

Project

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Data

From TidyTuesday URL:<https://github.com/rfordatascience/tidytuesday/tree/master/data/2020/2020-07-07>

```
coffee_ratings <- readr::read_csv('https://raw.githubusercontent.com/rfordatascience/tidytuesday/master/data/2020/2020-07-07/2020-07-07-coffee_ratings.csv')
```

```
##
## -- Column specification -----
## cols(
##   .default = col_character(),
##   total_cup_points = col_double(),
##   number_of_bags = col_double(),
##   aroma = col_double(),
##   flavor = col_double(),
##   aftertaste = col_double(),
##   acidity = col_double(),
##   body = col_double(),
##   balance = col_double(),
##   uniformity = col_double(),
##   clean_cup = col_double(),
##   sweetness = col_double(),
##   cupper_points = col_double(),
##   moisture = col_double(),
##   category_one_defects = col_double(),
##   quakers = col_double(),
##   category_two_defects = col_double(),
##   altitude_low_meters = col_double(),
##   altitude_high_meters = col_double(),
##   altitude_mean_meters = col_double()
## )
## i Use `spec()` for the full column specifications.
```

Quick overview

```
summary(coffee_ratings)
```

```
## total_cup_points species owner country_of_origin
## Min. : 0.00 Length:1339 Length:1339 Length:1339
## 1st Qu.:81.08 Class :character Class :character Class :character
## Median :82.50 Mode :character Mode :character Mode :character
## Mean :82.09
## 3rd Qu.:83.67
```

```

## Max.      :90.58
##
## farm_name      lot_number      mill      ico_number
## Length:1339    Length:1339    Length:1339    Length:1339
## Class :character    Class :character    Class :character    Class :character
## Mode  :character    Mode  :character    Mode  :character    Mode  :character
##
##
##
## company      altitude      region      producer
## Length:1339    Length:1339    Length:1339    Length:1339
## Class :character    Class :character    Class :character    Class :character
## Mode  :character    Mode  :character    Mode  :character    Mode  :character
##
##
##
## number_of_bags    bag_weight      in_country_partner    harvest_year
## Min.      : 0.0    Length:1339    Length:1339    Length:1339
## 1st Qu.: 14.0    Class :character    Class :character    Class :character
## Median : 175.0    Mode  :character    Mode  :character    Mode  :character
## Mean      : 154.2
## 3rd Qu.: 275.0
## Max.      :1062.0
##
## grading_date      owner_1      variety      processing_method
## Length:1339    Length:1339    Length:1339    Length:1339
## Class :character    Class :character    Class :character    Class :character
## Mode  :character    Mode  :character    Mode  :character    Mode  :character
##
##
##
## aroma      flavor      aftertaste      acidity      body
## Min.      :0.000    Min.      :0.00    Min.      :0.000    Min.      :0.000    Min.      :0.000
## 1st Qu.:7.420    1st Qu.:7.33    1st Qu.:7.250    1st Qu.:7.330    1st Qu.:7.330
## Median :7.580    Median :7.58    Median :7.420    Median :7.580    Median :7.500
## Mean      :7.567    Mean      :7.52    Mean      :7.401    Mean      :7.536    Mean      :7.517
## 3rd Qu.:7.750    3rd Qu.:7.75    3rd Qu.:7.580    3rd Qu.:7.750    3rd Qu.:7.670
## Max.      :8.750    Max.      :8.83    Max.      :8.670    Max.      :8.750    Max.      :8.580
##
## balance      uniformity      clean_cup      sweetness
## Min.      :0.000    Min.      : 0.000    Min.      : 0.000    Min.      : 0.000
## 1st Qu.:7.330    1st Qu.:10.000    1st Qu.:10.000    1st Qu.:10.000
## Median :7.500    Median :10.000    Median :10.000    Median :10.000
## Mean      :7.518    Mean      : 9.835    Mean      : 9.835    Mean      : 9.857
## 3rd Qu.:7.750    3rd Qu.:10.000    3rd Qu.:10.000    3rd Qu.:10.000
## Max.      :8.750    Max.      :10.000    Max.      :10.000    Max.      :10.000
##
## cupper_points      moisture      category_one_defects      quakers
## Min.      : 0.000    Min.      :0.00000    Min.      : 0.0000    Min.      : 0.0000
## 1st Qu.: 7.250    1st Qu.:0.09000    1st Qu.: 0.0000    1st Qu.: 0.0000
## Median : 7.500    Median :0.11000    Median : 0.0000    Median : 0.0000

```

```
## Mean : 7.503 Mean :0.08838 Mean : 0.4795 Mean : 0.1734
## 3rd Qu.: 7.750 3rd Qu.:0.12000 3rd Qu.: 0.0000 3rd Qu.: 0.0000
## Max. :10.000 Max. :0.28000 Max. :63.0000 Max. :11.0000
## NA's :1
## color category_two_defects expiration certification_body
## Length:1339 Min. : 0.000 Length:1339 Length:1339
## Class :character 1st Qu.: 0.000 Class :character Class :character
## Mode :character Median : 2.000 Mode :character Mode :character
## Mean : 3.556
## 3rd Qu.: 4.000
## Max. :55.000
##
## certification_address certification_contact unit_of_measurement
## Length:1339 Length:1339 Length:1339
## Class :character Class :character Class :character
## Mode :character Mode :character Mode :character
##
##
##
## altitude_low_meters altitude_high_meters altitude_mean_meters
## Min. : 1 Min. : 1 Min. : 1
## 1st Qu.: 1100 1st Qu.: 1100 1st Qu.: 1100
## Median : 1311 Median : 1350 Median : 1311
## Mean : 1751 Mean : 1799 Mean : 1775
## 3rd Qu.: 1600 3rd Qu.: 1650 3rd Qu.: 1600
## Max. :190164 Max. :190164 Max. :190164
## NA's :230 NA's :230 NA's :230
```

A few NA's.

1 within quakers, and 230 in Altitude low/high/mean

Check what is happening in the rest of the data set

Count of NA's per coloumn

```
apply(X=is.na(coffee_ratings), MARGIN = 2, FUN = sum)
```

```
## total_cup_points species owner
## 0 0 7
## country_of_origin farm_name lot_number
## 1 359 1063
## mill ico_number company
## 315 151 209
## altitude region producer
## 226 59 231
## number_of_bags bag_weight in_country_partner
## 0 0 0
## harvest_year grading_date owner_1
## 47 0 7
## variety processing_method aroma
## 226 170 0
## flavor aftertaste acidity
## 0 0 0
```

```
##          body          balance          uniformity
##          0              0              0
##      clean_cup          sweetness      cupper_points
##          0              0              0
##          moisture category_one_defects          quakers
##          0              0              1
##          color category_two_defects          expiration
##          218              0              0
##      certification_body certification_address certification_contact
##          0              0              0
##      unit_of_measurement altitude_low_meters altitude_high_meters
##          0              230              230
##      altitude_mean_meters
##          230
```

