



### **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



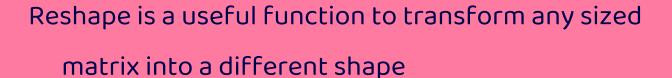
**corrcoef** — Returns the correlation between vectorized matrices

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



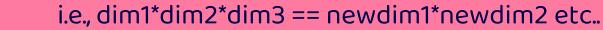


# Reminder: Reshape can help you



#### 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



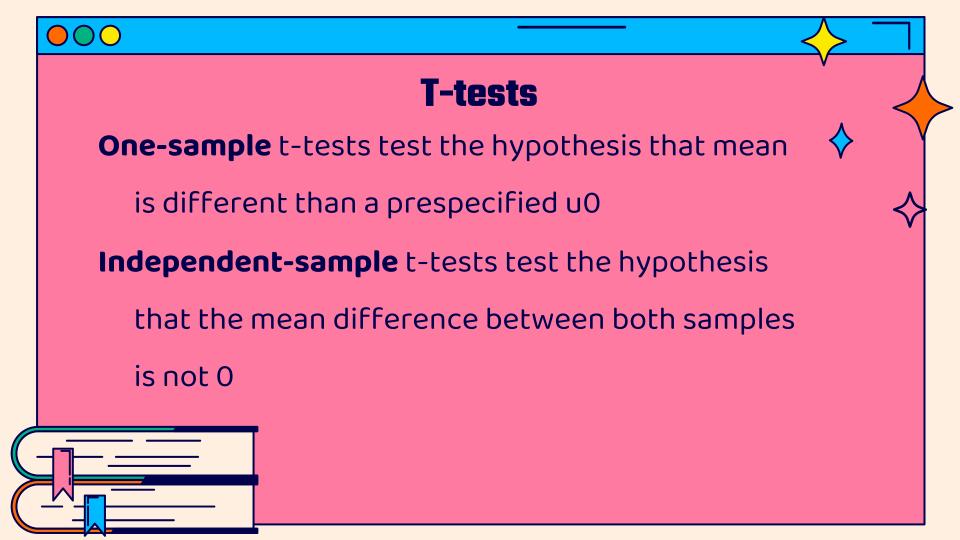


