

# Chapter 1 Study Guide: Exploring Data



**Individuals:** The objects described by a set of data. (ex: people, animals, things...)

**Variable:** Any characteristic of an individual.

**Categorical Variable:** Places an individual into groups or categories. (ex: colors)

**Quantitative Variable:** Variables that are numbers. (ex: height, weight)

**Describe Distribution:**

Shape: symmetric or skewed  
 Outliers: value that is outside pattern  
 Center: mean/median  
 Spread(Variability): range

} + CONTEXT

**Compare Distributions:**

Use comparative words  
 (similar, greater than, less than)

**SHAPE of a distribution:** Use -ly words (slightly, moderately, strongly)

Symmetric	Skewed Left	Skewed Right
Mean = Median	Mean < Median	Mean > Median

Other words to describe shape:

Bimodal	Uniform

**CENTER of a distribution:**

~Mean:  $\longrightarrow$  Use with symmetric data

~Median: middle point of a distribution ( $location = \frac{n+1}{2}$ )  $\longrightarrow$  Use with skewed data

**SPREAD/VARIABILITY of a distribution:**

~Range = max - min

~ Standard Deviation =  $\sqrt{variance}$   $\longrightarrow$  The average distance from the mean  
 $\longleftarrow$  The (context) typically varies by (SD) from the mean by ( $\bar{x}$ ).

~IQR = IQR =  $Q_3 - Q_1$

**Resistant:** A measure that is unaffected by extreme values (ex-median)

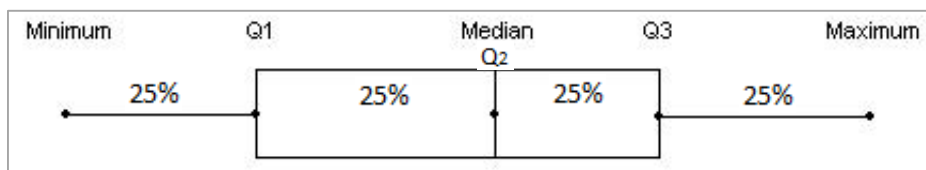
~Skewed data or outliers  $\rightarrow$  Use median and IQR

~Symmetric data  $\rightarrow$  Use mean and standard deviation

**5 Number Summary:** MIN  $Q_1$  MED  $Q_3$  MAX  $\longrightarrow$  Use to make boxplot

**Outliers:** way too small  $< Q_1 - 1.5IQR$  and way too big  $> Q_3 + 1.5IQR$

**Boxplots:**



# Chapter 2 Study Guide: Modeling Distributions of Data



**Percentile:** The value that has p% of data less than or equal to it

**Z-score:** The number of standard deviations above or below the mean.

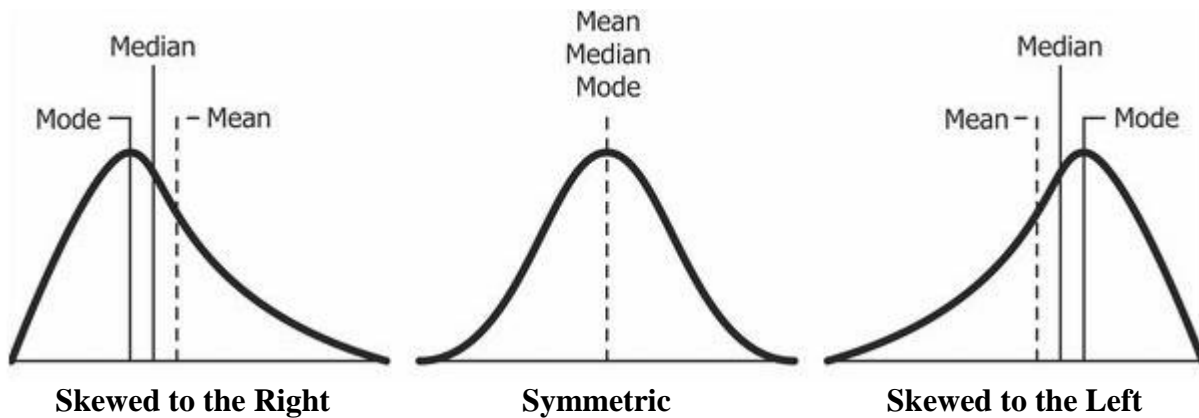
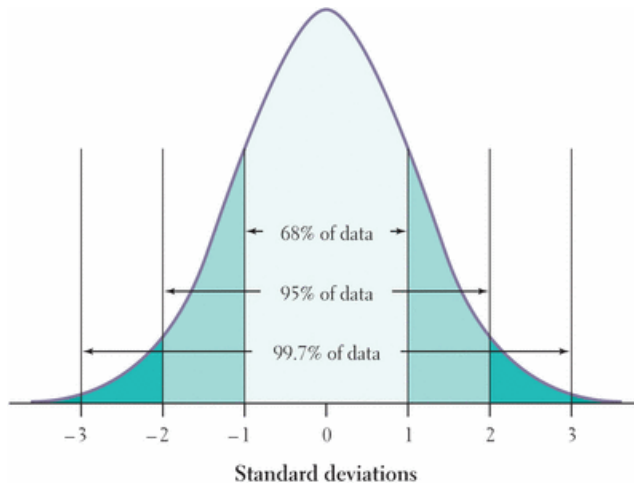
$$Z = \frac{x - \text{mean}}{\text{standard deviation}}$$

**Transforming Data:**

	+ / -	÷ / ×
S (shape)	S	S
C (center)	C	C
S (spread)	S	C

**Density Curve:** Area under the curve is 1 (100%).

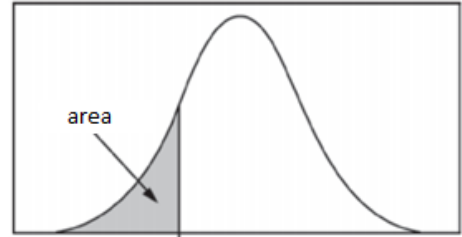
**Normal Distribution:** Symmetric and bell-shaped



