

Interpret Standard Deviation	Outlier Rule
Linear Transformations	Describe the Distribution OR Compare the Distributions
SOCS	<u>Using Normalcdf and Invnorm</u> (Calculator Tips)
Interpret a z-score	What is an Outlier?
Interpret LSRL Slope “ b ”	Interpret LSRL y -intercept “ a ”

<p>Upper Bound = $Q_3 + 1.5(IQR)$</p> <p>Lower Bound = $Q_1 - 1.5(IQR)$</p> <p>$IQR = Q_3 - Q_1$</p>	<p>Standard Deviation measures spread by giving the “typical” or “average” distance that the observations (context) are away from their (context) mean</p>
<p>SOCS! Shape, Outliers, Center, Spread Only discuss outliers if there are obviously outliers present. Be sure to address SCS in context!</p> <p>If it says “Compare” YOU MUST USE <u>comparison phrases</u> like “is greater than” or “is less than” for Center & Spread</p>	<p>Adding “a” to every member of a data set adds “a” to the measures of position, but does not change the measures of spread or the shape.</p> <p>Multiplying every member of a data set by “b” multiplies the measures of position by “b” and multiplies most measures of spread by b , but does not change the shape.</p>
<p>Normalcdf (min, max, mean, standard deviation)</p> <p>Invnorm (area to the left as a decimal, mean, standard deviation)</p>	<p>Shape – Skewed Left (Mean < Median) Skewed Right (Mean > Median) Fairly Symmetric (Mean \approx Median)</p> <p>Outliers – Discuss them if there are obvious ones Center – Mean or Median Spread – Range, <i>IQR</i>, or Standard Deviation</p> <p>Note: Also be on the lookout for gaps, clusters or other unusual features of the data set. Make Observations!</p>
<p>When given 1 variable data: An outlier is any value that falls more than $1.5(IQR)$ above Q_3 or below Q_1</p> <p>Regression Outlier: Any value that falls outside the pattern of the rest of the data.</p>	$z = \frac{\text{value} - \text{mean}}{\text{standard deviation}}$ <p>A z-score describes how many standard deviations a value or statistic (x, \bar{x}, \hat{p}, etc.) falls away from the mean of the distribution and in what direction. The further the z-score is away from zero the more “surprising” the value of the statistic is.</p>
<p>When the x variable (context) is zero, the y variable (context) is estimated to be <u>put value here</u>.</p>	<p>For every one unit change in the x variable (context) the y variable (context) is predicted to increase/decrease by _____ units (context).</p>

Interpret r^2	Interpret r
Interpret LSRL “ SE_b ”	Interpret LSRL “ s ”
Interpret LSRL “ \hat{y} ”	Extrapolation
Interpreting a Residual Plot	What is a Residual?
Sampling Techniques	Experimental Designs

Correlation measures the **strength** and **direction** of the **linear relationship** between x and y .

- r is always between -1 and 1 .
- Close to zero = very weak,
- Close to 1 or -1 = stronger
- Exactly 1 or -1 = perfectly straight line
- Positive r = positive correlation
- Negative r = negative correlation

___% of the variation in y (**context**) is accounted for by the LSRL of y (**context**) on x (**context**).

Or

___% of the variation in y (**context**) is accounted for by using the linear regression model with x (**context**) as the explanatory variable.

$s = \underline{\hspace{1cm}}$ is the standard deviation of the residuals.

It measures the typical distance between the actual y -values (**context**) and their predicted y -values (**context**)

SE_b measures the standard deviation of the estimated slope for predicting the y variable (**context**) from the x variable (**context**).

SE_b measures how far the estimated slope will be from the true slope, on average.

Using a LSRL to predict outside the domain of the explanatory variable.

(Can lead to ridiculous conclusions if the current linear trend does not continue)

\hat{y} is the “estimated” or “predicted” y -value (**context**) for a given x -value (**context**)

$$\text{Residual} = y - \hat{y}$$

A residual measures the difference between the actual (observed) y -value in a scatterplot and the y -value that is predicted by the LSRL using its corresponding x value.

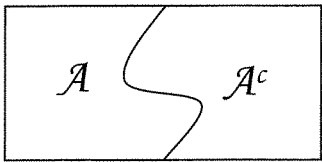
In the calculator: $L_3 = L_2 - Y_1(L_1)$

1. **Is there a curved pattern?** If so, a linear model may not be appropriate.
2. **Are the residuals small in size?** If so, predictions using the linear model will be fairly precise.
3. **Is there increasing (or decreasing) spread?** If so, predictions for larger (smaller) values of x will be more variable.

