

Week 2: Measurement and Distributions

Univariate Statistics and Methodology using R

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Today's Key Topics

- Histograms & Density Plots
- The Normal Distribution
- Populations & Samples
- Central Limit Theorem

Part 1



Measurement

- When we measure something, we attempt to identify its true measurement, or the ground truth
- The problem is that we don't have any way of measuring accurately enough
 - Our measurements are likely to be close to the truth
 - They will likely vary if we take multiple measures
- Let's run a quick experiment:



Measurement

We might expect values close to the true measurement to be more frequent if we take multiple measurements:





(Though there are still limits to our precision)

Considering this principle, it might be useful to create a histogram of all measurements taken.



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Note difference from a bar chart:

- Histogram represents continuous data
- Bar Chart represents categorical data



Considering this principle, it might be useful to create a histogram of all measurements taken.



- We know that there are 17 measurements around 7.55
- Strictly, between 7.548 & 7.567

Histograms in R

df <- data.frame(x=rnorm(200, mean = 7.55, sd = 0.1))
head(df)</pre>

x
1 7.609
2 7.614
3 7.471
4 7.578
5 7.491
6 7.524

hist(df\$x)



Histograms in R

df <- data.frame(x=rnorm(200, mean = 7.55, sd = 0.1))
head(df)</pre>

x
1 7.780
2 7.461
3 7.514
4 7.638
5 7.476
6 7.583

library(ggplot2)
ggplot(df, aes(x)) +
 geom_histogram(colour='black')



stat_bin() using bins = 30. Pick better value with binwidth.

Histograms in R

Note that the bin width of the histogram matters. Every figure below displays the same data.



- The Good
 - Way to examine the *distribution* of the data

 - Easy to interpret (y axis = counts)
 Sometimes helpful in spotting weird data (outliers)

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- Way to examine the *distribution* of the data
- Easy to interpret (y axis = counts)
- Sometimes helpful in spotting weird data (outliers)
- The Bad
 - $\circ~$ Only gives us information about distribution and mode; doesn't give us other information
 - E.g., the mean or median
 - Changing bin width can completely change the graph

Density Plots

- Similar to histogram in that it shows the distribution of the data
- However, the *y* axis is no longer a count, but represents a **proportion** of cases.
- The area under the curve is equal to 1 (or 100%, reflecting all cases)



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Density Plots in R

ggplot

ggplot(df, aes(x)) + geom_density()



base R

densDat <- density(df\$x) plot(densDat)



density.default(x = df\$x)

Part 2

The Normal Distribution

The Normal Distribution

- A hypothetical density plot
 - Probability distribution of a random variable
- Normal curves are unimodal, with values symmetrically distributed around the peak
 - $\circ~$ Centered around the mean
 - A higher proportion of cases near the mean and a lower proportion of cases with more extreme values



The Normal Distribution

Normal curves can be defined in terms of *two parameters*:

• The mean of the distribution (\bar{x} , or sometimes μ)

x <- c(22, 24, 21, 19, 22, 20) mean(x)

[1] 21.33

• The standard deviation of the distribution (sd, or sometimes σ)

sd(x)

[1] 1.751



A quick note on standard deviation

The standard deviation is the average distance of observations from the mean

$$\mathrm{sd} = \sqrt{rac{\sum{(x-ar{x})^2}}{n-1}}$$

x = individual observation

 \bar{x} = mean

n = sample size

 \sum = add it up





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Part 3

Sampling from a Population

Samples vs Populations

- **Population** all members of the group that you are hypothesizing about
- **Sample** the subset of the population that you're testing to find the answer
- If we repeatedly sample from a population and measure the mean of each sample, we'll get a normal distribution
 - $\circ\;$ the mean will be (close to) the population mean
 - the standard deviation ("width") of the distribution of sample means is referred to as the standard error of the distribution



Statistical Estimates

- so far, we've talked about sampling repeatedly from a population
- this might not be possible
- if we only have one sample we can make *estimates* of the mean and standard error
 - \circ the estimated *mean* is the sample mean (we have no other info)
 - the estimated *standard error* of the mean is defined in terms of the sample standard deviation

$$ext{se} = rac{\sigma}{\sqrt{n}} = rac{\sqrt{rac{\sum (x-ar{x})^2}{n-1}}}{\sqrt{n}}$$

Part 4

Towards Statistical Testing

Central Limit Theorem

- What we have just seen is a demonstration of Central Limit Theorem
- Lay version: sample means will be normally distributed about the true mean
- The more samples you take, the more normal the distribution should look, regardless of the variable's distribution in the population



The Standard Normal Curve

We can *standardize* any value on any normal curve by:

- subtracting the mean
 - the effective mean is now zero
- dividing by the standard deviation
 - $\circ~$ the effective standard deviation is now one

These new standardized values are called *z*-scores.

The standardized normal distribution is also known as the z-distribution.



The Standard Normal Curve

- ~68% of observations fall within one standard deviation of the mean.
- ~32% of observations fall **greater than one** standard deviation above or below the mean



The Standard Normal Curve

- ~95% of observations fall within 1.96 standard deviations of the mean
- ~5% of observations fall **greater than 1.96** standard deviations above or below the mean
- We can phrase it another way: *an area of .95* lies between -1.96 and 1.96 standard deviations from the mean
 - "95% of predicted observations" (the 95% confidence interval)



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- perhaps we're interested in the "average height of a young statistician" (!)

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Can we use the information from the sample of 386 responses we have to say anything about the population?

Looking at the class data

```
ggplot(hData, aes(height)) +
  geom_histogram(colour = 'black', binwidth = 1) +
  labs(x = 'height (cm)')
```



head(hData\$height)
[1] 171 149 173 159 157 177
mean(hData\$height)
[1] 167.9
sd(hData\$height)
[1] 8.23

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Remember, in normally distributed data, 95% of the data fall between $ar{x}-1.96\sigma$ and $ar{x}+1.96\sigma$

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mean(hData\$height) + 1.96*sd(hData\$height)

[1] 184

mean(hData\$height) - 1.96*sd(hData\$height)

[1] 151.7

Statistically Useful Information

Remember, in normally distributed data, 95% of the data fall between $ar{x}-1.96\sigma$ and $ar{x}+1.96\sigma$



If we measure the mean height of 386 people from the same population as the USMR class, we estimate that the answer we obtain will lie between 151.7cm and 184cm 95% of the time

The Aim of the Game

- As statisticians, a major goal is to infer from samples to populations
- More about how we do this next time

Today's Key Points

- The distribution of data can be visualized with histograms and density plots
- Normally distributed data are symmetrically distributed, with scores near the mean being measured more often than scores further away
- Populations include every member of a group of interest, while a sample includes only those members being observed or tested
- The Central Limit Theorem states that as sample size increases, a variable's distribution begins to approximate the normal distribution.