**Area under the curve:**

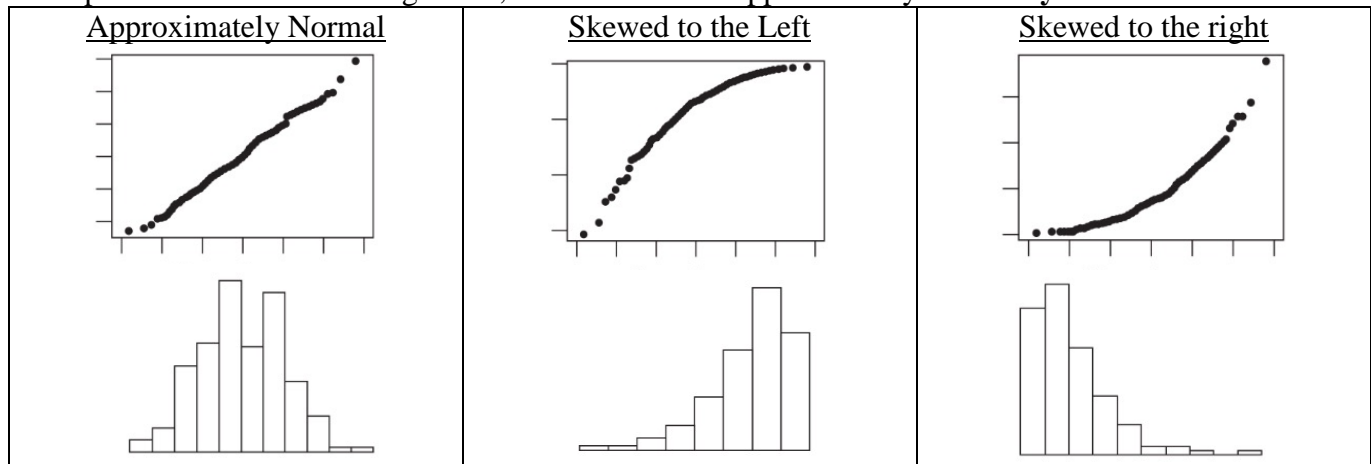
**TO FIND AREA** → 2<sup>nd</sup> → VARS → normalcdf(  
Use upper and lower as -10000 or 10000

**IF GIVEN AREA** → 2<sup>nd</sup> → VARS → invNorm(  
~Only finds area to the left



**Normal Probability Plot:**

If the points are close to a straight line, then the data are approximately **Normally distributed**.



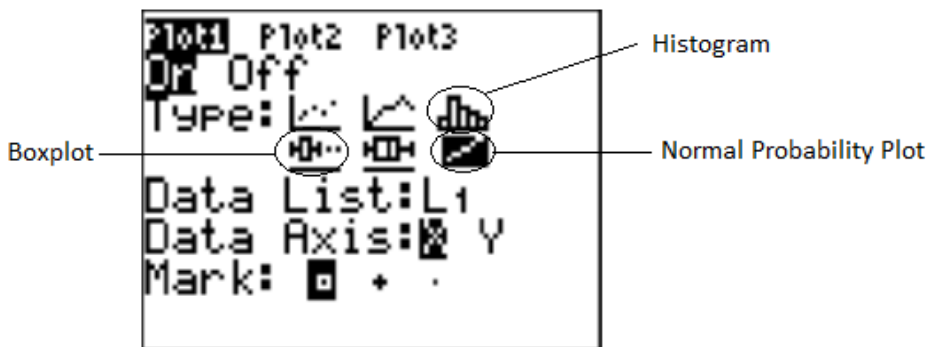
- 1) **Turn Plot1 On:** 2<sup>nd</sup> → STAT PLOT → ENTER → On
- 2) **Set Type:** Use arrow keys to highlight the Type to bottom right
- 3) **Graph:** ZOOM → 9



Does your graph look like an approximate straight line? \_\_\_\_\_

\*\*\* If the points of the NORMAL probability plot are close to a straight line, then the data are approximately **Normally distributed**.

**Graphing a Histogram, Boxplot, and Normal Probability Plot**



# Chapter 3 Study Guide: Exploring Two-Variable Data



**Explanatory Variable:** Used to predict. ( $x$ -values)

**Response Variable:** What you want to predict. Outcome, responds to explanatory. ( $y$ -values)

**Describe Relationship: (DUFS + context)**

Direction: $+/-$	} + CONTEXT
Unusual points: possible outliers	
Form: linear	
Strength: strong/moderate/weak	

**Template for Describing Relationship & Interpreting Correlation( $r$ ):**

There is a \_\_\_\_\_, \_\_\_\_\_ linear relationship between \_\_\_\_\_ and \_\_\_\_\_.

(strength)      (direction)

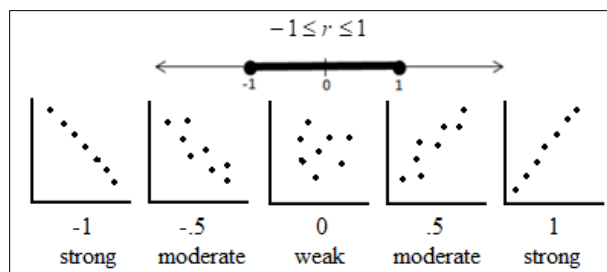
↳ ly words

**Correlation ( $r$ ):** Relationship/Association

~Direction:  $+/-$

~Strength: strong/moderate/weak

(use  $-ly$  words for in between)



**Facts/Cautions about Correlation:**

- $r$  doesn't have units  $\rightarrow$  it is just a number used to measure direction and strength!
- Switching axes doesn't change  $r \rightarrow$  any variable can be  $x$  and  $y$
- $r$  nonresistant  $\rightarrow$  strongly affected by outliers & high-leverage points
  - ~ high-leverage points in pattern strengthen
  - ~outliers out of pattern weaken
- **Correlation does NOT imply causation!!!**

**Outlier:** A point that does not follow the pattern and has a large residual. (far up/down)

**High-Leverage:** A point that has a much larger or smaller  $x$  value than the other points. (far left/right)

**Influential Point:** A point that, if removed, changes the relationship (slope/ $y$ -int/ $r/r^2$ , or  $s$ ) substantially.

\*Outliers and high-leverage points are often influential!

**Least-Squares Regression Line:** used to describe and/or predict the relationship between two variables.

$$\hat{y} = a + bx$$

↓ predicted
↓ y-int
↓ slope

~**Interpreting y-intercept:** When \_\_\_\_\_, the predicted \_\_\_\_\_ is \_\_\_\_\_.  
 (x context = 0) (y context) (y-intercept)

~**Interpreting Slope:** For every additional \_\_\_\_\_, the predicted \_\_\_\_\_ increases/decreases by \_\_\_\_\_.  
 (units in x context) (y context) (slope)

~**Extrapolation:** using the LSRL to make a prediction far outside the observed x values.