1. **CRD** (Completely Randomized Design) – All experimental units are allocated at random among all treatments
2. **RBD** (Randomized Block Design) – Experimental units are put into homogeneous blocks. The random assignment of the units to the treatments is carried out separately within each block.
3. **Matched Pairs** – A form of blocking in which each subject receives both treatments in a random order or the subjects are matched in pairs as closely as possible and one subject in each pair receives each treatment, determined at random.

1. **SRS** – Number the entire population, draw numbers from a hat (every set of n individuals has equal chance of selection)
2. **Stratified** – Split the population into homogeneous groups, select an SRS from each group.
3. **Cluster** – Split the population into heterogeneous groups called clusters, and randomly select whole clusters for the sample. Ex. Choosing a carton of eggs actually chooses a cluster (group) of 12 eggs.
4. **Census** – An attempt to reach the entire population
5. **Convenience** – Selects individuals easiest to reach
6. **Voluntary Response** – People choose themselves by responding to a general appeal.

<p>Goal of Blocking Benefit of Blocking</p>	<p>Advantage of using a Stratified Random Sample Over an SRS</p>
<p>Experiment Or Observational Study?</p>	<p>Does ___ CAUSE ___?</p>
<p>SRS</p>	<p>Why use a control group?</p>
<p>Complementary Events</p>	<p>$P(\text{at least one})$</p>
<p>Two Events are Independent If...</p>	<p>Interpreting Probability</p>

<p>Stratified random sampling guarantees that each of the strata will be represented. When strata are chosen properly, a stratified random sample will produce better (less variable/more precise) information than an SRS of the same size.</p>	<p>The goal of blocking is to create groups of homogeneous experimental units.</p> <p>The benefit of blocking is the reduction of the effect of variation within the experimental units. (context)</p>
<p>Association is NOT Causation!</p> <p>An observed association, no matter how strong, is not evidence of causation. Only a well-designed, controlled experiment can lead to conclusions of cause and effect.</p>	<p>A study is an experiment ONLY if researchers IMPOSE a treatment upon the experimental units.</p> <p>In an observational study researchers make no attempt to influence the results.</p>
<p>A control group gives the researchers a comparison group to be used to evaluate the effectiveness of the treatment(s). (context) (gauge the effect of the treatment compared to no treatment at all)</p>	<p>An SRS (simple random sample) is a sample taken in such a way that every set of n individuals has an equal chance to be the sample actually selected.</p>
<p>$P(\text{at least one}) = 1 - P(\text{none})$</p> <p>Ex. $P(\text{at least one 6 in three rolls}) = \underline{\hspace{2cm}}$ $P(\text{Get at least one six}) = 1 - P(\text{No Sixes})$ $= 1 - (5/6)^3$ $= 0.4213$</p>	<p>Two mutually exclusive events whose union is the sample space.</p>  <p>Ex: Rain/Not Rain, Draw at least one heart / Draw NO hearts</p>
<p>The probability of any outcome of a random phenomenon is the proportion of times the outcome would occur in a very long series of repetitions. Probability is a long-term relative frequency.</p>	<p>$P(B) = P(B A)$ Or $P(B) = P(B A^c)$</p> <p>Meaning: Knowing that Event A has occurred (or not occurred) doesn't change the probability that event B occurs.</p>

<p>Interpreting Expected Value/Mean</p>	<p>Mean and Standard Deviation of a Discrete Random Variable</p>
<p>Mean and Standard Deviation of a Difference of Two Random Variables</p>	<p>Mean and Standard Deviation of a Sum of Two Random Variables</p>
<p>Binomial Distribution (Conditions)</p>	<p>Geometric Distribution (Conditions)</p>
<p>Binomial Distribution (Calculator Usage)</p>	<p>Mean and Standard Deviation Of a Binomial Random Variable</p>
<p>Why Large Samples Give More Trustworthy Results... (When collected appropriately)</p>	<p>The Sampling Distribution of the Sample Mean (Central Limit Theorem)</p>

Also on the formula sheet!

Mean (Expected Value):

$$\mu_x = \sum x_i p_i$$

(Multiply & add across the table)

Standard Deviation:

$$\sigma_x = \sqrt{\sum (x_i - \mu_x)^2 p_i}$$

Square root of the sum of (Each x value – the mean)²(its probability)

The mean/expected value of a random variable is the long-run average outcome of a random phenomenon carried out a very large number of times.

