

Lecture 2: Univariate Data

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PO7001: Quantitative Methods I

Summarizing Categorical Data

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- In R, this data will typically appear as “factors”
- e.g., we import the correlates of war data

An Example from the CoW Data

```
url <- 'http://www.correlatesofwar.org/data-sets/COW-war/in
cow <- read.csv(url)
head(cow$StateName)
```

```
## [1] Spain                France                01
## [4] Russia                Mexico                Un
## 105 Levels: Afghanistan Angola Argentina Armenia Austrai
```

```
class(cow$StateName)
```

```
## [1] "factor"
```

```
levels(cow$StateName)
```

```
## [1] "Afghanistan"          "Angola"
## [3] "Argentina"            "Armenia"
## [5] "Australia"            "Austria"    3
## [7] "Austria-Hungary"      "Azerbaijan"
```

Summarizing Categorical Data

- Typically using a table
- E.g.:

```
table(cow$StateName)
```

Now that's not very pretty, is it?

Outcome	Frequency
1	155.00
2	119.00
3	4.00
4	28.00
6	30.00
8	1.00

Table 1: Frequency Distribution of war outcomes

You can convert data from one type to another

```
table(cow$Outcome)
```

```
##
```

```
##    1    2    3    4    6    8
```

```
## 155 119    4   28   30    1
```

```
cow$Outcomef <- factor(cow$Outcome,  
                        labels=c("Winner", "Loser", "Tied",  
                                "Different type", "Stalemate",  
                                "Changed sides"))
```

Outcome	Frequency
Winner	155.00
Loser	119.00
Tied	4.00
Different type	28.00
Stalemate	30.00
Changed sides	1.00

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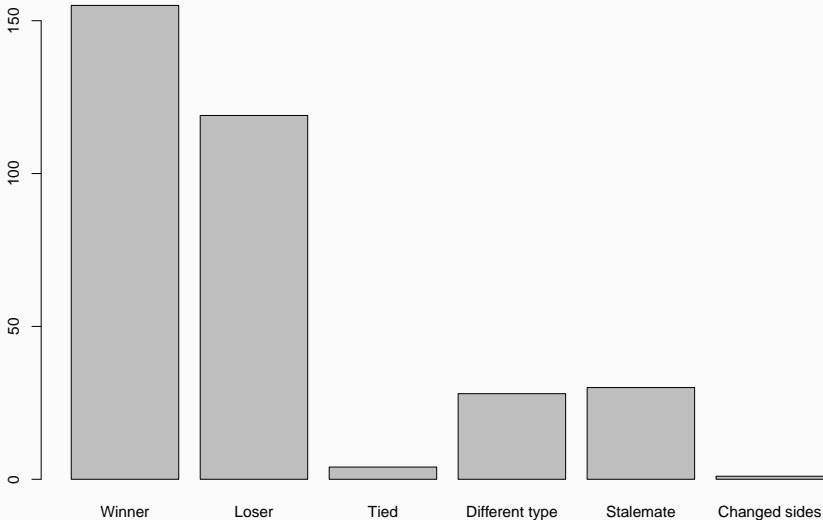
- Very intuitive and common way of representing data
- The height (or length if it is horizontal) of the bar corresponds to the frequency of a given category
- With some exceptions, height of bars should start at 0. Why?

Plotting Categorical Variables: The Barplot

- Very intuitive and common way of representing data
- The height (or length if it is horizontal) of the bar corresponds to the frequency of a given category
- With some exceptions, height of bars should start at 0. Why?
- But sometimes rules need to be broken. . .

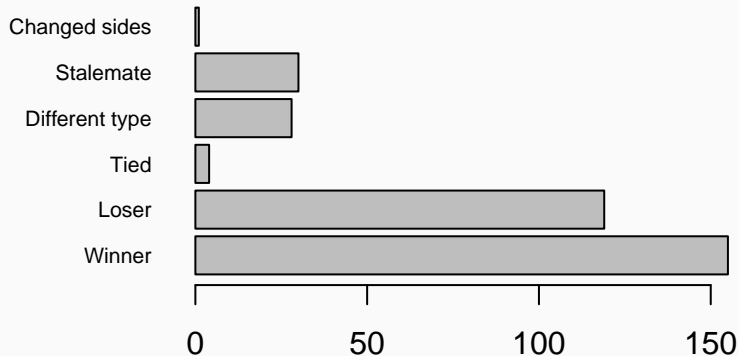
The Barplot (cont'd)

```
barplot(table(cow$Outcomef))
```



