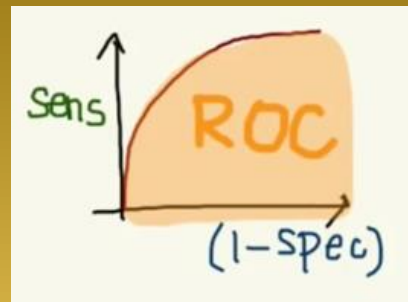


ML EVALUATION

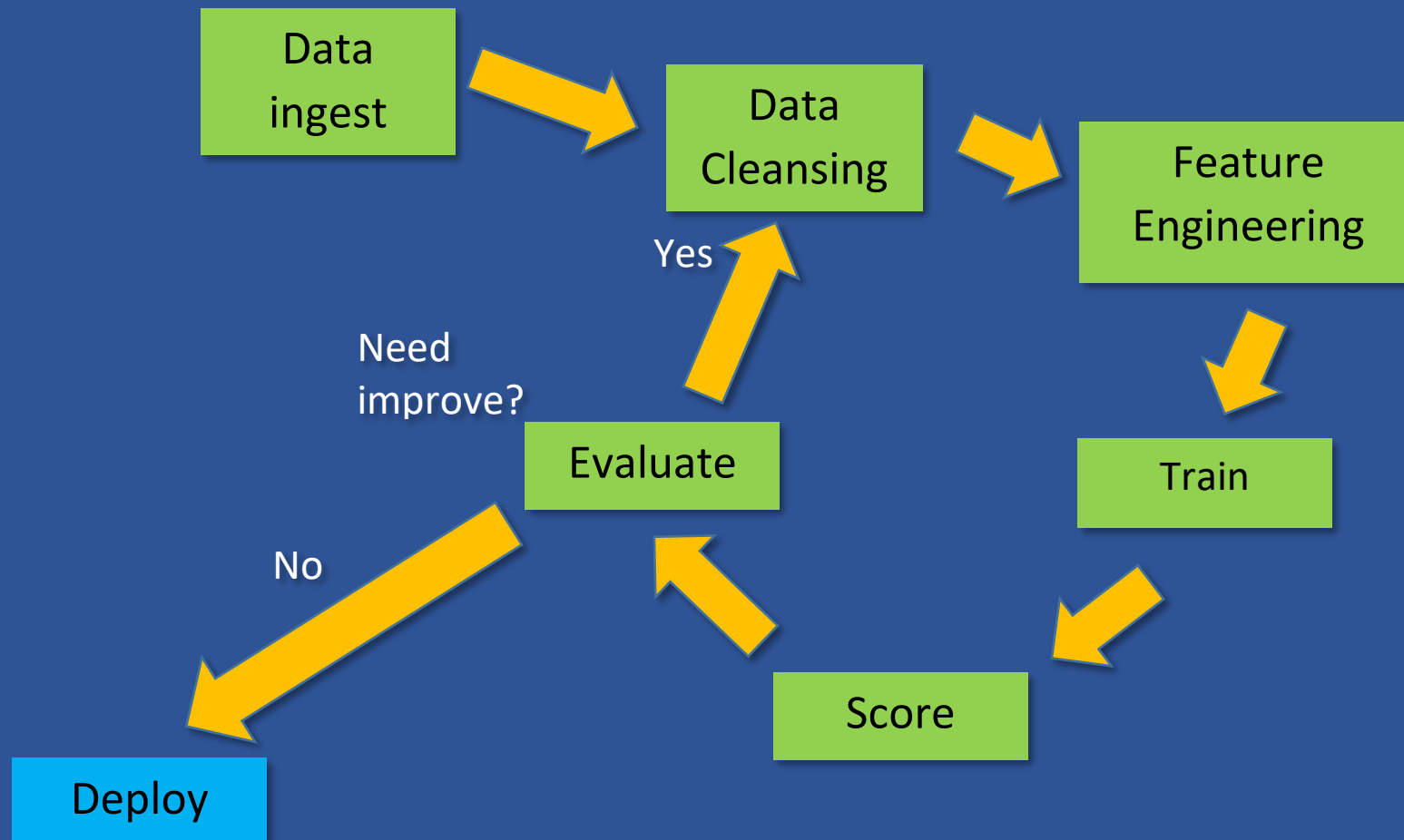


In this session

- ML train and evaluation circle
- How to read Histogram
- How to read Box Plot
- Adding Evaluate Model
- How to read ROC curve
- Area Under the Curve (AUC)
- How to read Evaluation metrics

ML Evaluation

ML evaluation circle





ML Evaluation

How to read Scoring results

Titanic Evaluate > Score Model > Scored dataset

rows 267 columns 10

view as  

Survived	PassengerClass	Gender	Age	SiblingSpouse	ParentChild	FarePrice	PortEmbarkation	Scored Labels	Scored Probabilities
1	3	male	20	1	1	15.7417	C	0	0.128143
1	2	female	25	1	1	30	S	1	0.999319
0	3	male	28	0	0	7.8958	C	0	0.40695
1	3	female	28	1	1	22.3583	C	1	0.993964
0	3	male	28	0	0	9.5	S	0	0.000195
0	1	male	29	0	0	30	S	1	0.97861
1	1	male	49	1	0	56.9292	C	1	0.932772