**Residual:** Actual y – Predicted  $\hat{y}$  ( $R = A - P$ )

~**Interpreting Residual:** The actual \_\_\_\_\_ is higher/lower than the predicted by \_\_\_\_\_.  
 (y-context) (residual)

**Standard Deviation (s) of the Residuals:** used to assess how well the line fits all the data. This value gives the approximate size of a “typical” prediction error (residual).

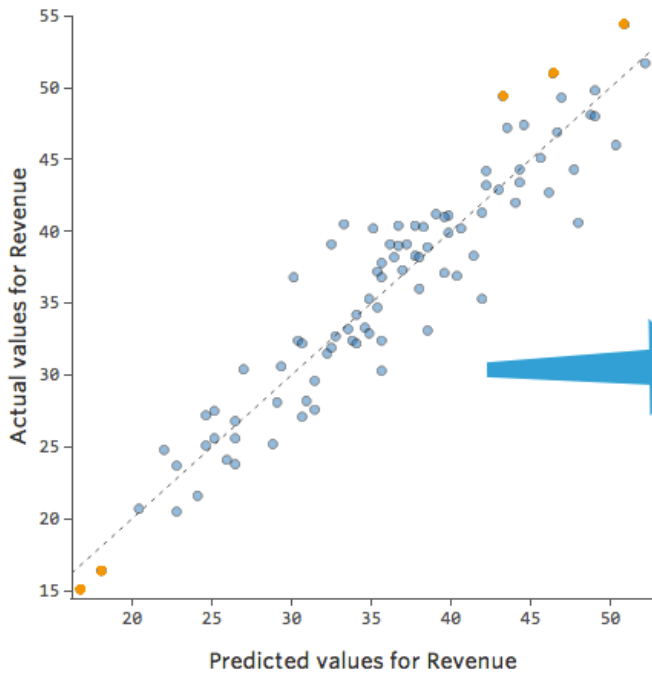
- Interpretation: The actual (y-context) is typically about (standard deviation) away from the predicted (y-context) by the LSRL.

**Coefficient of Determination ( $r^2$ ):** the fraction/percent (or piece) of the variation in the values of y that can be explained by the least-squares regression line.

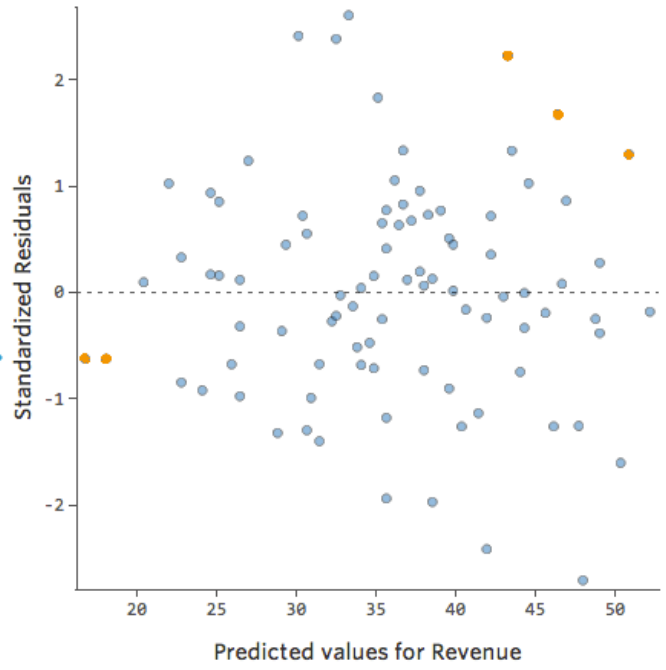
- Interpretation: About \_\_\_\_\_% of the variability in (y-context) can be explained by the LSRL.

<p><b>y-intercept (a)</b> ~When ___(x=0), the predicted ___(y) is ___.</p>	<p><b>slope (b)</b> ~For every one ___(unit in x), the predicted ___(y) increases/decreases by ___.</p>															
<table style="width: 100%; border-collapse: collapse;"> <thead> <tr> <th style="width: 25%;">Predictor</th> <th style="width: 15%;">Coef</th> <th style="width: 15%;">SE Coef</th> <th style="width: 15%;">T</th> <th style="width: 10%;">P</th> </tr> </thead> <tbody> <tr> <td>Constant</td> <td>1.201</td> <td><del>0.0074</del></td> <td>13.72</td> <td><del>0</del></td> </tr> <tr> <td>GPA (X)</td> <td>7.507</td> <td><del>1.29</del></td> <td>5.82</td> <td><del>0.0006514</del></td> </tr> </tbody> </table> <p>S = 3.252686      R-Sq = 82.8%      <del>R-Sq(adj) = 76.5%</del></p>		Predictor	Coef	SE Coef	T	P	Constant	1.201	<del>0.0074</del>	13.72	<del>0</del>	GPA (X)	7.507	<del>1.29</del>	5.82	<del>0.0006514</del>
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<p><b>SD of residuals (s)</b> ~The actual ___(y) is typically about ___(SD) away from the predicted ___(y) by the LSRL.</p>	<p><b>Coefficient of Determination (<math>r^2</math>)</b> ~ _____% of the variation in ___(y) can be explained by the LSRL.</p>															

