STATS8: Introduction to Biostatistics

Data Exploration

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Introduction

- After clearly defining the scientific problem, selecting a set of representative members from the population of interest, and collecting data (either through passively observing events or experiments), we usually begin our analysis with data exploration.
- We start by focusing on data exploration techniques for one variable at a time.
- Our objective is to develop a high-level understanding of the data, learn about the possible values for each characteristic, and find out how a characteristic varies among individuals in our sample.
- In short, we want to learn about the distribution of variables.

Variable types

- The visualization techniques and summary statistics we use for a variable depend on its type.
- Based on the values a variable can take, we can classify them into two general groups: numerical variables and categorical variables.
- Consider the Pima.tr data available from the MASS library.
- Variables npreg, age, and bmi in this data set are numerical variables since they take numerical values, and the numbers they take have their usual meaning.
- The type variable in this data set, on the other hand, is categorical since the set of values it can take consists of a finite number of categories.



Variable types

- Some numerical variables are count variables. For example, number of pregnancies and number of physician visits.
- For categorical variables, we typically use numerical codings.
- Categorical variables are either ordinal or nominal depending on the extent of information the numerical coding provides.
- For nominal variables, such as type, the numbers are simply labels, which are chosen arbitrarily.
- For ordinal variables, such as disease severity, although the numbers do not have their usual meaning, they preserve a rank ordering.

Frequency and relative frequency

- The number of times a specific category is observed is called **frequency**. We denote the frequency for category c by n_c .
- The relative frequency is the sample proportion for each possible category. It is obtained by dividing the frequencies n_c by the total number of observations n:

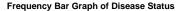
$$p_c = \frac{n_c}{n}$$

- Relative frequencies are sometimes presented as percentages after multiplying proportions p_c by 100.
- For a categorical variable, the mode of is the most common value, i.e., the value with the highest frequency.

Bar graph

- For categorical variables, bar graphs are one of the simplest ways of visualizing the data.
- Using a bar graph, we can visualize the possible values (categories) a categorical variable can take, as well as the number of times each category has been observed in our sample.
- The height of each bar in this graph shows the number of times the corresponding category has been observed.

Bar graphs and frequencies



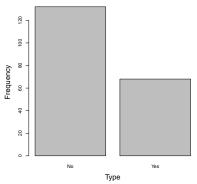
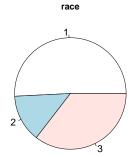


Figure: Using R-Commander to create and view a frequency bar graph for type in the Pima.tr data set. The heights of the bars sum to the sample size n. Overall, bar graphs show us how the observed values of a categorical variable in our sample are distributed

Pie chart

- We can use a pie chart to visualize the relative frequencies of different categories for a categorical variable.
- In a pie chart, the area of a circle is divided into sectors, each representing one of the possible categories of the variable.
- The area of each sector c is proportional to its frequency.



Exploring Numerical Variables

- For numerical variables, we are especially interested in two key aspects of the distribution: its location and its spread.
- The location of a distribution refers to the central tendency of values, that is, the point around which most values are gathered.
- The spread of a distribution refers to the dispersion of possible values, that is, how scattered the values are around the location.

Exploring Numerical Variables

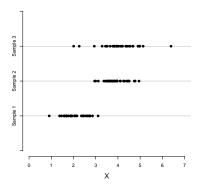
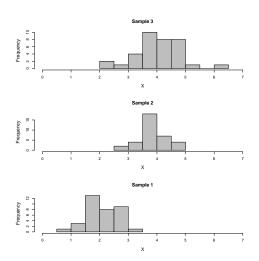


Figure: Three separate samples for variable X. Observations in Sample 1 are gathered around 2, whereas observations in Sample 2 and Sample 3 are gathered around 4. Observations in Sample 3 are more dispersed compared to those in Sample 1 and Sample 2

Histograms

- Histograms are commonly used to visualize numerical variables.
- A histogram is similar to a bar graph after the values of the variable are grouped (binned) into a finite number of intervals (bins).
- For each interval, the bar height corresponds to the frequency (count) of observation in that interval.

