

STAT 6500

Statistical Machine Learning

Land Use Cover - EDA

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Spring 2022
Wednesday, March 2

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Introduction

Dataset information for this project is obtained from UCI Machine Learning repository. The data includes training and testing data for classifying a high resolution aerial image into 9 types of urban land cover which are trees, grass, soil, concrete, asphalt, buildings, cars, pools, shadows. Features such as Multi-scale spectral, size, shape, and texture information are used for classification. There are a low number of training samples for each class (14-30) and a high number of classification variables or features (148 features in total). The testing data set is from a random sampling of the image. As the dataset contains a large number of features, it might be interesting to see their correlations between each pair of these variables in order to filter out some of them that have high correlation and use the remaining feature for prediction.

Details of Attributes

- Class : Land cover class (nominal)
- BrdIndx : Border Index (shape variable)
- Area : Area in m2 (size variable)
- Round : Roundness (shape variable)
- Bright : Brightness (spectral variable)
- Compact : Compactness (shape variable)
- ShpIndx : Shape Index (shape variable)
- Mean_G : Green (spectral variable)
- Mean_R : Red (spectral variable)
- Mean_NIR : Near Infrared (spectral variable)
- SD_G : Standard deviation of Green (texture variable)
- SD_R : Standard deviation of Red (texture variable)
- SD_NIR : Standard deviation of Near Infrared (texture variable)
- LW : Length/Width (shape variable)
- GLCM1 : Gray-Level Co-occurrence Matrix (texture variable)
- Rect : Rectangularity (shape variable)
- GLCM2 : Another Gray-Level Co-occurrence Matrix attribute (texture variable)
- Dens : Density (shape variable)
- Assym : Assymetry (shape variable)

- NDVI : Normalized Difference Vegetation Index (spectral variable)
- BordLngh: Border Length (shape variable)
- GLCM3 : Another Gray-Level Co-occurrence Matrix attribute (texture variable)
- Note: These variables repeat for each coarser scale (i.e. variable_40, variable_60, ... variable_140).

Loading Data

```
#load in training and test data
train = pd.read_csv('training.csv')
test = pd.read_csv('testing.csv')
print("Rows and Columns(Train): ",train.shape)
```

```
## Rows and Columns(Train): (168, 148)
```

```
print("Rows and Columns(Test) : ",test.shape)
```

```
## Rows and Columns(Test) : (507, 148)
```

Removing data with missing value

```
# check for missing values although it is clear there are none
train.isnull().any().any()
```

```
## False
```

```
# duplicated function of pandas returns a duplicate row as true and others as false
sum(train.duplicated())
```

```
## 0
```

Basic Statistics

```
# basic statistical details
fig = train.describe().T
fig = fig.round(5) # round to 5 decimal places
table = go.Table(
    columnwidth=[0.8]+[0.5]*8,
    header=dict(
        values=['Attribute'] + list(fig.columns),
        line = dict(color='darkslategray'),
```

```

        fill = dict(color='royalblue'),
    ),
    cells=dict(
        values=[fig.index] + [fig[k].tolist() for k in fig.columns[:]],
        line = dict(color='darkslategray'),
        fill = dict(color=['paleturquoise', 'white'])
    )
)
plot([table], filename='table-of-data.html')

```

```
## 'table-of-data.html'
```

```

# more general data exploration
print(train['class'].value_counts())

```

```

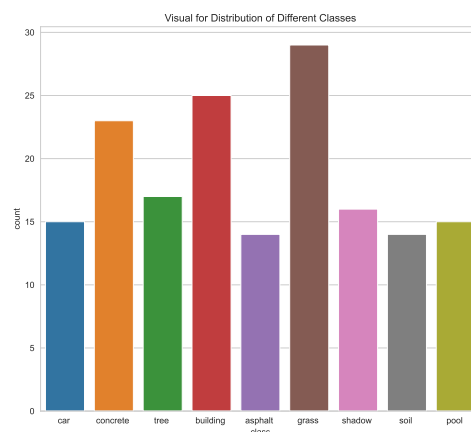
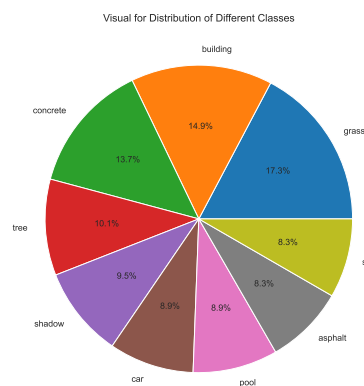
## grass      29
## building   25
## concrete   23
## tree       17
## shadow     16
## car        15
## pool       15
## asphalt    14
## soil       14
## Name: class, dtype: int64

```

```

f,axes=plt.subplots(1,2,figsize=(20,8))
train['class'].value_counts().plot.pie(autopct='%1.1f%%',ax=axes[0])
axes[0].set_title('Visual for Distribution of Different Classes')
axes[0].set_ylabel('')
sns.countplot('class',data=train,ax=axes[1]) # sns.countplot is used
                                              # like a histogram but for
                                              # categorical data
axes[1].set_title('Visual for Distribution of Different Classes')
plt.show()

```



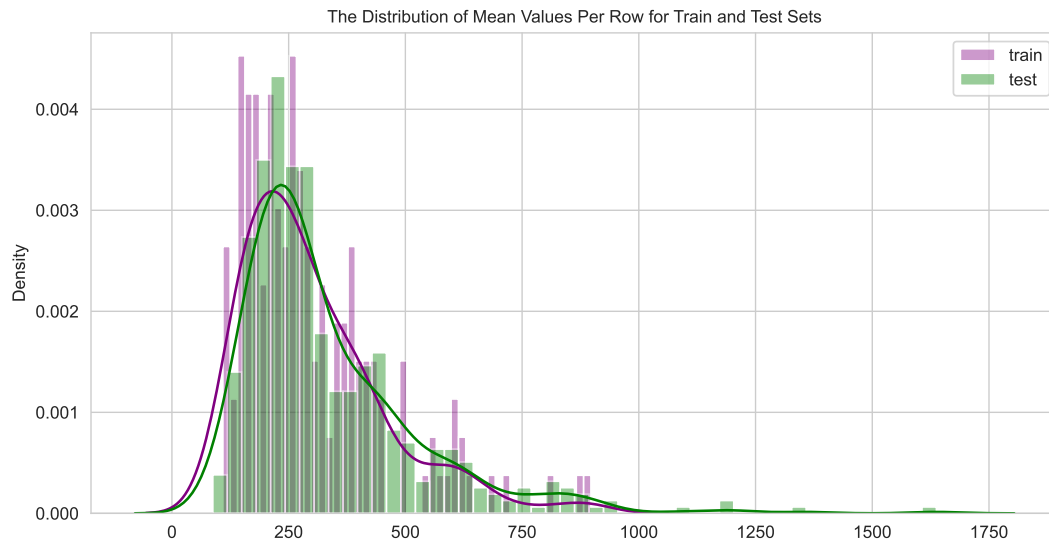
Outlier examination

```
# Lets take a look at any outliers that could be potential issues
from collections import Counter
def examine_outliers(train_data, n, features):
    outlier_indicator = []
    for out in features:
        Q1 = np.percentile(train_data[out], 25)
        Q3 = np.percentile(train_data[out], 75)
        IQR = Q3 - Q1
        outlier_step = 1.5 * IQR # IQR method of dealing with outliers, 1 of 2 methods
        outlier_list_out = train_data[
            (train_data[out] < Q1 - outlier_step) | (train_data[out] > Q3 +
            outlier_step)].index
        outlier_indicator.extend(outlier_list_out)
        outlier_indices = Counter(outlier_indicator)
        multiple_outliers = list(k for k, j in outlier_indices.items() if j > n)
    return multiple_outliers
# find outliers that should be removed
list_attributes = train.drop('class', axis=1).columns
outliers_to_remove = examine_outliers(train, 2, list_attributes)
len(outliers_to_remove)
#outliers_to_remove
#train.loc[outliers_to_remove]
```

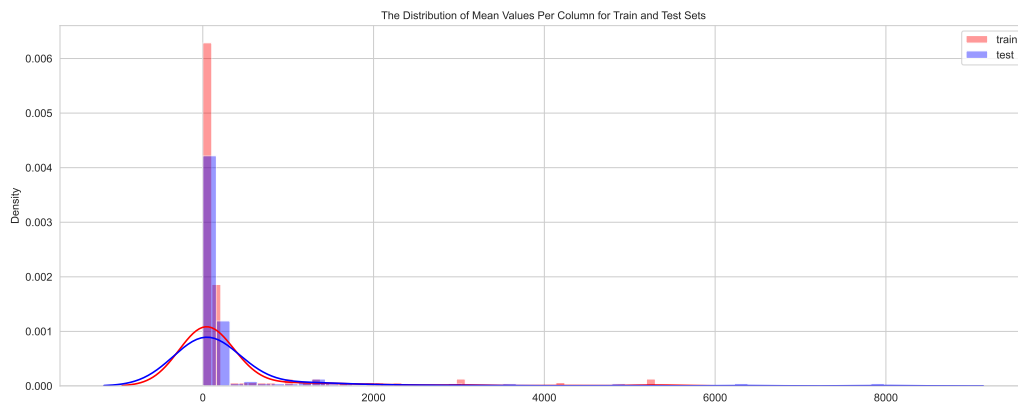
```
## 0
```

Distribution of Mean values

```
# lets look at mean values per row and column for train test data
# mean for rows
plt.figure(figsize=(10,5))
features = train.columns.values[1:148]
plt.title("The Distribution of Mean Values Per Row for Train and Test Sets",
          fontsize=10)
sns.distplot(train[features].mean(axis=1),color="purple", kde=True,bins=50,
              label='train') # kde is kernel density estimation
sns.distplot(test[features].mean(axis=1),color="green", kde=True,bins=50,
              label='test')
plt.legend()
plt.show()
```



```
# mean for columns
plt.figure(figsize=(16,6))
plt.title("The Distribution of Mean Values Per Column for Train and Test Sets",
          fontsize=10)
sns.distplot(train[features].mean(axis=0),color="red",kde=True,bins=50,
              label='train')
sns.distplot(test[features].mean(axis=0),color="blue",kde=True,bins=50,
              label='test')
plt.legend()
plt.show()
```