Mean of a Sum of 2 RV's:

$$\mu_{X+Y} = \mu_X + \mu_Y$$

Stdev of a Sum of 2 Independent RV's:

$$\sigma_{X+Y} = \sqrt{\sigma_X^2 + \sigma_Y^2}$$

Stdev of a Sum 2 Dependent RV's:

Cannot be determined because it depends on how strongly they are correlated.

Mean of a Difference of 2 RV's:

$$\mu_{X-Y} = \mu_X - \mu_Y$$

Stdev of a Difference of 2 Indep RV's:

$$\sigma_{X-Y} = \sqrt{\sigma_X^2 + \sigma_Y^2}$$

Stdev of a Difference of 2 Dependent RV's:

Cannot be determined because it depends on how strongly they are correlated.

1. **B**inary? Trials can be classified as success/failure
2. **I**ndependent? Trials must be independent.
3. **T**rials? The goal is to count the number of trials until the first success occurs
4. **S**uccess? The probability of success (p) must be the same for each trial.

1. **B**inary? Trials can be classified as success/failure
2. **I**ndependent? Trials must be independent.
3. **N**umber? The number of trials (n) must be fixed in advance
4. **S**uccess? The probability of success (p) must be the same for each trial.

Also on the formula sheet!

Mean: $\mu_x = np$

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

- Exactly 5: $P(X = 5) = \text{Binompdf}(n, p, 5)$
- At Most 5: $P(X \leq 5) = \text{Binomcdf}(n, p, 5)$
- Less Than 5: $P(X < 5) = \text{Binomcdf}(n, p, 4)$
- At Least 5: $P(X \geq 5) = 1 - \text{Binomcdf}(n, p, 4)$
- More Than 5: $P(X > 5) = 1 - \text{Binomcdf}(n, p, 5)$

Remember to define X , n , and p !

1. If the population distribution is Normal the sampling distribution will also be Normal with the same mean as the population. Additionally, as n increases the sampling distribution's standard deviation will decrease
2. If the population distribution is not Normal the sampling distribution will become more and more Normal as n increases. The sampling distribution will have the same mean as the population and as n increases the sampling distribution's standard deviation will decrease.

When collected appropriately, large samples yield more precise results than small samples because in a large sample the values of the sample statistic tend to be closer to the true population parameter.

<p>Unbiased Estimator</p>	<p>Bias</p>
<p>Explain a <i>P</i>-value</p>	<p>Can we generalize the results to the population of interest?</p>
<p>Finding the Sample Size (For a given margin of error)</p>	<p>Carrying out a Two-Sided Test from a Confidence Interval</p>
<p><u>4-Step Process</u> Confidence Intervals</p>	<p><u>4-Step Process</u> Significance Tests</p>
<p>Interpreting a Confidence Interval (Not a Confidence Level)</p>	<p>Interpreting a Confidence Level (The Meaning of 95% Confident)</p>

<p>The systematic favoring of certain outcomes due to flawed sample selection, poor question wording, undercoverage, nonresponse, etc.</p> <p>Bias deals with the center of a sampling distribution being “off”!</p>	<p>The data is collected in such a way that there is no systematic tendency to overestimate or underestimate the true value of the population parameter.</p> <p>(The mean of the sampling distribution equals the true value of the parameter being estimated)</p>
<p>Yes, if:</p> <p>A large random sample was taken from the same population we hope to draw conclusions about.</p>	<p>Assuming that the null is true (context) the <i>P</i>-value measures the chance of observing a statistic (or difference in statistics) (context) as large as or larger than the one actually observed.</p>
<p>We do/(do not) have enough evidence to reject $H_0: \mu = ?$ in favor of $H_a: \mu \neq ?$ at the $\alpha = 0.05$ level because $?$ falls outside/(inside) the 95% CI.</p> <p>$\alpha = 1 - \text{confidence level}$</p>	<p>For one mean: $m = z^* \left(\frac{\sigma}{\sqrt{n}} \right)$</p> <p>For one proportion: $m = z^* \sqrt{\frac{p(1-p)}{n}}$</p> <p>If an estimation of p is not given, use 0.5 for p. Solve for n.</p>
<p>STATE: What hypotheses do you want to test, and at what significance level? Define any parameters you use.</p> <p>PLAN: Choose the appropriate inference method. Check conditions.</p> <p>DO: If the conditions are met, perform calculations. Compute the test statistic and find the <i>P</i>-value.</p> <p>CONCLUDE: Interpret the result of your test in the context of the problem.</p>	<p>STATE: What parameter do you want to estimate, and at what confidence level?</p> <p>PLAN: Choose the appropriate inference method. Check conditions.</p> <p>DO: If the conditions are met, perform calculations.</p> <p>CONCLUDE: Interpret your interval in the context of the problem.</p>
<p>Intervals produced with this <u>method</u> will capture the true population _____ in about 95% of all possible samples of this same size from this same population.</p>	<p>I am ___% confident that the interval from ___ to ___ captures the true ____.</p>