I will be just removing some of the columns with many missing vlaues, for instance farm_name.

```
library(tidyverse)
```

```
## -- Attaching packages ----- tidyverse 1.3.1 --
```

```
## v ggplot2 3.3.4    v purrr  0.3.4
## v tibble  3.1.2    v dplyr  1.0.7
## v tidyr   1.1.3    v stringr 1.4.0
## v readr   1.4.0    v forcats 0.5.1
```

```
## -- Conflicts ----- tidyverse_conflicts() --
```

```
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()    masks stats::lag()
```

Removal of columns

```
coffee = coffee_ratings%>%
  select(-farm_name,-lot_number,-mill,-ico_number,-altitude,
         -altitude_low_meters,-altitude_high_meters,-producer,-company,
         -expiration,-certification_address,-owner_1,-grading_date,
         -certification_contact,-unit_of_measurement)
apply(X=is.na(coffee), MARGIN = 2, FUN = sum)
```

```
##      total_cup_points          species          owner
##          0              0              7
##      country_of_origin          region      number_of_bags
##          1              59              0
##          bag_weight in_country_partner      harvest_year
##          0              0              47
##          variety    processing_method          aroma
##          226              170              0
##          flavor          aftertaste          acidity
##          0              0              0
##          body          balance          uniformity
##          0              0              0
##      clean_cup          sweetness      cupper_points
##          0              0              0
##          moisture category_one_defects          quakers
##          0              0              1
##          color category_two_defects      certification_body
##          218              0              0
```

```

## altitude_mean_meters
##                230

#view(coffee)

coffee = na.omit(coffee)
#view(coffee)

coffee[grepl("lbs",coffee$bag_weight),]

## # A tibble: 18 x 28
##   total_cup_points species owner    country_of_origin region  number_of_bags
##   <dbl> <chr>    <chr>    <chr>          <chr>          <dbl>
## 1      87.2 Arabica the coff~ Costa Rica    san ram~      250
## 2      86.3 Arabica francisc~ Costa Rica    west an~      250
## 3      85.3 Arabica the coff~ Costa Rica    west va~      250
## 4      85.3 Arabica the coff~ Costa Rica    san ram~      250
## 5      84.7 Arabica fabian c~ Costa Rica    tarrazu      50
## 6      84.5 Arabica fabian c~ Costa Rica    tarrazu      250
## 7      83.8 Arabica german n~ United States (Pu~ yauco r~      18
## 8      83.8 Arabica the coff~ Guatemala     quetzal~      250
## 9      83.3 Arabica the coff~ Costa Rica    san ram~      250
## 10     83.3 Arabica itiah co~ Haiti          thiotte~       2
## 11      83 Arabica german n~ United States (Pu~ yauco r~      17
## 12     81.5 Arabica myriam k~ Haiti          dondon,~     300
## 13     81.2 Arabica essencec~ Guatemala     huehuet~      36
## 14     81.1 Arabica german n~ United States (Pu~ yauco r~      18
## 15     80.9 Arabica chris fi~ Nicaragua     matagal~     275
## 16     80.8 Arabica the coff~ Costa Rica    san ram~      250
## 17     79.3 Arabica the coff~ Colombia      pereira      250
## 18     79.1 Arabica german n~ United States (Pu~ yauco r~      18
## # ... with 22 more variables: bag_weight <chr>, in_country_partner <chr>,
## #   harvest_year <chr>, variety <chr>, processing_method <chr>, aroma <dbl>,
## #   flavor <dbl>, aftertaste <dbl>, acidity <dbl>, body <dbl>, balance <dbl>,
## #   uniformity <dbl>, clean_cup <dbl>, sweetness <dbl>, cupper_points <dbl>,
## #   moisture <dbl>, category_one_defects <dbl>, quakers <dbl>, color <chr>,
## #   category_two_defects <dbl>, certification_body <chr>,
## #   altitude_mean_meters <dbl>

coffee = separate(data = coffee, col = bag_weight, into = c("weight", "type"), sep = " ")

coffee$weight = as.numeric(coffee$weight)

for(i in 1:length(coffee)){
  if(coffee[i,8]=="kg"){
    coffee[i,7] = round(coffee[i,7] * 2.20462,0)
    coffee[i,8] = "lbs"
  }
}

coffee = coffee%>%
  select(-type)

coffee = coffee%>%
  rename(avg_altitude=altitude_mean_meters)

```

```
coffee$avg_altitude = round(coffee$avg_altitude * 3.28084,0)
```

```
coffee$harvest_year = substr(coffee$harvest_year,1,4)
coffee$harvest_year = as.numeric(coffee$harvest_year)
```

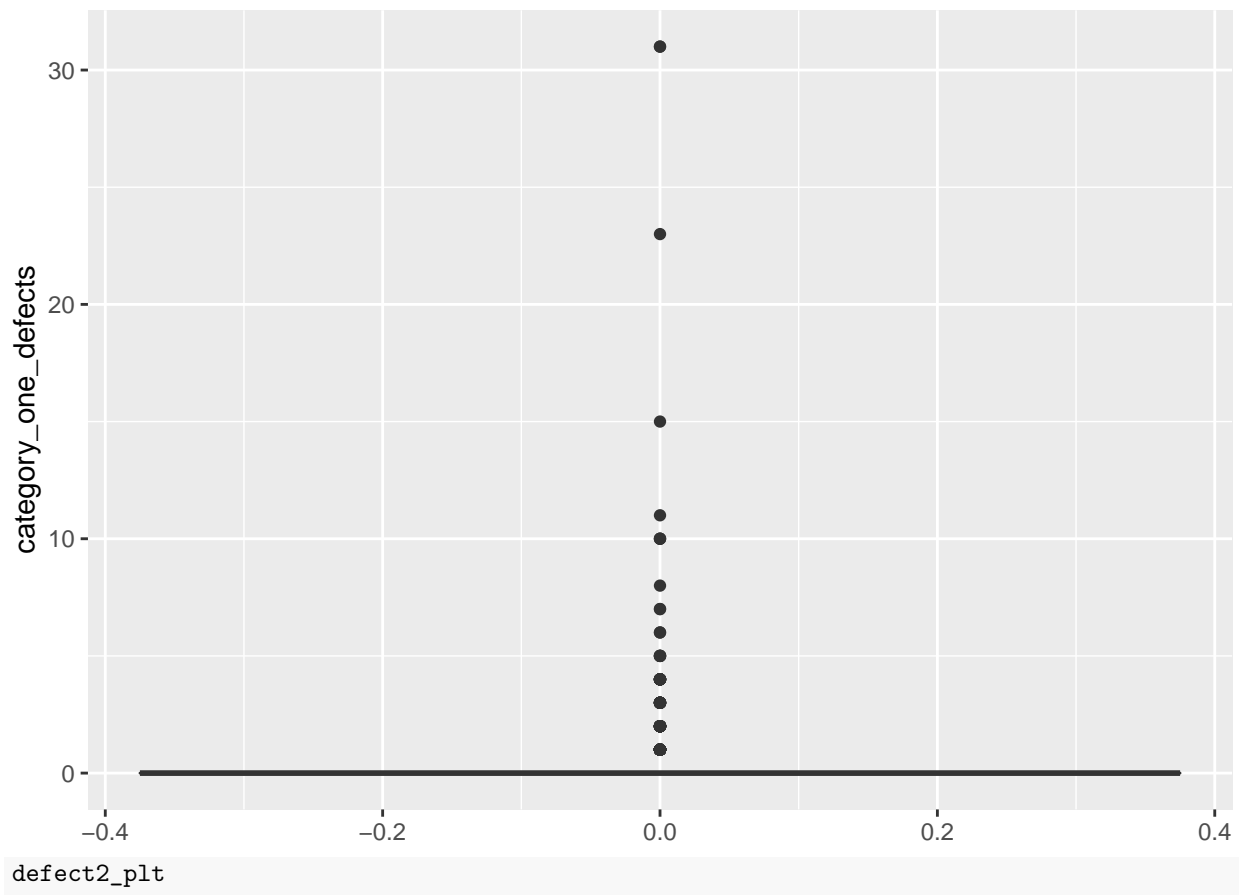
```
summary(coffee[,c(9,12:24,26,28)])
```

