

## RELATIONSHIPS BETWEEN VARIABLES

To answer research questions, data must be collected.

Analyses are motivated by **looking** for a **relationship between two or more variables**.

Examining **summary statistics** could provide insights for each of the research questions about the study.

A **summary statistics** is a **single number summarizing a large amount of data**. In other words, a **summary statistics** is a value computed from the **data**.

## EXAMINING NUMERICAL DATA

We will be introduced to techniques for **exploring** and **summarizing numerical** variables, working with two datasets : 'email50', 'county' and 'cars'.

```
In [ ]: # importing Libraries
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
```

## EXPLORING BIVARIATE VARIABLES WITH SCATTERPLOTS

A **Scatterplot** provides a case-by-case view of data for **two (bivariate) numerical variables**.

**Scatterplots** are helpful in quickly **spotting associations relating variables**, whether those associations come in the form of **simple trends** or whether those relationships are more **complex**.

We will use a **Scatterplot** to examine how *federal spending* and *poverty* are related in the *county* dataset.

```
In [ ]: # Open the chosen file
county = pd.read_csv('D:\\Documents\\EureCat\\Eurecat 2019\\BTS\\Datasets\\county.txt', sep='\\t',
encoding='utf-8')
```

```
In [ ]: county.shape
```

```
In [ ]: # Create data
x = county.fed_spend
y = county.poverty
colors = 'Blue'
area = np.pi*5

plt.axis([0, 100, 0, 60])

# Plot
plt.scatter(x, y, s=area, c=colors, alpha=0.4, edgecolors='black')

plt.title('Federal Spending vs Poverty by County')
plt.ylabel('Federal Spending per Capita')
plt.xlabel('Poverty Rate (Percent)')
plt.show()
```

In any [Scatterplot](#), each point represents a single **case/observation**. Since there are **3.143** cases in *county*, there are **3.143** points

Now, We will compare the [number of line breaks \(line\\_breaks\)](#) and [number of characters \(num\\_char\)](#) in emails for the *email50* dataset.

```
In [ ]: dbe = pd.read_csv('D:\\Documents\\EureCat\\Eurecat 2019\\BTS\\Datasets\\email50.txt',
                        encoding='utf-8', sep='\\t')
```

```
In [ ]: dbe.shape
```

```
In [ ]: # Create data
x = dbe.num_char
y = dbe.line_breaks

colors = "Blue"
area = np.pi*20
plt.axis([0, 70, 0, 1200])

# Plot
plt.scatter(x, y, s=area, c=colors, alpha=0.4, edgecolors='black')
plt.title('Spam email - # Lines vs # Characters')
plt.ylabel('Number of Lines')
plt.xlabel('Number of Characters (in thousands)')
plt.show()
```

To put the [number of characters](#) in perspective, this paragraph has **363** characters. Looking at [scatterplot](#), it seems that some emails are incredibly verbose!. Upon further investigation, we would actually find that most of the long emails use the **HTML format**, which means most of the characters in those emails are used to **format the email** rather than **provide text**.

```
In [ ]: dbcars = pd.read_csv('D:\\Documents\\EureCat\\Eurecat 2019\\BTS\\Datasets\\cars.txt',
                          encoding='utf-8', sep='\\t')
```

Let's consider a new dataset *cars* of 54 *cars* with 6 variables. Create scatterplot to examine how *vehicle price* and *weight* are related.

What can be said about the relationship between these variables?

```
In [ ]: dbcars.shape
```

```
In [ ]: # Checking dataset variables
dbcars.dtypes
```

```
In [ ]: dbcars.head()
```

```
In [ ]: # Create data
x = dbcars.weight
y = dbcars.price

colors = "Blue"
area = np.pi*30

# Plot
plt.scatter(x, y, s=area, c=colors, alpha=0.4, edgecolors='black')
plt.title('Cars - Price vs Weight')
plt.ylabel('Price ($1000s)')
plt.xlabel('Weight (Pounds)')
```