Histograms



Histograms

- The bar height for each interval could be set to its relative frequency $p_c = n_c/n$, or the percentage $p_c \times 100$, of observations that fall into that interval.
- For histograms, however, it is more common to use the density instead of the relative frequency or percentage.
- The density is the relative frequency for a unit interval. It is obtained by dividing the relative frequency by the interval width:

$$f_c = \frac{p_c}{w_c}. (1)$$

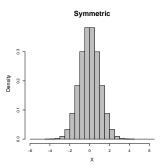
Here, $p_c = n_c/n$ is the relative frequency with n_c as the frequency of interval c and n as the total sample size.

The width of interval c is denoted w_c.



Shapes of histograms

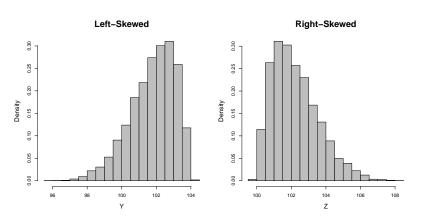
- Besides the location and spread of a distribution, the shape of a histogram also shows us how the observed values spread around the location.
- We say the following histogram is symmetric around its location (here, zero) since the densities are the [almost] same for any two intervals that are equally distant from the center.



Skewed histograms

- In many situations, we find that a histogram is stretched to the left or right.
- We call such histograms skewed.
- More specifically, we call them left-skewed if they are stretched to the left, or right-skewed if they are stretched to the right.

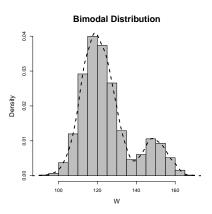
Skewed histograms



Unimodal vs. bimodal

- The above histograms, whether symmetric or skewed, have one thing in common: they all have one peak (or mode).
- We call such histograms (and their corresponding distributions) unimodal.
- Sometimes histograms have multiple modes.
- The bimodal histogram appears to be a combination of two unimodal histograms.
- Indeed, in many situations bimodal histograms (and multimodal histograms in general) indicate that the underlying population is not *homogeneous* and may include two (or more in case of multimodal histograms) subpopulations.

Unimodal vs. bimodal



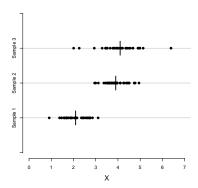
Sample mean

- Histograms are useful for visualizing numerical data and identifying their location and spread. However, we typically use summary statistics for more precise specification of the central tendency and dispersion of observed values.
- A common summary statistic for location is the sample mean.
- The sample mean is simply the average of the observed values. For observed values x_1, \ldots, x_n , we denote the sample mean as \bar{x} and calculate it by

$$\bar{x} = \frac{\sum_{i} x_{i}}{n},$$

where x_i is the *i*th observed value of X, and n is the sample size.

Sample mean



Sample mean

- Sample mean is sensitive to very large or very small values, which might be outliers (unusual values).
- For instance, suppose that we have measured the resting heart rate (in beats per minute) for five people.

$$x = \{74, 80, 79, 85, 81\},$$
 $\bar{x} = \frac{74 + 80 + 79 + 85 + 81}{5} = 79.8.$

- In this case, the sample mean is 79.8, which seems to be a good representative of the data.
- Now suppose that the heart rate for the first individual is recorded as 47 instead of 74.

$$x = \{47, 80, 79, 85, 81\},$$
 $\bar{x} = \frac{47 + 80 + 79 + 85 + 81}{5} = 74.4.$

Now, the sample mean does not capture the central tendency.



Sample median

- The **sample median** is an alternative measure of location, which is less sensitive to outliers.
- For observed values x_1, \ldots, x_n , the median is denoted \tilde{x} and is calculated by first sorting the observed values (i.e., ordering them from the lowest to the highest value) and selecting the middle one.
- If the sample size *n* is odd, the median is the number at the middle of the sorted observations. If the sample size is even, the median is the average of the two middle numbers.
- The sample medians for the above two scenarios are

$$x = \{74, 79, 80, 81, 85\},$$
 $\tilde{x} = 80;$
 $x = \{47, 79, 80, 81, 85\},$ $\tilde{x} = 80.$



- While summary statistics such as mean and median provide insights into the central tendency of values for a variable, they are rarely enough to fully describe a distribution.
- We need other summary statistics that capture the dispersion of the distribution.
- Consider the following measurements of blood pressure (in mmHg) for two patients:

Patient A:
$$x = \{95, 98, 96, 95, 96\}, \quad \bar{x} = 96, \quad \tilde{x} = 96.$$

Patient B: $y = \{85, 106, 88, 105, 96\}, \quad \bar{y} = 96, \quad \tilde{y} = 96.$

• While the mean and median for both patients are 96, the readings are more dispersed for Patient B.