- This table = Scored dataset
- Row = 267 / Columns = 10
- Total column = 10 / Left 8 = features / Right 2 = prediction results
- Scored Label 0 = dead 1 = survived
- Scored Probabilities (SP) $SP \leq 0.5$ == dead / $SP > 0.5$ == survived

ML Evaluation

How to read Scoring Statistics

Statistics	
Mean	28.8265
Median	28
Min	0.42
Max	80
Standard Deviation	12.3791
Unique Values	61
Missing Values	0
Feature Type	Numeric Feature

Show Statistics of the Scored dataset

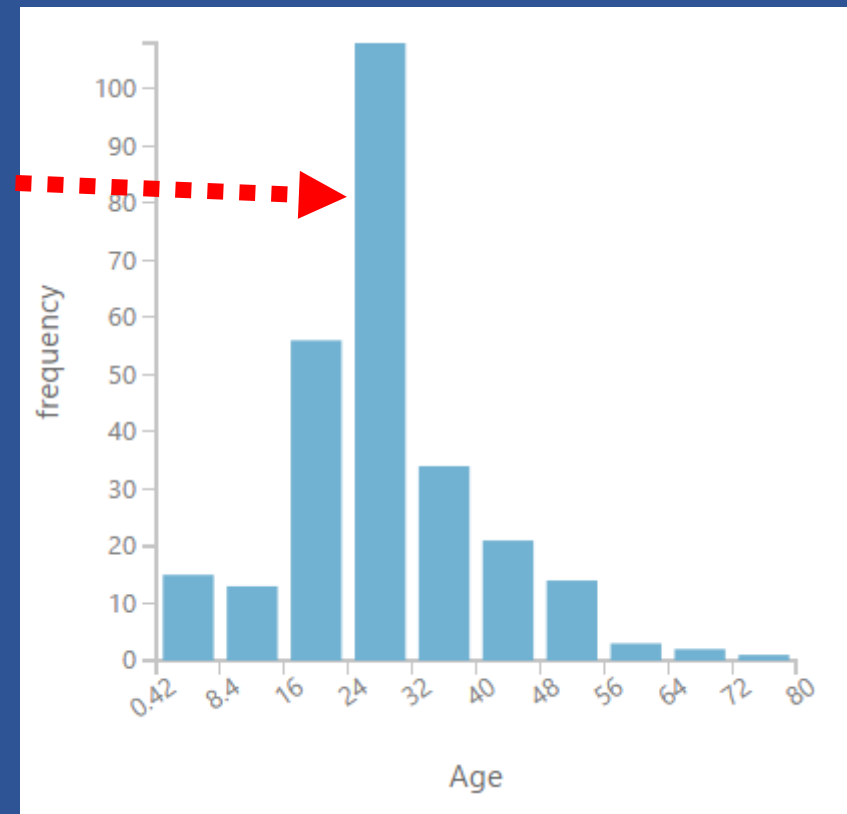
- Mean = Sum of all the values divided by the number of values
- Median = The midpoint of the data after being ranked
- Standard Deviation = The square root of the variance
- Unique Values
- Missing Value

ML Evaluation

How to read Score Histogram

Histogram

- Representation: distribution of numerical data
- Bin: series of intervals (bin)
- Count: values fall into each interval