The relationship is [evidently nonlinear](#).

```
In [ ]: fig = plt.figure(figsize=(12,4))

ax1 = fig.add_subplot(1, 3, 1)

# Create data
x = county.fed_spend
y = county.poverty
colors = 'Blue'
area = np.pi*5

plt.axis([0, 100, 0, 60])

# Plot
ax1.scatter(x, y, s=area, c=colors, alpha=0.4, edgecolors='black')

plt.title('County Dataset')
plt.ylabel('Federal Spending per Capita')
plt.xlabel('Poverty Rate (Percent)')

ax2 = fig.add_subplot(1, 3, 2)
# Create data
x = dbc.num_char
y = dbc.line_breaks

colors = "Blue"
area = np.pi*20
plt.axis([0, 70, 0, 1200])

# Plot
ax2.scatter(x, y, s=area, c=colors, alpha=0.4, edgecolors='black')
plt.title('Spam email Dataset')
plt.ylabel('# of Lines')
plt.xlabel('# of Characters (in thousands)')

ax3 = fig.add_subplot(1, 3, 3)
# Create data
x = dbcars.weight
y = dbcars.price

colors = "Blue"
area = np.pi*30
plt.axis([1500, 4300, 0, 65])

# Plot
ax3.scatter(x, y, s=area, c=colors, alpha=0.4, edgecolors='black')
plt.title('Cars Dataset')
plt.ylabel('Price ($1000s)')
plt.xlabel('Weight (Pounds)')

# plt.tight_layout(pad=0.4, w_pad=0.5, h_pad=1.0)
plt.tight_layout()
```

```
In [ ]: sns.pairplot(county, diag_kind='kde', plot_kws={'alpha': 0.2})
```

# HISTOGRAMS

Dot plots, like in [scatterplot](#), show the **exact value for each observation**. This is useful for **small datasets**, but they can become hard to read with **larger samples**.

Rather than showing the *value of each observation*, we prefer to think of the value as belonging to a **bin**.

These **bins** - (*counts*) are plotted as bars into what is called a [Histogram](#).

[Histogram](#) provide a view of the **data density**. Higher bars represent where the data are relatively more common.

[Histogram](#) are especially convenient for describing the *shape of the data distribution*.

- When data trail off to the right and have a longer right tail, the shape is said to be [Right Skewed](#) or also called [Skewed to the Positive End](#).
- Contrary, data with the reverse characteristic – a *long, thin tail to the left* – are said to be [Left Skewed](#). We also say that such a distribution has a **long left tail**.
- Data that show roughly equal trailing off in both directions are called [Symmetric](#).

```
In [ ]: dbe.hist(['num_char'], bins=14)
plt.title('Spam email - # Characters')
plt.ylabel('Frequency')
plt.xlabel('# Characters (in thousands)')
```

**Long tails to identify skew** When data trail off in one direction, the distribution has a long tail. If a distribution has a long left tail, it is Left Skewed. If a distribution has a long right tail, it is Right Skewed.

## Modal Distribution

In addition to looking at whether a distribution is [Skewed](#) or [Symmetric](#), histograms can be used to identify **Modes**.

A **mode** is the *value with the most occurrences*.

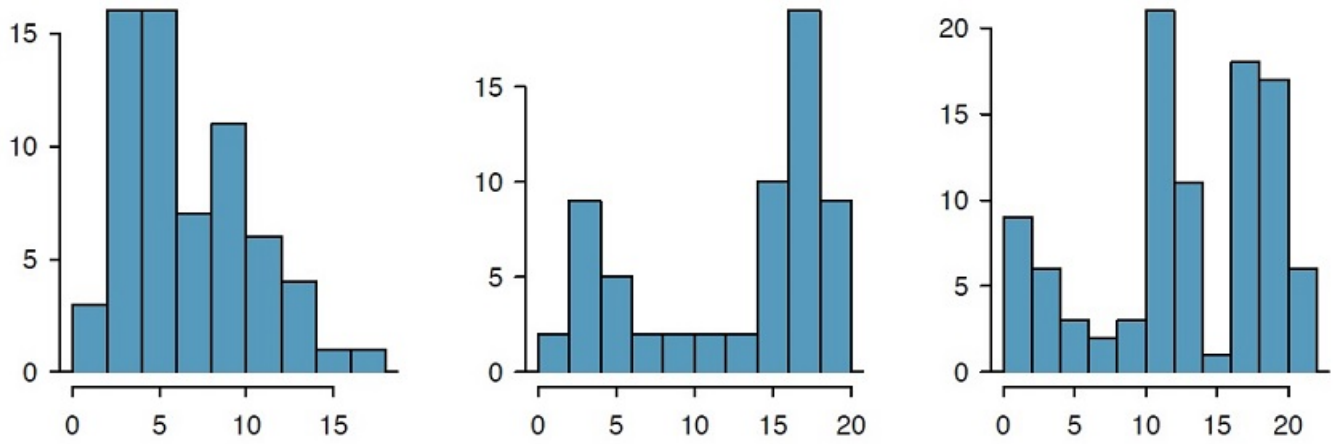
However, It is common to have **no observations** with the same value in a dataset, which makes, **mode**, **useless** for many real datasets.