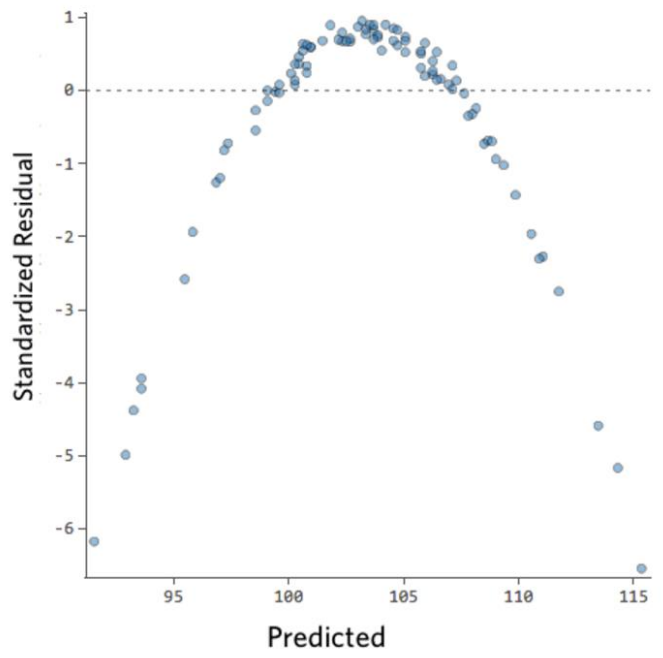
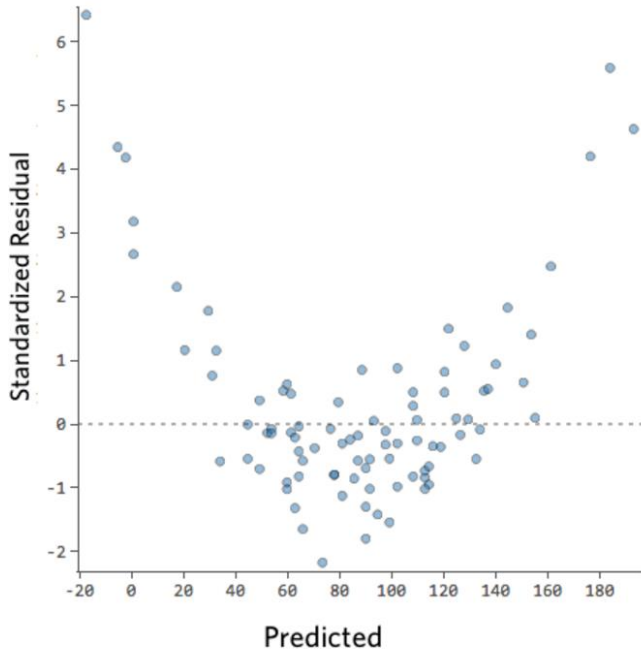
Predicted vs Actual



Residuals



## Nonlinear



# Chapter 4 Study Guide: Designing Studies



**Population:** All items or subjects.

**Sample:** Subset of population

**Census:** Selection of all items/subjects in a population

## Sampling Methods (DATA COLLECTION):

- **Simple Random Sample (SRS):** Every group of size  $n$  has an equal chance of being chosen.

<b>Describing a SRS Procedure:</b>	
<u>Hat Trick</u>	<u>Technology</u>
1) <b>Label</b> <u>individuals</u> on equal-sized pieces of paper. (or assign each individual a number) (or assign each <u>individual</u> a number) 2) <b>Randomize:</b> Put them in a hat and mix well. 3) <b>Select:</b> Randomly select _____ ( <u>individuals</u> ) (or those corresponding to the numbers) for the sample. Mix thoroughly after each selection.	1) <b>Label</b> each <u>individual</u> with a number. 2) <b>Randomize:</b> Using a random number generator, generate _____ random numbers from _____ to _____, ignoring repeats. 3) <b>Select:</b> The ( <u>individuals</u> ) corresponding to those numbers would be selected for the sample.
*Individuals: Who/what you are sampling	*In Calculator: Math → PRB → randInt(1,n)

- **Stratified Random Sample:** Split the population into groups of similarity (strata), then take an SRS of each group. Those chosen from each SRS will be selected for the sample.
  - **Homogeneous grouping**
- **Cluster** Split population into groups based on location (clusters), then randomly select clusters (1 SRS). All individuals within the cluster are selected for the sample.
  - **Heterogeneous grouping**
- **Systematic Random Sample:** Randomly select a starting point and every  $n$ th individual thereafter will be selected for the sample.

## Non-Random Sampling Methods:

- **Convenience Sample:** Choosing individuals easy to reach. (Bias)
- **Voluntary Response Sample:** People who choose themselves to be in the sample. (Bias)
  - Using people with strong opinions

**Bias:** Favoring a group (not fair).

- **Undercoverage Bias:** When members of the population are less likely to be chosen for a sample
- **Nonresponse Bias:** Individuals selected to be in a sample but can't be contacted or refuse.
- **Response Bias:** Inaccurate responses (lying or confusing questions).

**Observational Study:** No treatments imposed, only observed. (NO CAUSE/EFFECT)

- **Retrospective:** examine existing data
- **Prospective:** track individuals into the future

**Experiment:** Treatments imposed

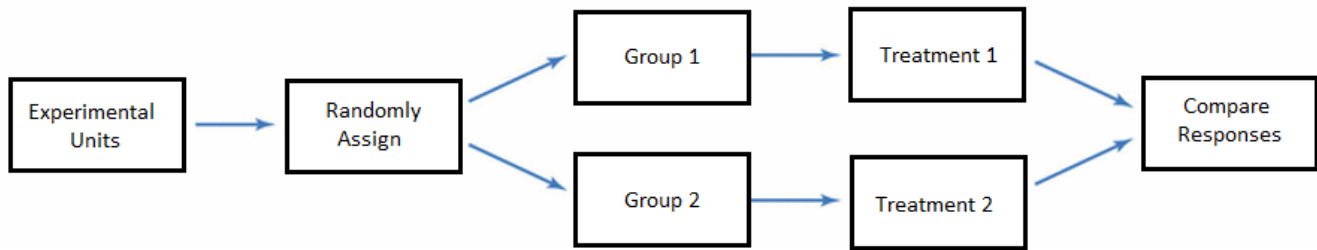
- **Experimental Units:** Individuals assigned treatments. (Humans often called **subjects**)
- **Explanatory Variable:** Variable whose levels are manipulated intentionally.
- **Response Variable:** Outcome from the treatments administered.
- **Confounding Variables:** Variables that influence the response variable.

**Well-Designed Experiment:**

- a) **Comparison:** Compares at least two treatment groups.
- b) **Random Assignment:** Experimental units randomly assigned to treatments (balances confounding variables).
- c) **Control:** Control potential confounding variables.
- d) **Replication:** Enough experimental units in each treatment group (more than one).