ML Evaluation

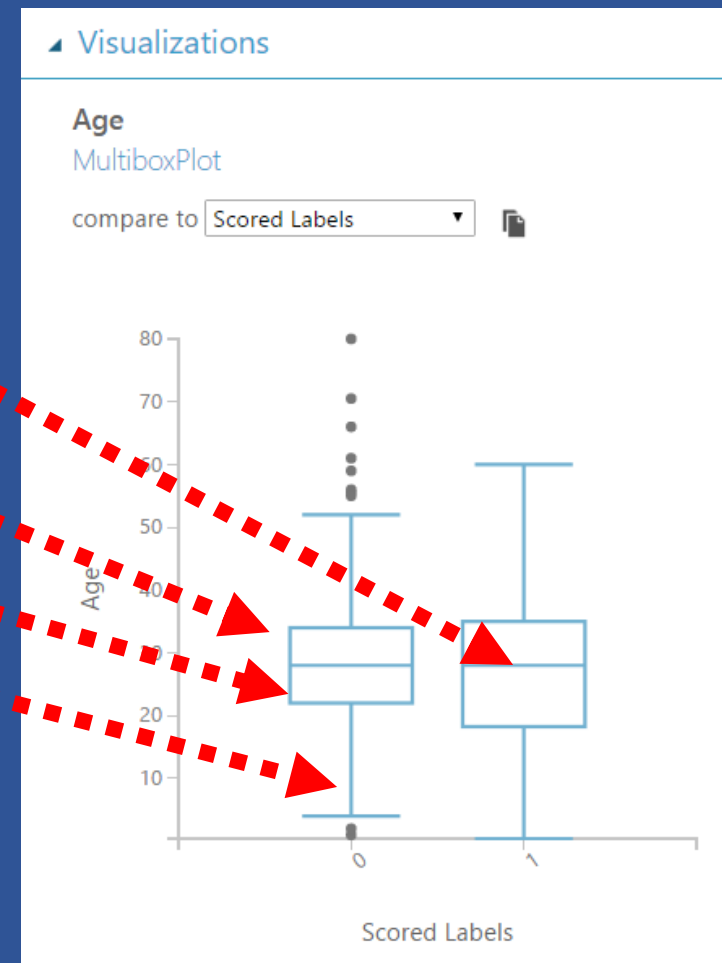
How to read Box Plot

Box Plot

Box Plot (whisker) is a standardized way of displaying the distribution of data

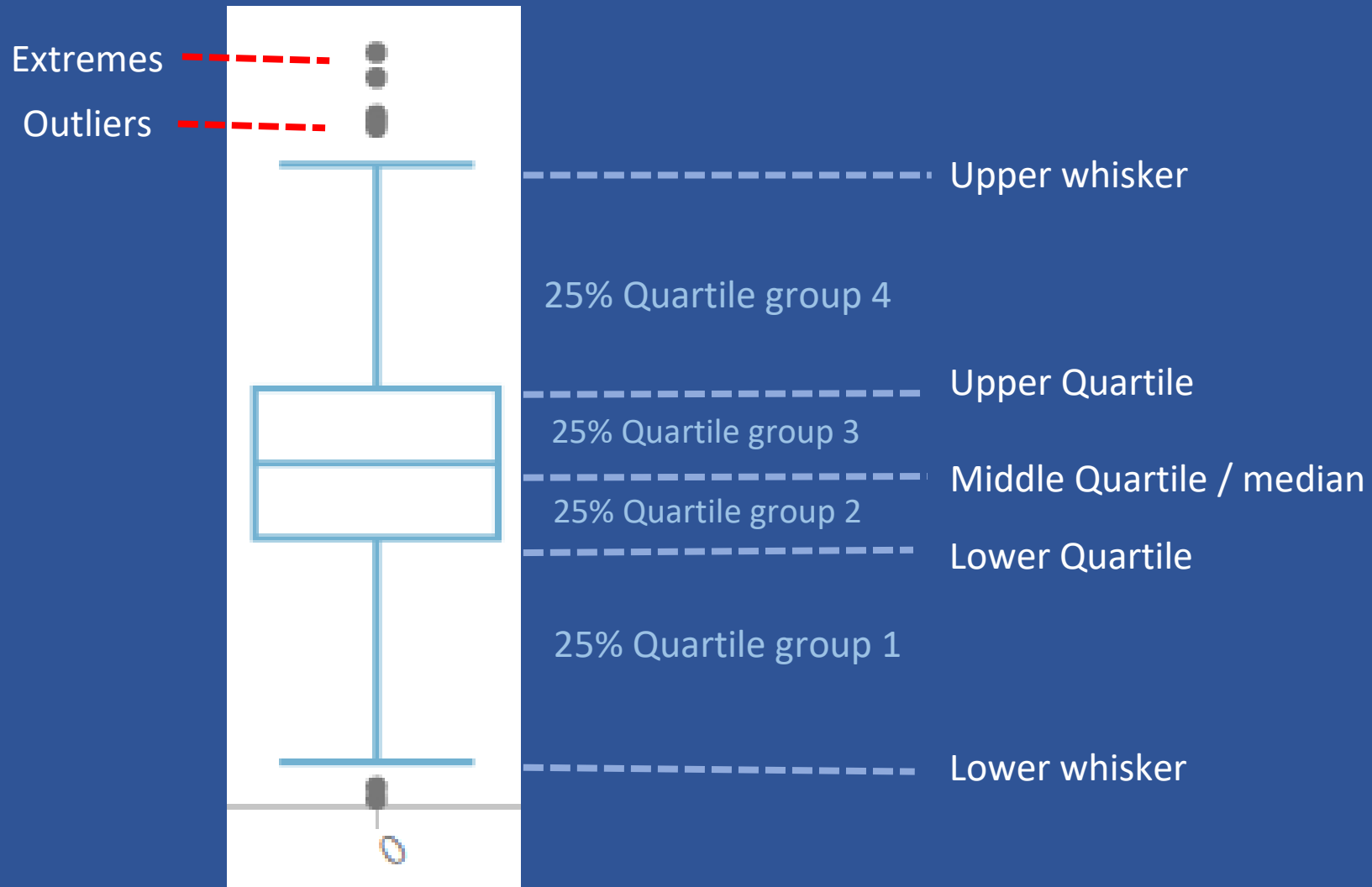
- **Median:** marks the mid-point of the data
- **Box:** middle 50% of scores for the group.
- **Upper quartile:** 75% of the scores fall below the upper quartile.
- **Lower quartile:** 25% of scores fall below the lower quartile.
- **Whiskers:** scores outside the middle 50%