Attributes Correlation

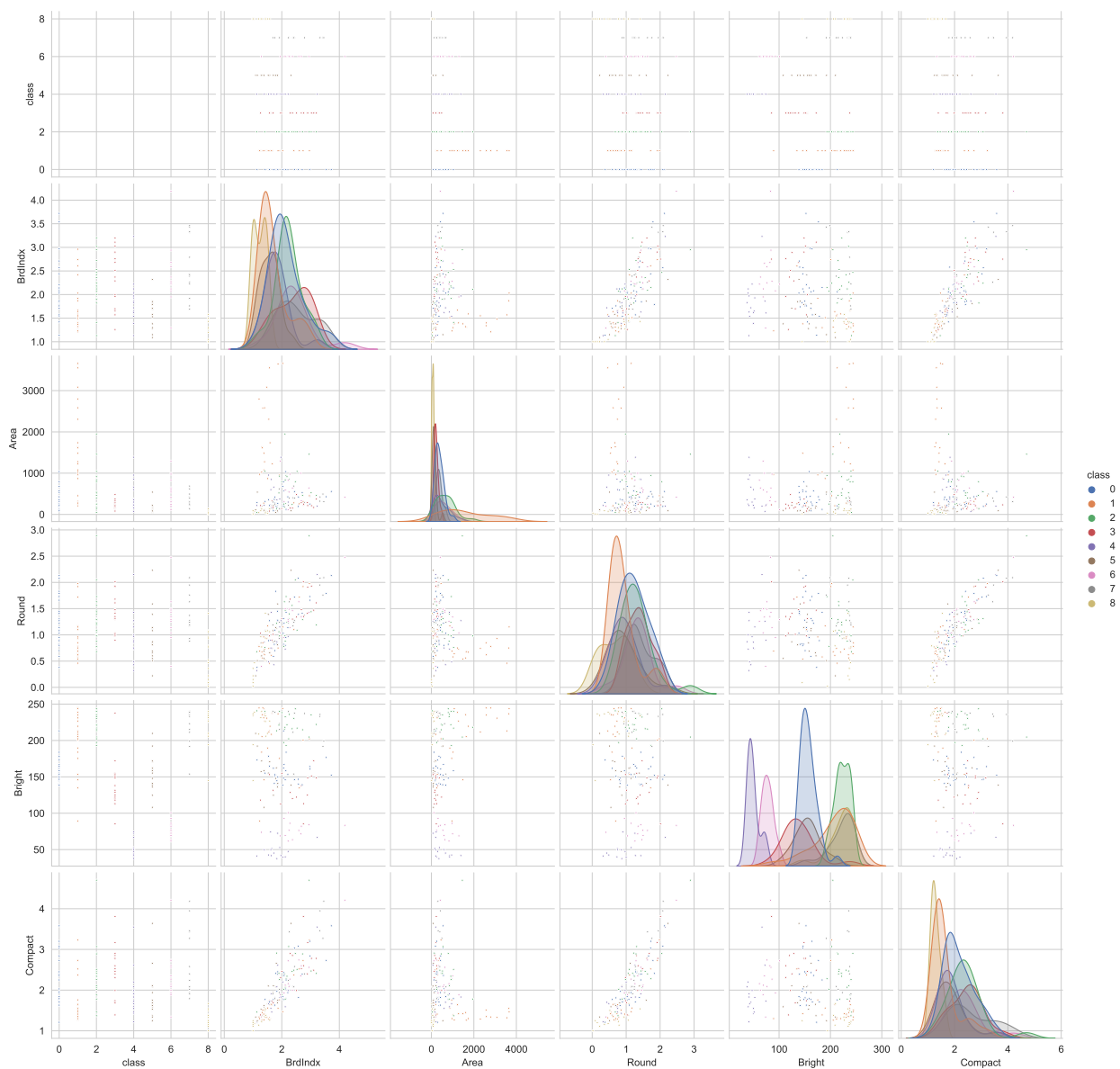
```
# lets examine correlations between features
# first the categorical variable 'class' needs to be changed to numerical variable so...
```

```
group_map = {"grass ":0,"building ":1,'concrete ':2,'tree ':3,'shadow ':4,'pool ':5,
             'asphalt ':6,'soil ':7,'car ':8}
train['class'] = train['class'].map(group_map)
test['class'] = test['class'].map(group_map)
train['class'].unique()
```

```
## array([8, 2, 3, 1, 6, 0, 4, 7, 5], dtype=int64)
```

```
# now lets look at the correlation between a few variables
```

```
sns.pairplot(train, vars=['class', 'BrdIndx', 'Area', 'Round', 'Bright', 'Compact'],
              hue='class', palette='deep', plot_kws={"s": 4})
#plt.show()
```




```
# correlation of features with target
corr = train.corr().abs().unstack().sort_values(kind="quicksort").reset_index()
corr = corr[corr['level_0'] != corr['level_1']]
corr.head()
```

```
##      level_0    level_1      0
## 0      SD_R    GLCM1_60 0.000052
## 1    GLCM1_60      SD_R 0.000052
## 2    Mean_NIR  BordLngth 0.000162
## 3  BordLngth    Mean_NIR 0.000162
## 4    Mean_R_40    Round_40 0.000190
```

```
correlations = corr.loc[corr[0] == 1]
features_to_be_removed = set(list(correlations['level_1']))
correlations.shape
```

```
## (42, 3)
```

```
# A first start would be to investigate each coarseness level
#create a list of dataframes
trainingList <- list()

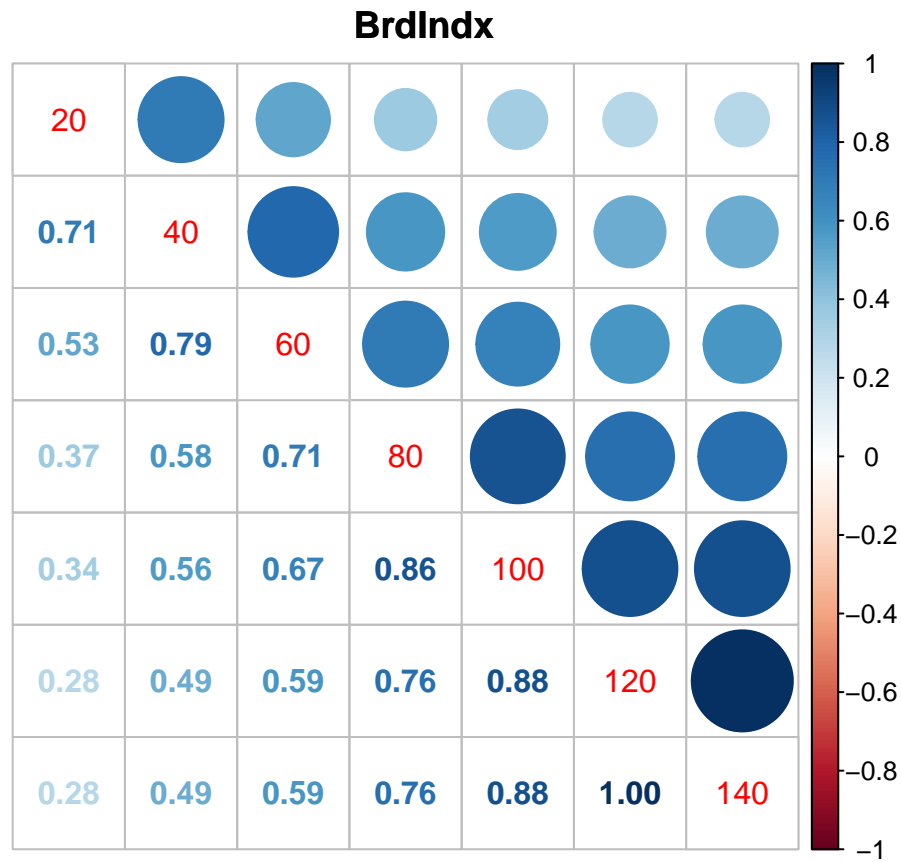
#create new data-frame for each coarseness level

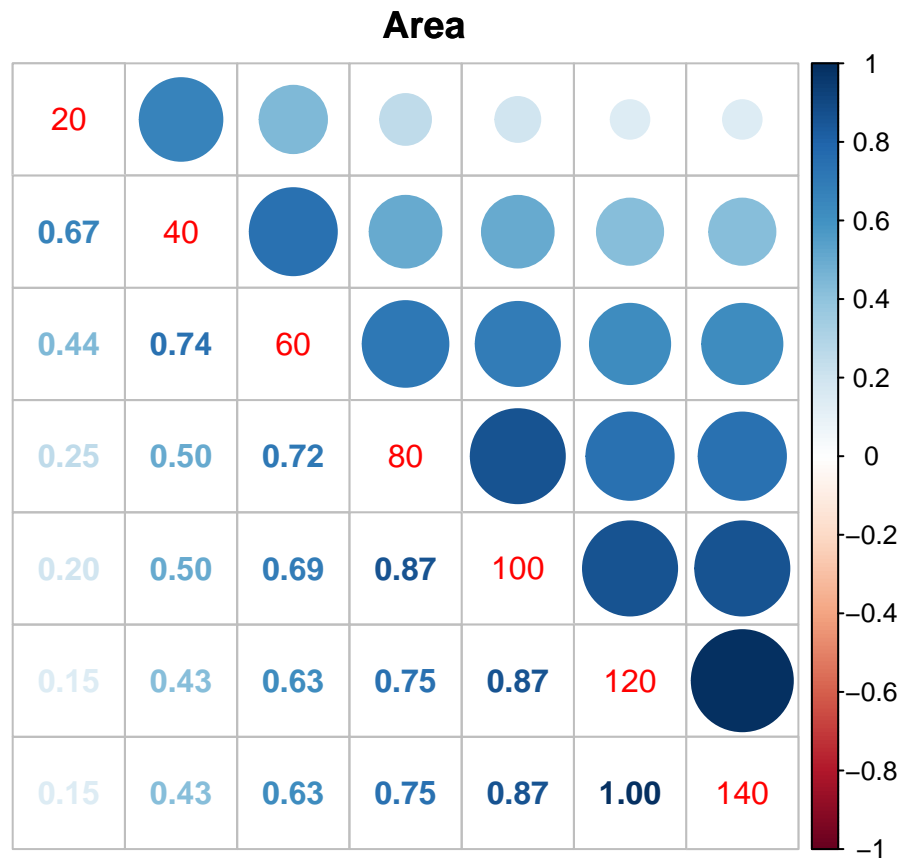
for( i in 1:7){
  #set the name to data + coarseness value
  name<- paste("data", i*20, sep = "")
  #first column index of the selected coarseness
  begin = (i-1)*21+2
  #last column index
  ending = begin+20
  #assign the first column and the columns between first and last
  assign(name, trainData[,c(1,begin:ending)])
  # add to the list
  trainingList[[i]] <- trainData[,c(1,begin:ending)]
}
```