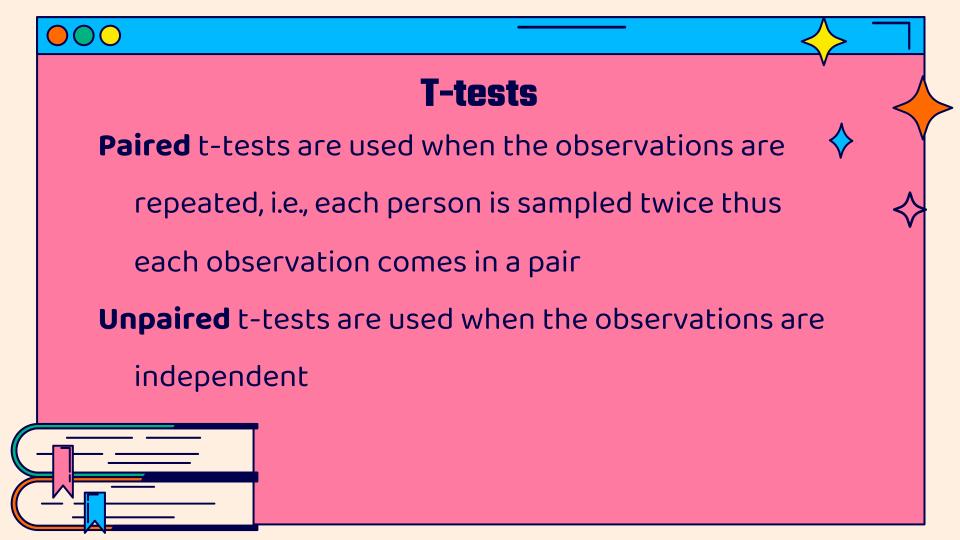


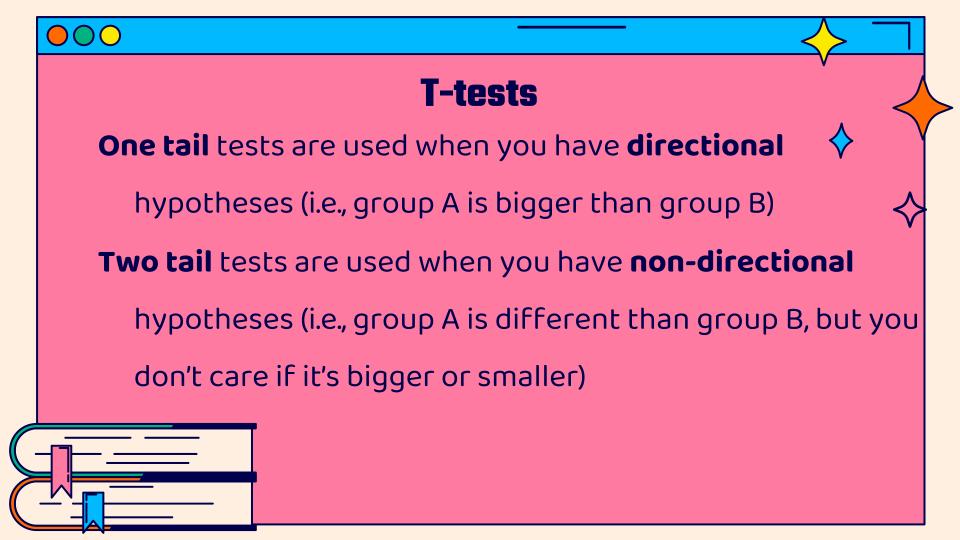


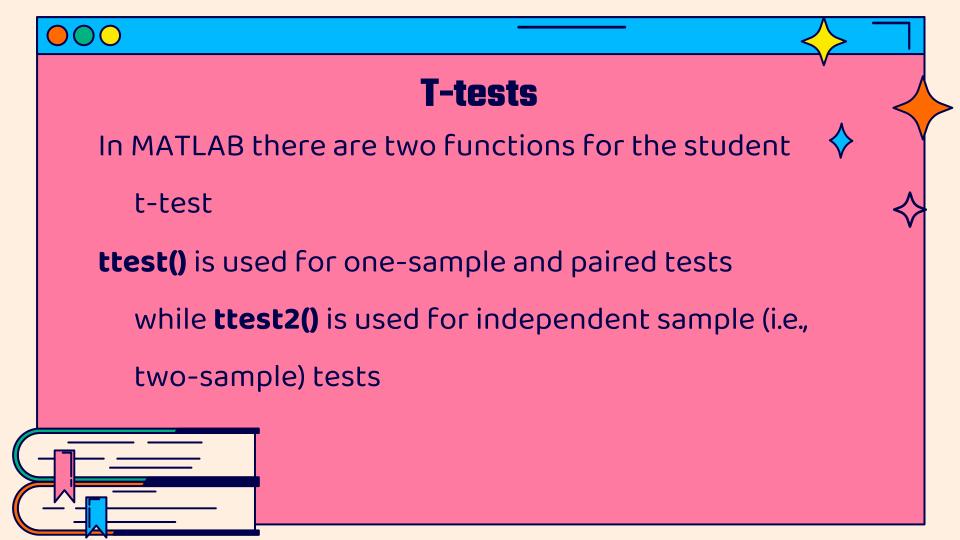
T-tests assume that your data come from a **normal** distribution and the observations are sampled **independently** from one another

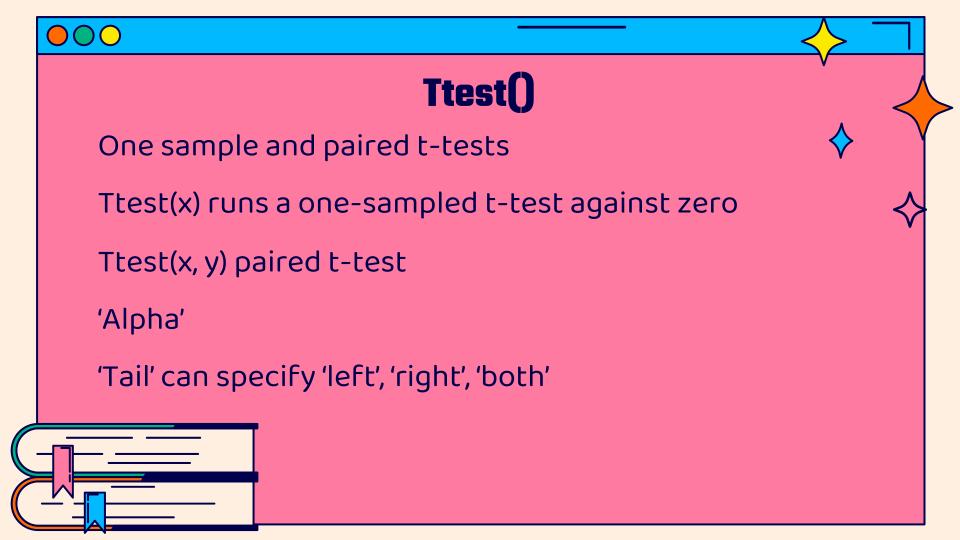
These assumptions apply for both paired and unpaired test