0 = dead



ML Evaluation

Box Plot Definitions

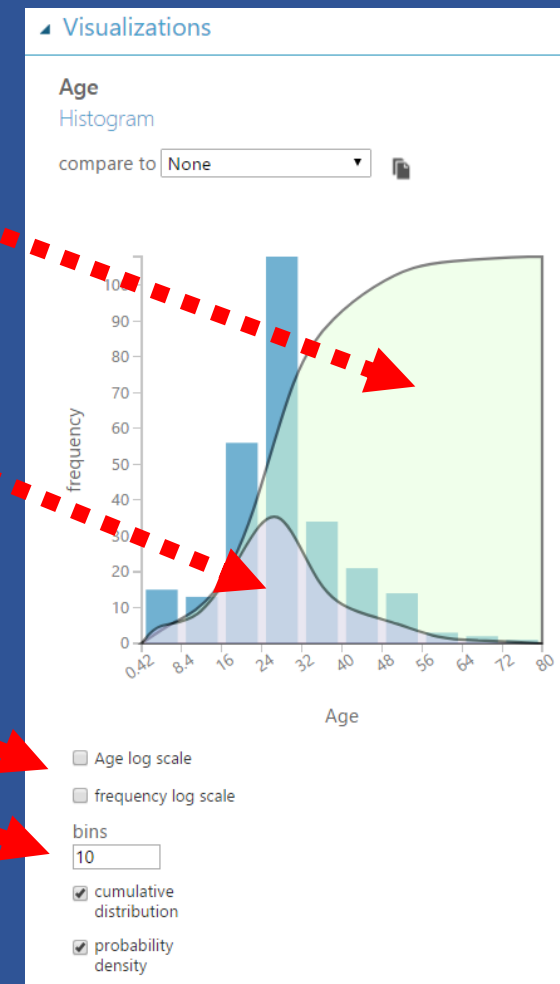


ML Evaluation

Histogram option

Histogram options

- **Cumulative distribution function (cdf)**: shows "How common are samples that are *less than or equal* to this value?"
- **Probability density function (pdf)**: shows "How common are samples at exactly this value?"
- **Scale**: scaling the distribution
- **bins**: number of bin

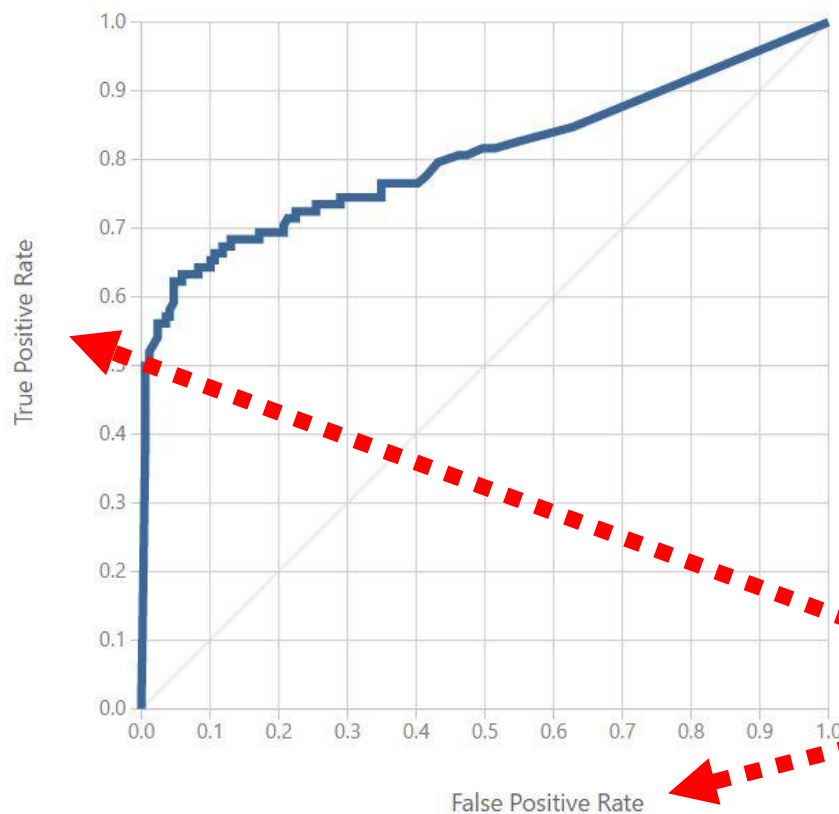


ML Evaluation

Receiver Operating Characteristic (ROC) Curve

Titanic Evaluate > Evaluate Model > Evaluation results

ROC PRECISION/RECALL LIFT



True Positive	False Negative	Accuracy	Precision
64	34	0.805	0.780
False Positive	True Negative	Recall	F1 Score
18	151	0.653	0.711
Positive Label	Negative Label		
1	0		

