# STAT 3355 Introduction to Data Analysis

Lecture 06: Summaries for Univariate Data I

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### Last Class

#### Summarize a bivariate data

	Discrete	Discrete	Continuous
	+	+	+
	Discrete	Continuous	Continuous
Numerical	F <- xtabs( $\sim x + y$ )		cor(x, y)
Graphical	barplot(F)		$plot(y \sim x)$
			$ ext{plot}(y \sim x) \  ext{abline}(( ext{lm}(y \sim x)))$

# Learning Goals

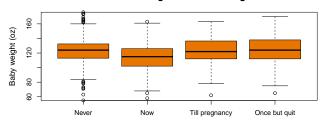
- Graphical summaries for discrete and continuous data
  - Side-by-side boxplots
  - Density plots

### Table of Contents

## Boxplot

- Displays the five-number summary of each group side-by-side
  - One axis arranges the levels of the discrete data in some order
  - The other axis represents the value of the data
  - Each box corresponds to a unique value in the discrete data
    - IQR: Box length
    - The median: Thick line through the box
    - Outliers: points out of Whiskers

#### Does smoking affects birth weight?



## Boxplot

- Implementation in R
  - lacktriangledown x is a numeric vector and y is an integer/factor/logical vector
    - lacksquare boxplot( $x \sim y$ )
  - x is a numeric variable and y is an integer/factor/logical variable in a data frame D
    - boxplot(x\_name  $\sim$  y\_name, data = D)

## Boxplot

```
# Load data
data("babies")
# Clean baby weight variable
babies$smoke[which(babies$smoke == 9)] <- NA
babies$smoke <- factor(babies$smoke, labels
   = c("Never", "Now", "Till pregnancy", "
   Once but quit"))
# Draw the boxplot
boxplot(wt ~ smoke, data = babies)
boxplot(wt \sim smoke, data = babies,
   horizontal = TRUE)
```

## The Function boxplot()

- Important arguments controlling the boxes
  - horizontal: A logical value for the orientation of the bars
  - range: A number that determines how far the whiskers extend out from the box
  - outline: A logical value for drawing the outlines defined by the range
  - col: A vector of colors for each bar
  - xlim or ylim: A numerical vector of two values indicating the limits for the axis that represents the values

## The Function boxplot()

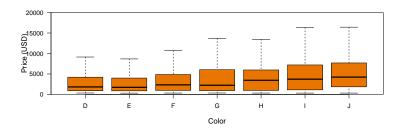
- Important arguments controlling the labels
  - main: A string of title
  - xlab and ylab: A string of label for the axis names
  - **las:** A numeric value of  $\{0,1,2,3\}$  for the orientation of axis labels

### The Function boxplot()

- Important arguments controlling the label size
  - **cex.main**: A numeric value for the title size
  - cex.lab: A numeric value for the size of axis labels
  - cex.axis: A numeric value for the size of x axis label

### Discussions

- Easy to compare the centers: The medians
- Easy to compare the spreads: The IQRs
- Easy to identify trend if the discrete data is ordinal



### Your Turn

- Continue to work on the dataset babies in the package UsingR, which contains a collection of variables taken for each new mother in a Child and Health Development Study
  - Draw a side-by-side boxplot to compare birth weight between mothers whose age were greater than or equal to 35 (advanced maternal age) and those whose age were below

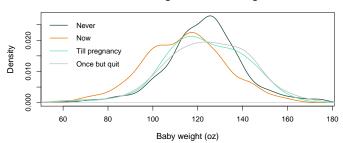
### Your Turn

#### Solutions

```
# Load data
data("babies")
# Clean baby age variable
babies$age[which(babies$age == 99)] <- NA
# Create a new variable indicating advanced
   maternal age
babies$ama <- babies$age >= 35
# Draw the side-by-side boxplot
boxplot(wt \sim ama, data = babies)
```

- Visualizes the probability distributions of a continuous data conditional on each value of the discrete data
  - x axis represents the value of the data
  - y axis represents the probability density
  - The colors indicate the levels of the discrete data in some order

#### Does smoking affects birth weight?



- Implementation in R
  - lacktriangledown x is a numeric vector and y is an integer/factor/logical vector
    - plot(density(x[which(y == )]), type = "1") and lines(density(x[which(y == )]))
  - x is a numeric variable and y is an integer/factor/logical variable in a data frame D
    - plot(density(D\$x\_name[which(D\$y\_name == )]), type
      = "1") and lines(density(D\$x\_name[which(D\$y\_name ==
      )]))

```
# Obtain the density
density_never <- density(babies$wt[which(
    babies$smoke == "Never")]);
density_now <- density(babies$wt[which(
    babies$smoke == "Now")]);
density_till <- density(babies$wt[which(
    babies$smoke == "Till pregnancy")]);
density_quit <- density(babies$wt[which(
    babies$smoke == "Once but quit")]);</pre>
```

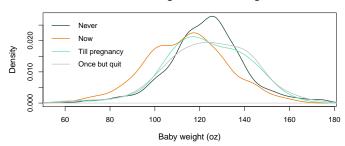
```
# Obtain the plot range
ymin <- min(density_never$y, density_now$y,
    density_till$y, density_quit$y)
ymax <- max(density_never$y, density_now$y,
    density_till$y, density_quit$y)
xmin <- min(babies$wt)
xmax <- max(babies$wt)</pre>
```

```
plot(density_never, xlim = c(xmin, xmax),
   ylim = c(ymin, ymax), xlab = "Baby weight
    (oz)", main = "", lwd = 1.5, col = 1)
lines(density_now, lwd = 1.5, col = 2)
lines(density_till, lwd = 1.5, col = 3)
lines(density_quit, lwd = 1.5, col = 4)
# Add legend
legend("topleft", c("Never", "Now", "Till
   pregnancy", "Once but quit"), col = 1:4,
   lwd = rep(1.5, 4), lty = c(1, 1, 1, 1),
   btv = 'n'
```

### Discussions

- Easy to compare the centers: The means and the modes
- Easy to compare the spreads
- Easy to identify the shapes: Symmetry, skewness, and multi-modality

#### Does smoking affects birth weight?



## After-class Reading

- Using R for Introductory Statistics (1st Ed.) by John Verzani
- Chapter 3 Bivariate data
  - Section 3.2 Comparing independent samples
    - Subsection 3.2.1 Side-by-side boxplots
    - Subsection 3.2.2 Densityplots

## After-class Reading

- Using R for Introductory Statistics (2nd Ed.) by John Verzani
- Chapter 3 Bivariate data
  - Section 3.1 Independent samples