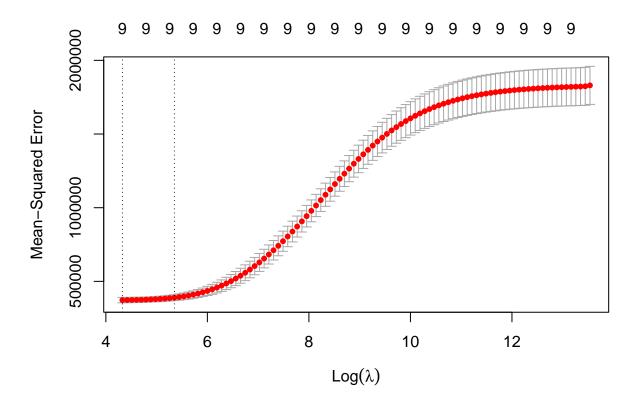
Ridge Regression

```
library(glmnet)
train.test = model.matrix(yield~., data = traindata)
x.test = model.matrix(yield~., data = testdata)
```

```
# Finding lambda chosen by cross-validation
set.seed(42)
ridge.cv = cv.glmnet(train.test, traindata$yield, alpha = 0)
lambda.ridge = ridge.cv$lambda.min
lambda.ridge
```

[1] 75.6745

plot(ridge.cv)



```
#Fitting to ridge regression
ridge.fit = glmnet(train.test, traindata$yield, alpha = 0, lambda = lambda.ridge)
coef(ridge.fit)
```

11 x 1 sparse Matrix of class "dgCMatrix"

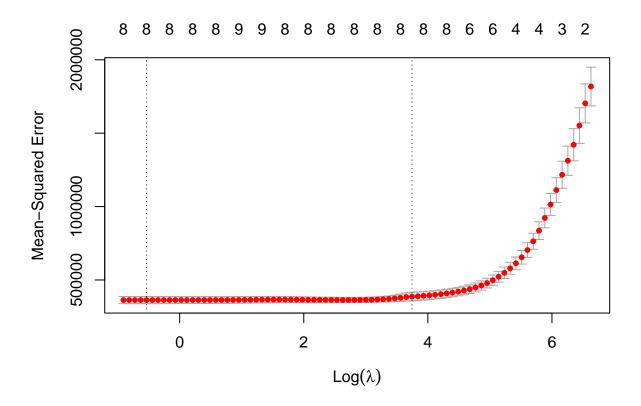
s0

```
## (Intercept)
                        7656.70319
## (Intercept)
## clonesize
                        -92.80157
## honeybee
                        113.78034
## bumbles
                        5423.50681
## andrena
                        500.51521
                        2244.36986
## osmia
## AverageOfUpperTRange -11.47164
## AverageOfLowerTRange -16.97510
                      -21.94500
## RainingDays
## AverageRainingDays -3012.19532
#Mean square error
ridge.pred = predict(ridge.fit, newx=x.test, s = lambda.ridge)
mean((testdata$yield - ridge.pred)^2)
```

Lasso Regression

[1] 359304.9

```
# Finding lambda chosen by cross-validation
set.seed(42)
lasso.cv = cv.glmnet(train.test, traindata$yield, alpha = 1)
lambda.lasso = lasso.cv$lambda.min
lambda.lasso
## [1] 0.5859202
```



```
#Fitting to lasso regression
lasso.fit = glmnet(train.test, traindata$yield, alpha = 1, lambda = lambda.lasso)
coef(lasso.fit)
```

```
## 11 x 1 sparse Matrix of class "dgCMatrix"
##
## (Intercept)
                         7880.01439
## (Intercept)
                          -98.15390
## clonesize
## honeybee
                          119.03992
## bumbles
                         5930.24157
## andrena
                           519.89848
## osmia
                         2419.36850
## AverageOfUpperTRange
                          -15.29857
## AverageOfLowerTRange
                           -12.36606
## RainingDays
                           44.92750
```

AverageRainingDays -7899.73466

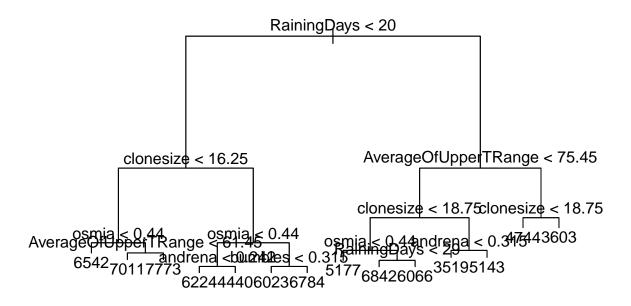
```
#Mean square error
lasso.pred = predict(lasso.fit, newx=x.test, s = lambda.lasso)
mean((testdata$yield - lasso.pred)^2)
```

```
## [1] 351616.5
```

There are 11 non-zero coefficient estimates in the lasso regression

Regression Tree

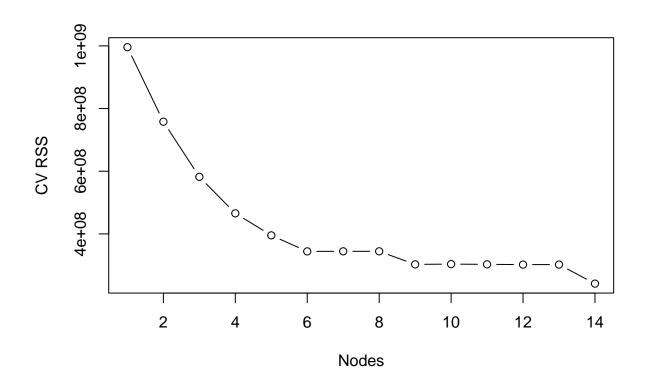
```
## creating regression tree with training data
library(tree)
set.seed(42)
tree.train = tree(yield ~ ., data = traindata)
plot(tree.train)
text(tree.train, pretty = 0)
```



```
##test MSE prior to CV + pruning
predict.yield1 = predict(tree.train, newdata = testdata)
mean((predict.yield1 - testdata$yield)^2)

## [1] 422800.7

## apply K-fold cross validation, K = 10 is standard size to use
## graph reveals that the training RSS is at its minimum at 14 nodes
cv.train = cv.tree(tree.train, K = 10)
plot(cv.train$size, cv.train$dev, xlab = "Nodes", ylab = "CV RSS", type = "b")
```



```
## prune tree

tree.prune = prune.tree(tree.train, best = 14)

##conclude pruning does not affect MSE

predict.yield2 = predict(tree.prune, newdata = testdata)

mean((predict.yield2 - testdata$yield)^2)
```

[1] 422800.7

Bagging

```
##Bagging tries to reduce variance which will in turn reduce MSE
library(randomForest)
set.seed(42)
tree.bag = randomForest(yield ~ ., data = traindata, mtry = 9, importance = T)
```

```
##Significantly smaller MSE relative to what we have done so far
predict.yield3 = predict(tree.bag, newdata = testdata)
mean((predict.yield3 - testdata$yield)^2)
```

[1] 145661.1

Random Forests

```
##Random Forests to see if it'll produce a even smaller MSE
set.seed(42)
tree.rf = randomForest(yield ~ ., data = traindata, mtry = 3, importance = T)
predict.yield4 = predict(tree.rf, newdata = testdata)
mean((predict.yield4 - testdata$yield)^2)
```

[1] 135009.7

##MSE produced by Random Forests is smallest out of all tree methods used