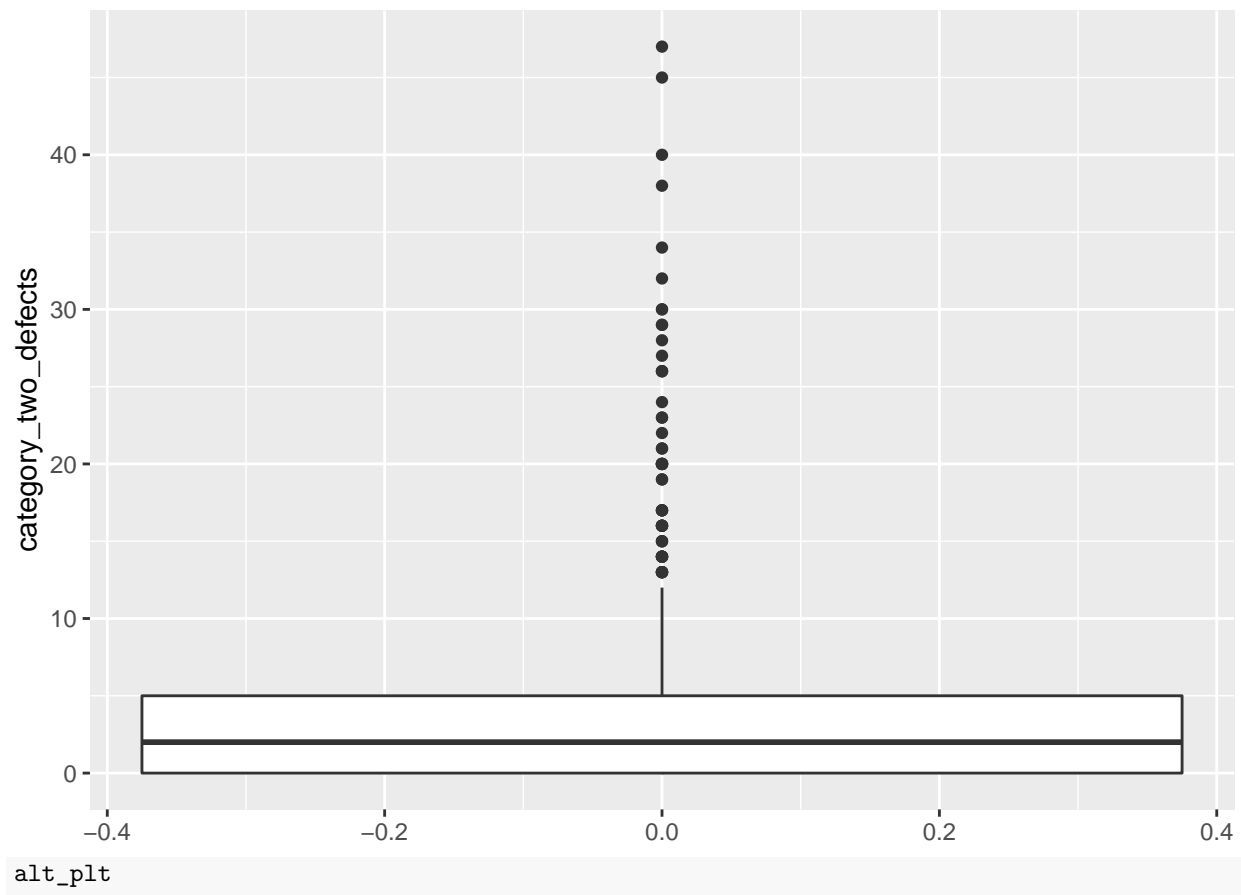
```
##  harvest_year      aroma      flavor      aftertaste      acidity
##  Min.   :2011   Min.   :5.080   Min.   :6.170   Min.   :6.170   Min.   :5.250
##  1st Qu.:2012   1st Qu.:7.420   1st Qu.:7.330   1st Qu.:7.170   1st Qu.:7.330
##  Median :2014   Median :7.580   Median :7.500   Median :7.420   Median :7.500
##  Mean   :2014   Mean   :7.559   Mean   :7.504   Mean   :7.374   Mean   :7.515
##  3rd Qu.:2015   3rd Qu.:7.750   3rd Qu.:7.670   3rd Qu.:7.580   3rd Qu.:7.670
##  Max.   :2018   Max.   :8.750   Max.   :8.670   Max.   :8.500   Max.   :8.580
##      body      balance      uniformity      clean_cup
##  Min.   :6.330   Min.   :6.080   Min.   : 6.000   Min.   : 0.000
##  1st Qu.:7.330   1st Qu.:7.330   1st Qu.:10.000   1st Qu.:10.000
##  Median :7.500   Median :7.500   Median :10.000   Median :10.000
##  Mean   :7.494   Mean   :7.488   Mean   : 9.871   Mean   : 9.849
##  3rd Qu.:7.670   3rd Qu.:7.670   3rd Qu.:10.000   3rd Qu.:10.000
##  Max.   :8.420   Max.   :8.580   Max.   :10.000   Max.   :10.000
##      sweetness      cupper_points      moisture      category_one_defects
##  Min.   : 1.33   Min.   :5.170   Min.   :0.00000   Min.   : 0.0000
##  1st Qu.:10.00   1st Qu.:7.250   1st Qu.:0.10000   1st Qu.: 0.0000
##  Median :10.00   Median :7.500   Median :0.11000   Median : 0.0000
##  Mean   : 9.93   Mean   :7.459   Mean   :0.09737   Mean   : 0.4262
##  3rd Qu.:10.00   3rd Qu.:7.670   3rd Qu.:0.12000   3rd Qu.: 0.0000
##  Max.   :10.00   Max.   :8.580   Max.   :0.17000   Max.   :31.0000
##      quakers      category_two_defects      avg_altitude
##  Min.   : 0.0000   Min.   : 0.000   Min.   :    3
##  1st Qu.: 0.0000   1st Qu.: 0.000   1st Qu.: 3609
##  Median : 0.0000   Median : 2.000   Median : 4300
##  Mean   : 0.1521   Mean   : 3.822   Mean   : 6145
##  3rd Qu.: 0.0000   3rd Qu.: 5.000   3rd Qu.: 5249
##  Max.   :11.0000   Max.   :47.000   Max.   :623898
```