## Completely Randomized Design:

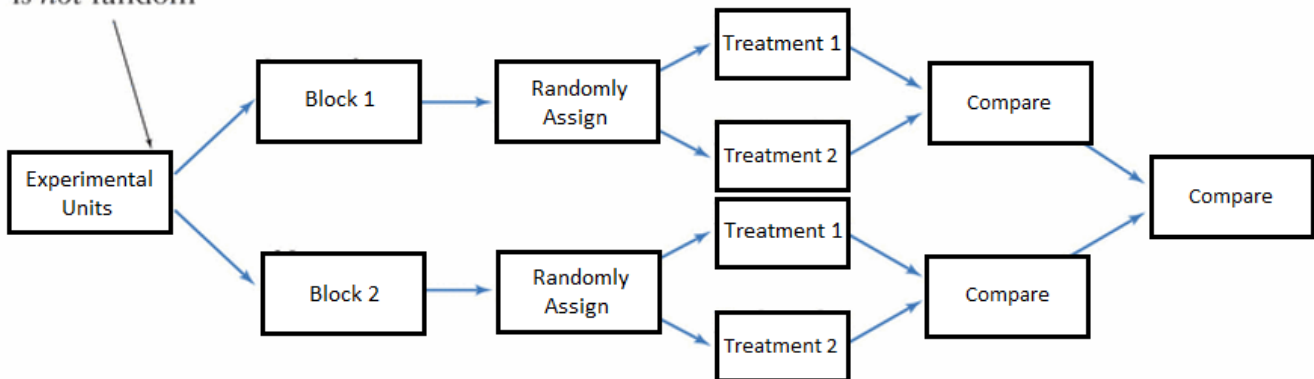


- **Random Selection:** Generalizes population.
- **Random Assignment:** Allows causation.
- **Single-blind Experiment:** Subjects do not know which treatment they are receiving.
- **Double-blind Experiment:** Neither the subjects nor the experimenters who interact with the subjects know which treatment a subject is receiving.
- **Control Group:** A group receiving an inactive treatment (**Placebo**).
- **Placebo Effect:** When experiment units have a response to a placebo (fake treatment works).

## Randomized Block Design: At the beginning of the experiment, units are divided.

- **Block:** Group of experimental units that are similar in some way.

Assignment to blocks  
is *not* random



## Matched Pairs Design: A pair or experimental units that are matched based off similarity then randomly assigned to each treatment.

- Sometimes a “pair” consists of a single unit that receives both treatments. The order of the treatments is randomly chosen.

## Statistically Significant: When results from a study are too unusual to have occurred purely by chance.

- Proportion of dots are  $\leq 5\%$  → STATISTICALLY SIGNIFICANT

# Chapter 5 Study Guide: Probability



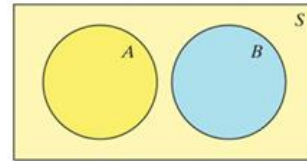
**Law of Large Numbers:** States that when an experiment is performed a large number of times, the relative frequency (proportion) of an event tends to become closer to the actual probability.

**Probability Model:** List showing all possible outcomes and their probabilities.

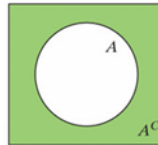
- Each probability must be between 0 and 1
- All probabilities (sample space) must add to 1

**Mutually Exclusive:** Events that cannot occur together

- If events are mutually exclusive  $\rightarrow P(A \text{ or } B) = P(A) + P(B)$

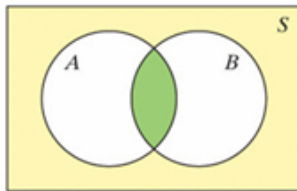


**Complement Rule:**  $P(A^C) = 1 - P(A)$

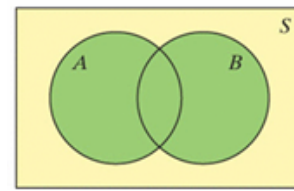


## Other Venn Diagrams and Probability Rules:

**A and B ( $A \cap B$ )**  
"both"



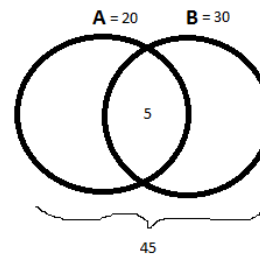
**A or B ( $A \cup B$ )**  
"either"



**General Addition Rule: (ON FORMULA SHEET)**

$$P(A \cup B) = P(A) + P(B) - P(A \cap B)$$

Example:  $P(A \text{ or } B) = 20 + 30 - 5 = 45$



**Given:**

$P(A|B) \rightarrow$  (out of B's, how many A's?)

**Independent:** One event has no effect on the probability of the other event

$$A|B = A$$

If events are independent  $\rightarrow P(A \text{ and } B) = P(A) \cdot P(B)$

# Chapter 6 Study Guide: Random Variables



**Random Variable:** Variable whose value is a numerical outcome of some chance process.

**Discrete Random Variable:** A random variable that takes a fixed set of possible values with gaps between.

**Probability Distribution:**

Value:	$x_1$	$x_2$	$x_3$	...
Probability:	$p_1$	$p_2$	$p_3$	...

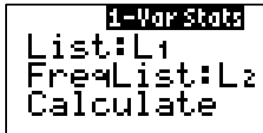
To find Expected Value (mean):  
Think “multiply and add”

Expected value (mean) =  $x_1 p_1 + x_2 p_2 + x_3 p_3 + \dots = \sum x_i p_i \rightarrow$  ON FORMULA SHEET:

$$\mu_X = E(X) = \sum x_i P(x_i)$$

\*You can also find mean and SD in calculator:

- 1) stat  $\rightarrow$  Edit
- 2) Input values into L<sub>1</sub> and L<sub>2</sub>
- 3) stat  $\rightarrow$  CALC  $\rightarrow$  1-VarStats  $\rightarrow$



**Continuous Random Variable:** A random variable that takes all values in an interval of numbers.

- To find continuous random variable probabilities  $\rightarrow$  Use normalcdf (2<sup>nd</sup>  $\rightarrow$  VARS  $\rightarrow$  normalcdf)

**Transforming and Combining Random Variables:**

**TRANSFORMATION RULES:**

Adding/Subtracting a Constant:

- Same: **shape** or **spread** (range, IQR, standard deviation)
- Change: **center** (mean, median, quartiles, percentiles)

	+/-	$\times/\div$
S	S	S
C	C	C
S	S	C

Multiplying/Dividing a Constant:

- Same: **shape**
- Changes: **center** (mean, median, quartiles, percentiles) or **spread** (range, IQR, standard deviation)

**Mean of the Sum/Difference of Random Variables**

$$\mu = \mu \pm \mu$$

\*The mean of the sum of *several* random variables is the sum or difference of their means.