A **mode** is represented by a prominent peak in the **distribution**. There is only one prominent peak in the histogram of num\_char.

[Histogram](#) that have one, two, or three prominent peaks are called [Unimodal](#), [Bimodal](#), and [Multimodal](#), respectively.

Any **distribution** with more than 2 prominent peaks is called **Multimodal**.

Notice that there was one prominent peak in the [Unimodal](#) distribution with a second less prominent peak that was not counted since it only differs from its neighboring **bins** by a few **observations**.



Looking for modes Looking for modes isn't about finding a clear and correct answer about the number of modes in a distribution. The important part of this examination is to better understand your data and how it might be structured.

## SUMMARY STATISTICS

### Mean - Average

The **mean**, sometimes called the **average**, is a common way to measure the **center of a distribution of data**.

To find the **mean number of characters (num\_char)** in the 50 emails, we add up all the character counts and divide by the number of emails.

For computational convenience, the number of characters is listed in the thousands and rounded to the first decimal.

$$\bar{x} = \frac{21.7 + 7.0 + \dots + 15.80}{50} = 11.6$$

```
In [ ]: dbe.num_char.mean().round(2)
```

The **sample mean** is often labeled  $\bar{x}$ . The letter  $x$  is being used as a generic placeholder for the variable of interest, *num\_char*, and the *bar over on the x* communicates that the average number of characters in the 50 emails is 11,6.

Mean The **sample mean**  $\bar{x}$  of a **numerical variable** is computed as the sum of all of the **observations** divided by the number of **observations**:

$$\bar{x} = \frac{x_1 + x_2 + \dots + x_n}{n}$$

where  $x_1, x_2, \dots, x_n$  represent the  $n$  observed values. It is useful to think of the mean as the **balancing point** of the distribution.

### EXERCISE - 3.1

Compare both Equations above.

- What does  $x_1$  correspond to ?,
- and  $x_2$  ?
- Can you infer a general meaning to what  $x_i$  might represent?
- What was  $n$  in this sample of emails?

**Population Mean** The **Population mean** has a special label :  $\mu$ . The symbol  $\mu$  is the *Greek* letter *mu* and represents the **average/mean of all observations in the Population**. Sometimes a subscript, such as  $x$ , is used to represent which variable the **population mean** refers to, e.g.  $\mu_x$

### EXERCISE - 3.2

The **average** number of characters across all emails (**population**) can be estimated using the **sample data**.

Based on the **sample** of 50 *emails*, what would be a reasonable estimate of  $\mu_x$ , the **mean** number of characters in all emails in the email dataset? (Recall that *email50* is a sample from *email*.)

## Variance and Standard Deviation

### Variance

The **mean** was introduced as a method to describe the **center of a data set**, but the **variability in the data** is also important.

We introduce **two measures of variability**: the **Variance** and the **Standard Deviation**. Both are very useful in data analysis.

The **Standard Deviation** describes how far away the typical observation is from the **mean**.

We call the **distance of an observation from its mean** its **Deviation**.