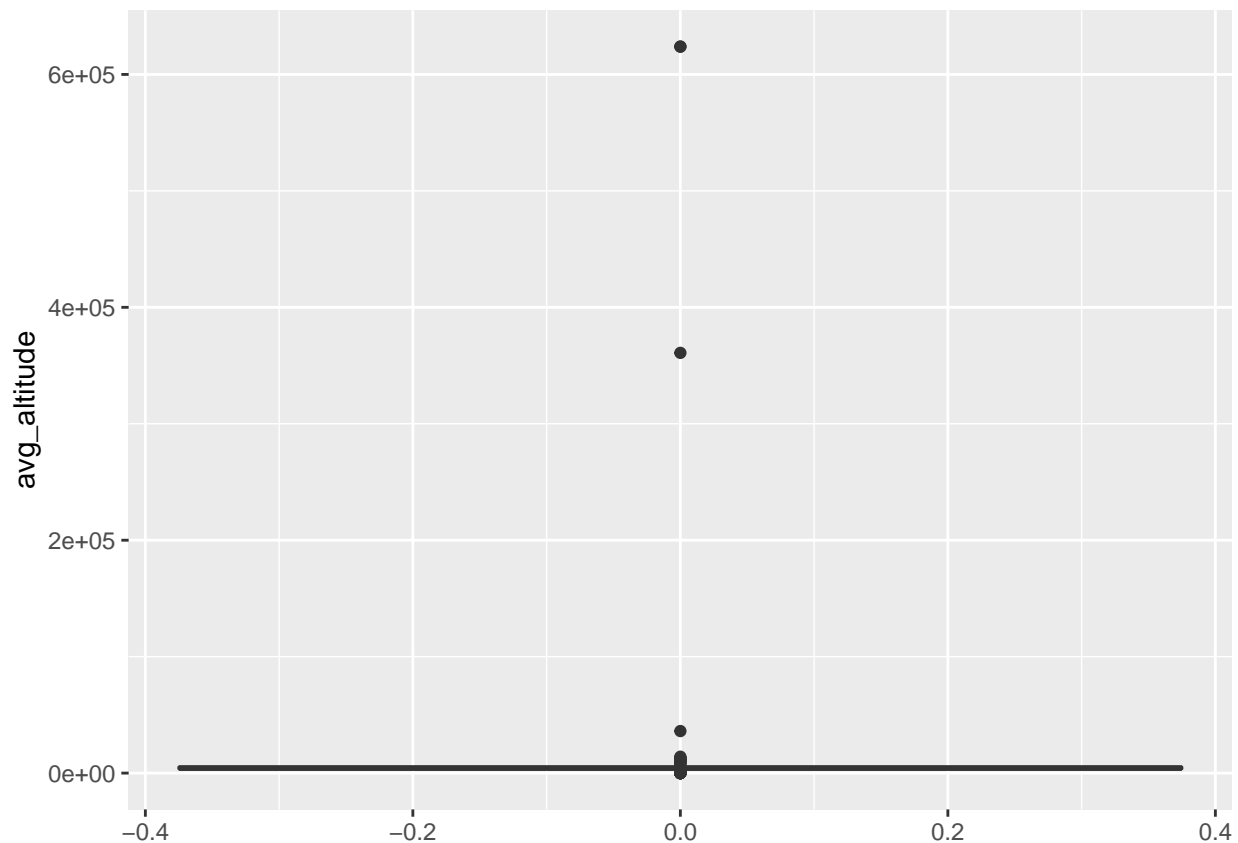
```
library(ggplot2)
```

check for outliers in some of the fields

```
defect1_plt = ggplot(coffee, aes(y=category_one_defects)) +
  geom_boxplot()
defect2_plt = ggplot(coffee, aes(y=category_two_defects)) +
  geom_boxplot()
alt_plt = ggplot(coffee, aes(y=avg_altitude)) +
  geom_boxplot()
defect1_plt
```







There are some outliers, but not that many that would result in a concern at this time.

Pick out some of the information that is not necessary at this point in exploration

```
c = coffee[,c(1:2,4,10:26,28)]
```

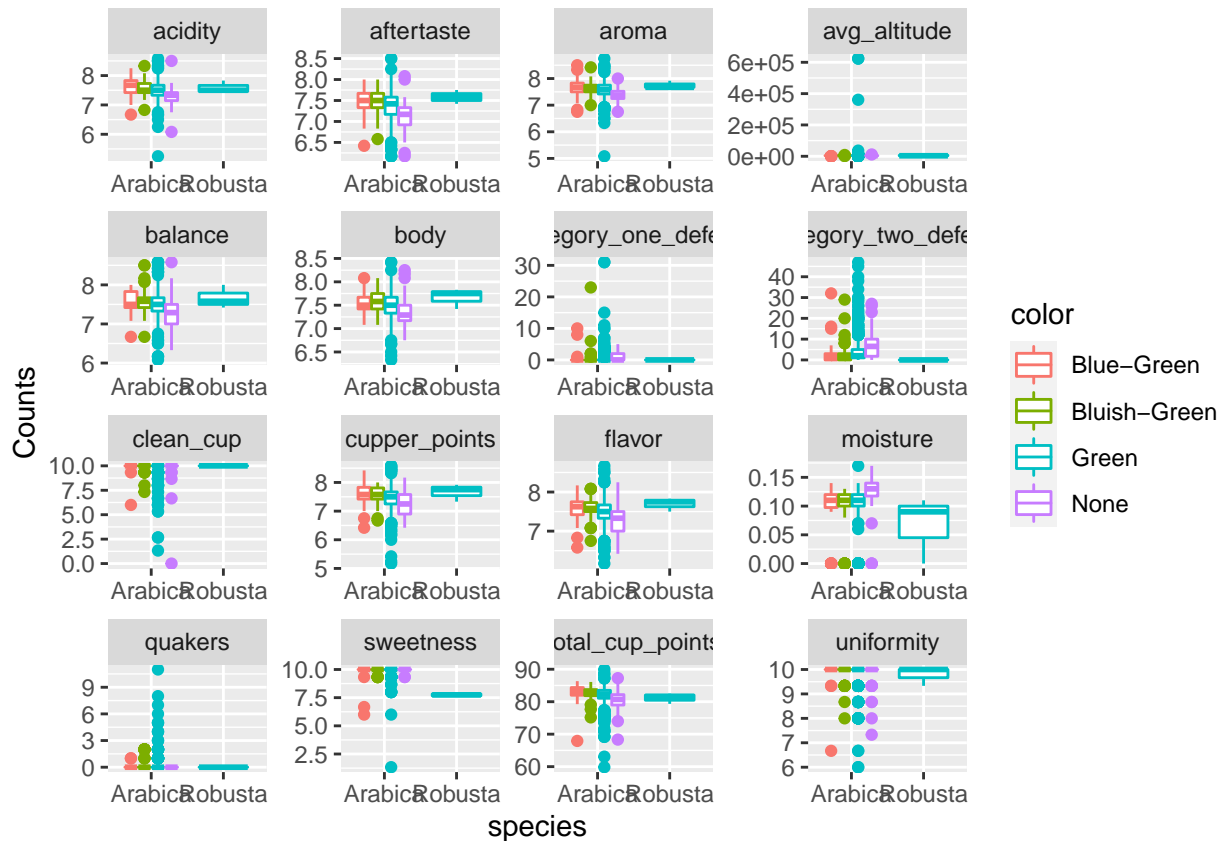
Condense the data

```
c.v1 = c%>%pivot_longer(
  cols = !c(species, country_of_origin,variety,processing_method,color),
  names_to = "Variables",
  values_to = "Counts")
c.v1
```

```
## # A tibble: 14,304 x 7
##   species country_of_origin variety processing_method color Variables    Counts
##   <chr>    <chr>          <chr>    <chr>          <chr> <chr>      <dbl>
## 1 Arabica Ethiopia      Other Washed / Wet    Green total_cup_p~ 89.9
## 2 Arabica Ethiopia      Other Washed / Wet    Green aroma        8.75
## 3 Arabica Ethiopia      Other Washed / Wet    Green flavor        8.67
## 4 Arabica Ethiopia      Other Washed / Wet    Green aftertaste    8.5
## 5 Arabica Ethiopia      Other Washed / Wet    Green acidity       8.58
## 6 Arabica Ethiopia      Other Washed / Wet    Green body          8.42
## 7 Arabica Ethiopia      Other Washed / Wet    Green balance       8.42
## 8 Arabica Ethiopia      Other Washed / Wet    Green uniformity    10
## 9 Arabica Ethiopia      Other Washed / Wet    Green clean_cup     10
## 10 Arabica Ethiopia      Other Washed / Wet    Green sweetness     10
## # ... with 14,294 more rows
```

