





## **Correlations**

Correlations is the relationship between any two variables

They can be described in different ways:

**Pearson**—linear relationship between continuous variables

**Spearman Rho**—nonparametric rank correlation, describing



two variables as a monotonic function







## **Correlations in MATLAB**



**corr** — returns a matrix of pairwise correlations between columns



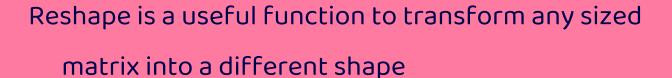
**corrcoef** — Returns the correlation between vectorized matrices

corr2 — returns correlation coefficient for matrices (i.e., one value for its 2-d inputs)



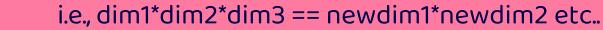


# Reminder: Reshape can help you



#### Reshape(X, [new dimensions])

Note that the new dimensions need to be consistent with the previous ones

















Student t-test allows you to test mean differences between normal distributions

There are many different **flavours** of t's

- one-sample vs two samples
- paired vs unpaired
- one tail vs two



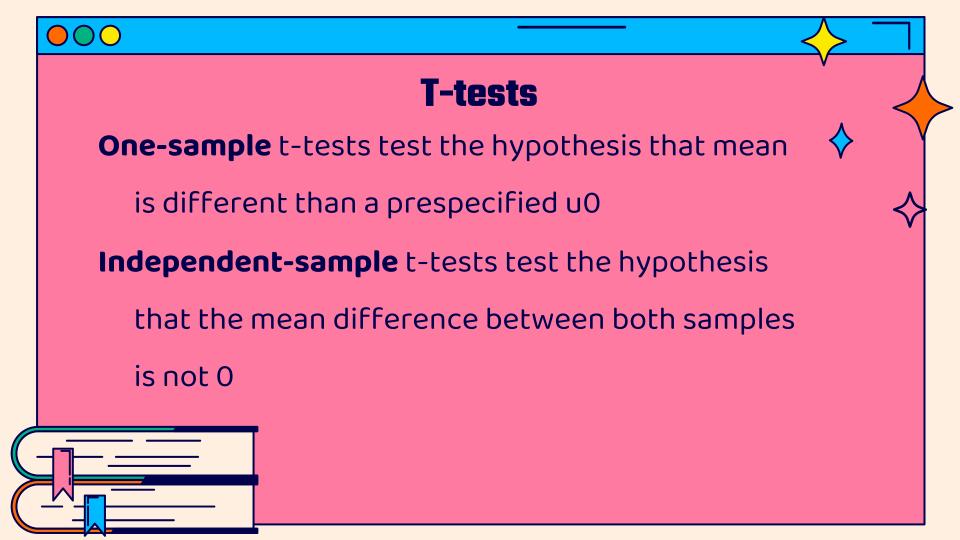


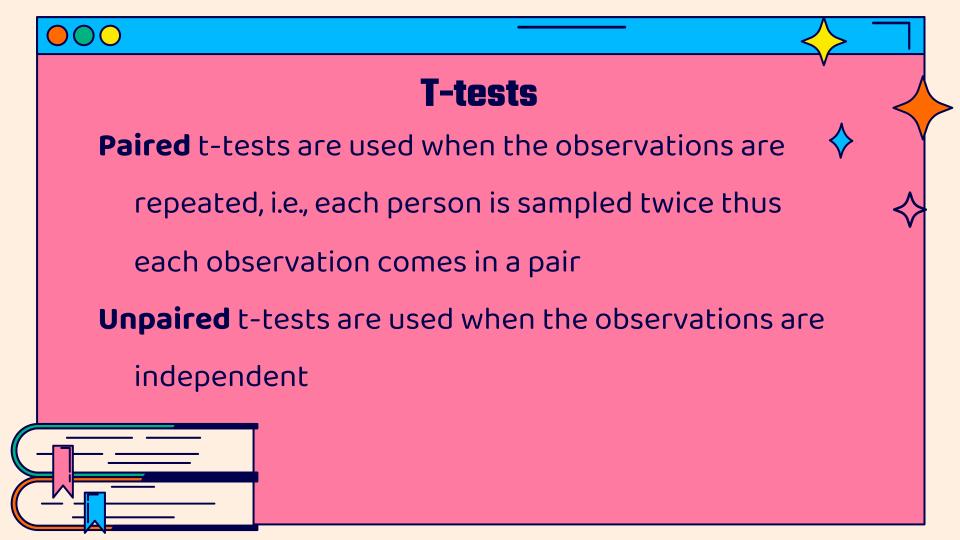


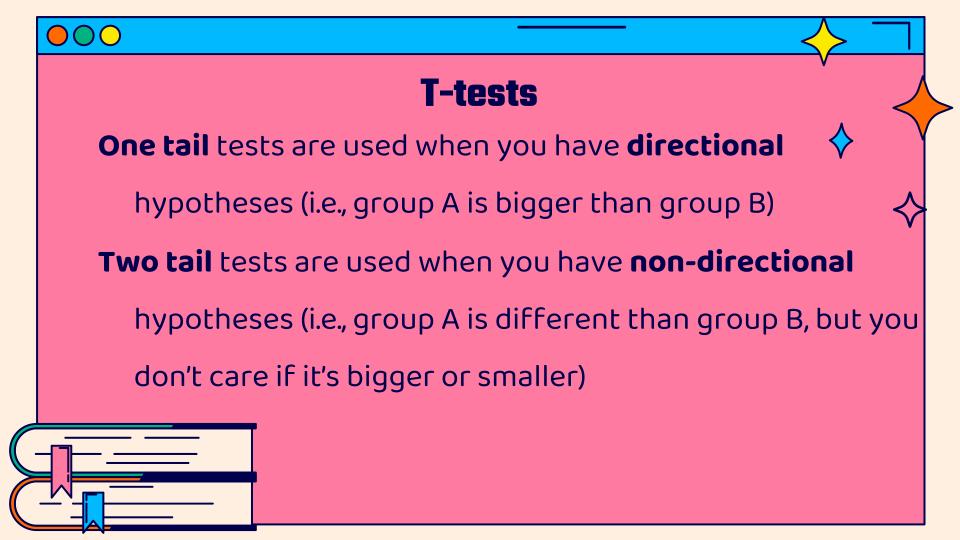


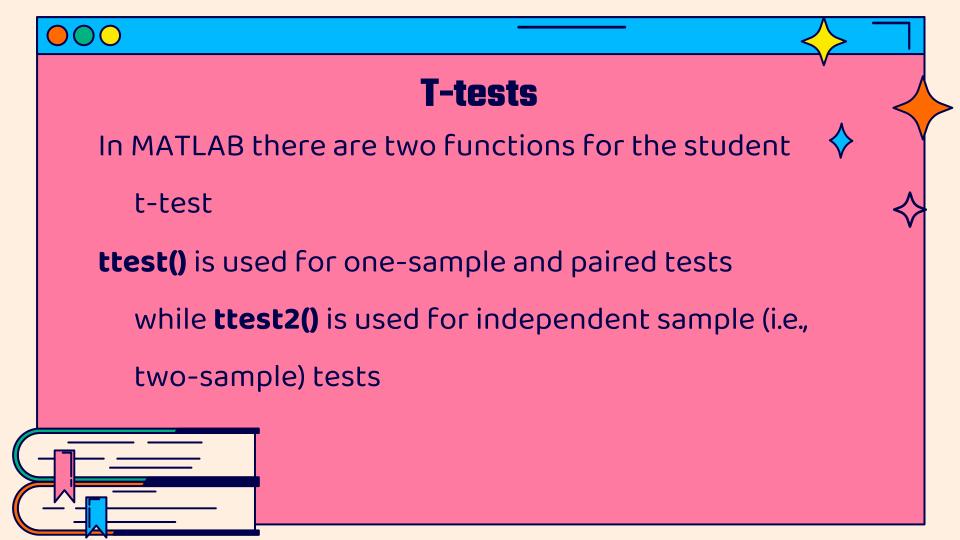
T-tests assume that your data come from a **normal** distribution and the observations are sampled **independently** from one another