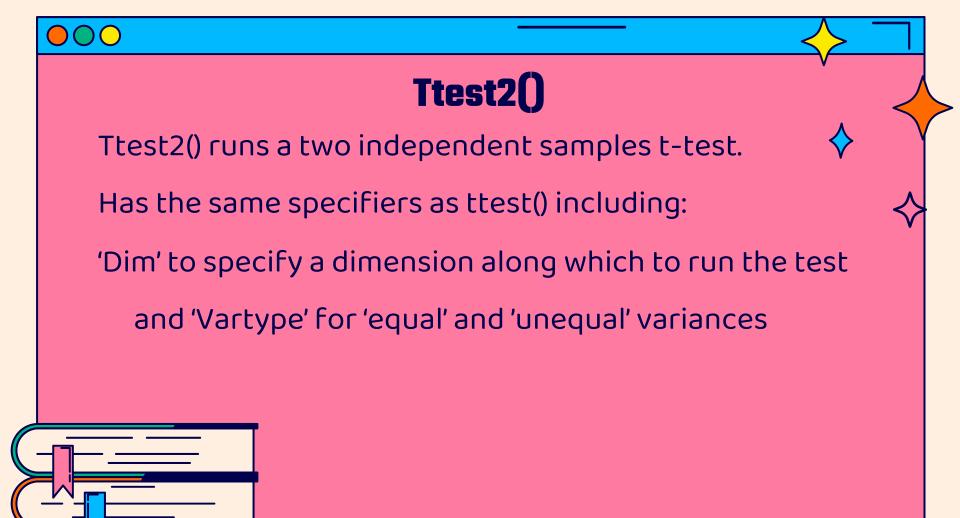


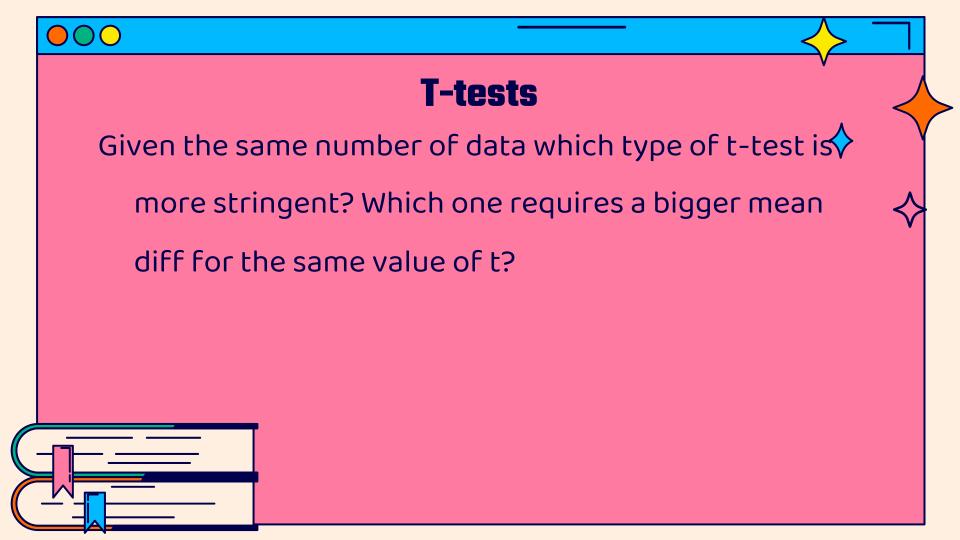














Let us assume we collect data from 20 people (paired)

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

Then the t value for a paired test would be:



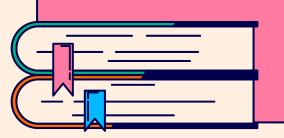
	$t = \frac{mean \ diff}{sd}$	- with df n — 1



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,



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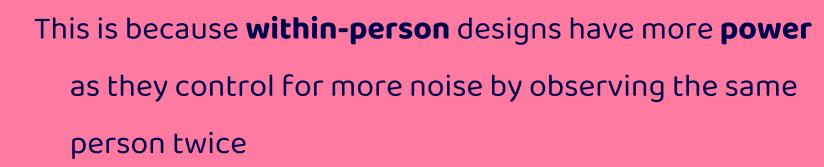
Then the t value for an unpaired test would be:

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



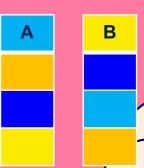
# permutations

**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

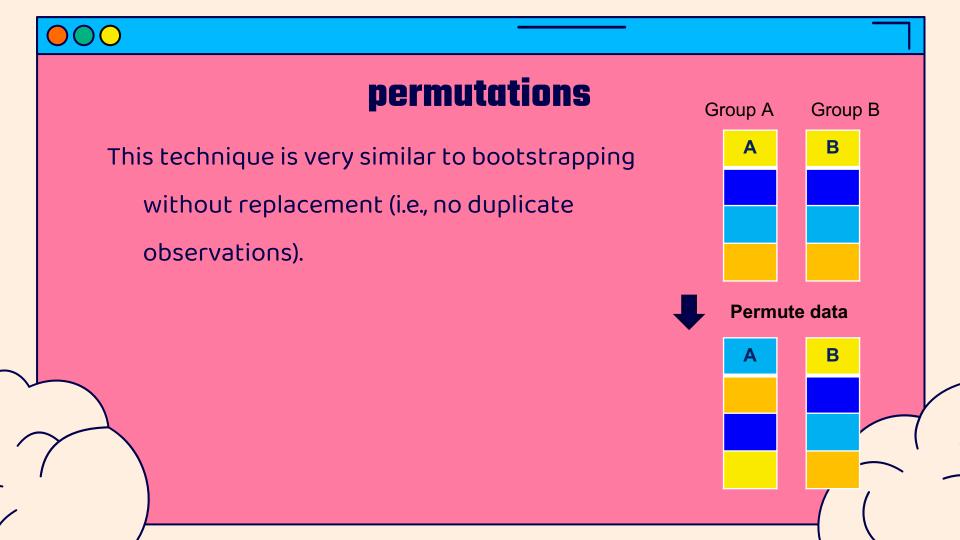
computed null distribution

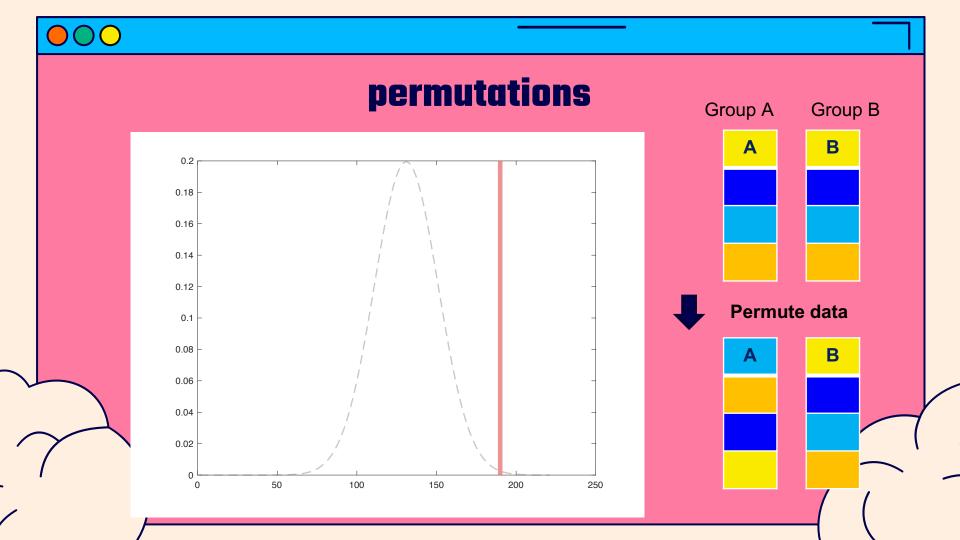
Group A Group B

A B



Permute data







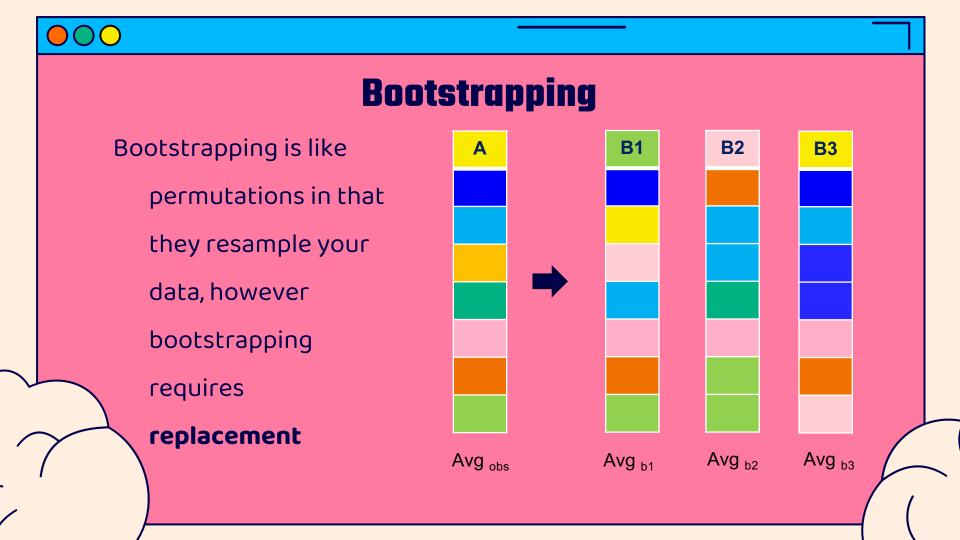
# **Bootstrapping**

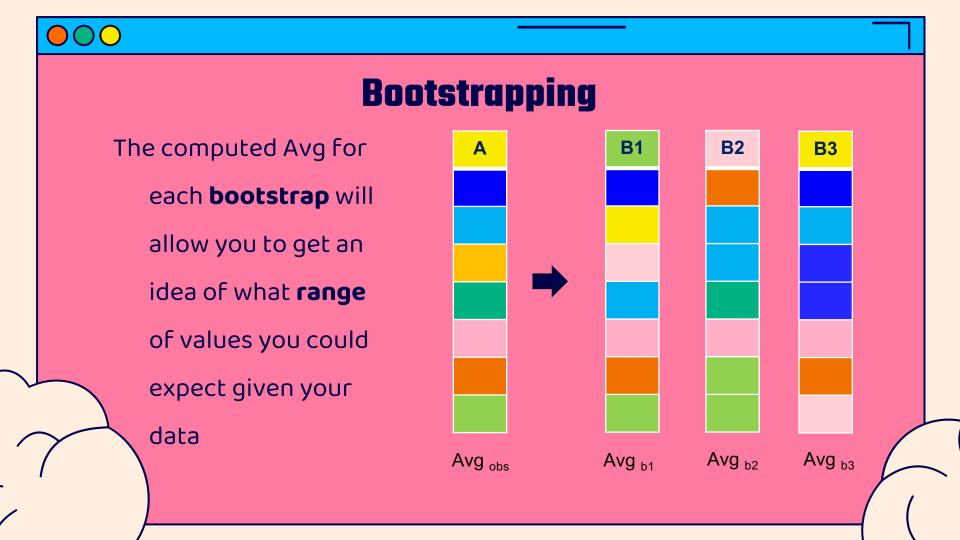