Plot the data to see overall behavior

```
ggplot(c.v1,aes(x=species,y=Counts,color=color))+geom_boxplot()+facet_wrap(~Variables,scales = "free")
```

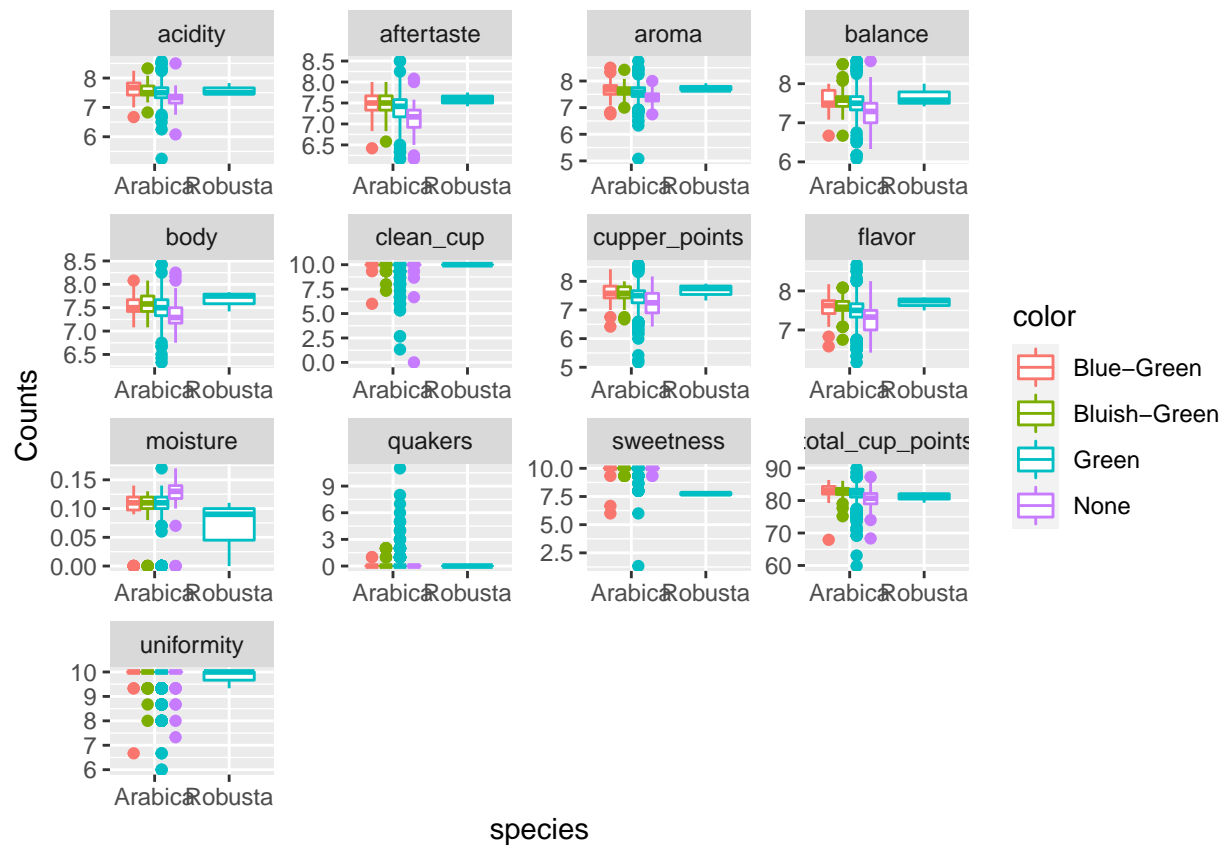


File-

ter out the items that have known outliers

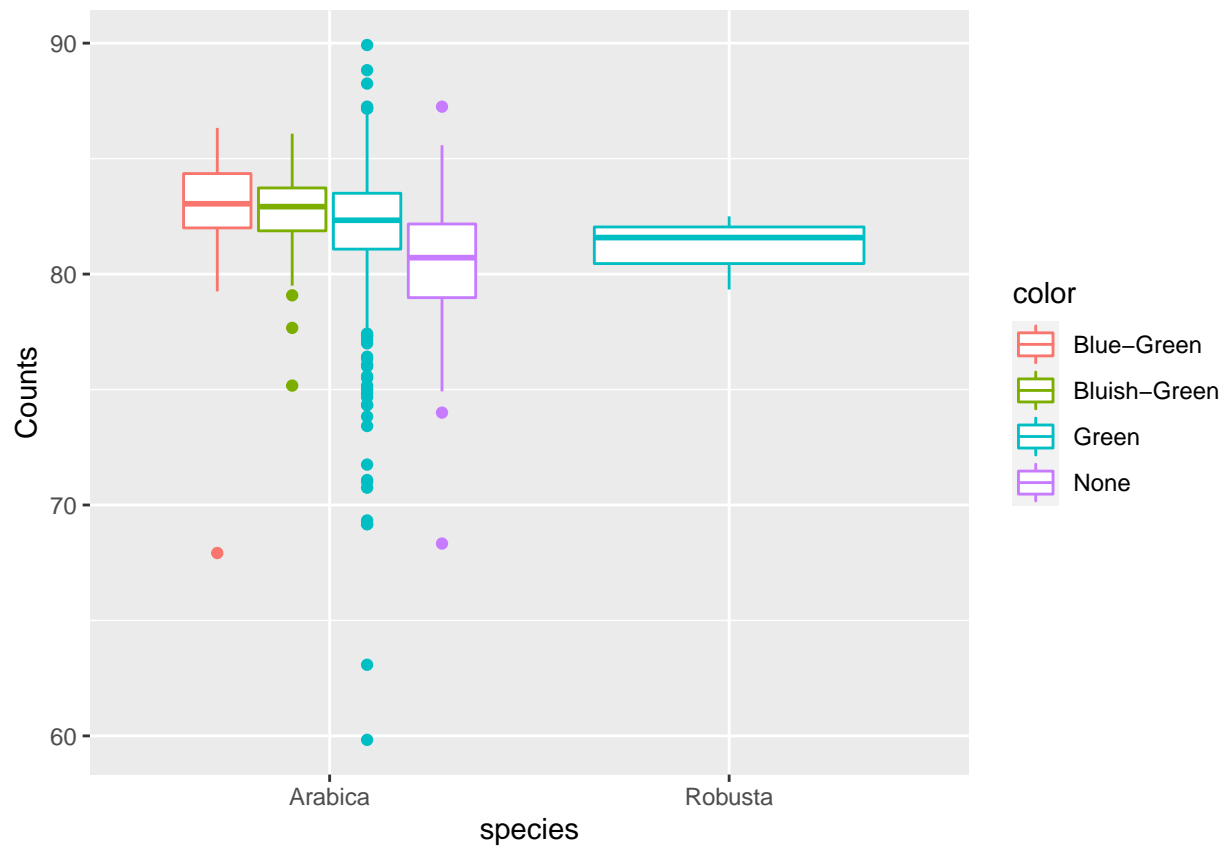
```
c.v2 = c.v1 %>%
  filter(Variables != 'avg_altitude' & Variables != 'category_one_defects' & Variables != 'category_two_defects')
```

```
ggplot(c.v2,aes(x=species,y=Counts,color=color))+geom_boxplot()+facet_wrap(~Variables,scales = "free")
```

