ML: Assignment 1 Report

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Tasks

Part 1: lecture script

This task was performed by the team members individually and there is no hand in required.

Part 2: Diamond Prices

- 1. Give an overview of the dataset structure:
- How many samples and features are in the dataset?

The dimensions of the diamonds dataset correspond to the amount of samples and features. By using the dim() command we get the information that there are 53940 samples and 10 features.

What are the feature data types

We can get information about the data types using the str() or structure command. While most of the features are numeric or integers the cut, color and clarity are labels or factors. The resulting types can be seen here:

Are diamonds balanced across color, cut and clarity?

Color: Fairly balanced, even though I and J are a bit fewer.

Cut: Not balanced. Most "ideal" with very few "fair".

Clarity: Not balanced. Very few I1 and IF and for example more SI1 and VS2.

- 2. Visualize diamond prices using a histogram, boxplot and densityplot:
- Is there a visible trend? If yes, which is it and from which plots can you derive it? There are more cheap diamonds than expensive ones (from hist and density plot). The lowest priced diamond is at 326, 25% of diamonds have a price below 950 (Q1), 50% below 2401 (Median), 75% below 5324 (Q3). While most diamonds cost below 12000 some outliers cost up to 18823 (from boxplot in combination with summary for exact numbers).

3. Calculate and state the mean, median, standard deviation, median absolute deviation, 1st and 3rd quantile, and inner quantile range of the diamond price:

Mean: 3933 Median: 2401

Standard Deviation: 3989.44

Median Absolute Deviation: 2475.942

1st Quantile: 950 3rd Quantile: 5321.25

Inner Quantile Range: 4374.25

- 4. Plot the diamond price against the carat values as an xy-plot:
- Is there a trend visible in the plot? If yes, which is it? The price of a diamond increases with its carat value.
- 5. Analyze the correlation between diamond price and diamond x, y, and z dimensions:
- Is there a trend visible between x, y, and z? If yes, which is it? Is there a trend visible between the dimensions and the price? If yes, which is it?

The plot shows logarithmic corellation between price and features x, y, z (bigger dimensions equal a higher price). The plot alsow shows linear corellation between the features x and y, x and z, as well as y and z.

- 6. Analyze diamond prices per diamond color:
- Create boxplots showing diamond price boxes for each diamond color. Use a single plot command to obtain this plot (hint: use the formula interface of boxplot with a "price ~ color" formula) Create densityplots showing diamond prices for each diamond color. Is there a trend visible? If yes, which is it?

Both plots shows that most of the diamonds in each color category have a price range between 0 and <= 5000.

- 7. Use vectorized commands to answer these questions:
- How many diamonds have a price above 9500?
 5734
- How many diamonds have a price above 9500 and have color "D"?
 461

• For all color "D" diamonds with cut "Fair" show the summary for price and carat

carat

Min. :0.2500 1st Qu.:0.7000 Median :0.9000 Mean :0.9201 3rd Qu.:1.0100 Max. :3.4000

price

Min.: 536 1st Qu.: 2204 Median: 3730 Mean: 4291 3rd Qu.: 4797 Max.: 16386

• For all color "J" diamonds with cut "Ideal" show the summary for price and carat

carat

Min. :0.230 1st Qu.:0.540 Median :1.030 Mean :1.064 3rd Qu.:1.410 Max. :3.010

price

Min.: 340 1st Qu.: 1132 Median: 4096 Mean: 4918 3rd Qu.: 6732 Max.: 18508

- 8. Utilize functions from the plyr library or the standard apply function on the diamonds data.frame to compute the following:
- **1.** Calculate the mean (?mean) for all numerics features using a single command. sapply(diamonds, function(x) if (!is.numeric(x)) NA else mean(x))
- 2. Do the same with median (?median). sapply(diamonds, function(x) if (!is.numeric(x)) NA else median(x))
- **3. Do the same with standard deviation (?sd).** sapply(diamonds, function(x) if (!is.numeric(x)) NA else sd(x))

4. Do the sa	me with	median	absolute	deviation	(?mad) a	and set the	constant	parameter
of the mad f	unction	to 1.						

sapply(diamonds, function(x) if (!is.numeric(x)) NA else mad(x, constant = 1))

Part 3: Cell Body Segmentation Data

1. Which classes exist? Are they (roughly) balanced?

PS and WS.

They are not balanced (1300:719).

2. Which noteworthy trends of features and relations between features as well as features and Class do you see?

AwgIntenCh1 is linearly related to DiffIntenDensityCh1. Between all other attributes there were no clear relations visible.

3. If you would need to distinguish between classes, which features do you think would be most helpful? Why?

There is no real good attribute to distinguish the classes. One of the more fitting attributes is AvgIntenCh2. Further attributes that may help distinguish classes are AvgIntenCh1 and ConvexHullPerimRatioCh1. These attributes show fewer overlaps in the density-, box- and hist-plot.

However these are still not an optimal indicator. We would recommend collecting other features.