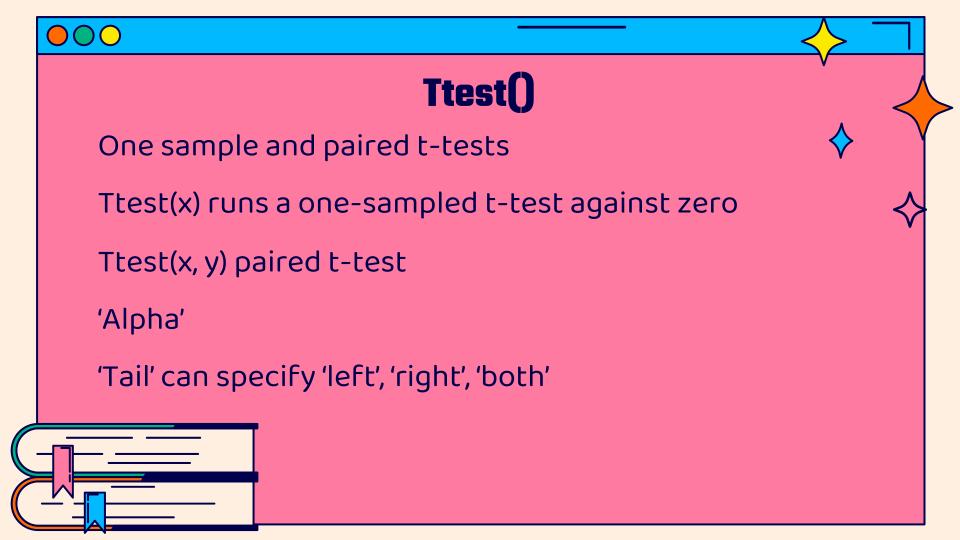
These assumptions apply for both paired and unpaired test

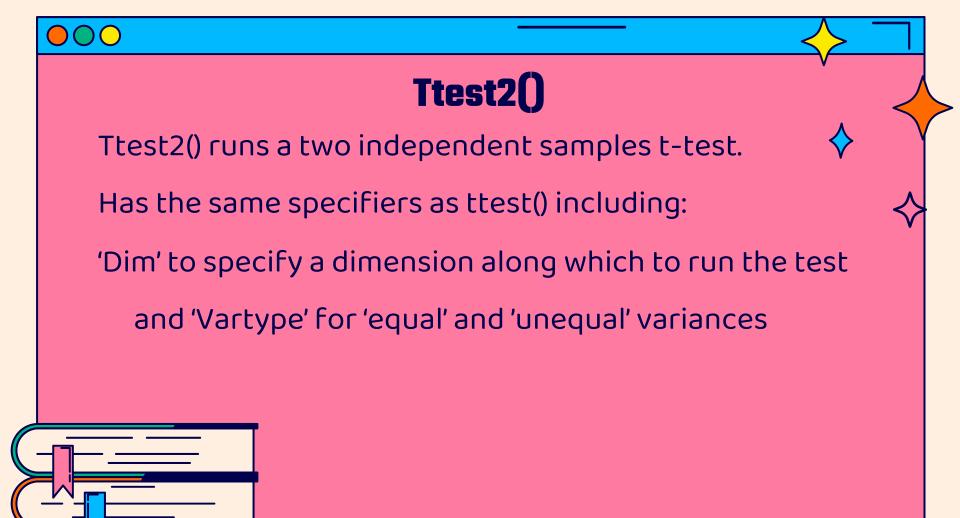


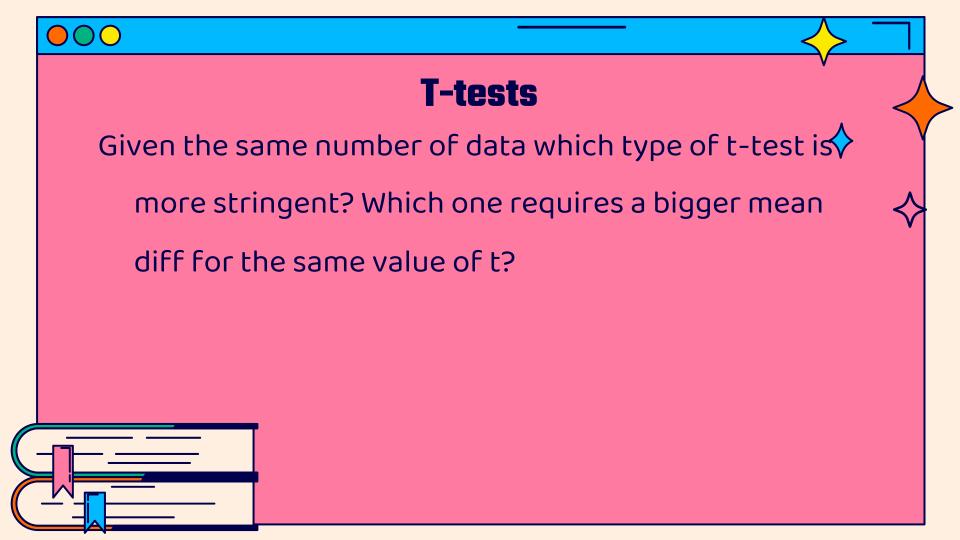














Let us assume we collect data from 20 people (paired)

and observe a mean diff of 3.75, and a sd of 1

Then the t value for a paired test would be:



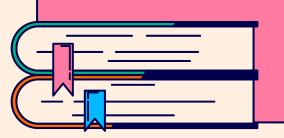
	$t = \frac{mean \ diff}{sd}$	- with df n — 1



Let us assume we collect data from 20 people (paired)

and observe a mean diff of 3.75, and a sd of 1

Then the t value for a paired test would be:



$$t = \frac{3.75}{\frac{1}{\sqrt{20}}}$$
 = with df 19



Let us assume we collect data from 20 people



(independent groups) and observe a mean diff of 3.75,



and a sd of 1 (assuming equal variance)



Then the t value for an unpaired test would be:

$$t = \frac{mean \ diff}{sp * \sqrt{\frac{1}{n1} + \frac{1}{n2}}} \text{ with } sp = \sqrt{\frac{(n1-1)*std1^2 + (n2-1)*std2^2}{n1 + n2 - 2}} \ df \ n1 + n2$$



Let us assume we collect data from 20 people



(independent groups) and observe a mean diff of 3.75,



and a sd of 1 (assuming equal variance)



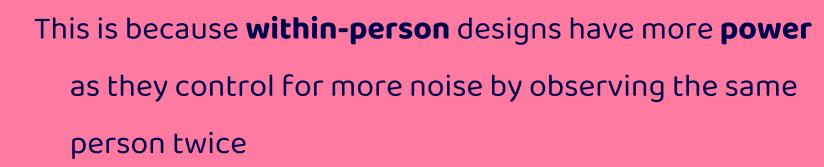
Then the t value for an unpaired test would be:

$$sp = \sqrt{\frac{(20-1)*1 + (20-1)*1}{20+20-2}} \qquad t = \frac{3.75}{sp*\sqrt{\frac{1}{20} + \frac{1}{20}}}.$$
 df 38





T value for paired is larger, in comparison to t values of unpaired or independent tests







## **Non-parametric tests**

In stats there are some tests that are **non-parametric,** by this statisticians mean that the test does not require assuming a specific model or distribution for your data

These are sometimes referred to as **non-distributional** tests

These tests are used when you do not want to assume a specific distribution (e.g., violation), or do not know the distribution of your data



## **Non-parametric t-tests**

Regular t-tests assume your data is normally distributed

There is a non-parametric equivalent of a t-test called the

permutation t-test. This test works on the premise that you

can observe a *null distribution* from your own data by

randomly permuting groups.