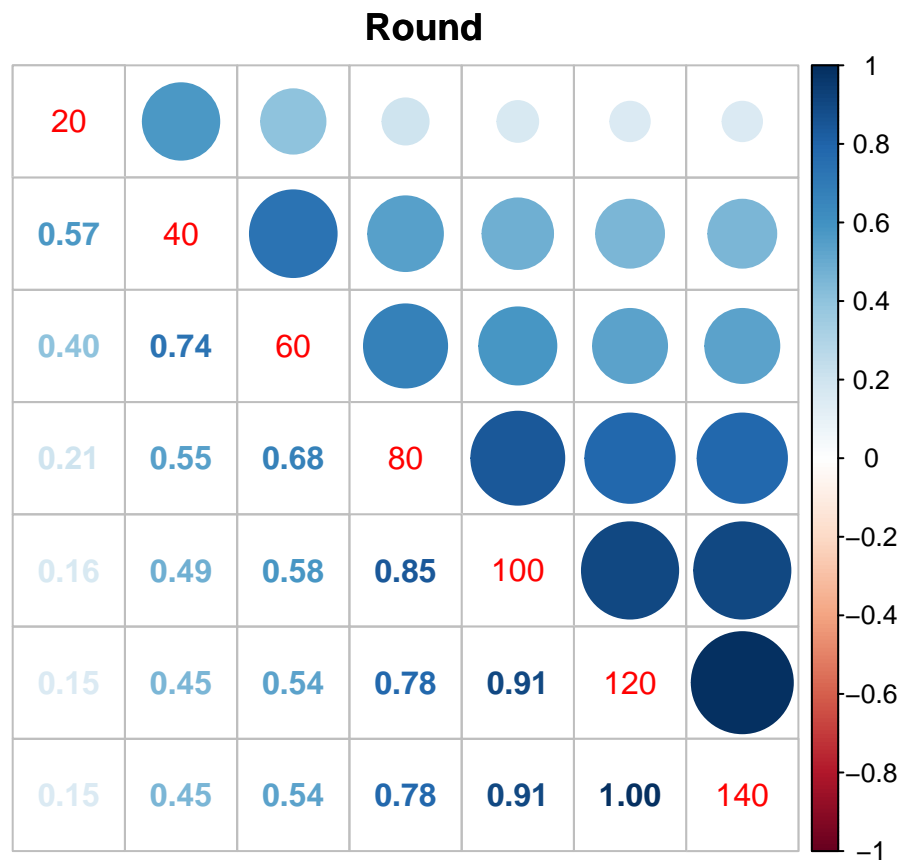
```
# check collinearity among repeated variables

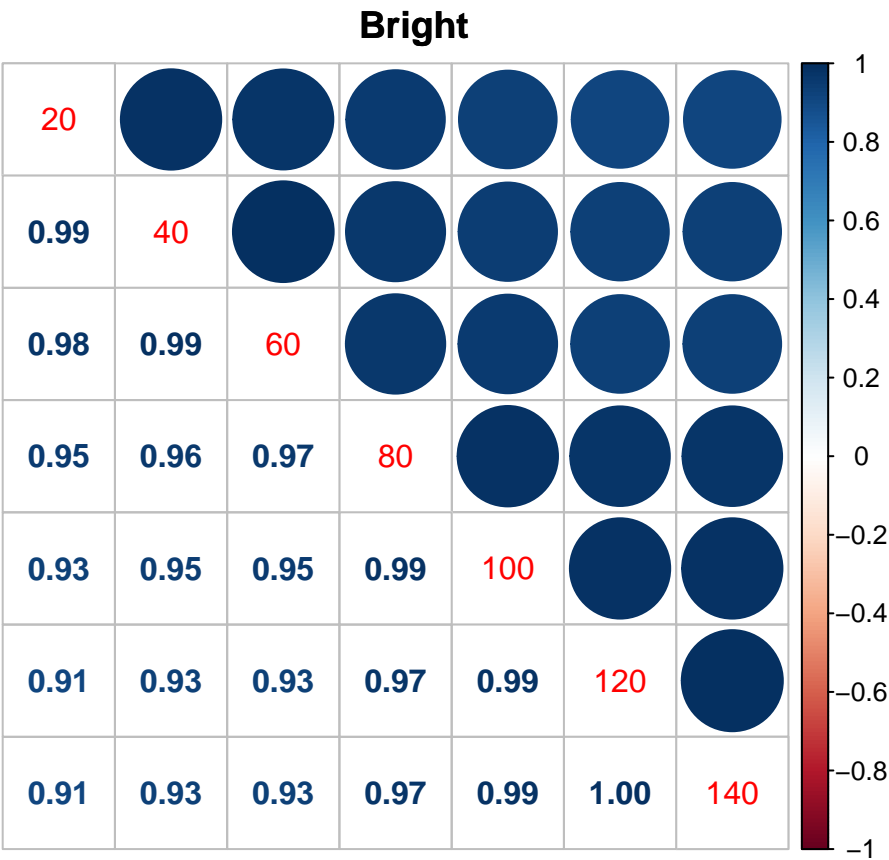
corList <- list()
#looping through all repeated variables
for(i in 2:22){
  #creating a dataframe with coarseness 20, and attribute i
  df <- trainingList[[1]][,i]
  for(j in 2:7){
    #adding the same attribute from a different coarseness level
    df <- cbind(df,trainingList[[j]][,i] )
    colnames(df)[j]<- paste(j*20)
  }
  colnames(df)[1]<- "20"
  #adding a correlation matrix to the list
  corList[[i]] <- cor(df)
```

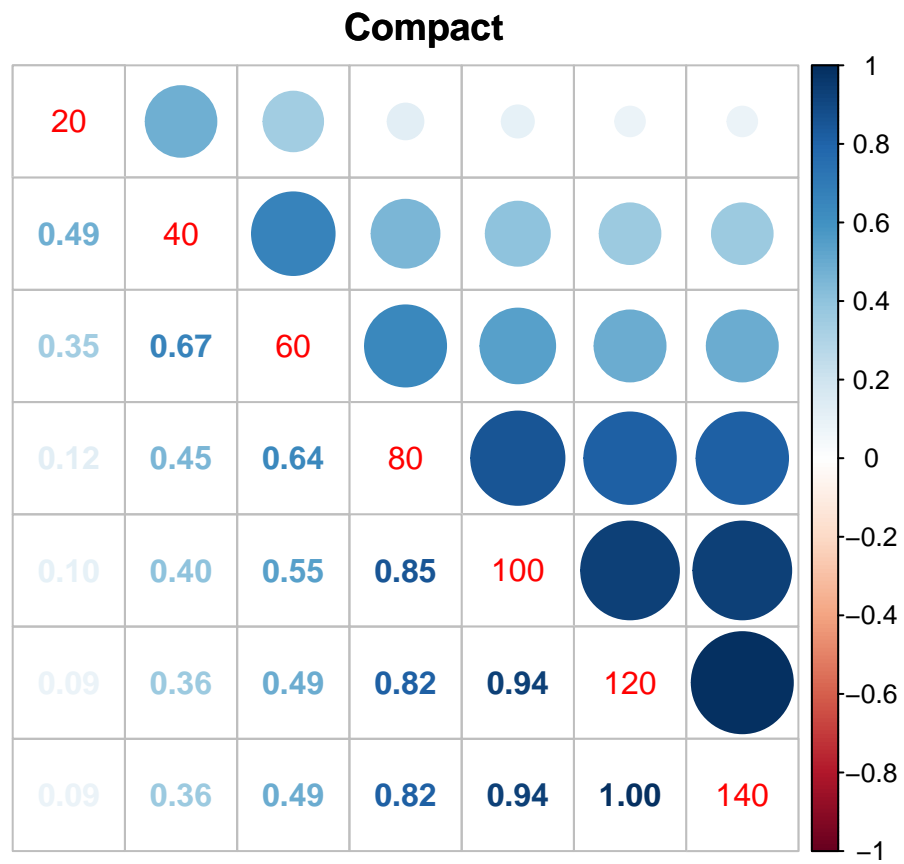
```
#plotting 21 correlation matrices
corrplot.mixed(corList[[i]], upper = 'circle', lower = 'number', title =
               colnames(data20[i]), mar=c(0,0,1,0))
}
```