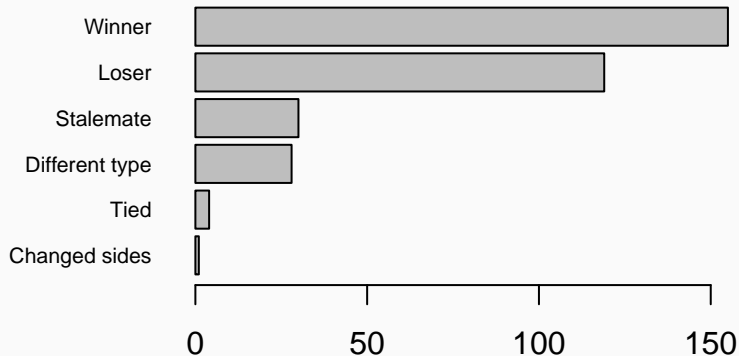
Horizontal Barplot (same thing)

```
par(mar=c(3,5,2,1))  
barplot(table(cow$Outcomef),  
        horiz=TRUE,  
        las=1,  
        cex.names=0.7)
```



Horizontal Barplot, ordered

```
par(mar=c(3,5,2,1))  
barplot(sort(table(cow$Outcomef)),  
        horiz=TRUE,  
        las=1,  
        cex.names=0.7)
```

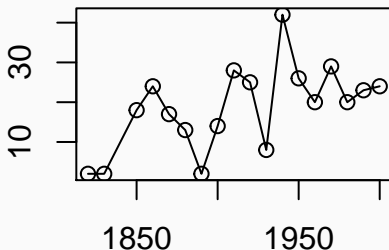
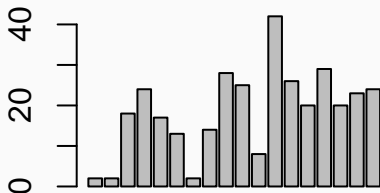


Barplots are not the best for time series

- For example, let us calculate the number of wars per decade:

```
par(mfrow = c(1,2), mar=c(2,2,1,1)) # tells R to print 2 p
cow$decade <- round(cow$StartYear1/10)*10
wars.by.year <- aggregate(cow$WarName,
                           by=list(cow$decade),
                           FUN = length)

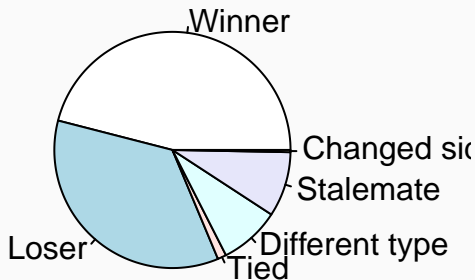
barplot(wars.by.year$x)
plot(wars.by.year, type='o')
```



Pie charts

- I rarely, if ever, see these graphs in publications. They don't look professional and are not particularly useful. If you insist on using them, though:

```
pie(table(cow$Outcomef))
```

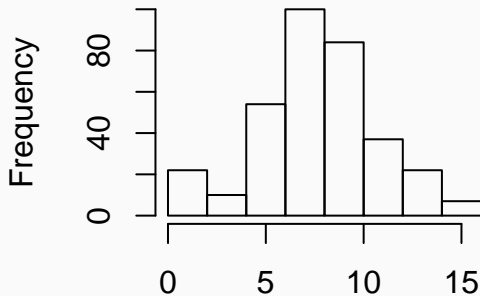


Histograms

A graphical display of tabulated frequencies shown as bars, showing the proportion of cases that fall into non-overlapping intervals of a variable

```
x <- log1p(cow$BatDeath)
hist(x)
```

Histogram of x



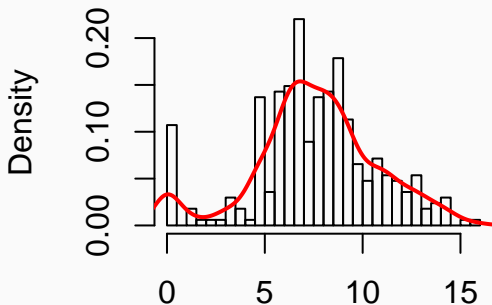
Histograms with density

```
x <- log1p(cow$BatDeath)

## Warning in log1p(cow$BatDeath): NaNs produced

hist(x, breaks=50, freq = FALSE)
lines(density(x, na.rm = TRUE), col=2, lwd=2)
```

Histogram of x

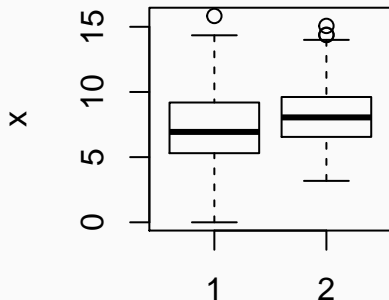
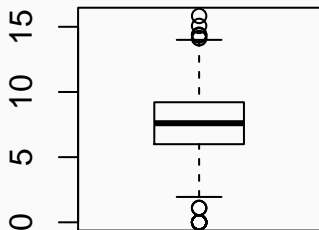