Reminder: a null distribution is the distribution when the null

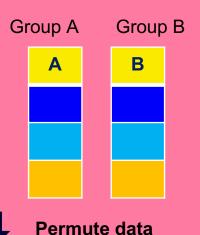
hypothesis is true

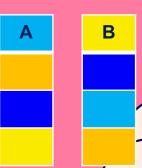


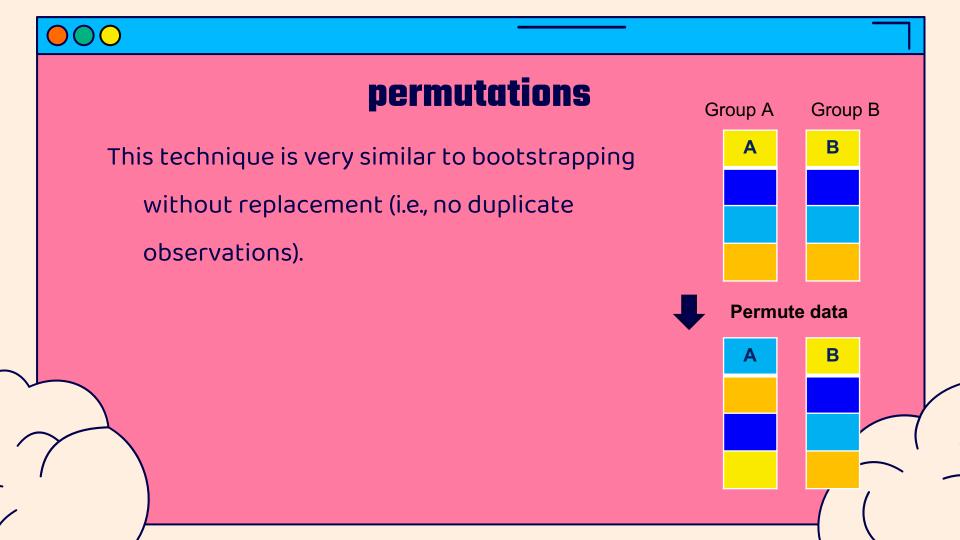
## permutations

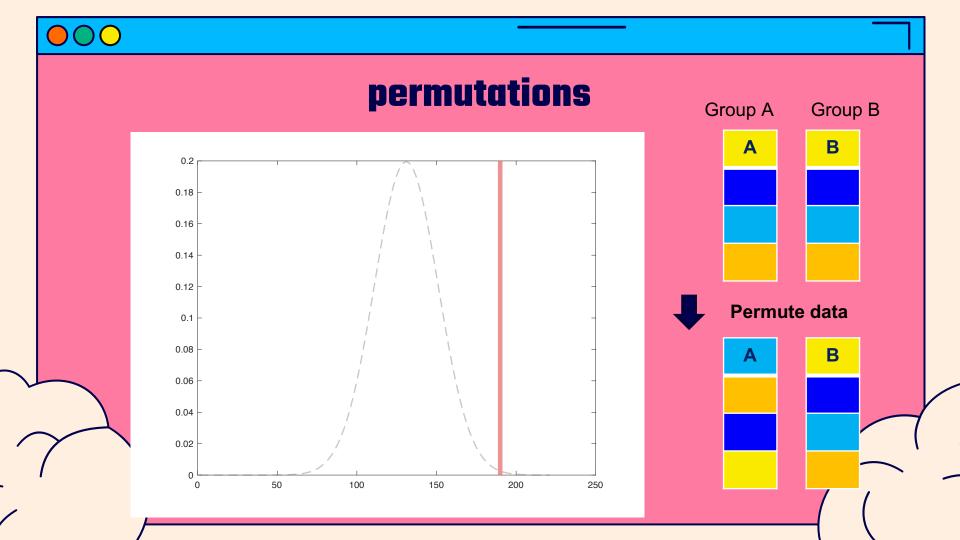
**Permutations** build a *null distribution* of data based on your observations under the assumption that randomly shuffling your data will void the effect of interest. Thus, you can measure how surprising the effect you observe is given your data based on the