**Variance of the Sum/Difference of Independent Random Variables**

$$\sigma^2 = \sigma^2 + \sigma^2$$

\*The variance of the sum of *several independent* random variables is the sum of their variances

## Binomial Probability:

Binomial Setting: (BINS)

- **Binary:** “success” or “failure”
- **Independent:** One trial’s outcome does not affect any other trial
- **Number:** Set number of trials  $n = \underline{\hspace{2cm}}$
- **Success:** Same probability  $p = \underline{\hspace{2cm}}$

To find an EXACT probability → use **binompdf**

➤ 2<sup>nd</sup> → VARS → binompdf(

**binompdf**  
trials:  
p:  
x value:  
Paste

To find a NO MORE THAN probability → use **binomcdf**

➤ 2<sup>nd</sup> → VARS → binomcdf(

To find an AT LEAST probability → use **binomcdf**

➤ 2<sup>nd</sup> → VARS → 1- binomcdf(

**binomcdf**  
trials:  
p:  
x value:  
Paste

## Mean and Standard Deviation of a Binomial Distribution: ON FORMULA SHEET:

$$\text{Mean} \rightarrow \mu_x = np$$

$$\text{Standard Deviation: } \sigma_x = \sqrt{np(1-p)}$$

## Geometric Probability:

Geometric Setting: (BINS without the N)

- **Binary:** “success” or “failure”
- **Independent:** One trial’s outcome does not affect any other trial
- **Success:** Same probability  $p = \underline{\hspace{2cm}}$

To find an EXACT probability → use **geometpdf**

➤ 2<sup>nd</sup> → VARS → geometpdf(

**geometpdf**  
p:  
x value:  
Paste

To find a NO MORE THAN probability → use **geometcdf**

➤ 2<sup>nd</sup> → VARS → geometcdf(

To find an AT LEAST probability → use **binomcdf**

➤ 2<sup>nd</sup> → VARS → 1- geometcdf(

**geometcdf**  
p:  
x value:  
Paste

## Mean and Standard Deviation of a Geometric Distribution: ON FORMULA SHEET:

$$\text{Mean} \rightarrow \mu_x = \frac{1}{p}$$

$$\text{Standard Deviation: } \sigma_x = \frac{\sqrt{1-p}}{p}$$

# Chapter 7 Study Guide: Sampling Distributions



**Sampling Distribution:** The distribution of values from all possible samples of size  $n$ .

**Parameter:** A number that describes a population.





- $p$  = population proportion and  $\mu$  = population mean

**Statistic:** A number that describes a sample.

- $\hat{p}$  = sample proportion and  $\bar{x}$  = sample mean

**Unbiased Estimator:** The mean of the sampling distribution is equal to the population parameter.

- $\hat{p} = p$  and  $\bar{x} = \mu$
- A larger sample size will decrease variability (Larger sample = Less variability)

<u>Proportions:</u>	<u>Means:</u>
 $\mu_{\hat{p}} = p$ $\sigma_{\hat{p}} = \sqrt{\frac{p(1-p)}{n}}$	 $\mu_{\bar{x}} = \mu$ $\sigma_{\bar{x}} = \frac{\sigma}{\sqrt{n}}$
 $\mu_{\hat{p}_1 - \hat{p}_2} = p_1 - p_2$ $\sigma_{\hat{p}_1 - \hat{p}_2} = \sqrt{\frac{p_1(1-p_1)}{n_1} + \frac{p_2(1-p_2)}{n_2}}$	 $\mu_{\bar{x}_1 - \bar{x}_2} = \mu_1 - \mu_2$ $\sigma_{\bar{x}_1 - \bar{x}_2} = \sqrt{\frac{\sigma_1^2}{n_1} + \frac{\sigma_2^2}{n_2}}$
<u>Conditions:</u>	<u>Conditions:</u>
<p><b>Independent:</b> <math>n \leq 10\%N</math></p> <p><b>Approximately Normal</b> <math>\rightarrow np \geq 10 \quad n(1-p) \geq 10</math></p>	<p><b>Independent:</b> <math>n \leq 10\%N</math></p> <p><b>Normal</b> <math>\rightarrow</math> Population Normal</p> <p><b>Approximately Normal</b> <math>\rightarrow n \geq 30</math></p>

# Chapter 8 Study Guide: Confidence Intervals



**Statistic  $\pm$  (critical value)(standard error of statistic)**  
(point estimate)

## One-Sample z Interval for a Population Proportion

In Calculator:

STAT  $\rightarrow$  TESTS  $\rightarrow$  1-PropZint (A)  $\rightarrow$

**1-PropZint**  
x:  
n:  
C-Level:

By Hand:

$$\hat{p} \pm z^* \sqrt{\frac{\hat{p}(1-\hat{p})}{n}}$$

Find  $z$ : 2<sup>nd</sup>  $\rightarrow$  VARS  $\rightarrow$

**invNorm(**  
area:  
 $\mu$ :  
 $\sigma$ :

## One-Sample t Interval for a Population Mean: (When Population SD is unknown)

In Calculator:

STAT  $\rightarrow$  TESTS  $\rightarrow$  TInterval (8)  $\rightarrow$

**TInterval**  
Input: Stats  
 $\bar{x}$ :  
Sx:  
n:  
C-Level:

By Hand:

$$\bar{x} \pm t^* \frac{s_x}{\sqrt{n}}$$

Find  $t$ : 2<sup>nd</sup>  $\rightarrow$  VARS  $\rightarrow$

**invT(**  
area:  
df:

(df = n-1)

Width of confidence interval:

- Confidence interval  $\downarrow$  as sample size  $n \uparrow$
- Confidence interval  $\uparrow$  as confidence level  $\uparrow$

statistic  $\pm$  (critical value)(standard error of statistic)

ME

To find sample size:  $z^* \sqrt{\frac{\hat{p}(1-\hat{p})}{n}} \leq ME$

$\rightarrow$  If  $\hat{p}$  is not given  $\rightarrow$  use 0.5

**Interpreting Confidence Level:**

\*In repeated random samples of size \_\_\_\_\_ (n) from this population, about \_\_\_\_\_% of confidence intervals created will capture the true population (proportion or mean) of...