Below are the deviations for the 1st, 2nd, 3rd, and 50th observations in the `num_char` variable. For computational convenience, the number of characters is listed in the thousands and rounded to the first decimal.

$$\begin{aligned}x_1 - \bar{x} &= 21.7 - 11.6 = 10.1 \\x_2 - \bar{x} &= 7.0 - 11.6 = -4.6 \\x_3 - \bar{x} &= 0.6 - 11.6 = -11.0 \\&\vdots \\x_{50} - \bar{x} &= 15.8 - 11.6 = 4.2\end{aligned}$$

If we square these **deviation** and then take an average, the result is about equal to the **sample variance**, denoted by  $s^2$ :

$$s^2 = \frac{10.1^2 + (-4.6)^2 + (-11.0)^2 + \dots + 4.2^2}{50 - 1} = 172,44$$

Sample Variance  $s^2$  We divide by  $n - 1$ , rather than dividing by  $n$ , when computing the **Variance**. **squaring the deviations** does two things: - First, it **makes large values much larger**, seen by comparing  $10.1^2$ ,  $(-4.6)^2$ ,  $(-11.0)^2$ , and  $4.2^2$ . - Second, it **gets rid of any negative signs**. The variance is roughly the **average squared distance** from the mean.

## Standard Deviation

Standard Seviation The Standard Deviation is defined as the square root of the **Variance** :

$$s = \sqrt{172.44} = 13.13$$

The Standard Deviation is useful when considering how close the data are to the Mean. Formulas and methods used to compute the Variance and Standard Deviation for a Population are similar to those used for a sample (*The only difference is that the Population Variance has a division by  $n$  instead of  $n - 1$* ). However, like the Mean, the Population values have special symbols : -  $\sigma^2$  for the Variance and -  $\sigma$  for the Standard Deviation. The symbol  $\sigma$  is the *Greek letter sigma*.

```
In [ ]: dbe.num_char.iloc[[0], ].std()
```

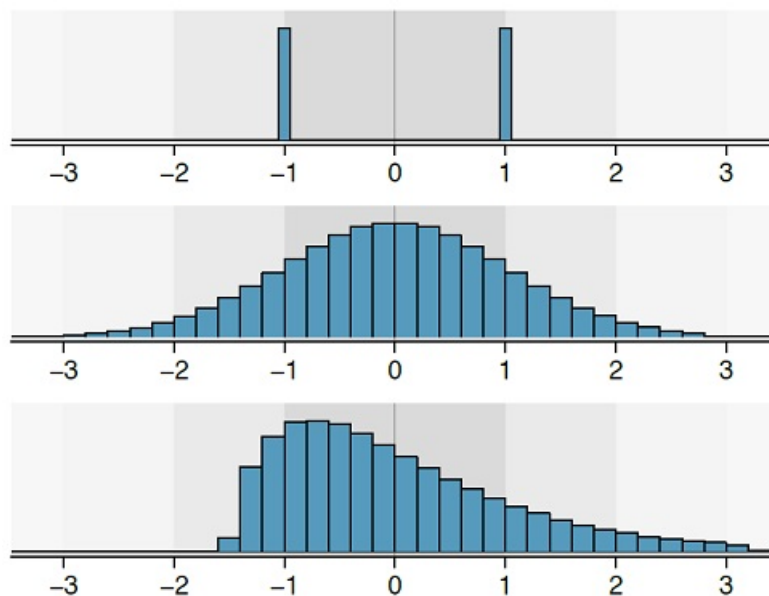
**Standard Deviation** describes **Variability**, so focus on the conceptual meaning of the **Standard Deviation** as a descriptor of **Variability** rather than the formulas.

Usually 70% of the data will be within **one standard deviation of the mean** and about 95% will be within **two standard deviations** two standard deviations. However, these percentages are not strict rules.

### EXERCISE - 3.6

A good description of the shape of a distribution should include modality and whether the distribution is symmetric or skewed to one side.

Explore the figure as an example, explain why such a description is important :



```
In [ ]: dbe.hist(['num_char'], bins=14)
plt.title('Spam email - # Characters')
plt.ylabel('Frequency')
plt.xlabel('# Characters (in thousands)')
```

### EXERCISE - 3.7

Describe the distribution of the `num_char` variable using the histogram display above.