computed null distribution











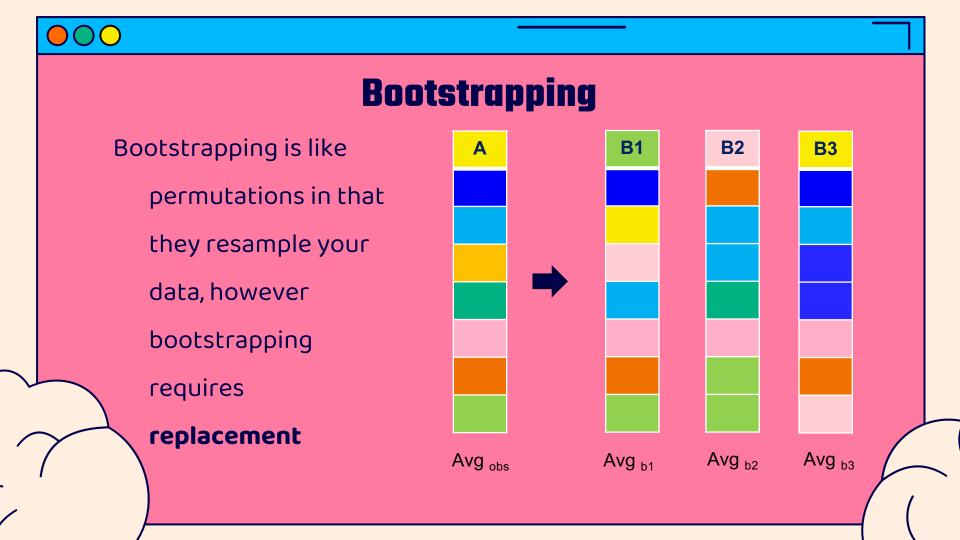
## **Bootstrapping**

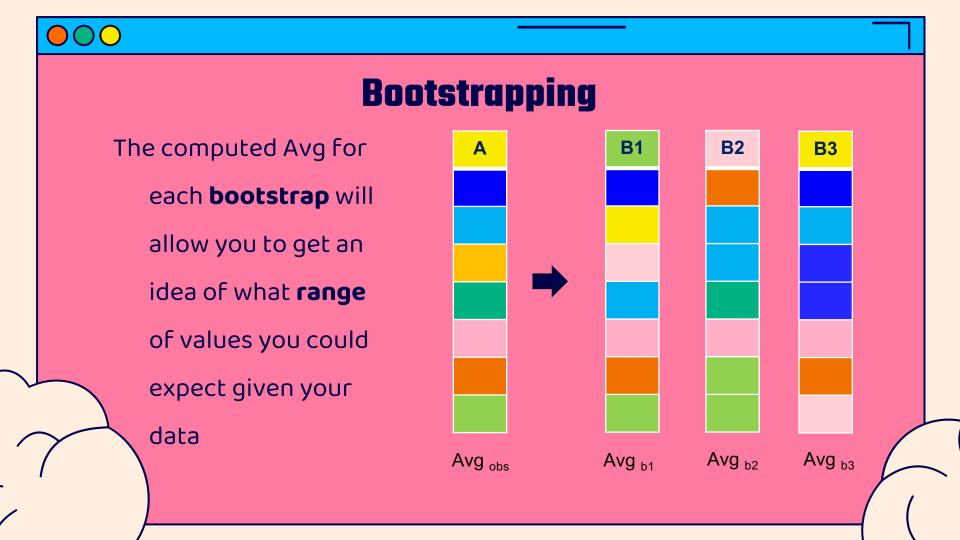
A very similar concept in statistics is the idea of **bootstrapping** to get a measure of uncertainty around an estimate (e.g., CI)

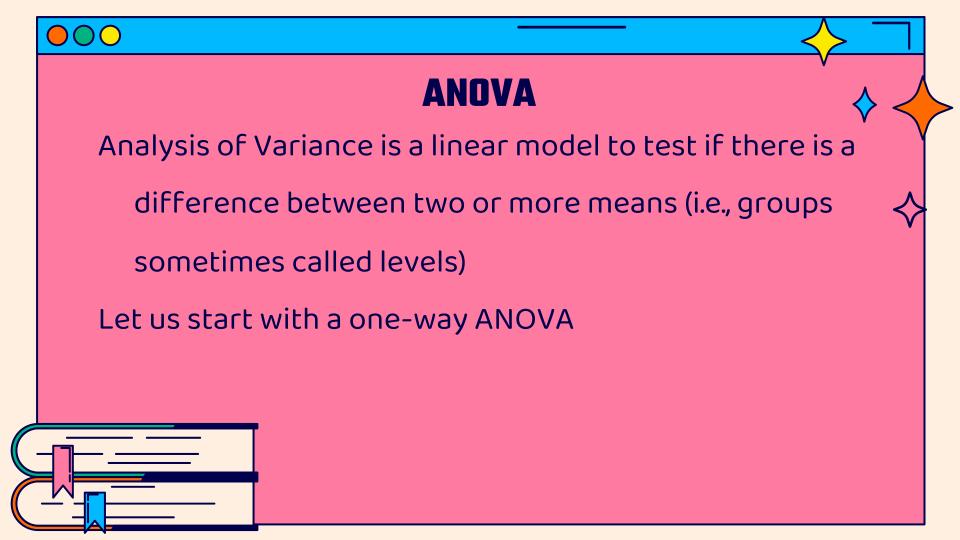
Bootstrapping is like permutations in that they resample your data, however bootstrapping requires **replacement** 

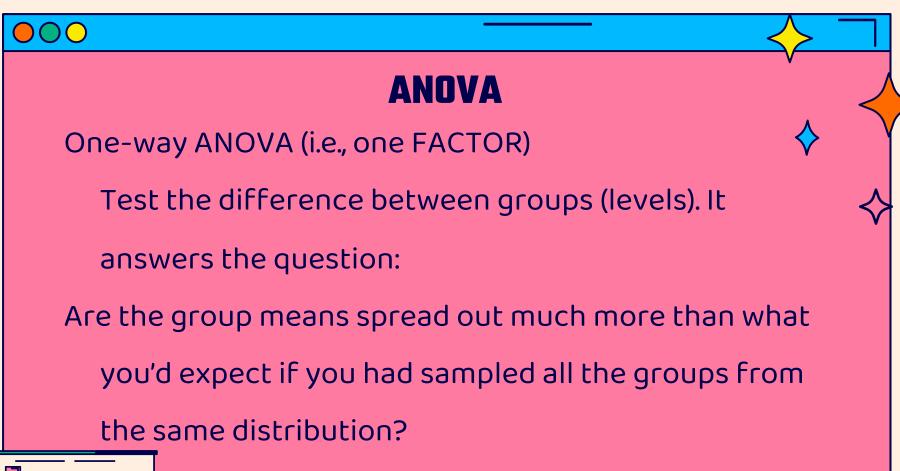
It is often used to calculate the error associated to an estimate, effect, or performance of an algorithm and allows you to

know if one given data point is driving the effect you see













## ANOVA

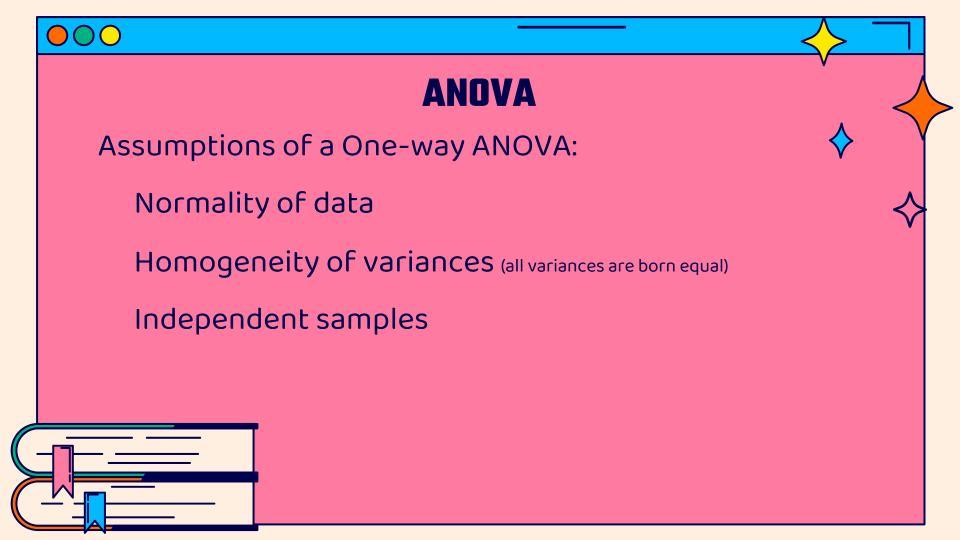


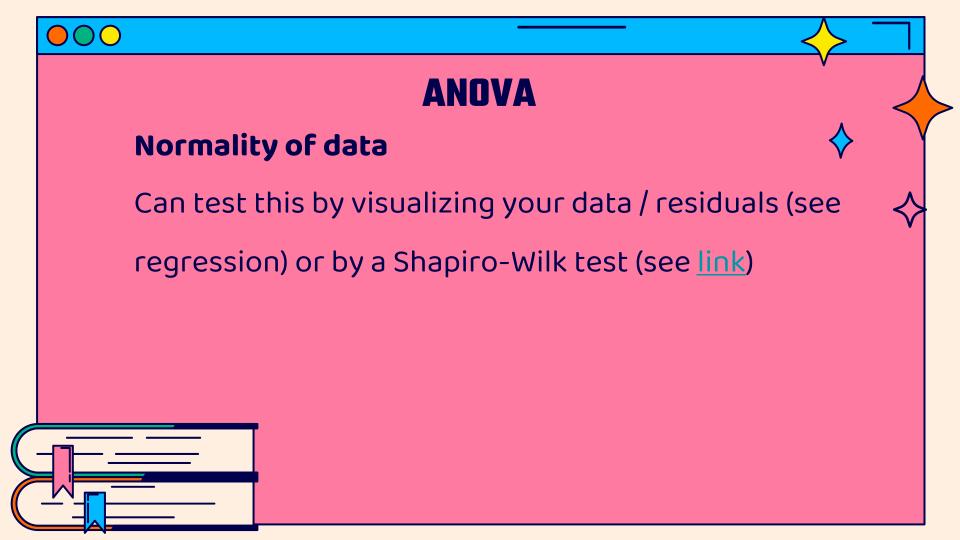
you'd expect if you had sampled all the groups from