A very similar concept in statistics is the idea of **bootstrapping** to get a measure of uncertainty around an estimate (e.g., CI)

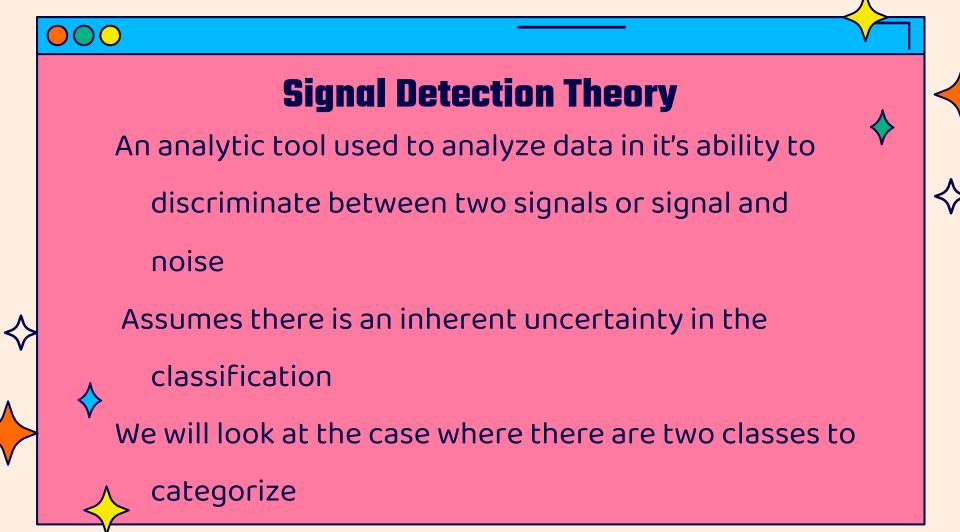
Bootstrapping is like permutations in that they resample your data, however bootstrapping requires **replacement** 

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











detection theory

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



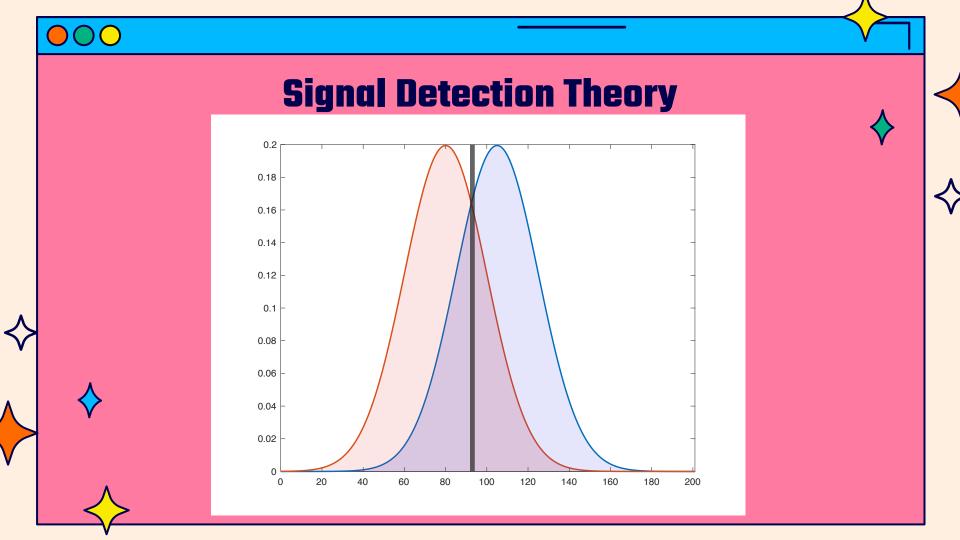
Sensitivity: one's ability to discriminate between signal and noise