Threshold AUC 0.817

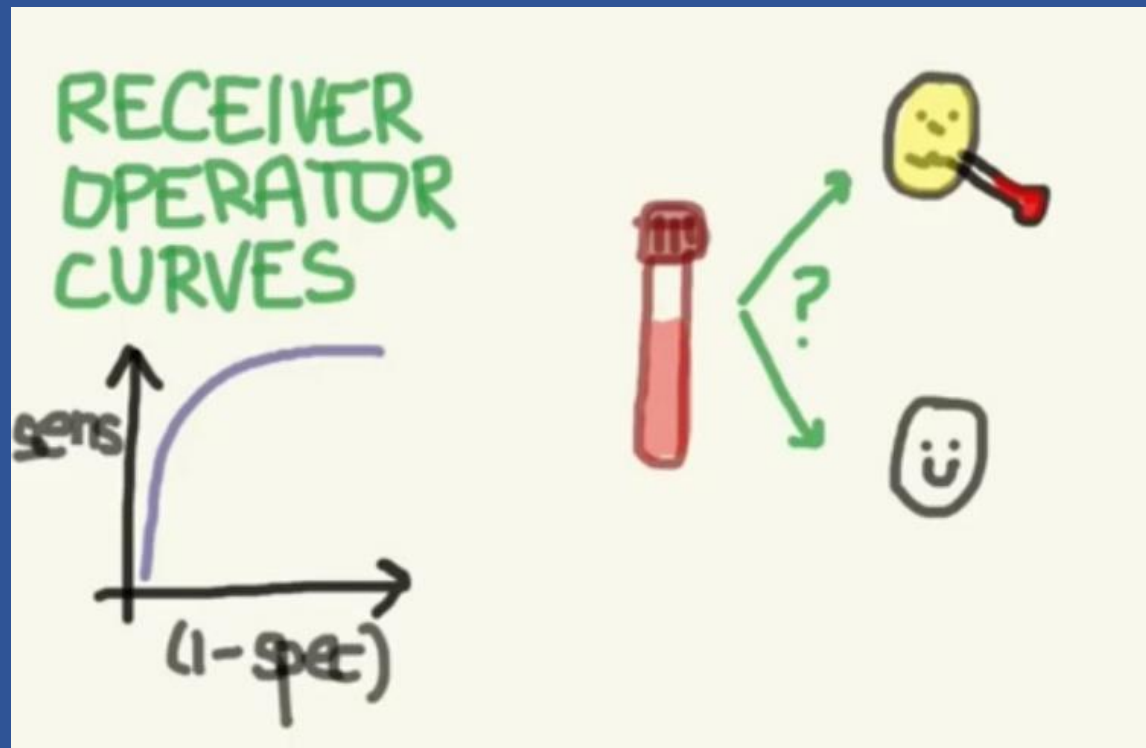
True Positive Rate (TPR)

False Positive Rate (FPR)

ML Evaluation

How to read ROC curve

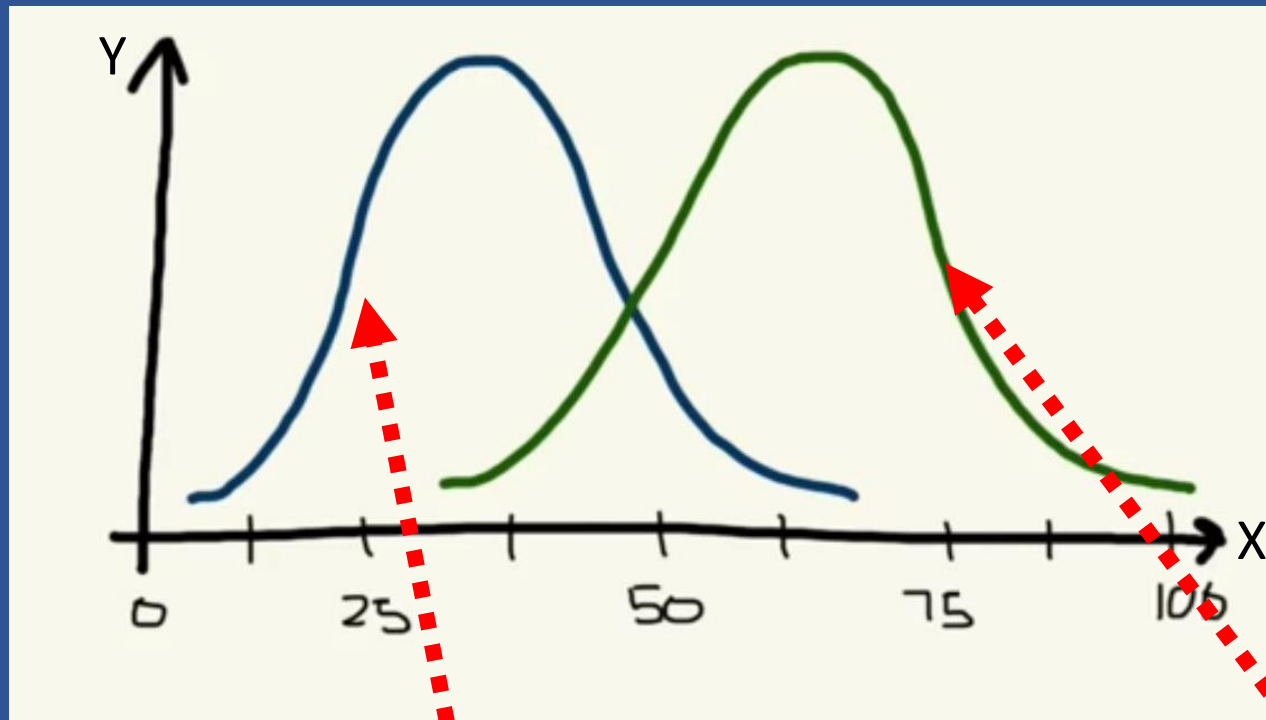
ROC curve is a graphical plot that illustrates the diagnostic ability of a **binary classifier** system as its discrimination threshold is varied.