The description should incorporate the center, variability, and shape of the distribution, and it should also be placed in context: the number of characters in emails. Also note any especially unusual cases.

We will use the **Variance** and **Standard Deviation** to assess how close the **Sample Mean** ( $\bar{x}$ ) is to the **Population Mean** ( $\mu$ ).

variable	description
<code>name</code>	County name
<code>state</code>	State where the county resides (also including the District of Columbia)
<code>pop2000</code>	Population in 2000
<code>pop2010</code>	Population in 2010
<code>fed_spend</code>	Federal spending per capita
<code>poverty</code>	Percent of the population in poverty
<code>homeownership</code>	Percent of the population that lives in their own home or lives with the owner (e.g. children living with parents who own the home)
<code>multiunit</code>	Percent of living units that are in multi-unit structures (e.g. apartments)
<code>income</code>	Income per capita
<code>med_income</code>	Median household income for the county, where a household's income equals the total income of its occupants who are 15 years or older
<code>smoking_ban</code>	Type of county-wide smoking ban in place at the end of 2011, which takes one of three values: <code>none</code> , <code>partial</code> , or <code>comprehensive</code> , where a <code>comprehensive</code> ban means smoking was not permitted in restaurants, bars, or workplaces, and <code>partial</code> means smoking was banned in at least one of those three locations

```
In [ ]: fig = plt.figure(figsize=(10,8))

ax1 = fig.add_subplot(2, 2, 1)

ax1.hist(county['multiunit'], bins=25)
plt.title('County - 2010 Population')
plt.ylabel('Frequency')
plt.xlabel('multi unit (%)')

ax2 = fig.add_subplot(2, 2, 2)

ax2.hist(county['income'], bins=25)

plt.title('2010 County Population')
plt.ylabel('Frequency')
plt.xlabel('Per Capita Income')

ax3 = fig.add_subplot(2, 2, 3)

ax3.hist(county['homeownership'], bins=25)
plt.title('2010 County Population')
plt.ylabel('Frequency')
plt.xlabel('Homeownership (%)')

ax4 = fig.add_subplot(2, 2, 4)

ax4.hist(county['med_income'], bins=25)

plt.title('2010 County Population')
plt.ylabel('Frequency')
plt.xlabel('Median Household Income')

plt.tight_layout()
```



```
In [ ]: fig = plt.figure(figsize=(20,5))

ax1 = fig.add_subplot(1, 4, 1)

ax1.hist(county['multiunit'], bins=25)
plt.title('2010 County Population')
plt.ylabel('Frequency')
plt.xlabel('multi unit (%)')

ax2 = fig.add_subplot(1, 4, 2)

ax2.hist(county['income'], bins=25)

plt.title('2010 County Population')
plt.ylabel('Frequency')
plt.xlabel('Per Capita Income')

ax3 = fig.add_subplot(1, 4, 3)

ax3.hist(county['homeownership'], bins=25)
plt.title('2010 County Population')
plt.ylabel('Frequency')
plt.xlabel('Homeownership (%)')

ax4 = fig.add_subplot(1, 4, 4)

ax4.hist(county['med_income'], bins=25)

plt.title('2010 County Population')
plt.ylabel('Frequency')
plt.xlabel('Median Household Income')

# plt.tight_layout(pad=0.4, w_pad=0.5, h_pad=1.0)
plt.tight_layout()
```

## BOX PLOTS

A **Box Plot** summarizes a dataset using *five statistics* while also plotting **unusual observations - Anomalies or Outliers**.

### Quartiles, and the Median

```
In [ ]: (dbe['num_char']).describe()
```

The median (6,890), splits the data into the bottom 50% and the top 50%, marked in the dot plot by horizontal dashes and open circles, respectively.

```
In [ ]: (dbe['num_char']).median().round(3)
```

The first step in building a **box plot** is drawing a dark line denoting the **median**, which splits the data in half. 50% of the data falling below the **median** and other 50% falling above the **median**.