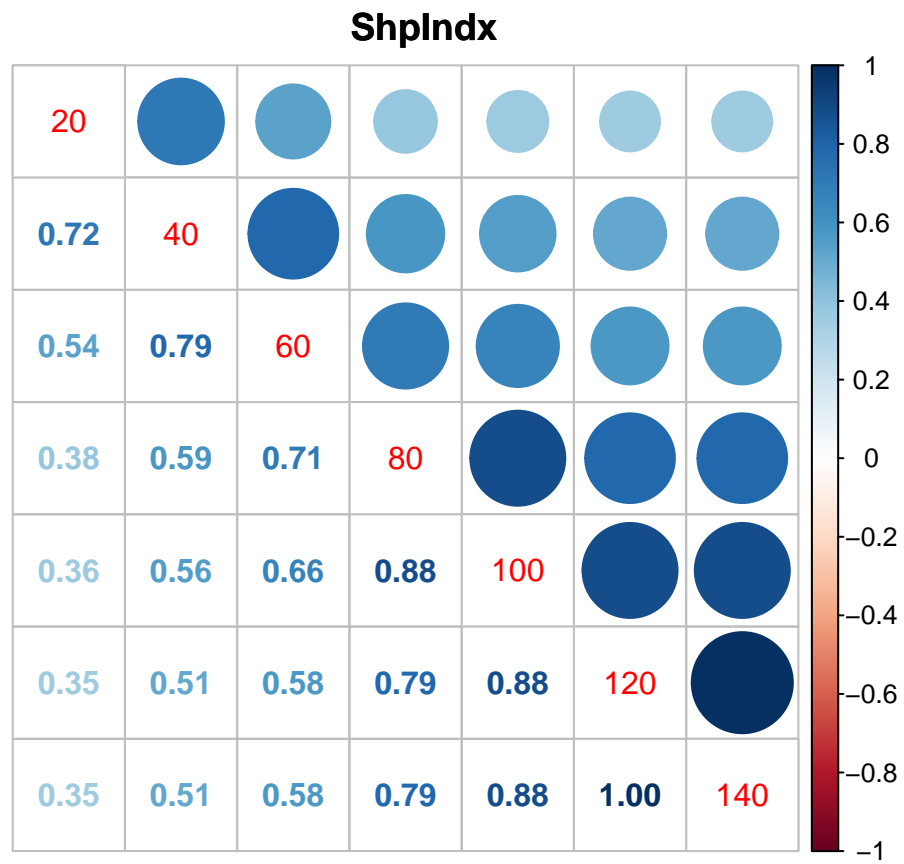


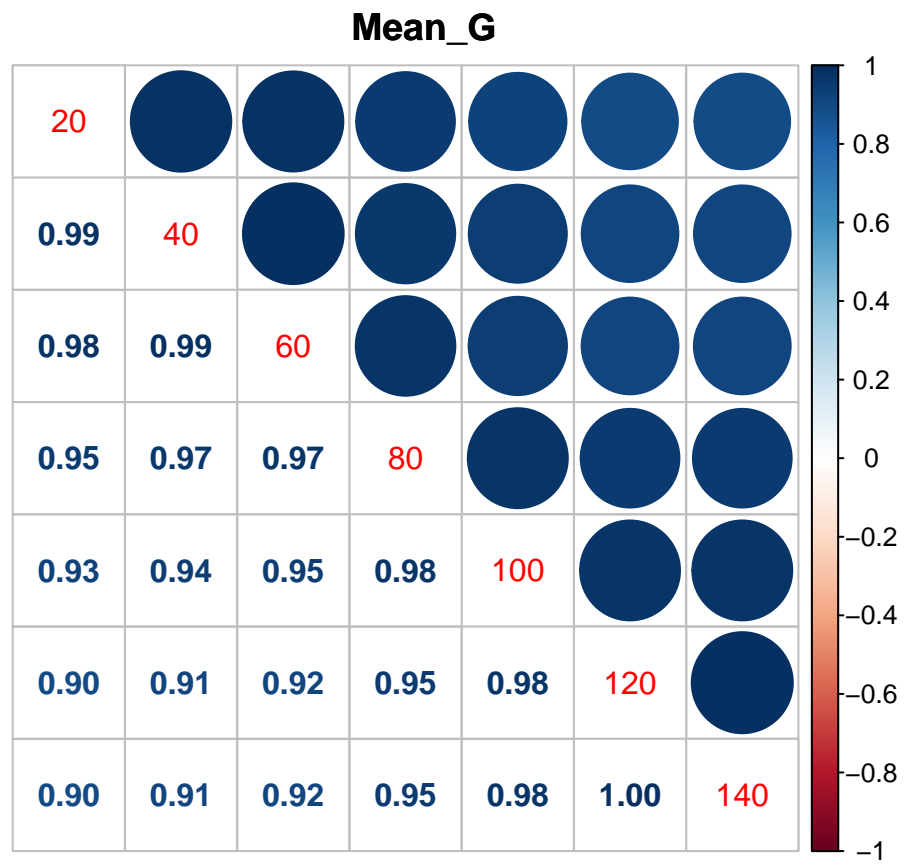


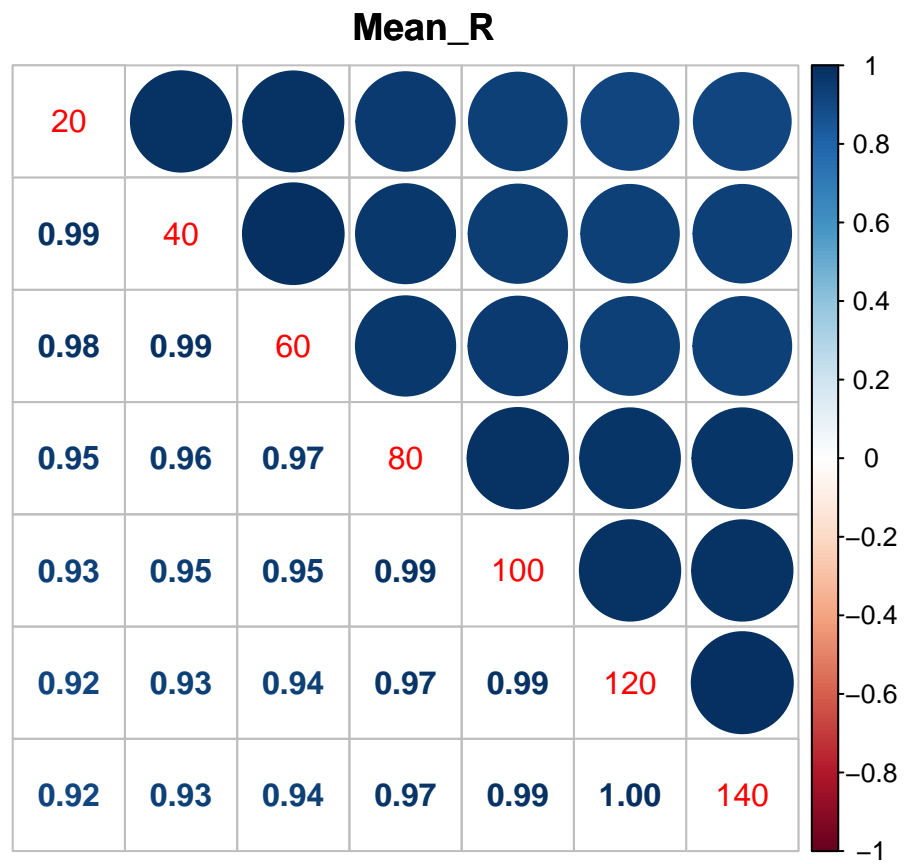


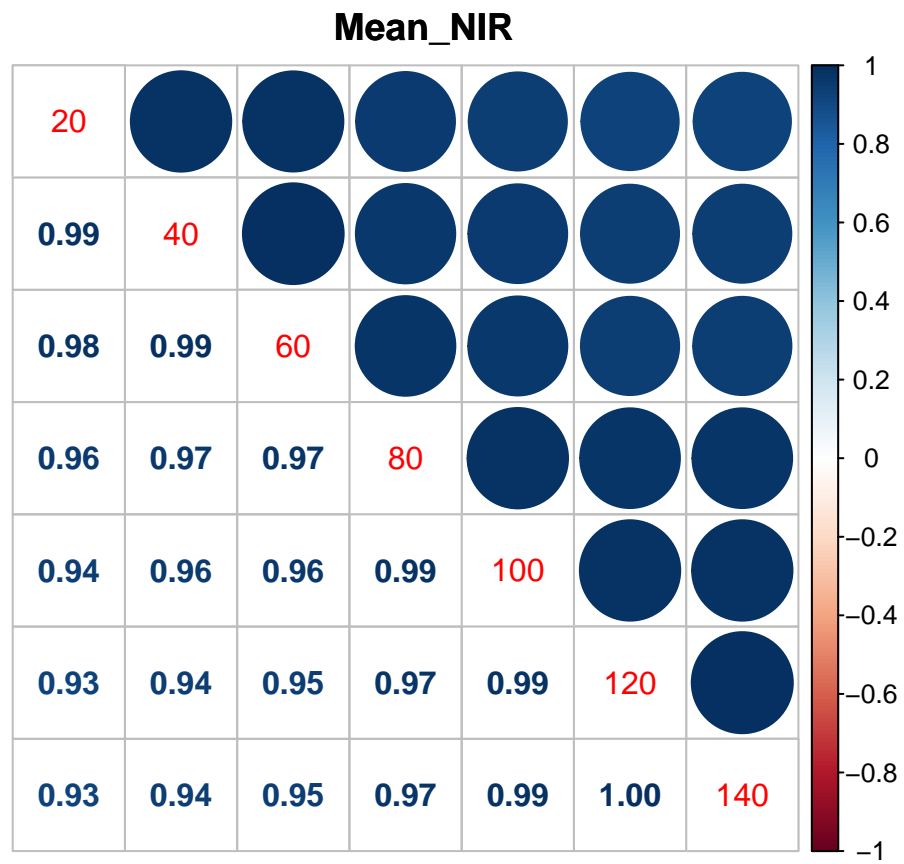


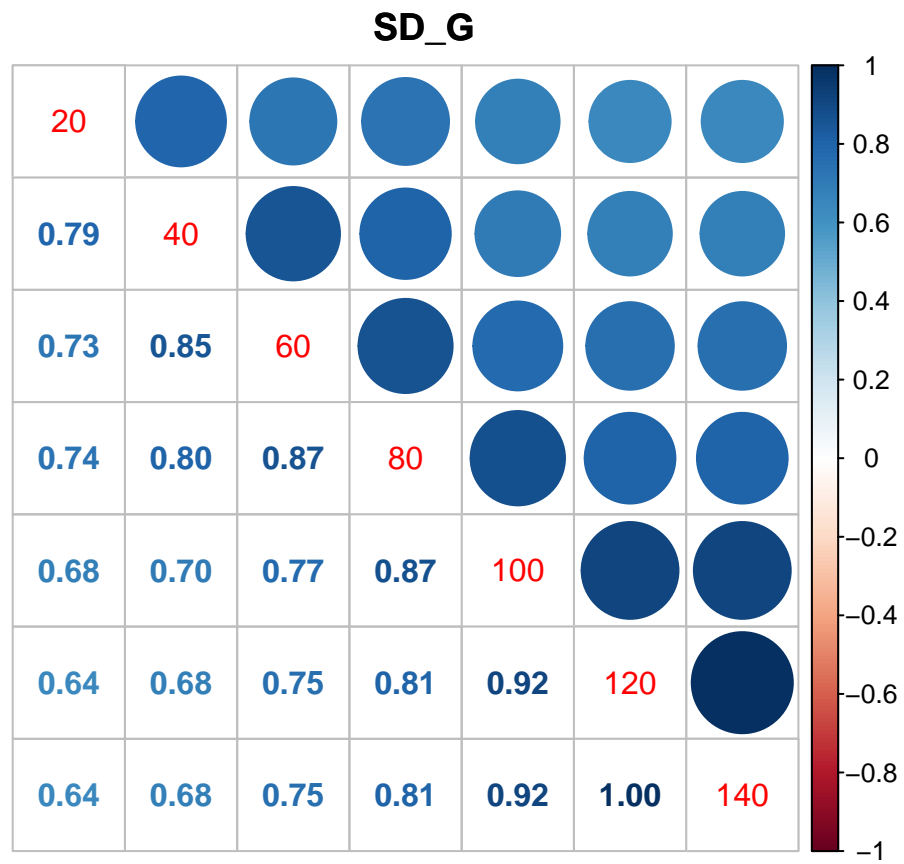


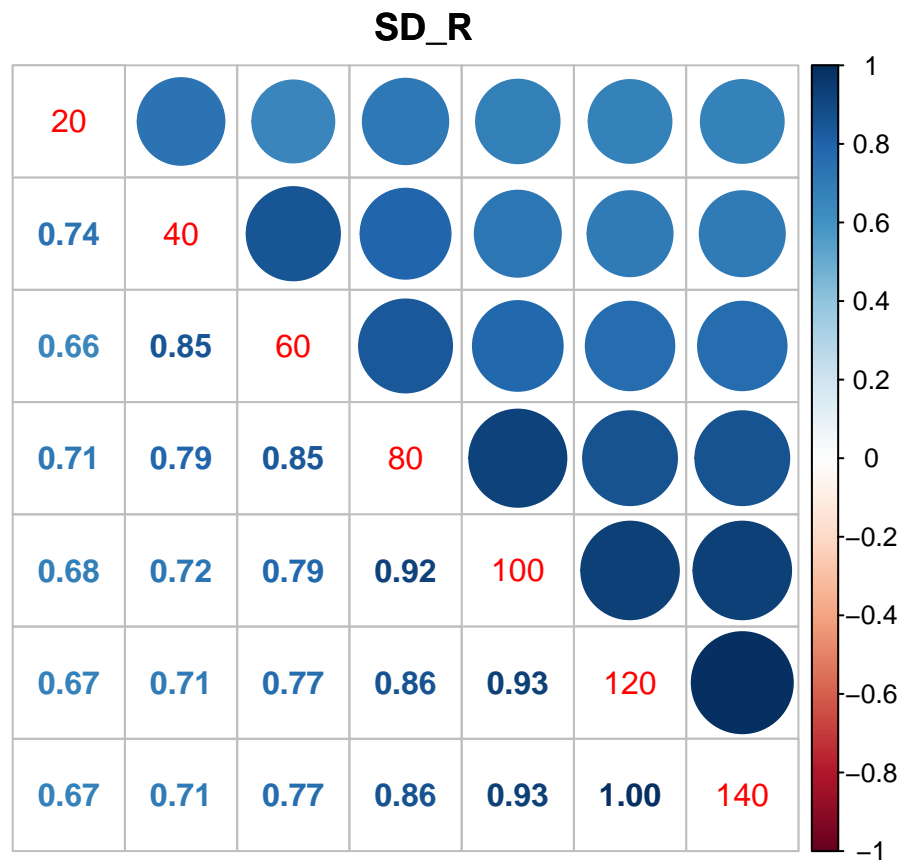


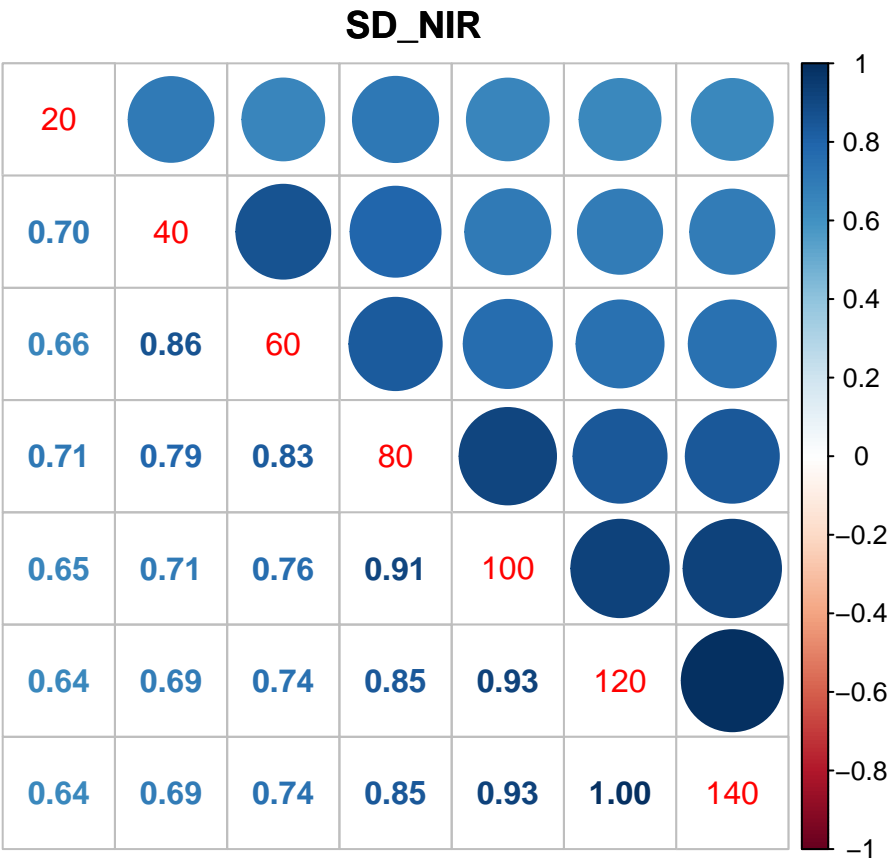


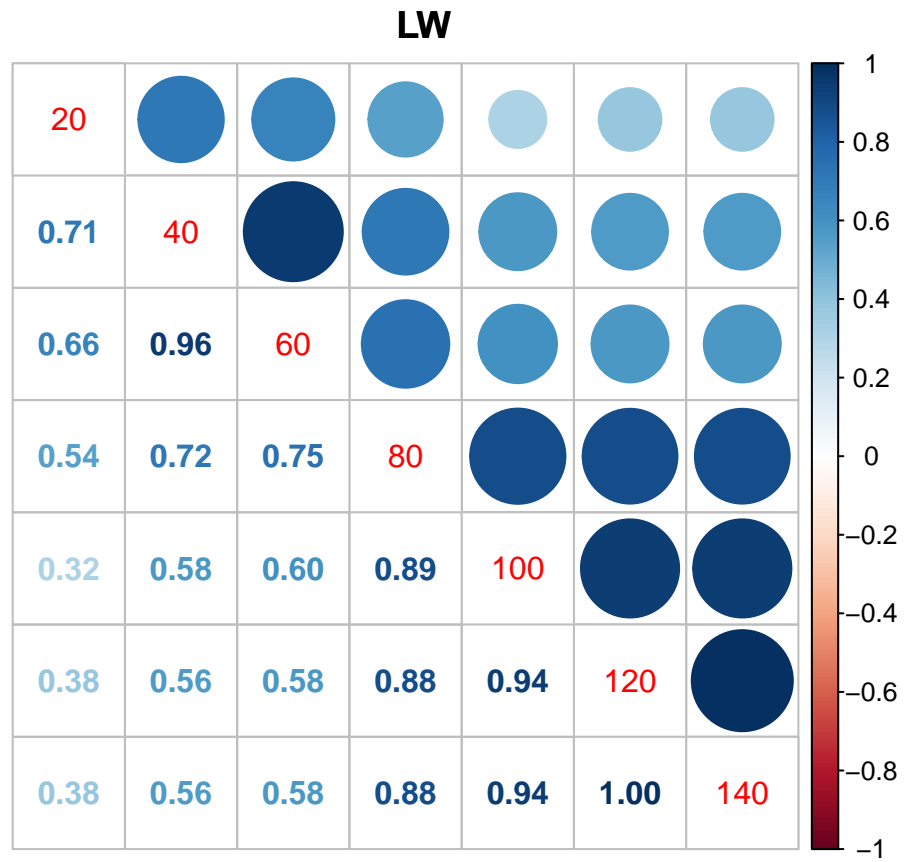


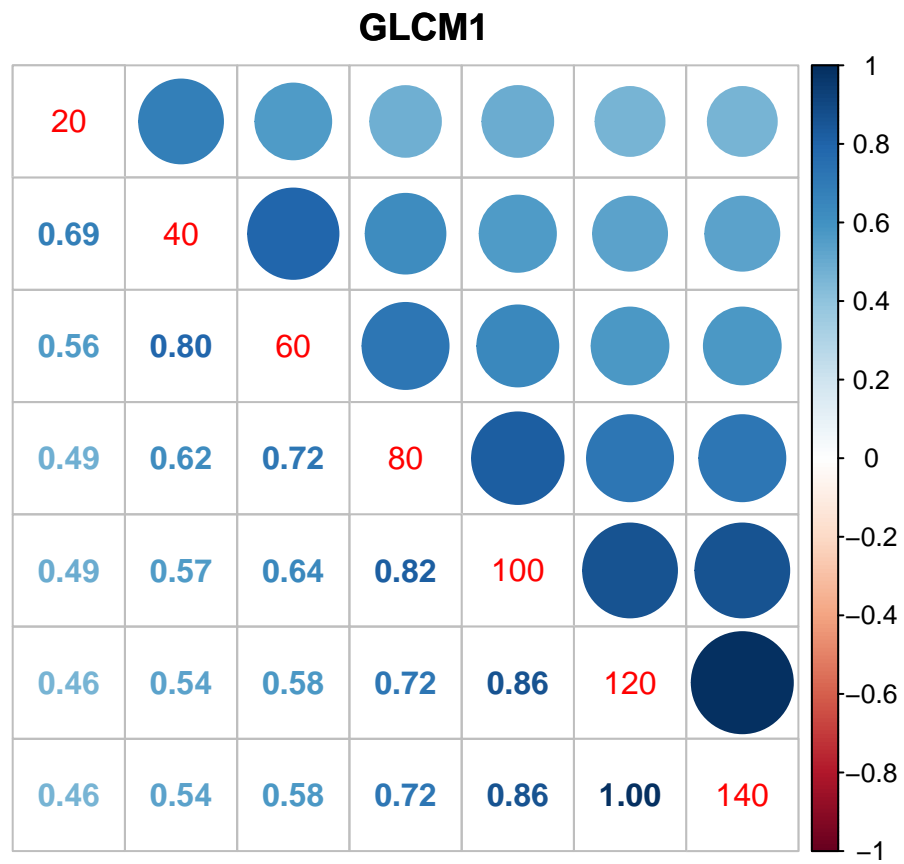


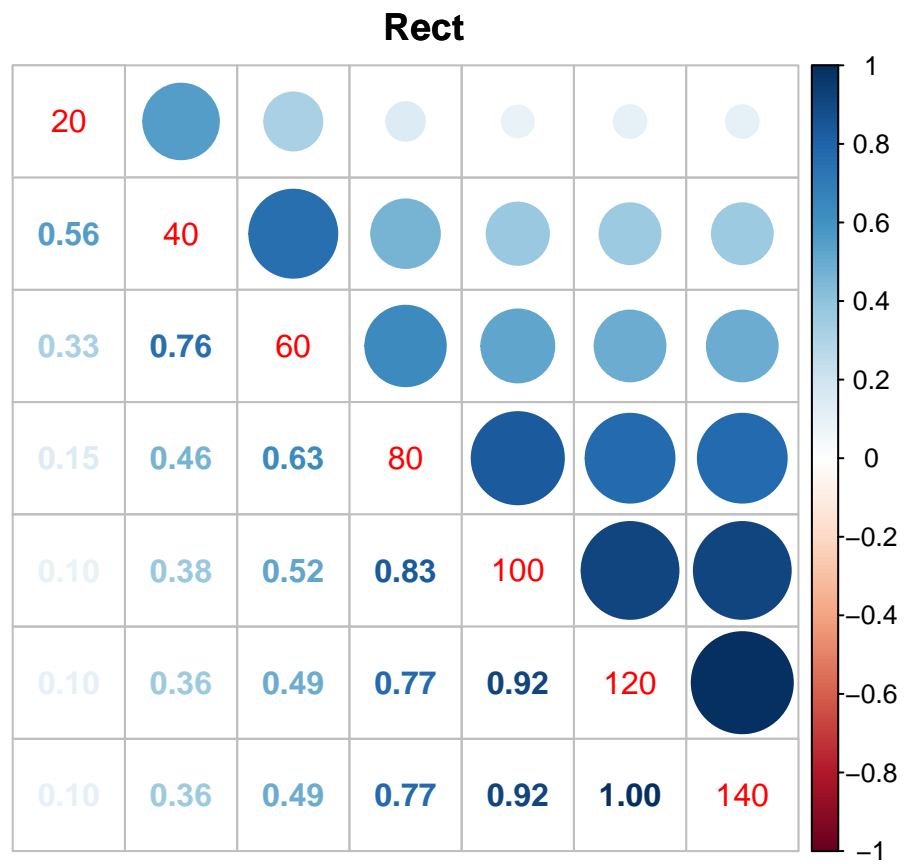