\*KEMP DEF: In the long run, \_\_\_\_\_ % of intervals generated capture the population proportion/mean

# Confidence Intervals: A Four-Step Process

	<b>Proportions</b>	<b>Means</b>
<b>S T A T E</b>	<p><math>p</math> = true <b>proportion</b> of...</p> <p>Confidence Level = _____</p>	<p><math>\mu</math> = true <b>mean</b> of...</p> <p>Confidence Level = _____</p>
<b>P L A N</b>	<p><u>Name</u>: One Sample <math>z</math> interval for <math>p</math></p> <p><u>Conditions</u>:</p> <ol style="list-style-type: none"> <li>1) Random sample</li> <li>2) Independent if <math>n \leq 10\%N</math></li> <li>3) App Normal if <math>n\hat{p} \geq 10</math> and <math>n(1 - \hat{p}) \geq 10</math></li> </ol>	<p><u>Name</u>: One Sample <math>t</math> interval for <math>\mu</math></p> <p><u>Conditions</u>:</p> <ol style="list-style-type: none"> <li>1) Random sample</li> <li>2) Independent if <math>n \leq 10\%N</math></li> <li>3) Approximately Normal if:                             <ul style="list-style-type: none"> <li>➤ <math>n \geq 30</math></li> <li>➤ Sample shows no strong skewness and outliers</li> </ul> </li> </ol>
<b>D O</b>	<p><math>\hat{p} \pm z^* \sqrt{\frac{\hat{p}(1 - \hat{p})}{n}}</math>    OR    1-PropZInt (A)</p>	<p><math>\bar{x} \pm t^* \frac{s_x}{\sqrt{n}}</math>    OR    TInterval (8)</p>
<b>C O N C L U D E</b>	<p>We are _____% confident that the interval from _____ to _____ captures the true <b>proportion</b> of...</p>	<p>We are _____% confident that the interval from _____ to _____ captures the true <b>mean</b> of...</p>

# Chapter 9 Study Guide: Significance Tests



$$\text{Standardized test statistic} = \frac{\text{statistic} - \text{parameter}}{\text{standard error of the statistic}} = \frac{\text{stat} - \text{null}}{SD}$$

## One-Sample z Test for a Population Proportion

In Calculator:

(5)  
STAT → TESTS → 1-PropZTest →

```

1-PropZTest
P0:
X:
n:
PROP≠P0 <P0 >P0
Calculate Draw
    
```

By Hand:

$$z = \frac{\hat{p} - p_0}{\sqrt{\frac{p_0(1-p_0)}{n}}}$$

2<sup>nd</sup> → VARS →

```

normalcdf
lower:
upper:
μ: 0
σ: 1
Paste
    
```

## One-Sample t Test for a Population Mean

In Calculator:

STAT → TESTS → T-Test (2) →

```

T-Test
Inpt: Data Stats
μ0:
X:
Sx:
n:
μ: ≠μ0 <μ0 >μ0
Calculate Draw
    
```

By Hand:

$$t = \frac{\bar{x} - \mu_0}{\frac{s}{\sqrt{n}}}$$

2<sup>nd</sup> → VARS →

```

tcdf
lower:
upper:
df:
Paste
    
```

(df = n-1)

### Errors (HOT HAT):



- **Type I Error (Hot):** Reject  $H_0$  when  $H_0$  is true
- **Type II Error (Hat):** Fail to reject  $H_0$  when  $H_a$  is true

	$H_0$ true	$H_a$ true
Reject $H_0$	Type I error $\alpha$	Correct conclusion → POWER
Fail to reject $H_0$	Correct conclusion	Type II error $\beta$

**Power:** Probability that the test will correctly reject  $H_0$

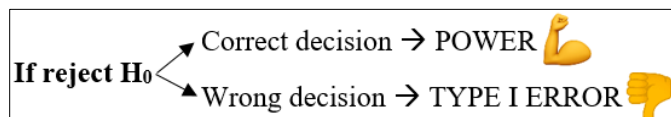
**P-value:** Assuming the  $H_0$  is true, the probability of one sample occurring

$\alpha$ : Probability of making a type I error

$\beta$ : Probability of making a type II error

$$P = 1 - \beta$$

↓  
prob of type II error





# Significance Tests: A Four-Step Process

<b>Proportions</b>		<b>Means</b>	
<b>S T A T E</b>	<p><math>p</math> = true <b>proportion</b> of...</p> <p><math>H_0: p =</math></p> <p><math>H_a: p \neq, &lt;, \text{ or } &gt;</math></p>	<b>S T A T E</b>	<p><math>\mu</math> = true <b>mean</b> of...</p> <p><math>H_0: \mu =</math></p> <p><math>H_a: \mu \neq, &lt;, \text{ or } &gt;</math></p>
<b>P L A N</b>	<p><u>Name</u>: One Sample <math>z</math> Test for <math>p</math></p> <p><u>Conditions</u>:</p> <ol style="list-style-type: none"> <li>1) Random sample</li> <li>2) Independent if <math>n \leq 10\%N</math></li> <li>3) App Normal if <math>np_0 \geq 10</math> and <math>n(1 - p_0) \geq 10</math></li> </ol>	<b>P L A N</b>	<p><u>Name</u>: One Sample <math>t</math> Test for <math>\mu</math></p> <p><u>Conditions</u>:</p> <ol style="list-style-type: none"> <li>1) Random sample</li> <li>2) Independent if <math>n \leq 10\%N</math></li> <li>3) Approximately Normal if:                             <ul style="list-style-type: none"> <li>➤ <math>n \geq 30</math></li> <li>➤ Sample shows no strong skewness and outliers</li> </ul> </li> </ol>
<b>D O</b>	$z = \frac{\hat{p} - p_0}{\sqrt{\frac{p_0(1 - p_0)}{n}}}$ <p style="text-align: center;">OR    1-PropZTest (5)</p>	<b>D O</b>	$t = \frac{\bar{x} - \mu_0}{\frac{s}{\sqrt{n}}}$ <p style="text-align: center;">OR    T-Test (2)</p>
<b>C O N C L U D E</b>	<p><math>p\text{-value} \leq \alpha</math>, reject <math>H_0</math>. We have convincing evidence that the true <b>proportion</b> of...(H<sub>a</sub>)</p> <p><math>p\text{-value} &gt; \alpha</math>, fail to reject <math>H_0</math>. We do not have convincing evidence that the true <b>proportion</b> of...(H<sub>a</sub>)</p>	<b>C O N C L U D E</b>	<p><math>p\text{-value} \leq \alpha</math>, reject <math>H_0</math>. We have convincing evidence that the true <b>mean</b> of...(H<sub>a</sub>)</p> <p><math>p\text{-value} &gt; \alpha</math>, fail to reject <math>H_0</math>. We do not have convincing evidence that the true <b>mean</b> of...(H<sub>a</sub>)</p>