ROC curve prediction result who have disease who don't

ML Evaluation

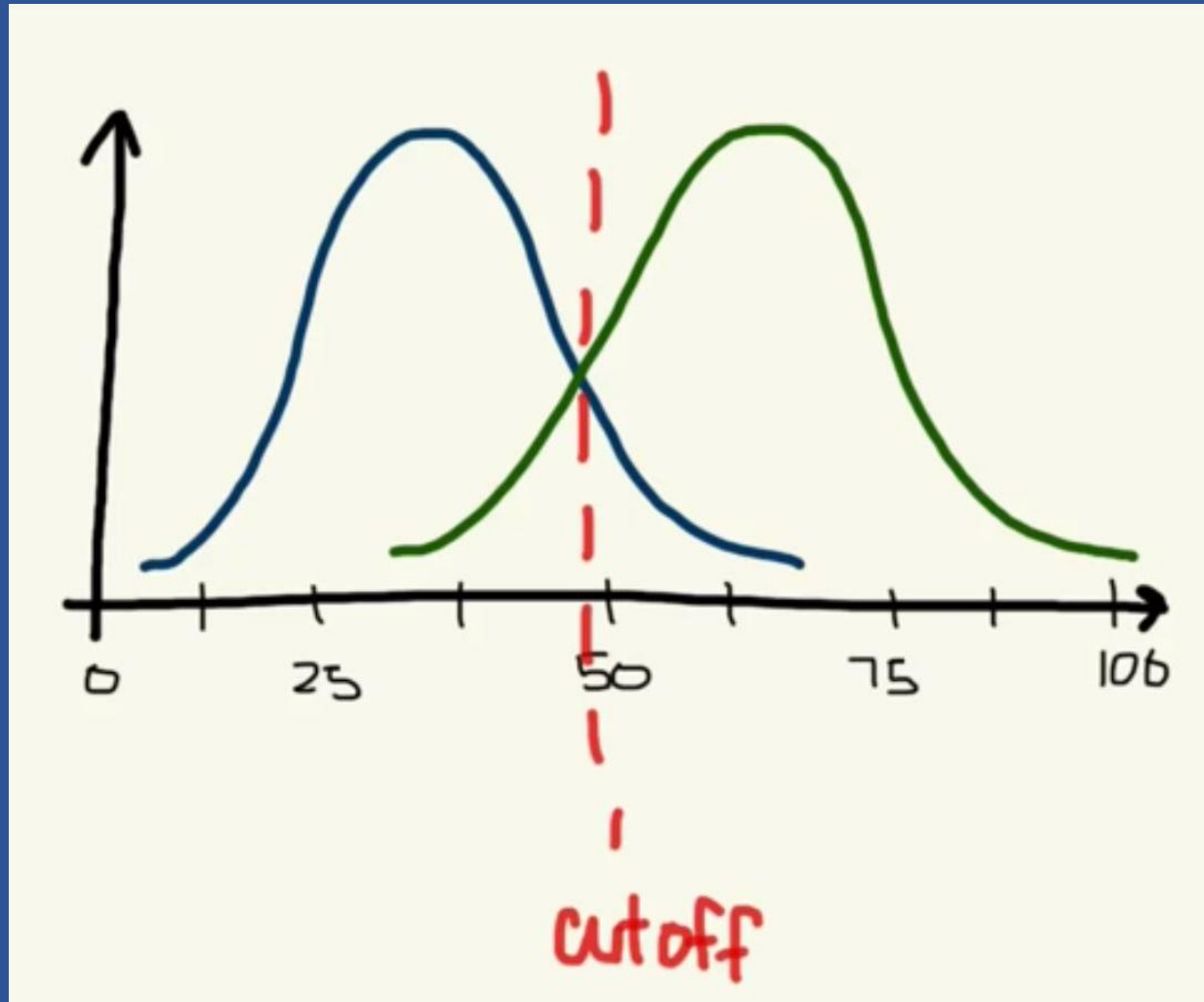
Distribution score



Left distribution = patient who do **NOT** have disease (survived) / Right = have disease (dead)