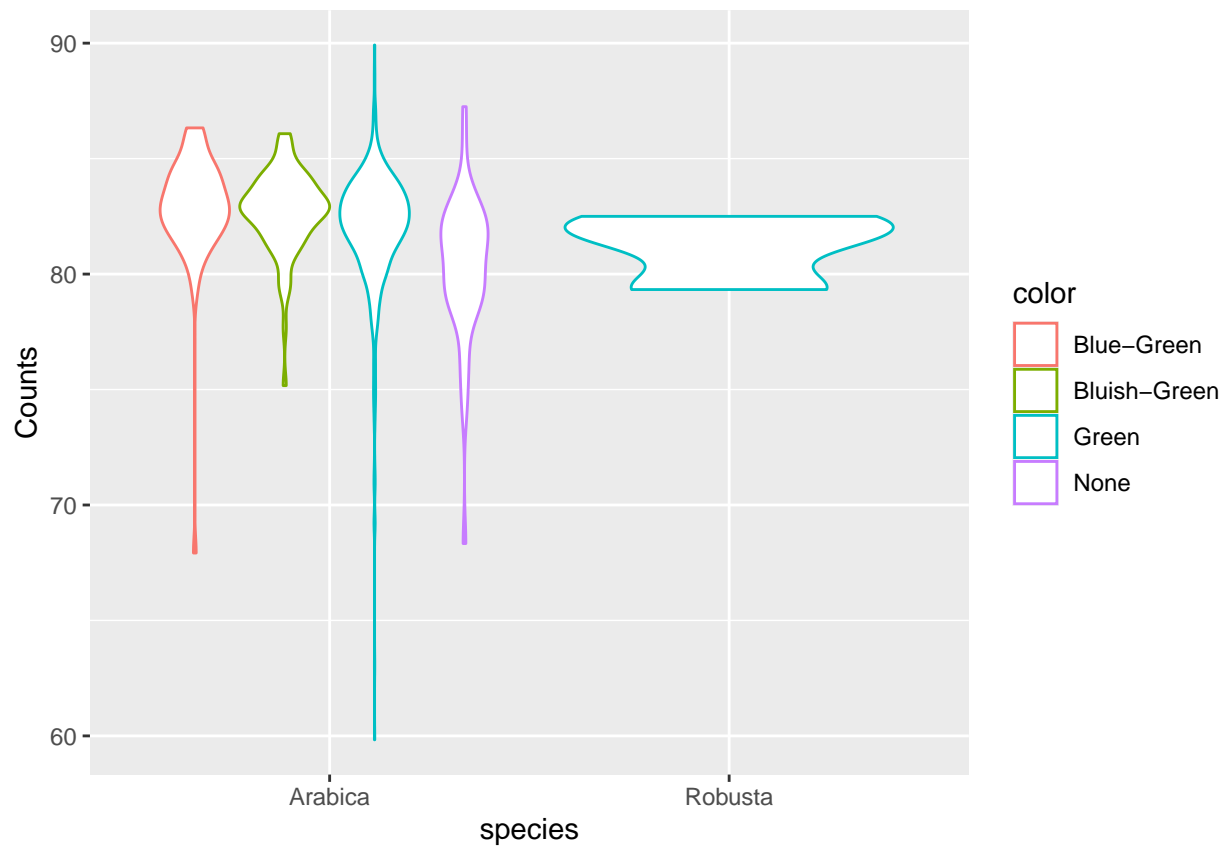


How are the cup points distributed and where the weight it is at

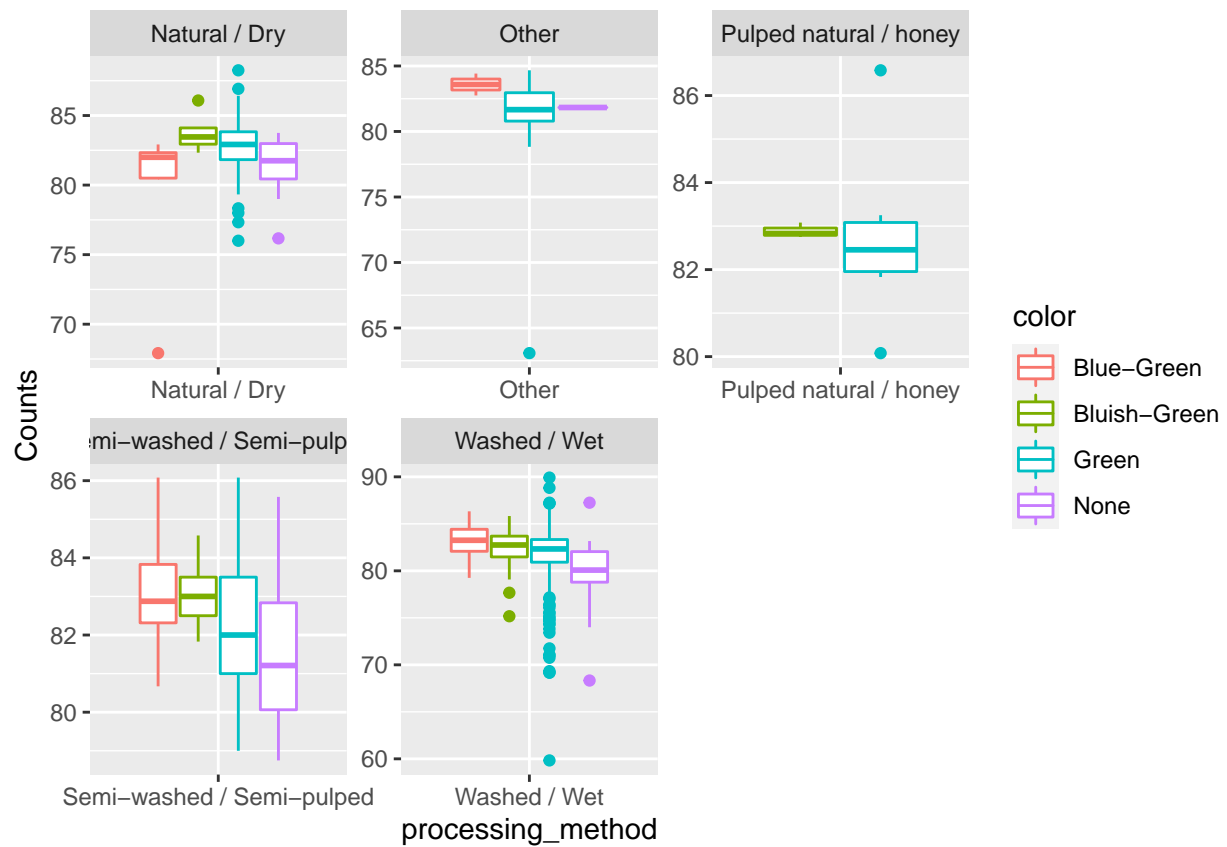
```
c.v2 %>%
  filter(Variables == 'total_cup_points')%>%
  ggplot(aes(x=species,y=Counts,color=color))+geom_boxplot()
```