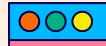
the same distribution?

$$H_0: a_1 = a_2 = a_3 = \dots a_n \mid H_1: a_1 \neq a_2 \neq a_3 \neq \dots a_n$$









### Homogeneity of variances



with a Bartlett test (see code)

This is a more 'serious' assumption to break when running an ANOVA







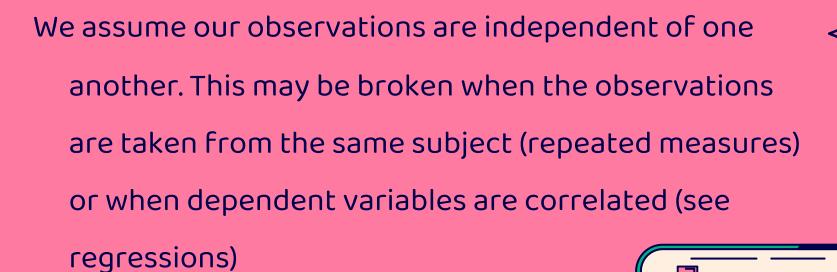






#### **ANOVA**

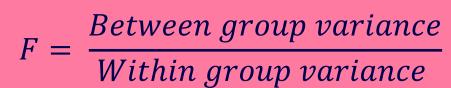
#### Independence





#### ANOVA

We test the significance of an ANOVA with an F test: 💠



We asses each using Sum of squares (i.e., SS) weighted by the number of observations (i.e., n) in each group