# Chapter 10 Study Guide: Comparing Two Populations or Groups



## Proportions

### Two Sample z Interval for $p_1 - p_2$

In Calculator:

STAT → TESTS → 2-PropZint (B) →

```

2-PropZInt
x1:
n1:
x2:
n2:
C-Level:
Calculate
    
```

By Hand

$$(\hat{p}_1 - \hat{p}_2) \pm z^* \sqrt{\frac{\hat{p}_1(1-\hat{p}_1)}{n_1} + \frac{\hat{p}_2(1-\hat{p}_2)}{n_2}}$$

```

invNorm
area:
μ:0
σ:1
Paste
    
```

### Two Sample z Test for $p_1 - p_2$

In Calculator: STAT → TESTS → 2-PropZTest (6) →

```

2-PropZTest
x1:
n1:
x2:
n2:
P1: ≠P2 <P2 >P2
Calculate Draw
    
```

## Means

### Two Sample t Interval for $\mu_1 - \mu_2$

In Calculator:

STAT → TESTS → 2-SampTInt (0) →

```

2-SampTInt
Inpt:Data Stats
x1:
Sx1:
n1:
x2:
Sx2:
n2:
C-Level:
Pooled: [ ] Yes
Calculate
    
```

By Hand

$$(\bar{x}_1 - \bar{x}_2) \pm t^* \sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}}$$

```

invT
area:
df:
Paste
    
```

### Two Sample t Test for $\mu_1 - \mu_2$

In Calculator: STAT → TESTS → 2-SampTTest (4) →

```

2-SampTTest
Inpt:Data Stats
x1:
Sx1:
n1:
x2:
Sx2:
n2:
μ1: ≠μ2 <μ2 >μ2
Pooled: [ ] Yes
Calculate Draw
    
```

# Confidence Intervals: A Four-Step Process

	Proportions	Means
<b>S T A T E</b>	$p_1 - p_2 =$ true difference in <b>proportions</b> of...  Confidence Level = _____	$\mu_1 - \mu_2 =$ true difference in <b>means</b> of...  Confidence Level = _____
<b>P L A N</b>	<p><u>Name</u>: Two Sample <math>z</math> interval for <math>p_1 - p_2</math></p> <p><u>Conditions</u>:</p> 1) Random samples 2) Independent if: $n_1 \leq 10\%N$ $n_2 \leq 10\%N$ 3) Approximately Normal if: $n_1 \hat{p}_1 \geq 10$ and $n_1(1 - \hat{p}_1) \geq 10$ $n_2 \hat{p}_2 \geq 10$ and $n_2(1 - \hat{p}_2) \geq 10$	<p><u>Name</u>: Two Sample <math>t</math> interval for <math>\mu_1 - \mu_2</math></p> <p><u>Conditions</u>:</p> 1) Random samples 2) Independent if: $n_1 \leq 10\%N$ $n_2 \leq 10\%N$ 3) Approximately Normal if: ➤ $n_1 \geq 30$ and $n_2 \geq 30$ ➤ Sample shows no strong skewness and outliers
<b>D O</b>	$(\hat{p}_1 - \hat{p}_2) \pm z^* \sqrt{\frac{\hat{p}_1(1 - \hat{p}_1)}{n_1} + \frac{\hat{p}_2(1 - \hat{p}_2)}{n_2}}$ OR 2-PropZint (B)	$(\bar{x}_1 - \bar{x}_2) \pm t^* \sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}}$ OR 2-SampTInt (0)
<b>C O N C L U D E</b>	We are _____% confident that the interval from _____ to _____ captures the true difference in <b>proportions</b> of...	We are _____% confident that the interval from _____ to _____ captures the true difference in <b>means</b> of...

# Significance Tests: A Four-Step Process

	Proportions	Means
<b>S T A T E</b>	$H_0: p_1 = p_2$ $H_a: \neq, <, \text{ or } >$  $p_1 = \text{true proportion of...}$ $p_2 = \text{true proportion of...}$	$H_0: \mu_1 = \mu_2$ $H_a: \mu \neq, <, \text{ or } >$  $\mu_1 = \text{true mean of...}$ $\mu_2 = \text{true mean of...}$
<b>P L A N</b>	<p style="text-align: center;"><u>Name:</u> Two Sample <math>z</math> Test for <math>p_1 - p_2</math></p> <p><u>Conditions:</u></p> <ol style="list-style-type: none"> <li>1) Random samples</li> <li>2) Independent if:  <math>n_1 \leq 10\%N</math>  <math>n_2 \leq 10\%N</math></li> <li>3) Approximately Normal if:  <math>n_1 \hat{p}_1 \geq 10</math> and <math>n_1(1 - \hat{p}_1) \geq 10</math>  <math>n_2 \hat{p}_2 \geq 10</math> and <math>n_2(1 - \hat{p}_2) \geq 10</math></li> </ol>	<p style="text-align: center;"><u>Name:</u> Two Sample <math>t</math> Test for <math>\mu_1 - \mu_2</math></p> <p><u>Conditions:</u></p> <ol style="list-style-type: none"> <li>1) Random samples</li> <li>2) Independent if:  <math>n_1 \leq 10\%N</math>  <math>n_2 \leq 10\%N</math></li> <li>3) Approximately Normal if: <ul style="list-style-type: none"> <li>➤ <math>n_1 \geq 30</math> and <math>n_2 \geq 30</math></li> <li>➤ Sample shows no strong skewness and outliers</li> </ul> </li> </ol>
<b>D O</b>	2-propZTest (6)	2-SampTTest (4)
<b>C O N C L U D E</b>	<p>p-value <math>\leq \alpha</math>, reject <math>H_0</math>. We have convincing evidence that...(H<sub>a</sub>)</p> <p>p-value <math>&gt; \alpha</math>, fail to reject <math>H_0</math>. We do not have convincing evidence...(H<sub>a</sub>)</p>	<p>p-value <math>\leq \alpha</math>, reject <math>H_0</math>. We have convincing evidence that...(H<sub>a</sub>)</p> <p>p-value <math>&gt; \alpha</math>, fail to reject <math>H_0</math>. We do not have convincing evidence that...(H<sub>a</sub>)</p>