Boxplots

```
par(mfrow=c(1,2))  
x <- log1p(cow$BatDeath)
```

```
## Warning in log1p(cow$BatDeath): NaNs produced
```

```
boxplot(x)  
boxplot(x ~ cow$Side)
```



Measures of Central Tendency

Measures of Central Tendency

- Central Tendency: a single number that characterizes the “typical” unit in a set of data
- Several measures:
 - Mode
 - Median
 - Mean
- Choose depending on nature of data, what you need to convey, and the distribution of the data

The Mode

- The most *frequently* occurring value in a distribution. I.e, the category with the largest frequency.
- E.g.:
 - The mode of {1, 2, 1, 3, 4, 5 } is one
 - The mode of {Republican, Republican, Democrat, Republican, Libertarian} is Republican

The mode in R

- Unfortunately, 'mode' does not work as expected:

```
mode(cow$Outcomef)
```

```
## [1] "numeric"
```

- Luckily it's easy enough from the table:

```
table(cow$Outcomef)
```

```
##
```

```
##           Winner           Loser           Tied Different t
```

```
##           155           119           4
```

```
## Changed sides
```

```
##           1
```

- Or, if you're lazy/have too many categories:

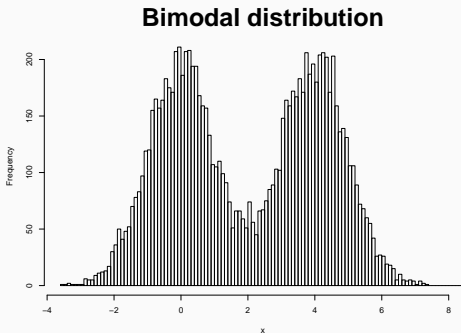
```
which(table(cow$Outcomef) == max(table(cow$Outcomef)))
```


Multiple modes

- There may be more than one mode in a sample
- For example, $\{1,2,3,1,2,4,5\}$ has two modes: 1 and 2. The uniform distribution has an infinity of modes
- In practice, large datasets make it unlikely that you have exactly two modes.

Bimodal Distributions

A *distribution* has two modes even if one of the modes is smaller than the others. What we mean then is that these are local maxima.
E.g.:



The median

- The median divides the sample in two groups of equal size. So 50% of the data will be below the median, 50% will be above.
- Find the median by ordering the data and looking for the $(N+1)/2$ point.
 - e.g.: median of $\{1,2,3,4,5\}$ is 3
 - e.g.: median of $\{1,2,3,4,5,6\}$ is 3.5
- In R:

```
x <- c(1,2,3,4,5,6)
```

```
median(x)
```

```
## [1] 3.5
```

The Mean (arithmetic)

- Same thing as the *average*
- Often written as \bar{X} or μ
- Calculated as $\frac{1}{N} \sum_{i=1}^N x_i$

```
mean(c(1,2,3,4,5))
```

```
## [1] 3
```

```
mean(c(1,2,3,4,5, 1000))
```

```
## [1] 169.1667
```

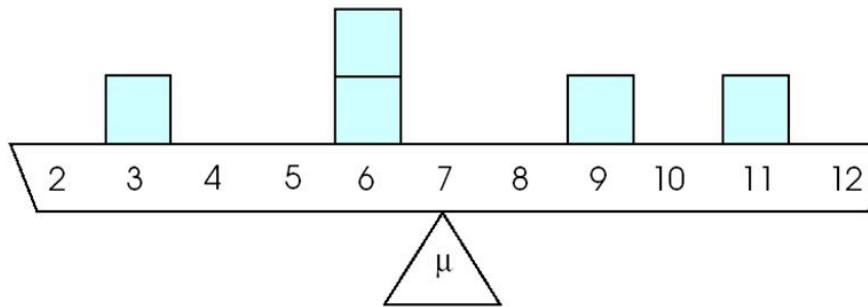
```
mean(c(1,2,3,4,5, 1000, NA))
```

```
## [1] NA
```

```
mean(c(1,2,3,4,5, 1000, NA), na.rm=TRUE)
```

```
## [1] 169.1667
```

Mean as a center of gravity



An aside on summation signs

- $\sum_{i=1}^N x_i = x_1 + x_2 + x_3 + \dots + x_n$

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- $\sum_{i=1}^N 1 = N$
- $\sum_{i=4}^6 \frac{1}{i} = \frac{1}{4} + \frac{1}{5} + \frac{1}{6}$