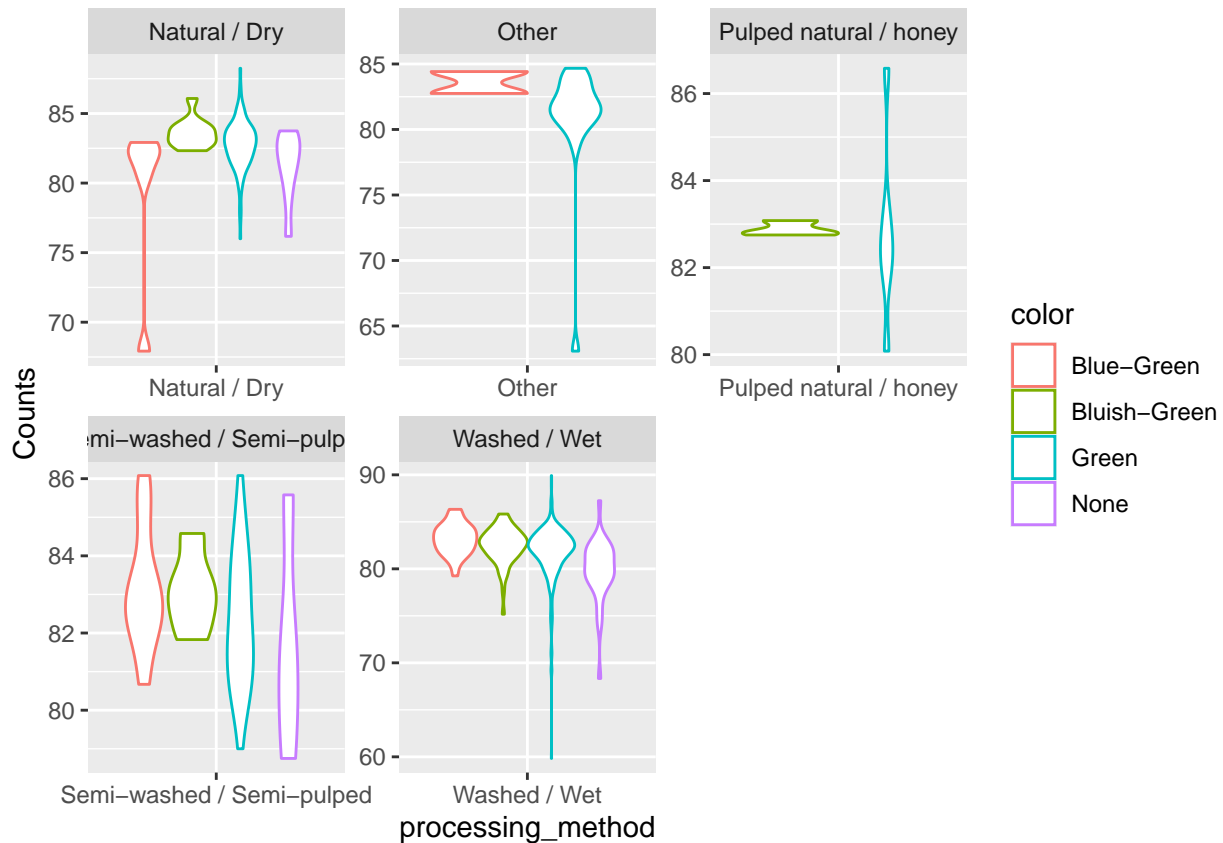
```
c.v2 %>%
  filter(Variables == 'total_cup_points')%>%
  ggplot(aes(x=species,y=Counts,color=color))+geom_violin()
```



```
c.v2 %>%
  filter(Variables == 'total_cup_points')%>%
  ggplot(aes(x=processing_method,y=Counts,color=color))+geom_boxplot()+
  facet_wrap(~processing_method,scales = "free")
```



```
c.v2 %>%
  filter(Variables == 'total_cup_points')%>%
  ggplot(aes(x=processing_method,y=Counts,color=color))+geom_violin()+
  facet_wrap(~processing_method,scales = "free")
```



See the data make-up in a different view

```
library(formattable)
```

```
#want to make this cleaner
```

```
c.v1 %>%
```

```
  group_by(Variables, Counts) %>%
```

```
  summarise(count = n()) %>%
```

```
  mutate(freq = formattable::percent(count / sum(count)))
```

```
## `summarise()` has grouped output by 'Variables'. You can override using the `.groups` argument.
```

```
## # A tibble: 611 x 4
```

```
## # Groups:   Variables [16]
```

```
##   Variables Counts count freq
```

```
##   <chr>      <dbl> <int> <formttbl>
```

```
## 1 acidity    5.25     1 0.11%
```

```
## 2 acidity    6.08     1 0.11%
```

```
## 3 acidity    6.25     1 0.11%
```

```
## 4 acidity    6.5      1 0.11%
```

```
## 5 acidity    6.67     3 0.34%
```

```
## 6 acidity    6.75     2 0.22%
```

```
## 7 acidity    6.83     6 0.67%
```

```
## 8 acidity    6.92     7 0.78%
```

```
## 9 acidity    7       23 2.57%
```

```
## 10 acidity   7.08    25 2.80%
```

```
## # ... with 601 more rows
```

Format the label (total_cup_points) to be categorical

```

coffee$tcp = coffee$total_cup_points

for(i in 1:894){
  if(coffee[i,29] >= 80){
    coffee[i,29] = 80
  }
  else if(coffee[i,29] >= 70 & coffee[i,29] < 80){
    coffee[i,29] = 70
  }
  else if(coffee[i,29] >= 60 & coffee[i,29] < 70){
    coffee[i,29] = 60
  }
  else{
    coffee[i,29] = 50
  }
}
coffee$tcp = round(coffee$tcp,0)