#### ANOVA

Between-group SS

How spread apart are your group means

Within-group SS

How spread apart is each distribution

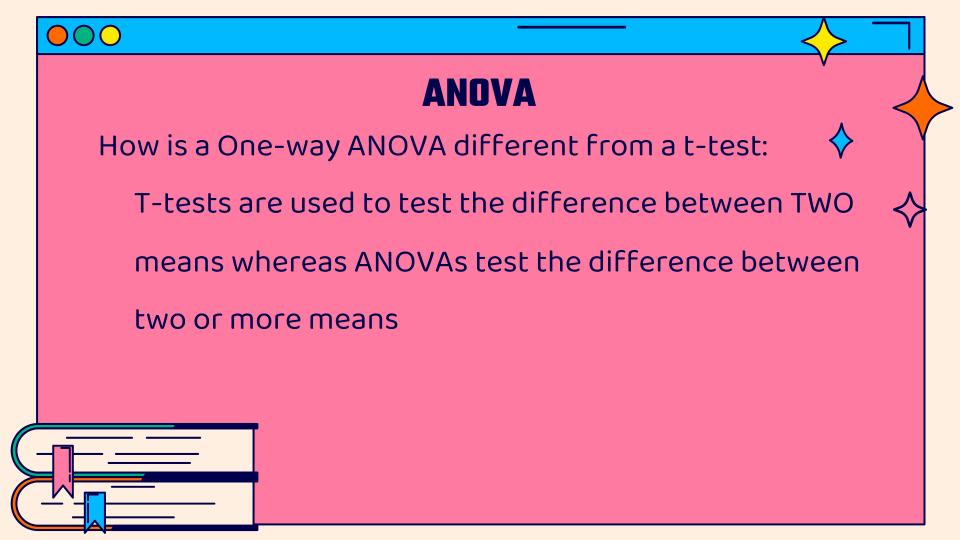


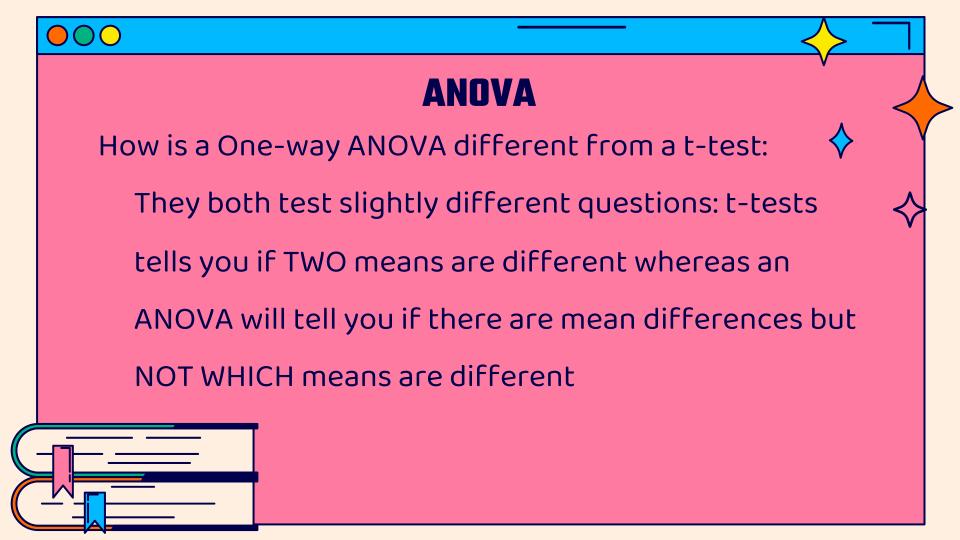


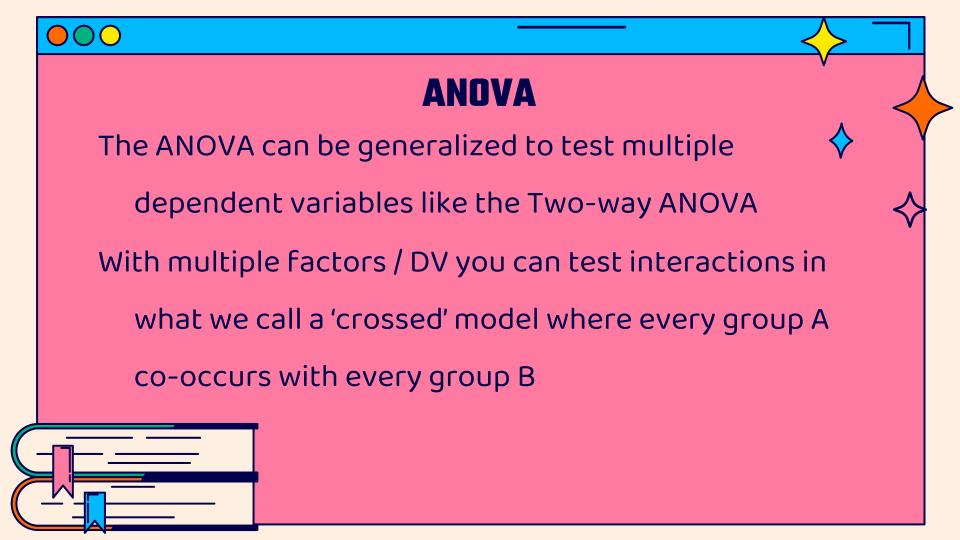


<sup>\*\*\*</sup> Note we always normalize these by their degrees of freedom

<sup>\*\*\*</sup> See code for visual depiction of an example with three means









## **Crossed Two-way ANOVA**





Factor / DV B

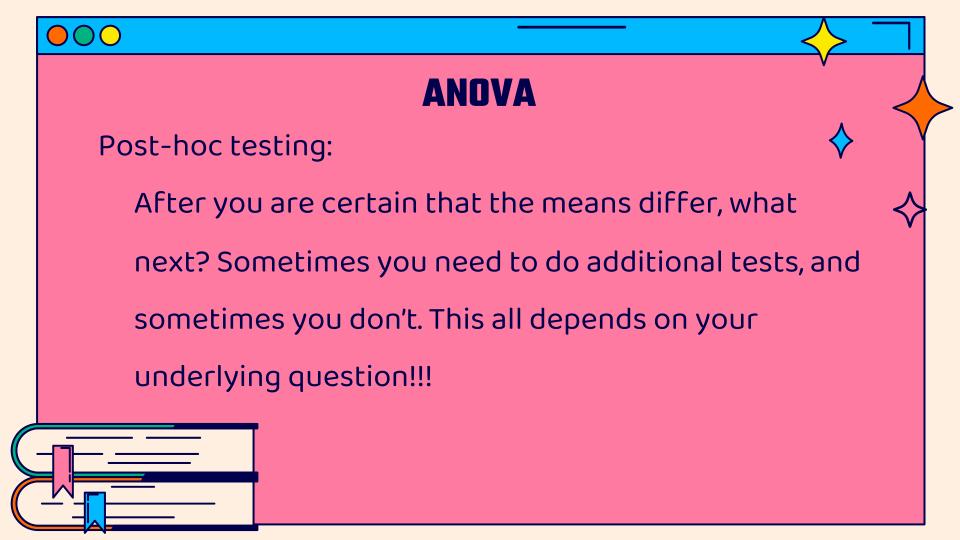
Group / Level 1A Group / Level 2A Group / Level 3A



**Group / Level 1B** 

**Group / Level 2B** 

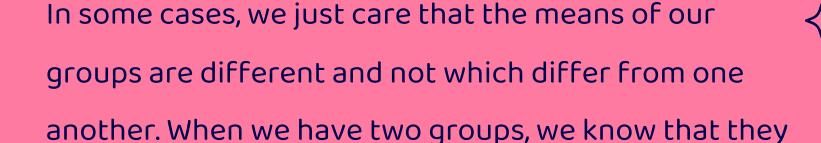
Mariah	Jlo	Lady Gaga
Halloween	Halloween	Halloween
Mariah	Jlo	Lady Gaga
Christmas	Christmas	Christmas





Post-hoc testing: NOT ALWAYS NECESSARY

are different, no additional test is needed.







#### **ANCOVA**

There are many iterations of an ANOVA including the  $\diamondsuit$  analysis of covariance!

Here we test the INDEPENDENT effect of dependent

variables (factors) regardless of COVARIATES of no

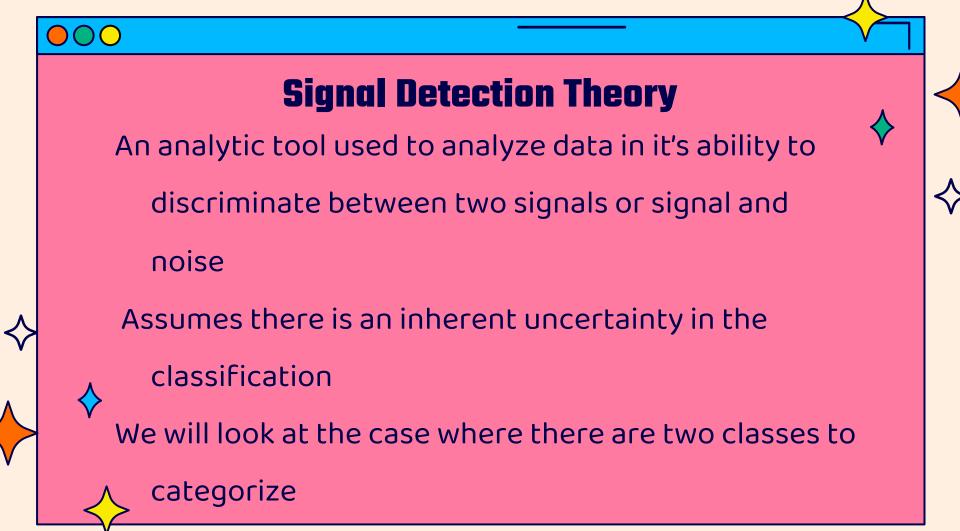
interest (referred to as nuisance variables)



\*\*\* we will cover this idea in more detail in the regression slides!











detection theory

The criterion: where you draw the boundary between signal and noise



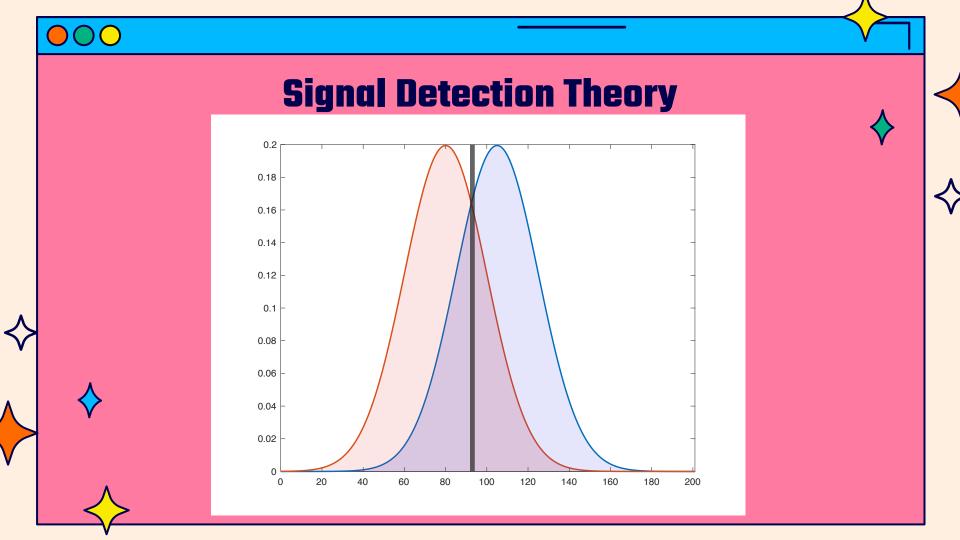
Sensitivity: one's ability to discriminate between signal and noise