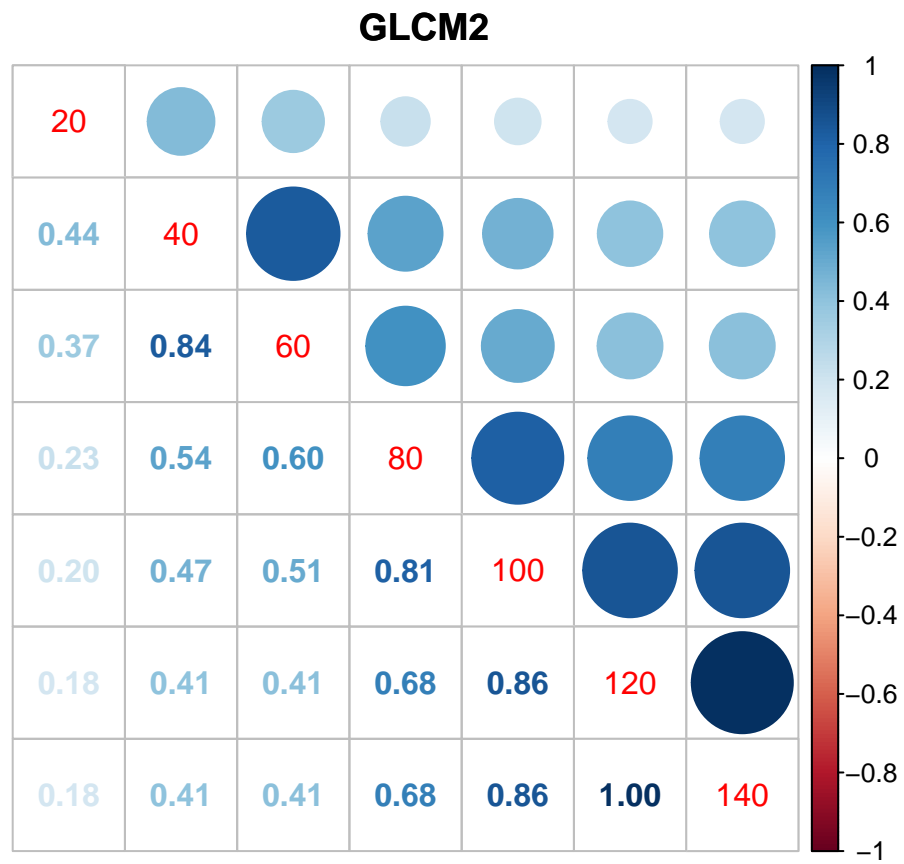


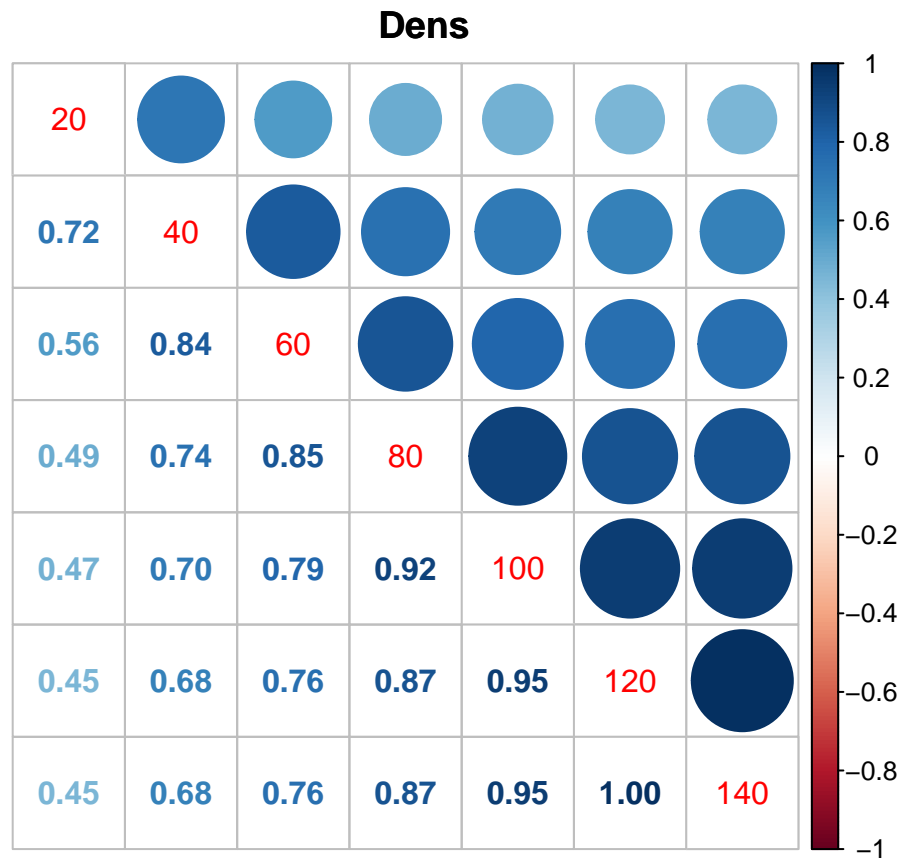


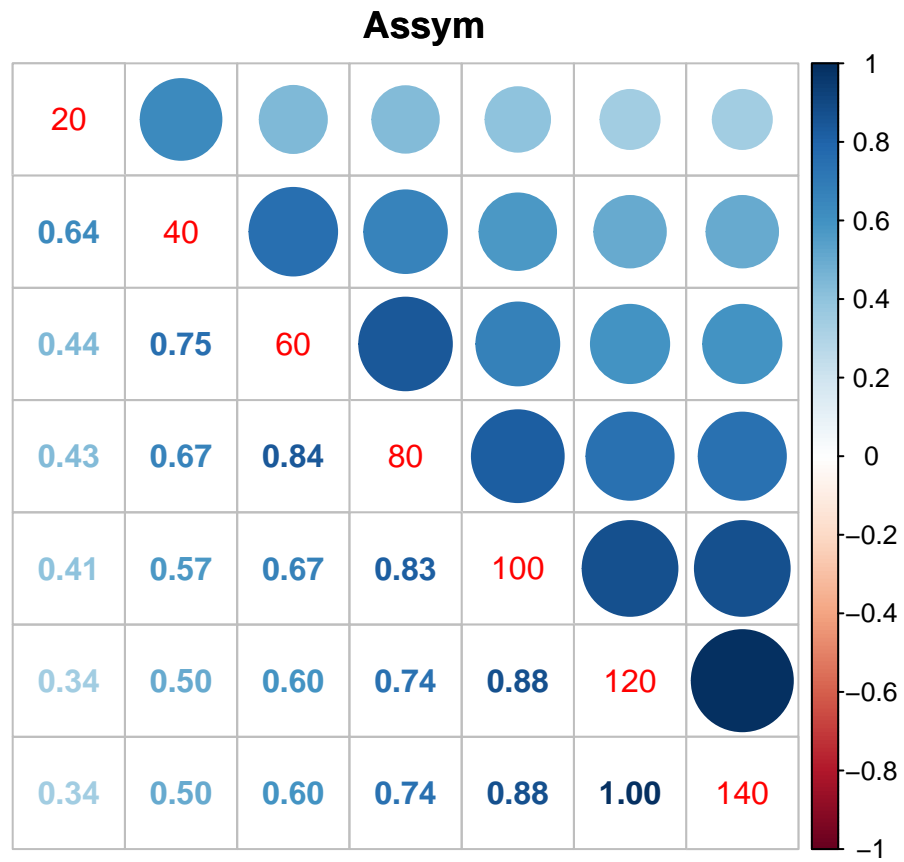


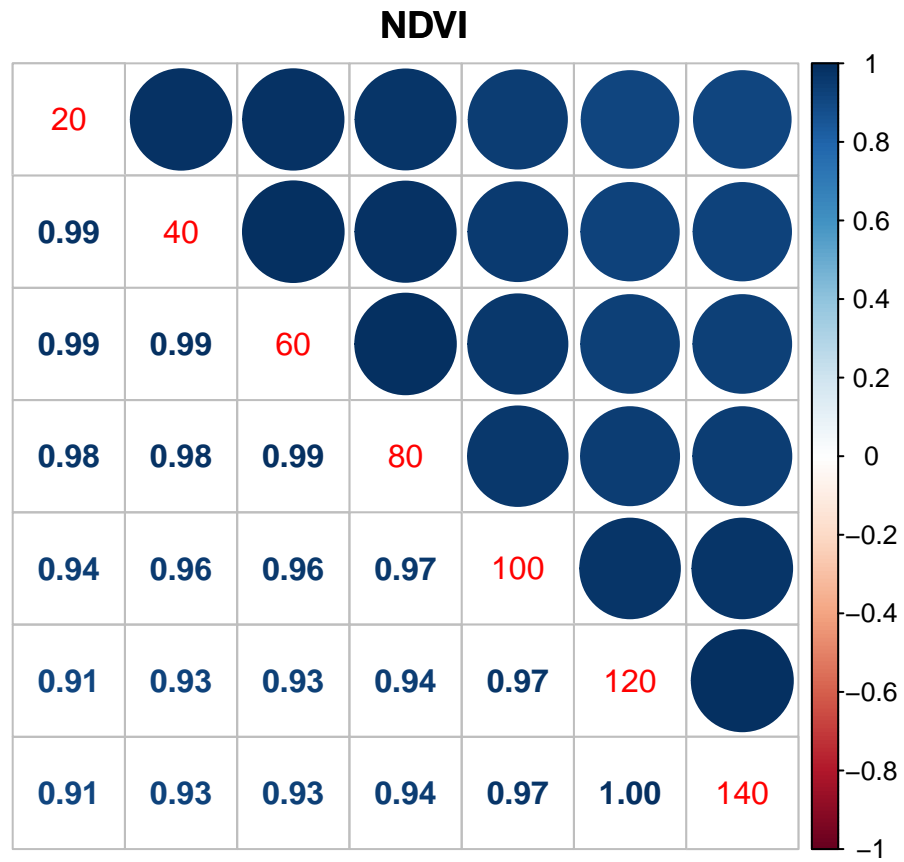


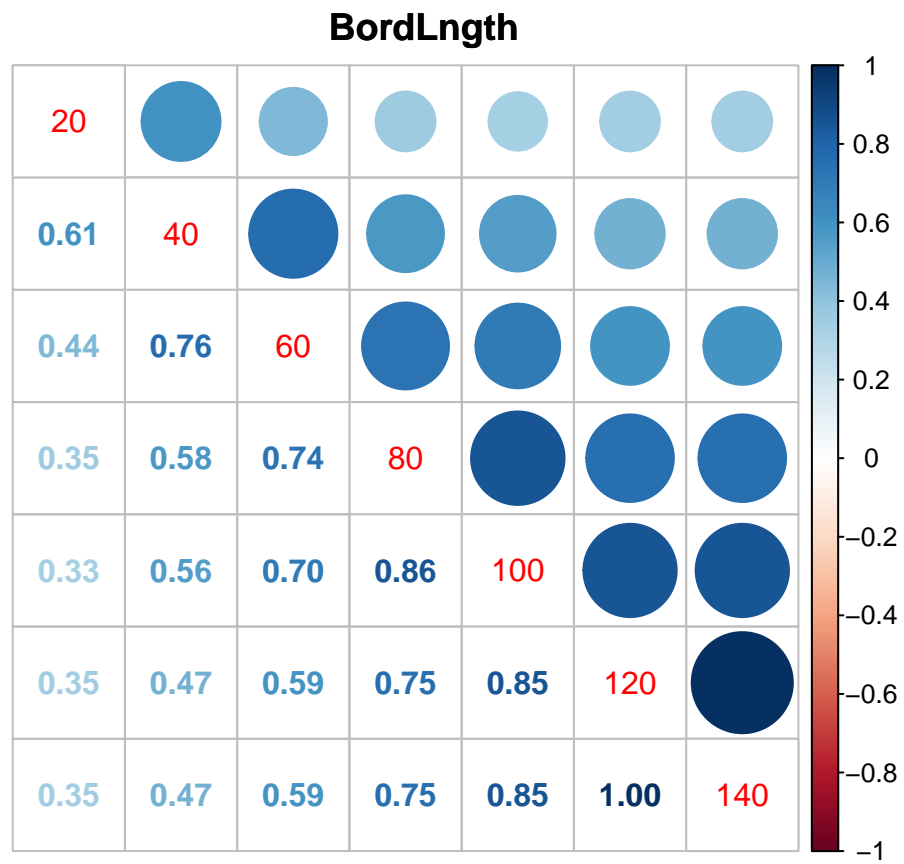


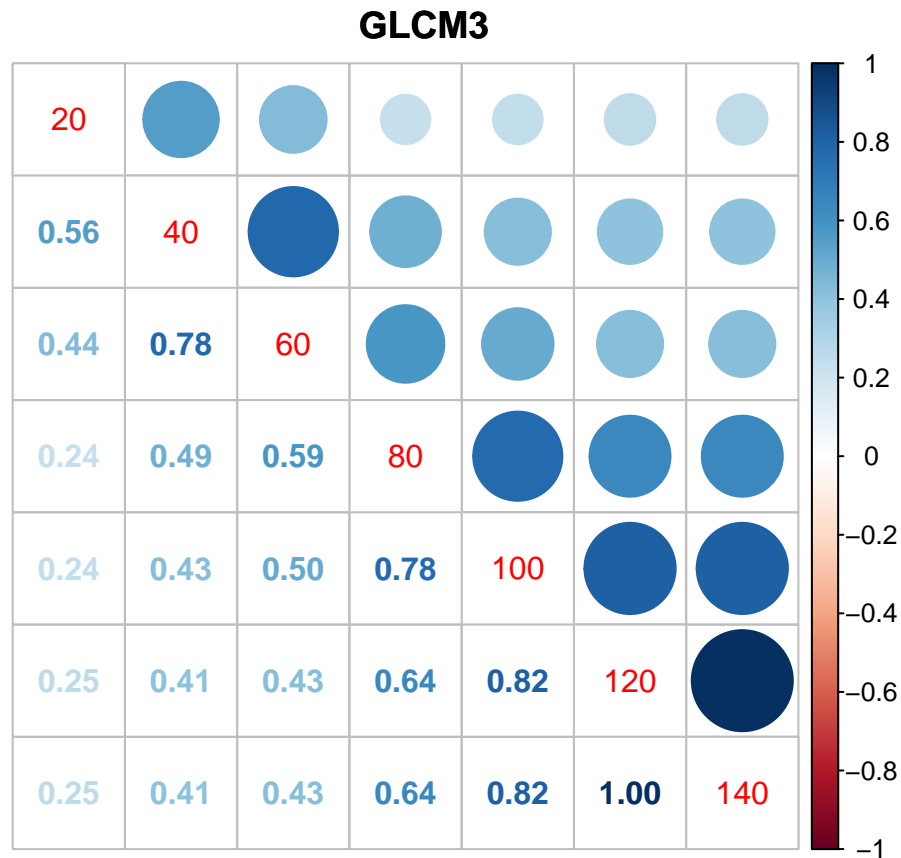












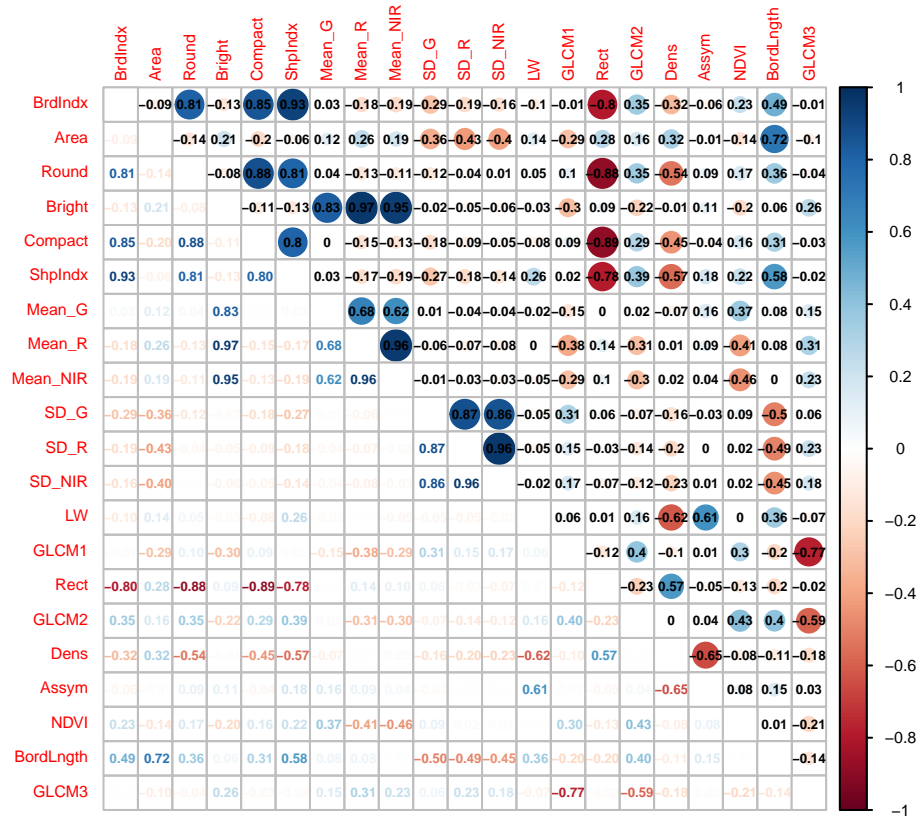
From the correlation plots: 1. In all coarseness cases, 120 and 140 are very highly