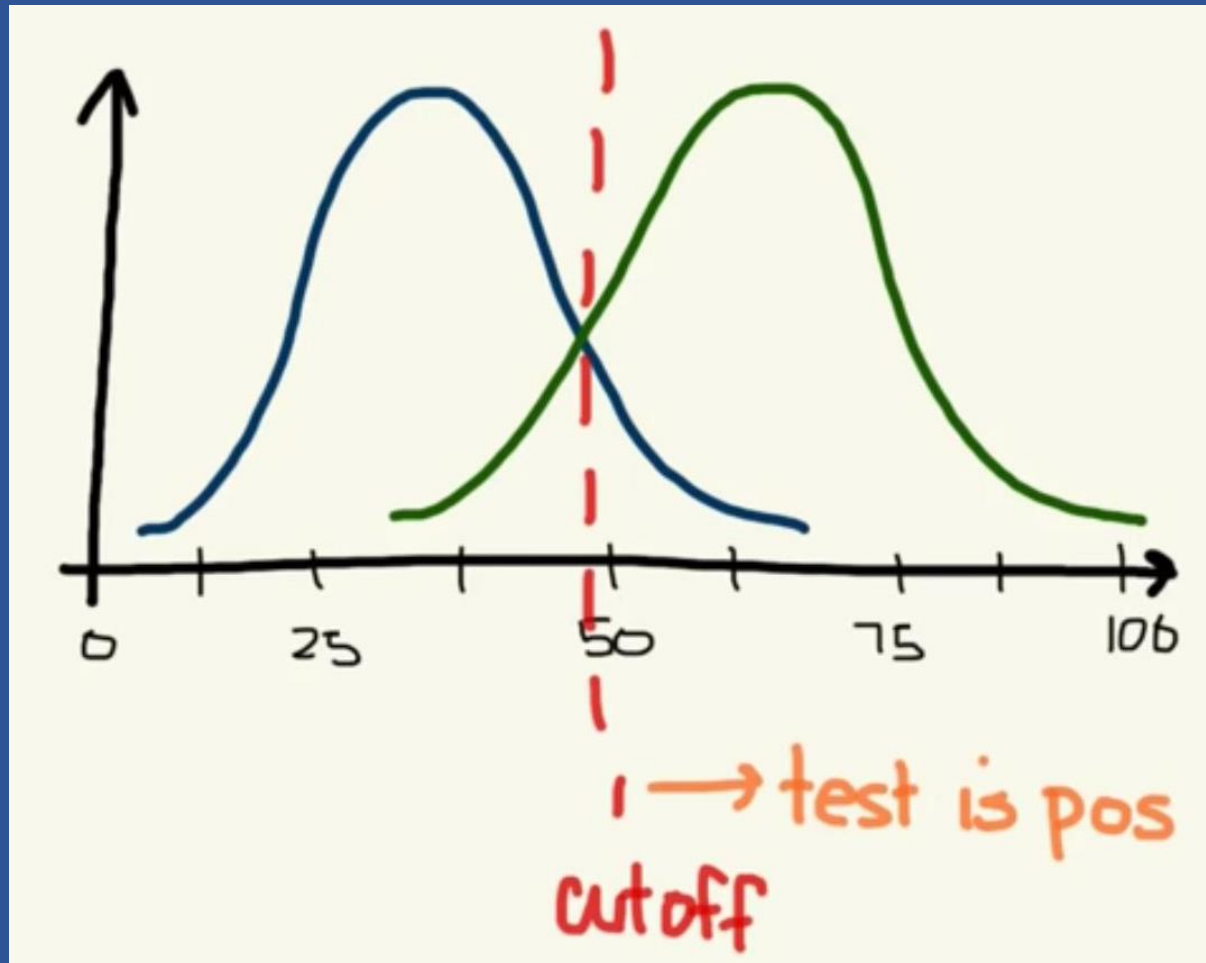
x axis = score / y axis = number of patient