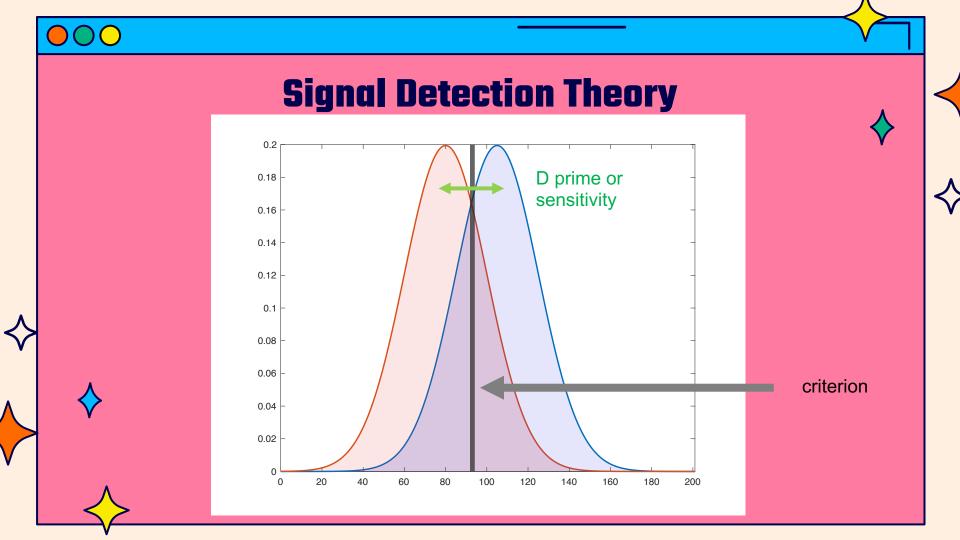






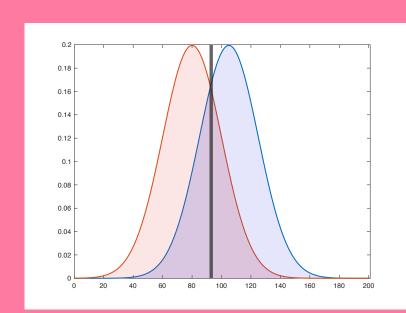












	Signal	Noise
Present	Hits	False Alarms
Absent	Misses	Correct Rejection







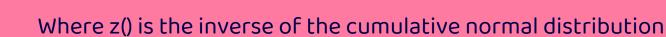






$$c = -\frac{1}{2}(z(Hits) + z(False\ Alarms))$$













What to do when you get values of 0 or 1 as

probabilities?

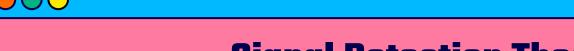
You cannot take the inverse cumulative normal distribution of 0 or 1 as it returns infinite values.