```

For analysis

```

df = coffee[,c(9:22,25,29)]
for(i in 4 : 13){
  df[,i]=round(df[,i],1)
}

```

Accuracy table for comparison between models

```

table_accuracy = matrix(nrow=6,ncol=1)
colnames(table_accuracy) = c('Accuracy')
rownames(table_accuracy) = c('DTree','NB','SVM-Linerar','SVM-Polynomial','ANN','KNN')
table_accuracy

```

```

##           Accuracy
## DTree           NA
## NB              NA
## SVM-Linerar     NA
## SVM-Polynomial  NA
## ANN            NA
## KNN            NA

```

Set seed so analysis is repeatable

```

set.seed(1)

```

Simple k-fold cross validation(cv) function

```

cv = function(data,k){
  n = nrow(data)
  folds = k
  tail = n%%folds

  set.seed(1)

  rnd = runif(n)
  rank = rank(rnd)

  #block/chunck from cv is blk

```



```

blk = (rank-1)%/%tail+1
blk = as.factor(blk)

#to see formation of folds
print(summary(blk))
}

n = nrow(df)
folds = 10
tail = n%/%folds

set.seed(1)

rnd = runif(n)
rank = rank(rnd)

#block/chunck from cv is blk
blk = (rank-1)%/%tail+1
blk = as.factor(blk)

#to see formation of folds
print(summary(blk))

## 1 2 3 4 5 6 7 8 9 10 11
## 89 89 89 89 89 89 89 89 89 89 4

df$tcp = as.factor(df$tcp)
df$moisture = round(df$moisture,1)

df = df[,c(4:13,16)]

library(rpart)
#dtree
set.seed(1)
all.acc = numeric(0)
for(i in 1:folds){
  tree = rpart(tcp~.,df[blk != i,],method="class")
  pred = predict(tree,df[blk==i,],type="class")
  confMat = table(pred,df$tcp[blk==i])
  acc = (confMat[1,1]+confMat[2,2]+confMat[3,3]+confMat[4,4])/sum(confMat)
  all.acc = rbind(all.acc,acc)
}

print(mean(all.acc))

## [1] 0.947191
table_accuracy[1,1] = mean(all.acc)

```

I re-formatted the label/target field and went from a binary (good/bad) grading and could not figure out why the accuracy was so low (0.003) and then looked into what the accuracy was calculating...

```

confMat

##
## pred 50 60 70 80
##    50  0  0  0  0
##    60  0  0  0  0

```

```
## 70 0 0 13 0
## 80 0 0 3 73

# naive Bayes (gaussian data)
library(e1071)
set.seed(1)

all.acc = numeric(0)
for(i in 1:folds){
  model = naiveBayes(tcp~.,df[blk != i,],method="class")
  pred = predict(model,df[blk==i,],type="class")
  confMat = table(pred,df$tcp[blk==i])
  acc = (confMat[1,1]+confMat[2,2]+confMat[3,3]+confMat[4,4])/sum(confMat)
  all.acc = rbind(all.acc,acc)
}

print(mean(all.acc))
```

```
## [1] 0.952809
table_accuracy[2,1] = mean(all.acc)
```

```
#svm linear
set.seed(1)

all.acc = numeric(0)
for(i in 1:folds){
  model = svm(tcp~.,df[blk != i,],kernel="linear",type="C")
  pred = predict(model,df[blk==i,],type="class")
  confMat = table(pred,df$tcp[blk==i])
  acc = (confMat[1,1]+confMat[2,2]+confMat[3,3]+confMat[4,4])/sum(confMat)
  all.acc = rbind(all.acc,acc)
}

print(mean(all.acc))
```

```
## [1] 0.9842697
table_accuracy[3,1] = mean(all.acc)
```

```
#svm poly
set.seed(1)

all.acc = numeric(0)
for(i in 1:folds){
  model = svm(tcp~.,df[blk != i,],kernel="polynomial",type="C")
  pred = predict(model,df[blk==i,],type="class")
  confMat = table(pred,df$tcp[blk==i])
  acc = (confMat[1,1]+confMat[2,2]+confMat[3,3]+confMat[4,4])/sum(confMat)
  all.acc = rbind(all.acc,acc)
}

print(mean(all.acc))
```

```
## [1] 0.9730337
```

```
table_accuracy[4,1] = mean(all.acc)
```

```
df$tcp = as.factor(df$tcp)
#ann
library(nnet)
set.seed(1)

all.acc = numeric(0)
for(i in 1:folds){
  model = nnet(tcp~.,df[blk != i,], size = 5, trace=FALSE, wgts=.05)
  pred = predict(model, df[blk==i,],type="class")
  confMat = table(factor(pred,levels=1:4),factor(df$tcp[blk==i],levels=1:4))
  acc = (confMat[1,1]+confMat[2,2]+confMat[3,3]+confMat[4,4])/sum(confMat)
  all.acc = rbind(all.acc,acc)
}
print(mean(all.acc))
```

```
## [1] NaN
```

```
table_accuracy[5,1] = mean(all.acc)
```

There was an (un)interesting issue with NN table, as it was dropping the first two rows as it was not forward feeding into those nodes. The following is the work around to resolve this issue.

```
#Before#
```

```
set.seed(1)
i=1
model = nnet(tcp~.,df[blk != i,], size = 5, trace=FALSE, wgts=.05)
```

```
## Warning in nnet.formula(tcp ~ ., df[blk != i, ], size = 5, trace = FALSE, :
## group '50' is empty
```

```
pred = predict(model, df[blk==i,],type="class")
confMat = table(pred,df$tcp[blk==i])
confMat
```

```
##
## pred 50 60 70 80
##    50  1  0  1  1
##    60  0  0 11  2
##    70  0  0  4 69
```

```
#After#
```

```
set.seed(1)
i=1
model = nnet(tcp~.,df[blk != i,], size = 5, trace=FALSE, wgts=.05)
```

```
## Warning in nnet.formula(tcp ~ ., df[blk != i, ], size = 5, trace = FALSE, :
## group '50' is empty
```

```
pred = predict(model, df[blk==i,],type="class")
confMat = table(factor(pred,levels=1:4),factor(df$tcp[blk==i],levels=1:4))
confMat
```

```
##
##      1 2 3 4
##    1 0 0 0 0
```

```
## 2 0 0 0 0
## 3 0 0 0 0
## 4 0 0 0 0

df$tcp = as.factor(df$tcp)
library (caret)

## Loading required package: lattice
##
## Attaching package: 'caret'
## The following object is masked from 'package:purrr':
##
## lift
trControl <- trainControl(method = "cv", number = 10)

knn = df[,]

model <- train(tcp ~ .,
               method = "knn",
               tuneGrid = expand.grid(k = 1:10),
               trControl = trControl,
               data = knn)

acc = mean(model$results$Accuracy)
table_accuracy[6,1] = acc

tab = round(table_accuracy,4)
tab
```

	Accuracy
## DTree	0.9472
## NB	0.9528
## SVM-Linerar	0.9843
## SVM-Polynomial	0.9730
## ANN	NaN
## KNN	0.9734