There are 50 character counts in the dataset (an even number) so the data are perfectly split into two groups of 25. We take the **median** in this case to be the **average** of the two observations closest to the 50th percentile:

$$(6,768 + 7,012)/2 = 6,890.$$

When there are an odd number of observations, there will be exactly one observation that splits the data into two halves, and in such a case that observation is the **median** (no average needed).

**Median** If the data are ordered from smallest to largest, the median is the observation right in the middle. If there are an even number of observations, there will be two values in the middle, and the median is taken as their average.

The second step in building a box plot is drawing a rectangle to represent the middle 50 of the data. The total length of the box, is called the **interquartile range (IQR)**. It, like the **Standard Deviation**, is a measure of Variability in data. The more variable the data, the larger the **Standard Deviation** and **IQR**.

The two boundaries of the box are called the **first quartile** (the 25<sup>th</sup> percentile), i.e. 25 of the data fall below this value and the **third quartile** (the 75<sup>th</sup> percentile), and these are often labeled **Q1** and **Q3**, respectively.

**Interquartile range (IQR)** The IQR is the length of the box in a box plot. It is computed as

$$IQR = Q3 - Q1$$

where **Q1** and **Q3** are the 25<sup>th</sup> and 75<sup>th</sup> percentiles.

```
In [ ]: sns.set(style="whitegrid")
ax = sns.boxplot(x=db["num_char"], color='lightblue', fliersize=5, orient='v', linewidth=1, width=0.3)
```

```
In [ ]: sns.stripplot(x=db["num_char"], orient='v', color='darkblue')
```

```
In [ ]: ax = sns.boxplot(y="num_char", data=db, color='lightblue', fliersize=5, orient='v', linewidth=1, width=0.3)
ax = sns.stripplot(x=db["num_char"], orient='v', color='darkblue')
```

```
In [ ]: sns.set(style="whitegrid")
ax = sns.boxplot(x=db["num_char"], color='lightblue', fliersize=5, orient='v', linewidth=1, width=0.3)
```

```
In [ ]: sns.swarmplot(x=db["num_char"], orient='v', color='darkblue')
```

```
In [ ]: ax = sns.boxplot(y="num_char", data=db, color='lightblue', fliersize=5, orient='v', linewidth=1, width=0.3)
ax = sns.swarmplot(y="num_char", data=db, color="darkblue", orient="v", size=4)
```

### EXERCISE - 3.8

1. What percent of the data fall between Q1 and the median?
2. What percent is between the median and Q3?

Extending out from the box, the **whiskers** attempt to capture the data outside of the box, however, their reach is never allowed to be more than  $1.5 \times IQR$

They capture everything within this reach. The **upper whisker** does not extend to the last three points, which is beyond  $Q3 + 1.5 \times IQR$ , and so it extends only to the last point below this limit.

The **lower whisker** stops at the lowest value, 33, since there is no additional data to reach; the **lower whisker's limit** is not shown in the figure because the plot does not extend down to  $Q1 - 1.5 \times IQR$ . In a sense, the box is like the body of the box plot and the **whiskers** are like its arms trying to reach the rest of the data.

Any observation that lies beyond the **whiskers** is labeled with a **dot**. The purpose of labeling these points – instead of just extending the whiskers to the minimum and maximum observed values – is to help identify any observations that appear to be **unusually** distant from the rest of the data. **Unusually** distant observations are called **Outliers**.

In this case, it would be reasonable to classify the emails with character counts of 41,623, 42,793, and 64,401 as **outliers** since they are numerically distant from most of the data.

Outlier An **outlier** is an *observation* that appears **extreme** relative to the rest of the **data**. Why it is important to look for outliers Examination of data for possible **outliers** serves many useful purposes, including : 1. Identifying strong **skew** in the distribution. 2. Identifying data collection or **entry errors**. For instance, we re-examined the email purported to have 64,401 characters to ensure this value was accurate. 3. Providing **insight** into interesting **properties** of the **data**.

### **EXERCISE - 3.9**

estimate the following values for num\_char in the *email50* dataset:

- a).-  $Q_1$ ,
- b).-  $Q_3$ , and
- c).-  $IQR$