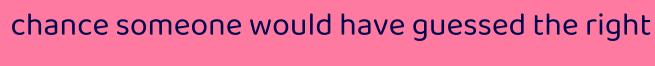








We assume that if we double the number of trials, by



answer i.e., if pHIT or pFA =0, then we correct to









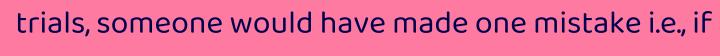




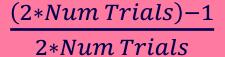




Similarry we assume that if we double the number of



pHIT or pFA =1, then correct with



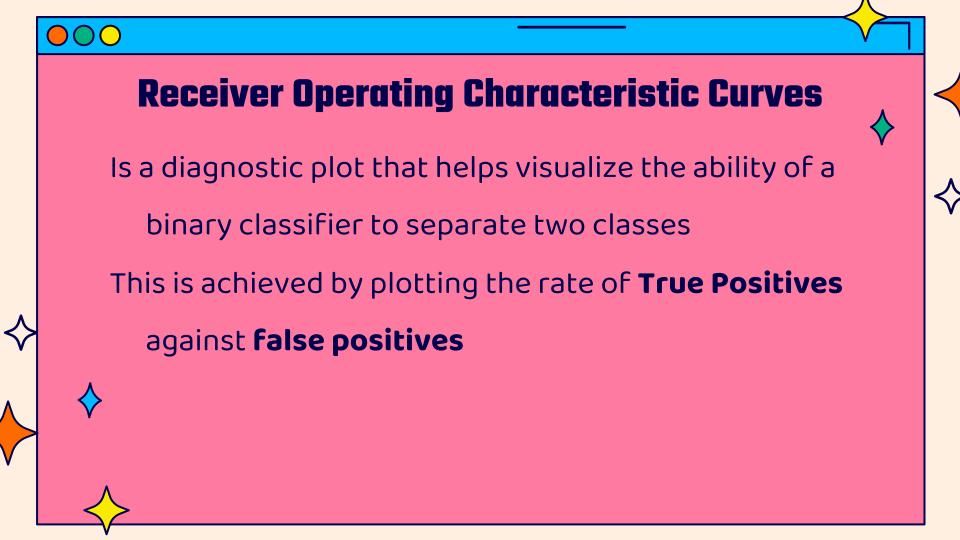




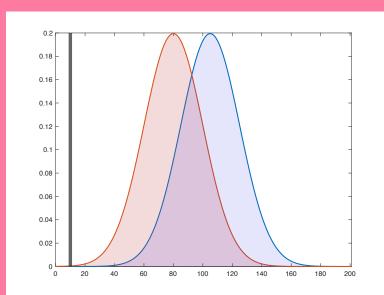


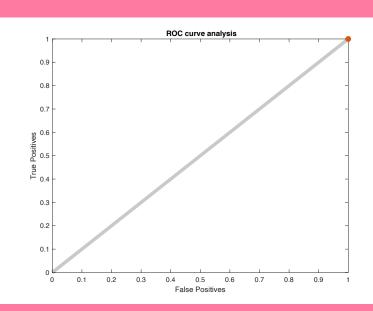










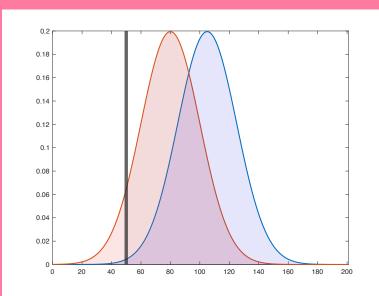


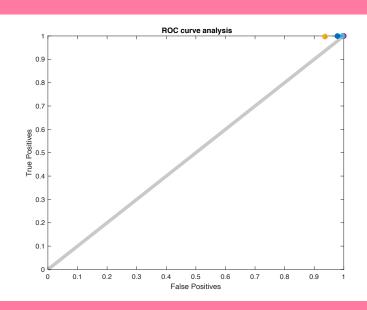










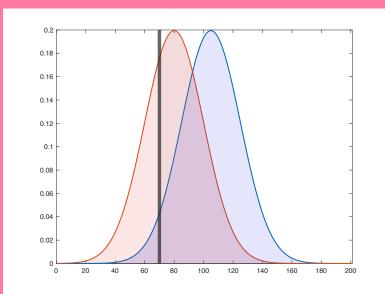


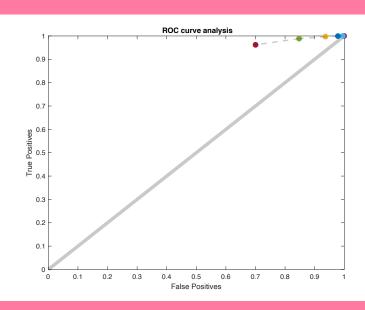










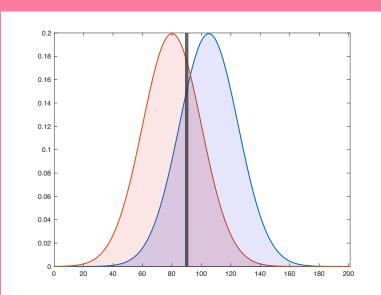


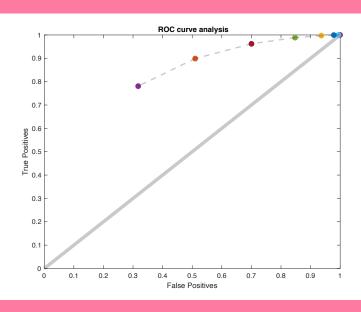










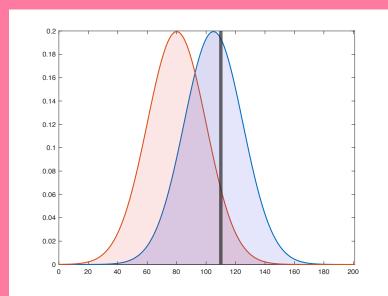


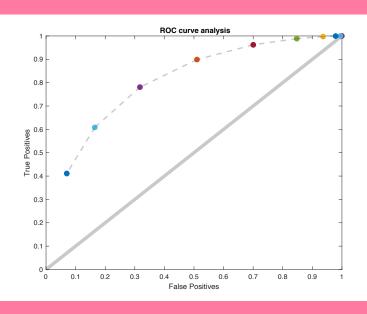






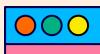


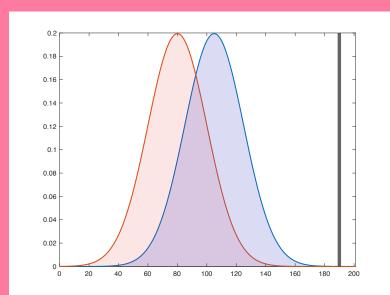


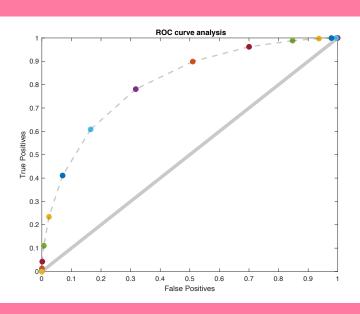








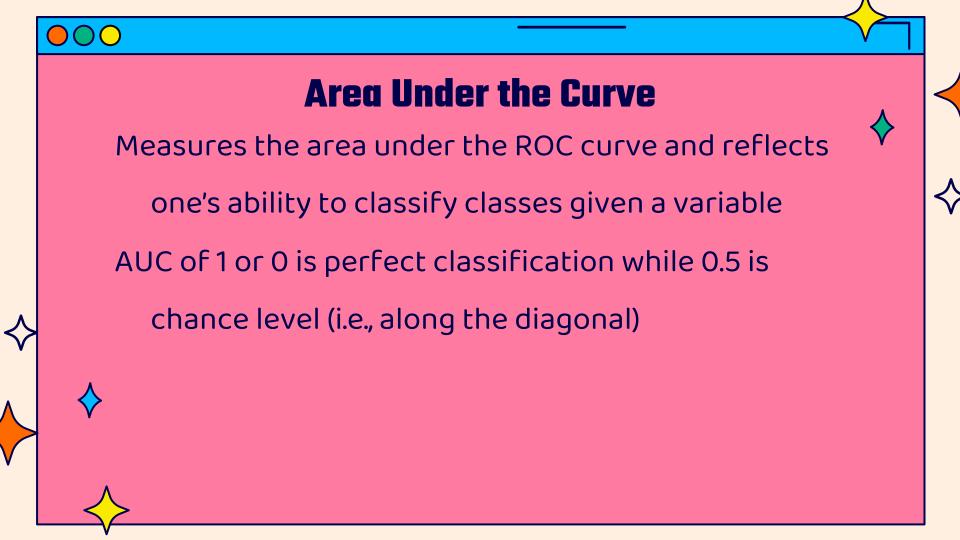


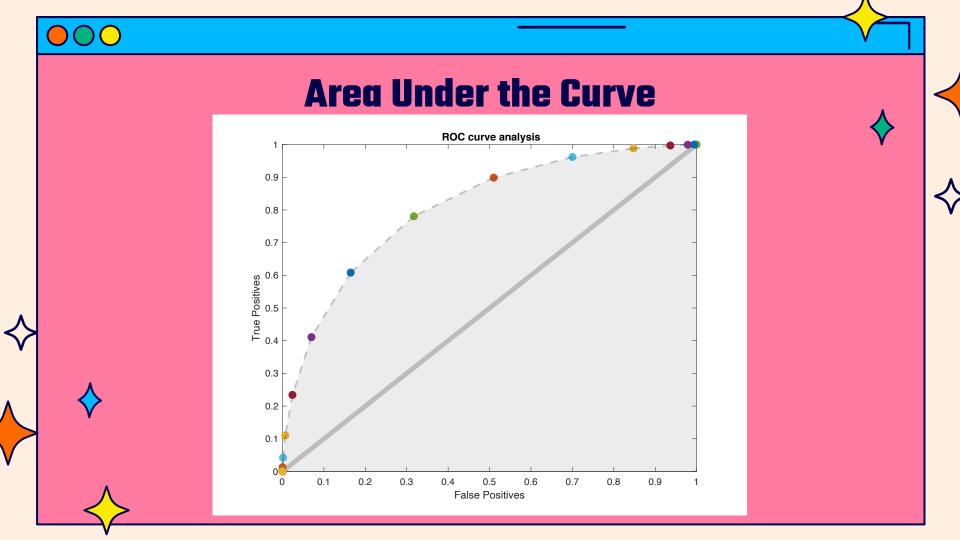


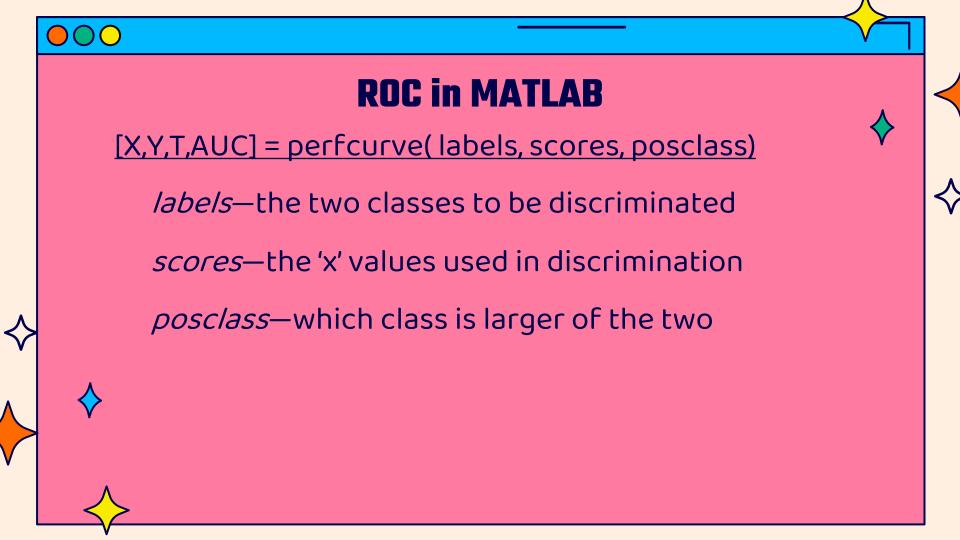












**ROC in MATLAB** 

[X,Y,T,AUC] = perfcurve( labels, scores, posclass)

X— x values of ROC curve

Y— y values of ROC curve

7—array of thresholds used



