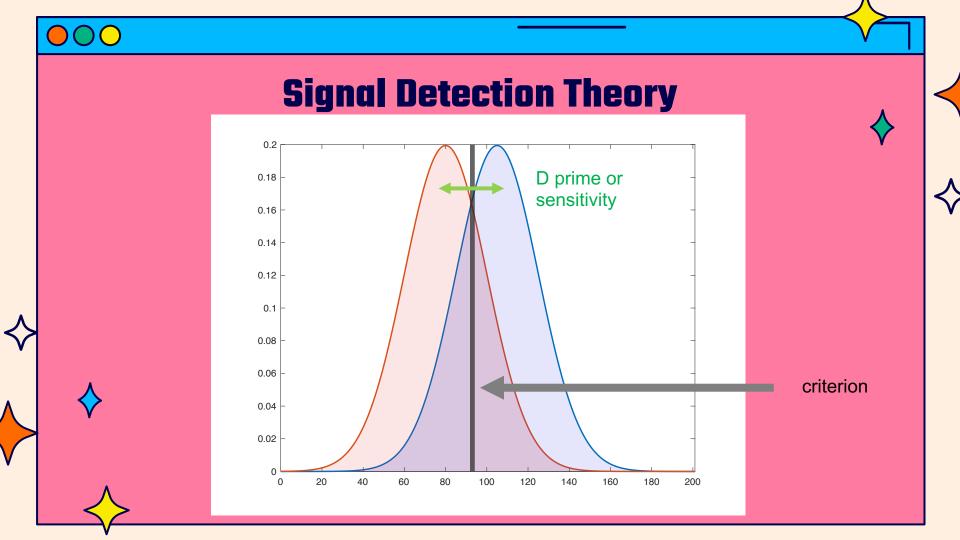






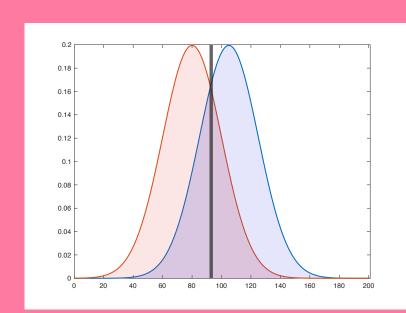












	Signal	Noise
Present	Hits	False Alarms
Absent	Misses	Correct Rejection







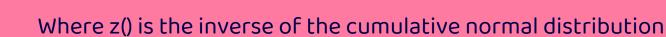






$$c = -\frac{1}{2}(z(Hits) + z(False\ Alarms))$$













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











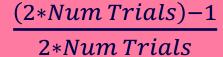




Similarry we assume that if we double the number of