- Two common summary statistics for measuring dispersion are the sample variance and sample standard deviation.
- These two summary statistics are based on the deviation of observed values from the mean as the center of the distribution.
- For each observation, the deviation from the mean is calculated as $x_i \bar{x}$.

 The sample variance is a common measure of dispersion based on the squared deviations

$$s^2 = \frac{\sum_{i=1}^{n} (x_i - \bar{x})^2}{n-1}.$$

The square root of the variance is called the sample standard deviation

$$s = \sqrt{\frac{\sum_{i=1}^{n}(x_i - \bar{x})^2}{n-1}},$$

Patient A			Patient B		
x_i	$x_i - \bar{x}$	$(x_i - \bar{x})^2$	Уi	$y_i - \bar{y}$	$(y_i - \bar{y})^2$
95	-1	1	85	-11	121
98	2	4	106	10	100
96	0	0	88	-8	65
95	-1	1	105	9	81
96	0	0	96	0	0
Σ	0	6	Σ	0	366
$s^2 = 6/4 = 1.5$			$s^2 = 366/4 = 91.5$		
$s=\sqrt{1.5}=1.22$			$s = \sqrt{91.5} = 9.56$		

Quantiles

- Informally, the sample median could be interpreted as the point that divides the ordered values of the variable into two equal parts.
- That is, the median is the point that is greater than or equal to at least half of the values and smaller than or equal to at least half of the values.
- The median is called the 0.5 quantile.
- Similarly, the 0.25 quantile is the point that is greater than or equal to at least 25% of the values and smaller than or equal to at least 75% of the values.
- In general, the q quantile is the point that is greater than or equal to at least 100q% of the values and smaller than or equal to at least 100(1-q)% of the values.

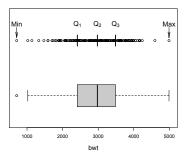
Quartiles

- We can divide the ordered values of a variable into four equal parts using 0.25, 0.5, and 0.75 quantiles.
- The corresponding points are denoted Q_1 , Q_2 , and Q_3 , respectively.
- We refer to these three points as **quartiles**, of which Q_1 is called the *first quartile* or the *lower quartile*, Q_2 (i.e., median) is called the *second* quartile, and Q_3 is called the *third quartile* or *upper quartile*.
- The interval from Q_1 (0.25 quantile) to Q_3 (0.75 quantile) covers the middle 50% of the ordered data.

- The **minimum** (min), which is the smallest value of the variable in our sample, is in fact the 0 quantile.
- On the other hand, the **maximum** (max), which is the largest value of the variable in our sample, is the 1 quantile.
- The minimum and maximum along with quartiles (Q_1 , Q_2 , and Q_3) are known as **five-number summary**.
- These are usually presented in the increasing order: min, first quartile, median, third quartile, max.
- This way, the five-number summary provides 0, 0.25, 0.50, 0.75, and 1 quantiles.

- The five-number summary can be used to derive two measures of dispersion: the range and the interquartile range.
- The range is the difference between the maximum observed value and the minimum observed value.
- The interquartile range (IQR) is the difference between the third quartile (Q_3) and the first quartile (Q_1) .

 To visualize the five-number summary, the range and the IQR, we often use a boxplot (a.k.a. box and whisker plot).



· Very often, boxplots are drawn vertically.

- The thick line at the middle of the "box" shows the median.
- The left side of the box shows the lower quartile.
- Likewise, the right side of the box is the upper quartile.
- The dashed lines are known as the whiskers.
- The whisker on the right of the box extends to the largest observed value or $Q_3 + 1.5 \times IQR$, whichever it reaches first.
- The whisker on the left extends to the lowest value or $Q_1 1.5 \times IQR$, whichever it reaches first.
- Data points beyond the whiskers are shown as circles and considered as possible outliers.