In class exercise

- Write a function called 'mymean', which will take a vector of numbers and return the mean

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- Create a function that will report both the mean and the median

In class exercise

- Write a function called `mymean`, which will take a vector of numbers and return the mean, without actually using the `mean` function

```
mymean <- function(x){  
  return(sum(x)/length(x))  
}
```

```
mymean(1:10)
```

```
## [1] 5.5
```

```
mymeanAndMedian <- function(x){  
  this.mean <- sum(x)/length(x)  
  midpoint <- (length(x)+1)/2  
  if(length(x)%2!=0){ #we have an odd number of observations  
    this.median <- sort(x)[midpoint]  
  }  
}
```

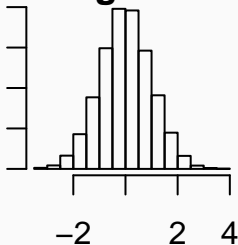
Weighted mean

- E.g., to calculate grades with different weights
- Or surveys to count observations differently
- $\bar{X}_{weighted} = \sum_i w_i X_i$

Mean, Median, and skewness

```
par(mar=c(2,1,1,1))  
x <- rnorm(10000) # Symmetric distribution (standard normal)  
hist(x)
```

Histogram of x



```
mean(x)
```

```
## [1] 0.003090373
```

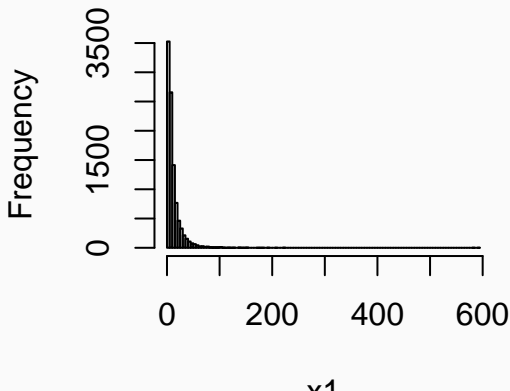
```
median(x)
```

nonsymmetric distribution (lognormal here for illustration)

```
x1 = rlnorm(10000, 2, 1)
```

```
hist(x1,  
     main=paste('mean = ', round(mean(x1)),  
                'median = ', round(median(x1))),  
     breaks=100)
```

mean = 12 median = 7



The range

- Simply the difference between largest and smallest observation
- I.e., $\text{range} = \max(x) - \min(x)$
- Dependent on extreme values

Percentiles

- The percentage of the data that is below a certain level.
- E.g., the 5th percentile means that 5% of the data is below that level
- Given an ordered variable with 100 observations, the x^{th} percentile is simply the x^{th} value
- Some percentiles have special designations:
 - the 25th percentile is the 1st *quartile*
 - the 50th percentile is the median
 - the 75th percentile is the 3rd *quartile*
 - deciles refer to every 10th percentile. E.g., 9th decile is the 90th percentile

Percentiles in R

```
x <- rnorm(1000)
# 25th percentiles
quantile(x, 0.25)
```

```
##           25%
## -0.6356508
```

```
quantile(x, 0.5) == median(x)
```

```
## 50%
## TRUE
```

```
quantile(x, probs = seq(0,1,0.1))
```

```
##           0%           10%           20%           30%
## -2.89082816 -1.18155106 -0.81261117 -0.49673094 -0.21224
##           60%           70%           80%           90%           32 1
```

Interquartile Range

- The difference between the 3rd and 1st quartile

```
x <- rnorm(1000)
```

```
IQR(x)
```

```
## [1] 1.352439
```

```
quantile(x, 0.75) - quantile(x, 0.25) # same thing
```

```
##      75%
```

```
## 1.352439
```

Variance

- $Var(x) = \sigma^2 = \frac{1}{N} \sum_{i=1}^N (x_i - \bar{x})^2$

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- Why $N - 1$? Because that is an unbiased estimate of the population variance. No need to worry too much about why, but do remember it. . .
- In R, very simple: `var(x)`

Standard Deviation

- Simply the square root of the variance
- $\text{sd}(x) = \sigma = \sqrt{\sigma^2}$
- *sample* standard deviation is denoted by s

A word about logs

- $\log_{10}(10) = 1$
- $\log_{10}(100) = 2$
- $\log_{10}(1000) = 3$
- $\ln(2.718) = 1 = \log_e(2.718)$
- $\ln(100) = 4.6$
- $\ln(1000) = 6.9$

Recommended:

- Tufte, Edward R. The visual display of quantitative information. Cheshire, CT: Graphics press, 1983. (esp. ch. 6)