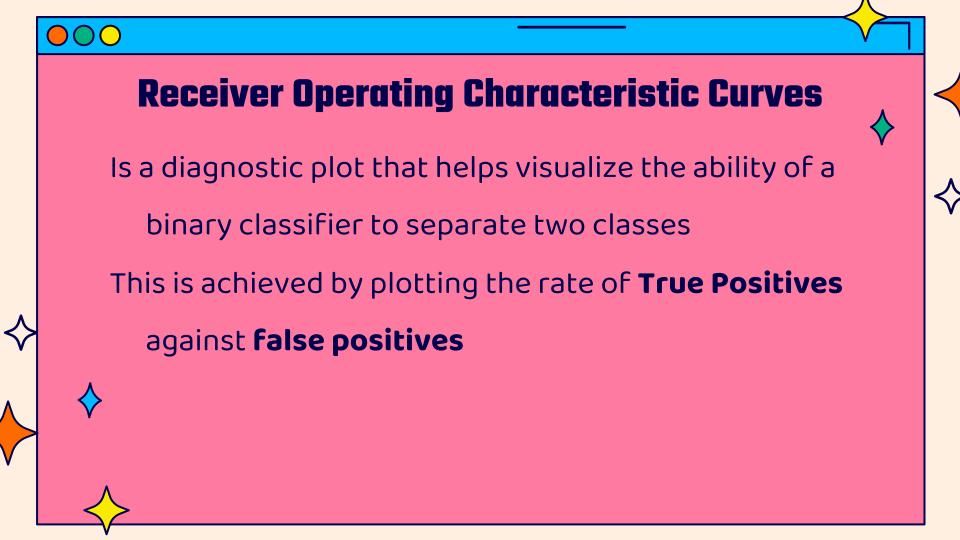
pHIT or pFA =1, then correct with



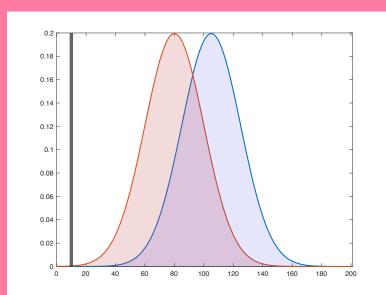


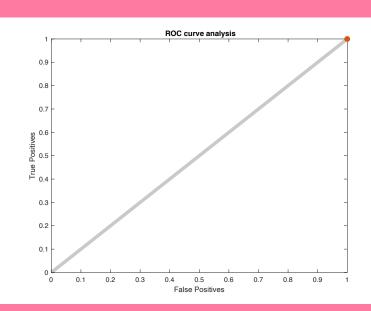










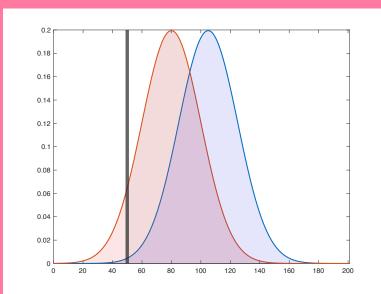


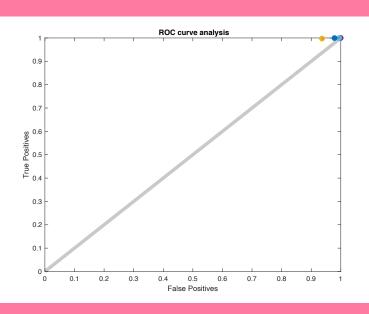










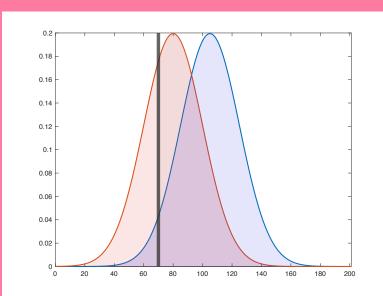


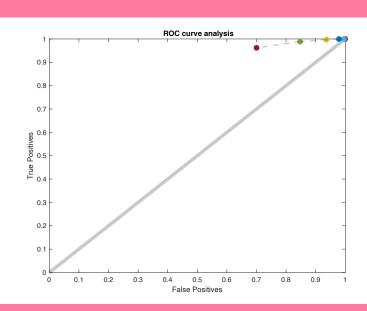










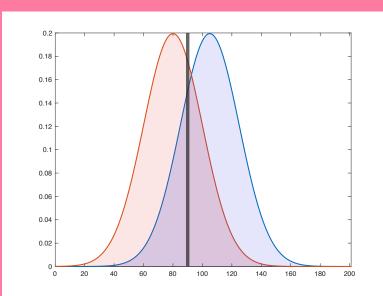


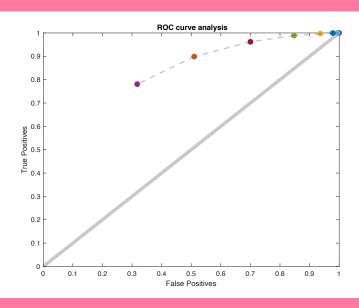