<p><u>Paired t-test</u> Phrasing Hints, H_0 and H_a, Conclusion</p>	<p><u>Two Sample t-test</u> Phrasing Hints, H_0 and H_a, Conclusion</p>
<p>Type I Error, Type II Error, & Power</p>	<p>Factors that Affect Power</p>
<p><u>Inference for Means</u> (Conditions)</p>	<p><u>Inference for Proportions</u> (Conditions)</p>
<p>Types of Chi-Square Tests</p>	<p><u>Chi-Square Tests</u> df and Expected Counts</p>
<p><u>Inference for Counts</u> (Chi-Squared Tests) (Conditions)</p>	<p><u>Inference for Regression</u> (Conditions)</p>

<p>Key Phrase: DIFFERENCE IN THE MEANS</p> <p>$H_0: \mu_1 - \mu_2 = 0$ OR $\mu_1 = \mu_2$</p> <p>$H_a: \mu_1 - \mu_2 < 0, > 0, \neq 0$</p> <p>$\mu_1 - \mu_2 =$ The difference between the mean ___ for all ___ and the mean ___ for all ___.</p> <p>We do/(do not) have enough evidence at the 0.05 level to conclude that the difference between the mean ___ for all ___ and the mean ___ for all ___ is ___.</p>	<p>Key Phrase: MEAN DIFFERENCE</p> <p>$H_0: \mu_{Diff} = 0$</p> <p>$H_a: \mu_{Diff} < 0, > 0, \neq 0$</p> <p>$\mu_{Diff} =$ The mean difference in ___ for all ___.</p> <p>We do/(do not) have enough evidence at the 0.05 level to conclude that the mean difference in ___ for all ___ is ___.</p>
<ol style="list-style-type: none"> Sample Size: To increase power, increase sample size. Increase α: A 5% test of significance will have a greater chance of rejecting the null than a 1% test. Consider an alternative that is farther away from μ_0: Values of μ that are in H_a, but lie close to the hypothesized value are harder to detect than values of μ that are far from μ_0. 	<ol style="list-style-type: none"> Type I Error: Rejecting H_0 when H_0 is actually true. (Ex. Convicting an innocent person) Type II Error: Failing to (II) reject H_0 when H_0 should be rejected. (Ex. Letting a guilty person go free) Power: Probability of rejecting H_0 when H_0 should be rejected. (Rejecting Correctly)
<p>Random: Data from a random sample(s) or randomized experiment</p> <p>Normal: At least 10 successes and failures (in both groups, for a two sample problem)</p> <p>Independent: Independent observations and independent samples/groups; 10% condition if sampling without replacement</p>	<p>Random: Data from a random sample(s) or randomized experiment</p> <p>Normal: Population distribution is normal or large sample(s) ($n_1 \geq 30$ or $n_1 \geq 30$ and $n_2 \geq 30$)</p> <p>Independent: Independent observations and independent samples/groups; 10% condition if sampling without replacement</p>
<ol style="list-style-type: none"> Goodness of Fit: df = # of categories - 1 Expected Counts: Sample size times hypothesized proportion in each category. Homogeneity or Association/Independence: df = (# of rows - 1)(# of columns - 1) Expected Counts: $\frac{(\text{row total})(\text{column total})}{\text{table total}}$ 	<ol style="list-style-type: none"> Goodness of Fit: Use to test the distribution of one group or sample as compared to a hypothesized distribution. Homogeneity: Use when you you have a sample from 2 or more independent populations or 2 or more groups in an experiment. Each individual must be classified based upon a single categorical variable. Association/Independence: Use when you have a single sample from a single population. Individuals in the sample are classified by two categorical variables.
<p>Linear: True relationship between the variables is linear.</p> <p>Independent observations, 10% condition if sampling without replacement</p> <p>Normal: Responses vary normally around the regression line for all x-values</p> <p>Equal Variance around the regression line for all x-values</p> <p>Random: Data from a random sample or randomized experiment</p>	<p>Random: Data from a random sample(s) or randomized experiment</p> <p>Large Sample Size: All expected counts are at least 5.</p> <p>Independent: Independent observations and independent samples/groups; 10% condition if sampling without replacement</p>