correlated 2. across all attributes, correlation among consecutive coarseness level is highly correlated 3. Bright, Mean_G, Mean_R, Mean_NIR, and NVDI are extremely correlated across all coarseness levels

Hypothesis: perhaps modelling the data using a single coarseness level is sufficient or using coarseness levels 20 and 140 as they are they least correlated

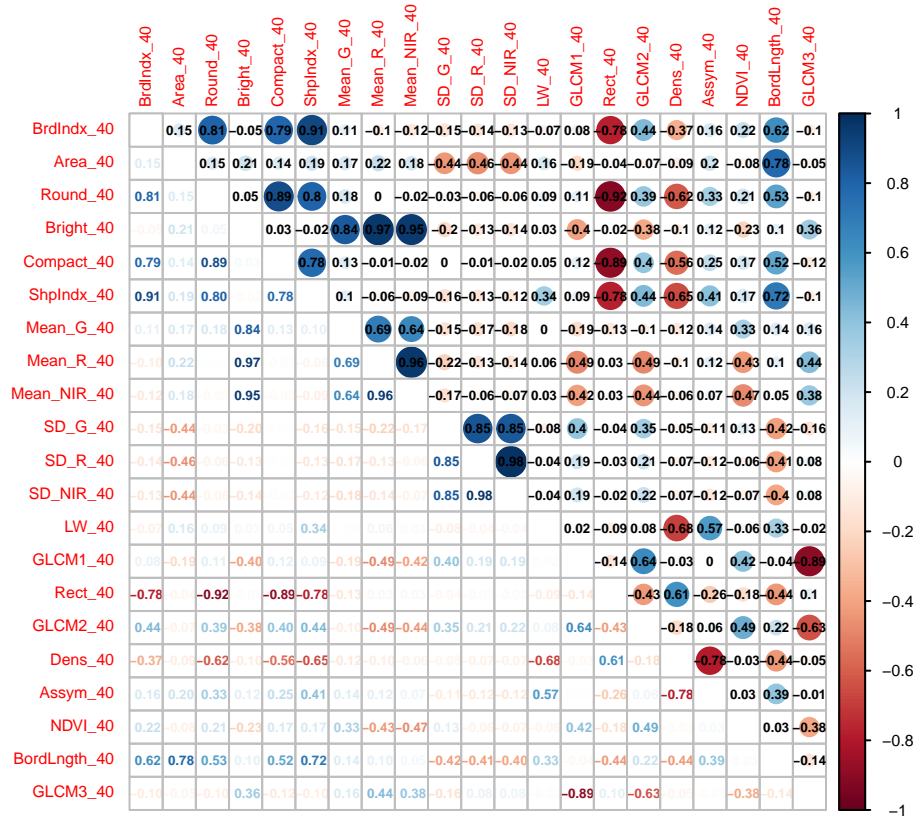
checking the correlation among different attributes within the same coarseness

```
corList2 <- list()
for(i in 1:7){
  corList2[[i]] <- cor(trainingList[[i]][,2:22])
  corrplot.mixed(corList2[[i]],addCoef.col = 'black', number.cex = 0.4, upper = 'circle',
    tl.cex=0.5, lower = 'number', title = paste("Coarseness Level",
    i*20, sep=" "), mar=c(0,0,1,0),
    cl.cex = 0.5, diag = "n", tl.pos = "lt")
}
```