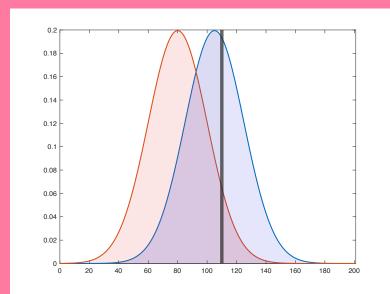


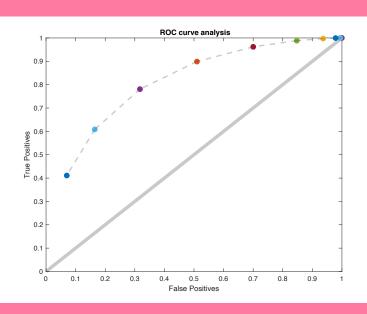










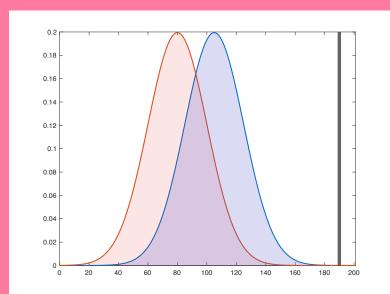


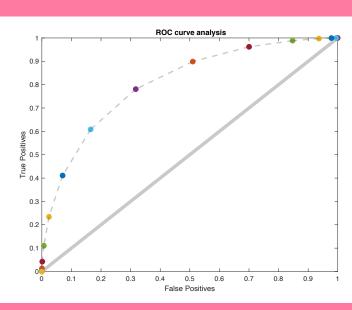






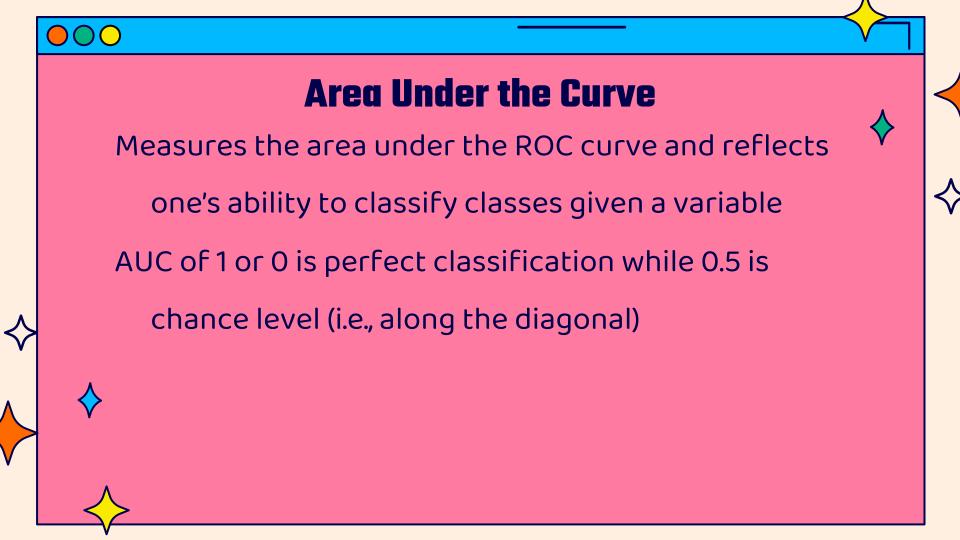


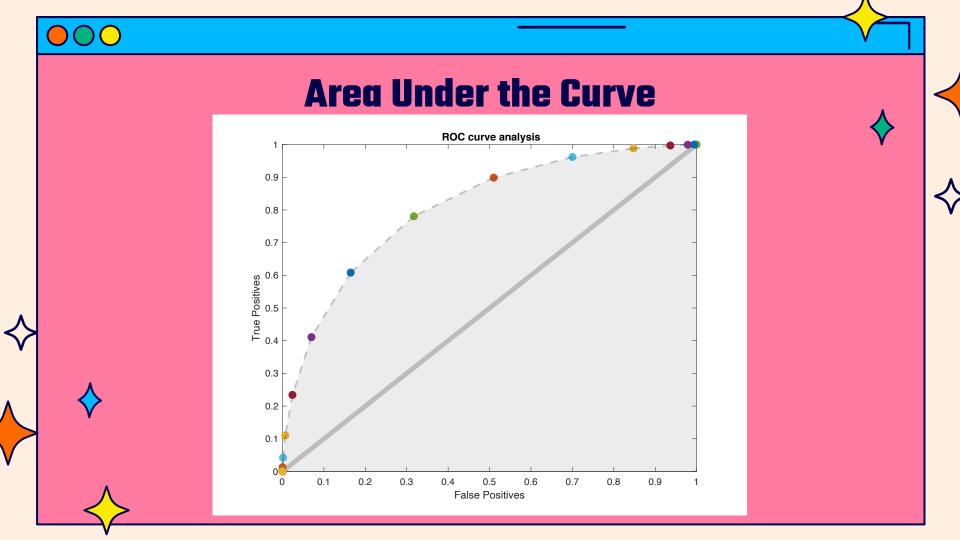


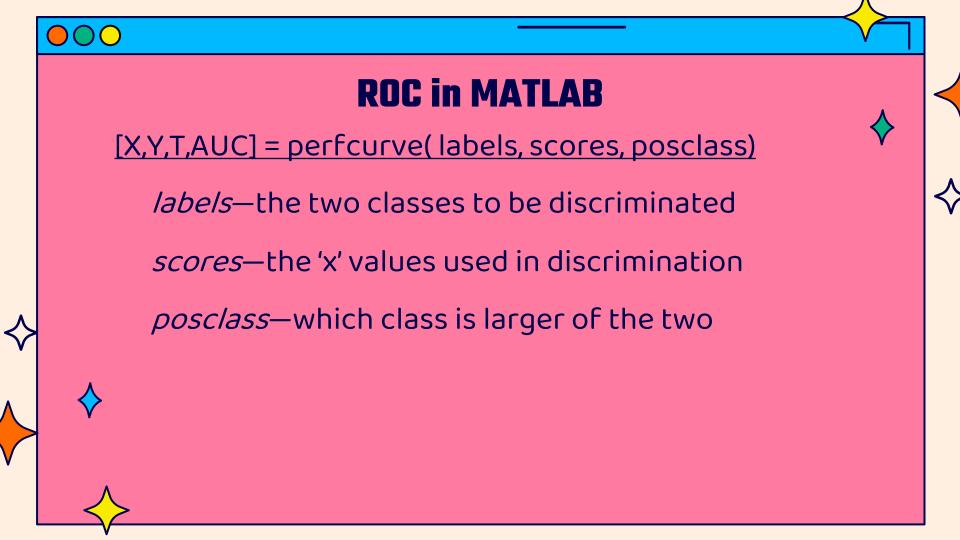












**ROC in MATLAB** 

[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





