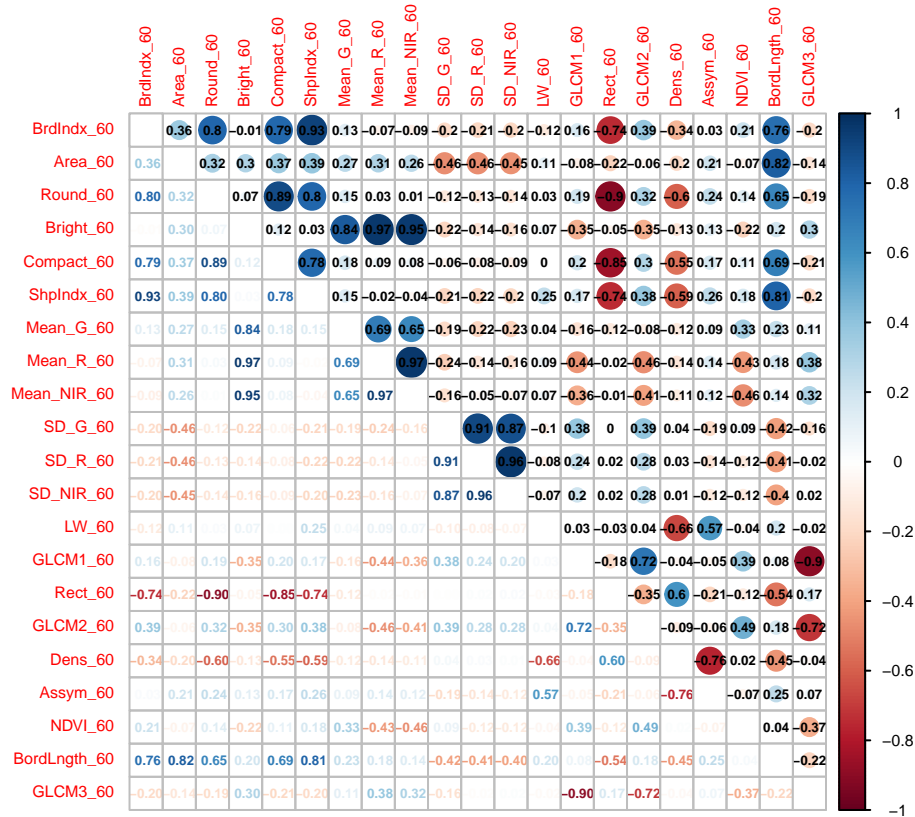
Coarseness Level 20



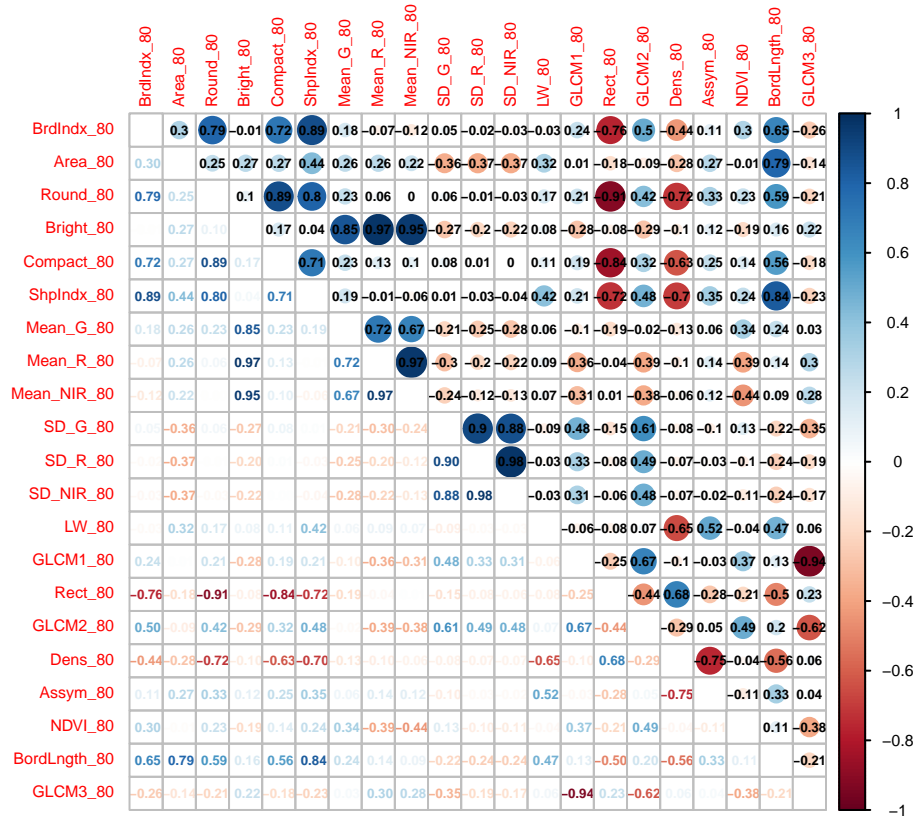
Coarseness Level 40



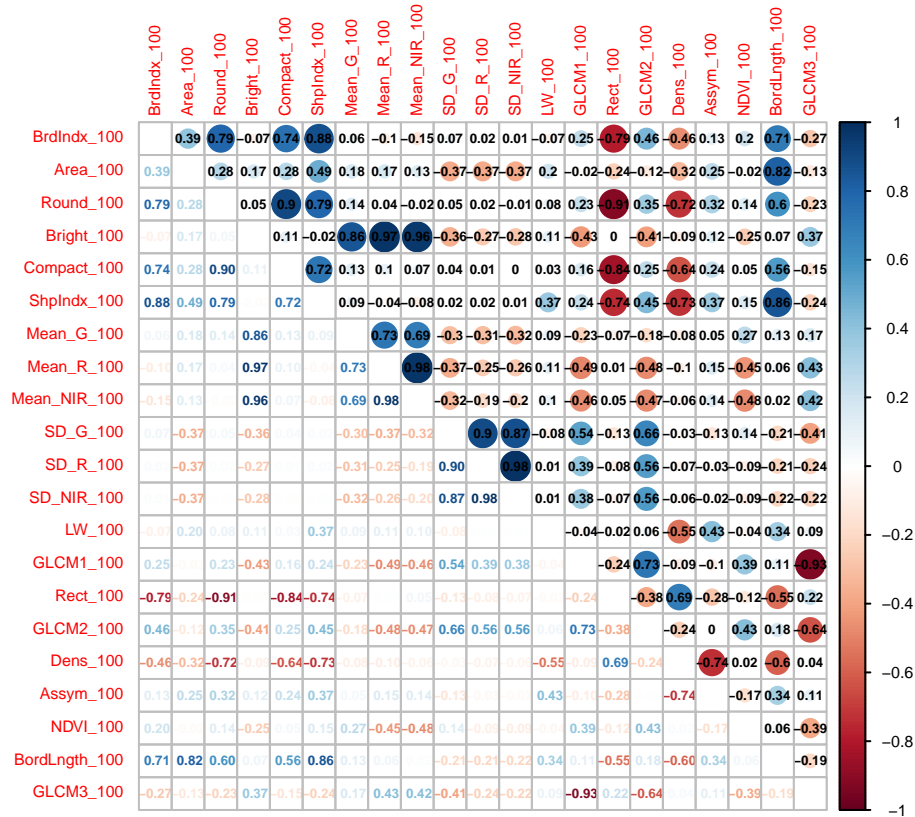
Coarseness Level 60



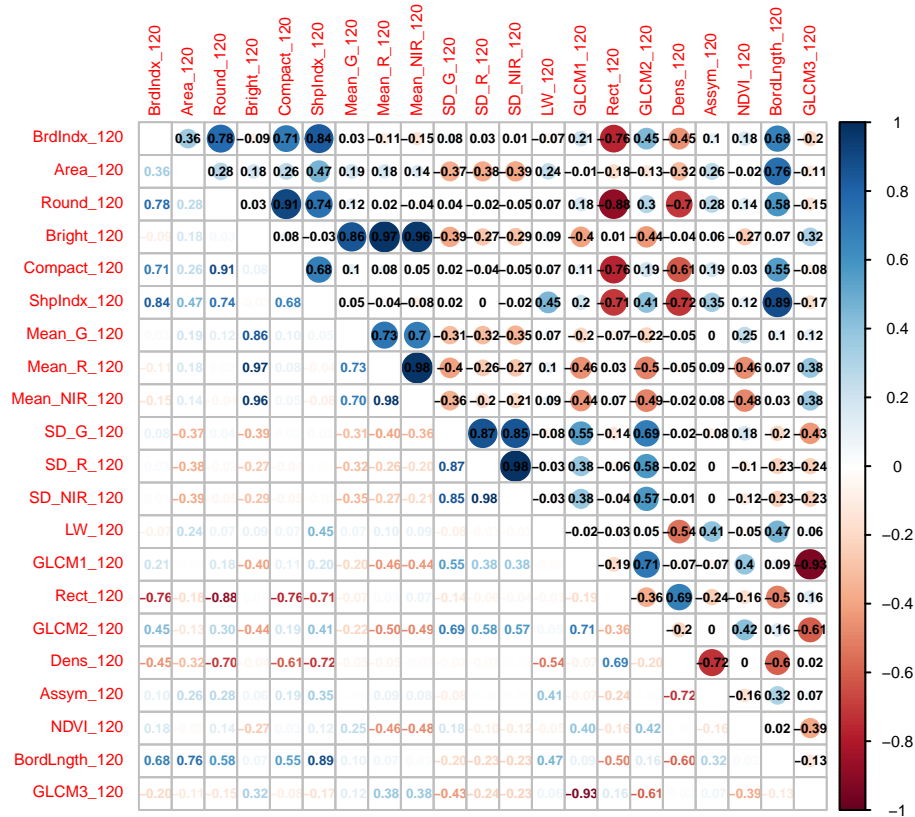
Coarseness Level 80



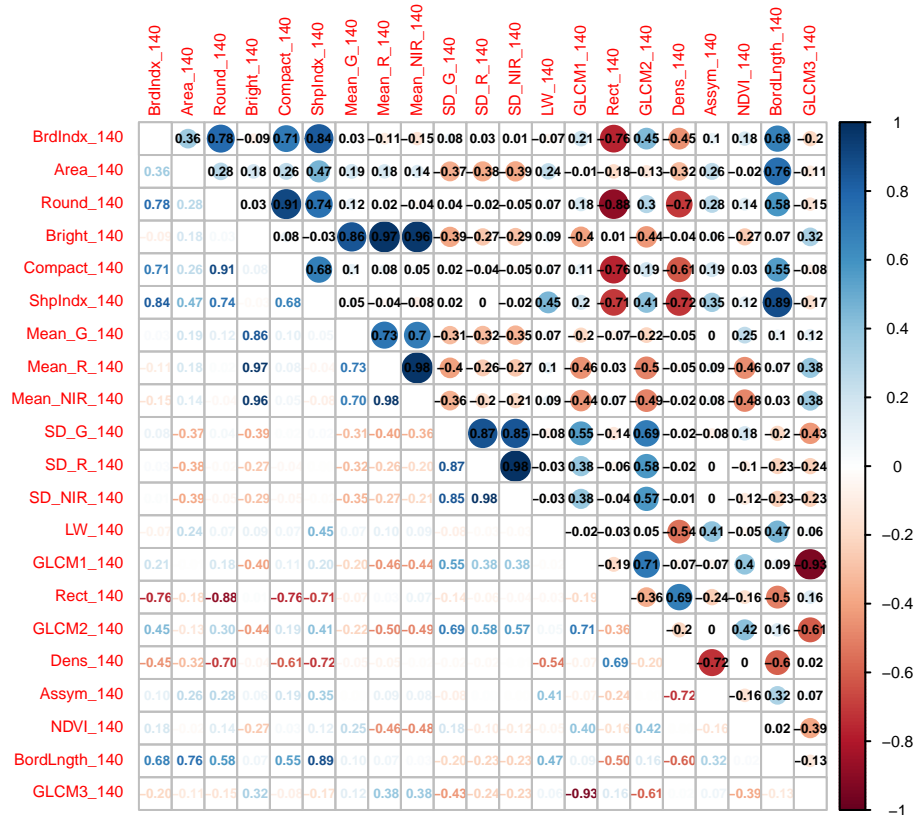
Coarseness Level 100



Coarseness Level 120



Coarseness Level 140



Analysis: 1. brdindx and shpindx are highly correlated across all coarseness levels 2. bright is highly correlated with mean_G, mean_R, and mean_NIR 3. SD_G, SD_R, SD_NIR are highly correlated among each other 4. rect is inversely correlated with shpindx, compact, round, and brdindx

Conclusions

1. classification can be attempted with only one coarseness level, or levels 20 and 140
2. Brdindx, Shpindx, compact, and round can be omitted from the model since rect is highly correlated with all of them
3. Mean_G, Mean_R, and Mean_NIR can be omitted from the model since bright is highly correlated with all of them
4. SD_G, SD_R, or SD_NIR should be used in the model; whichever produces the best results