ML Evaluation Cutoff line



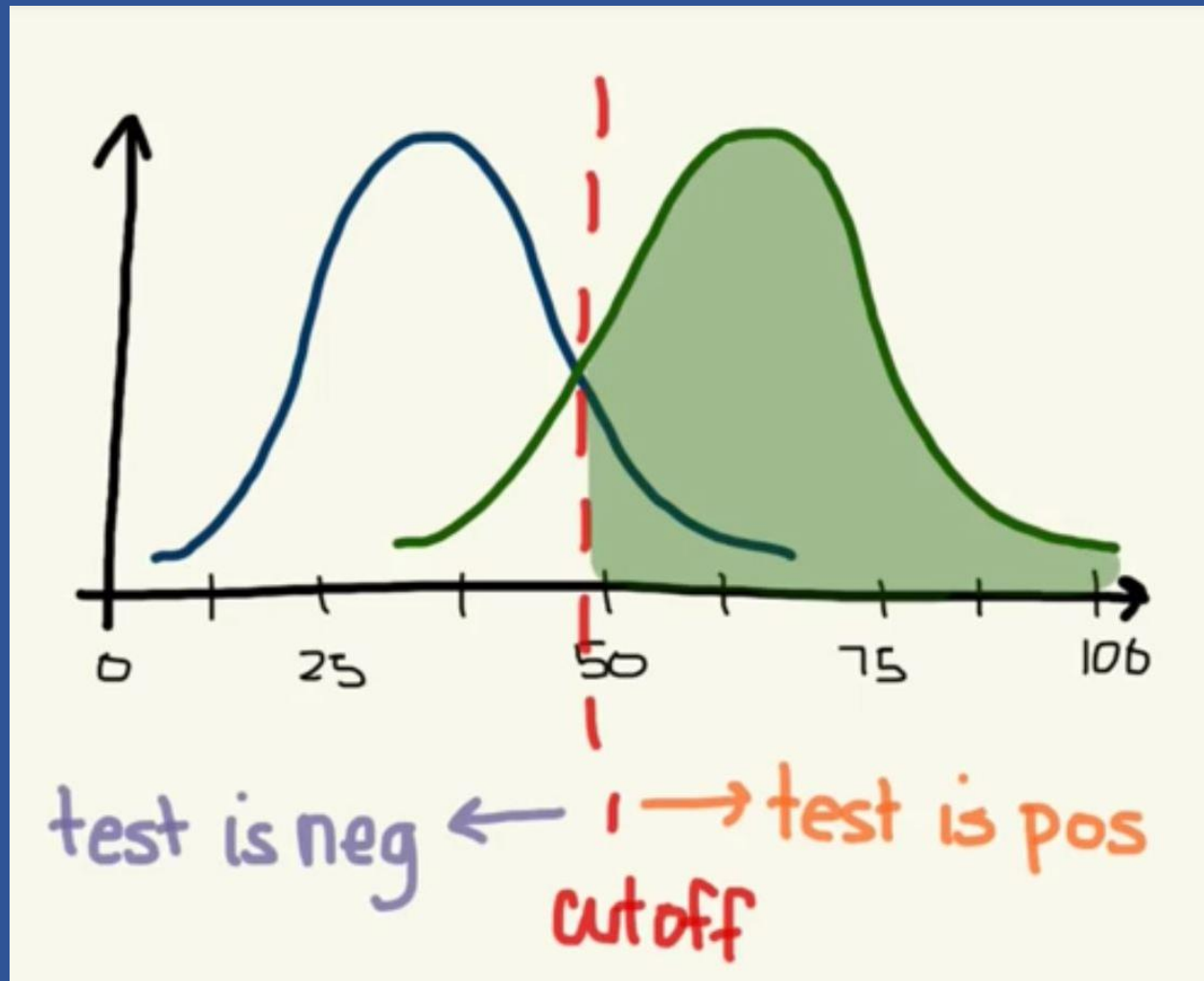
ML Evaluation

Area where the test is positive



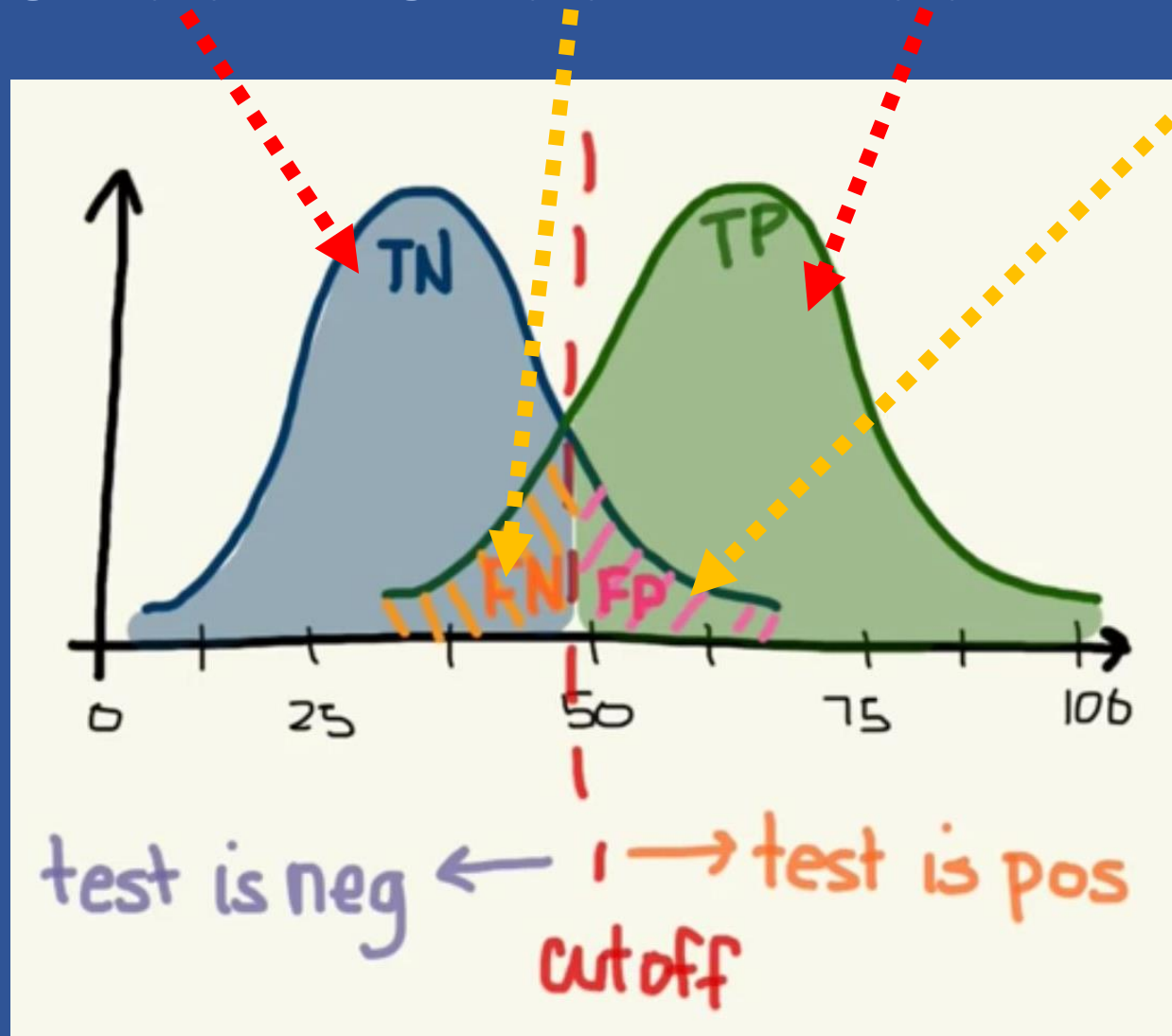
ML Evaluation

Area where the test is negative



ML Evaluation

True Negative (TN), False Negative (FN) / True Positive (TP), False Positive (FP)

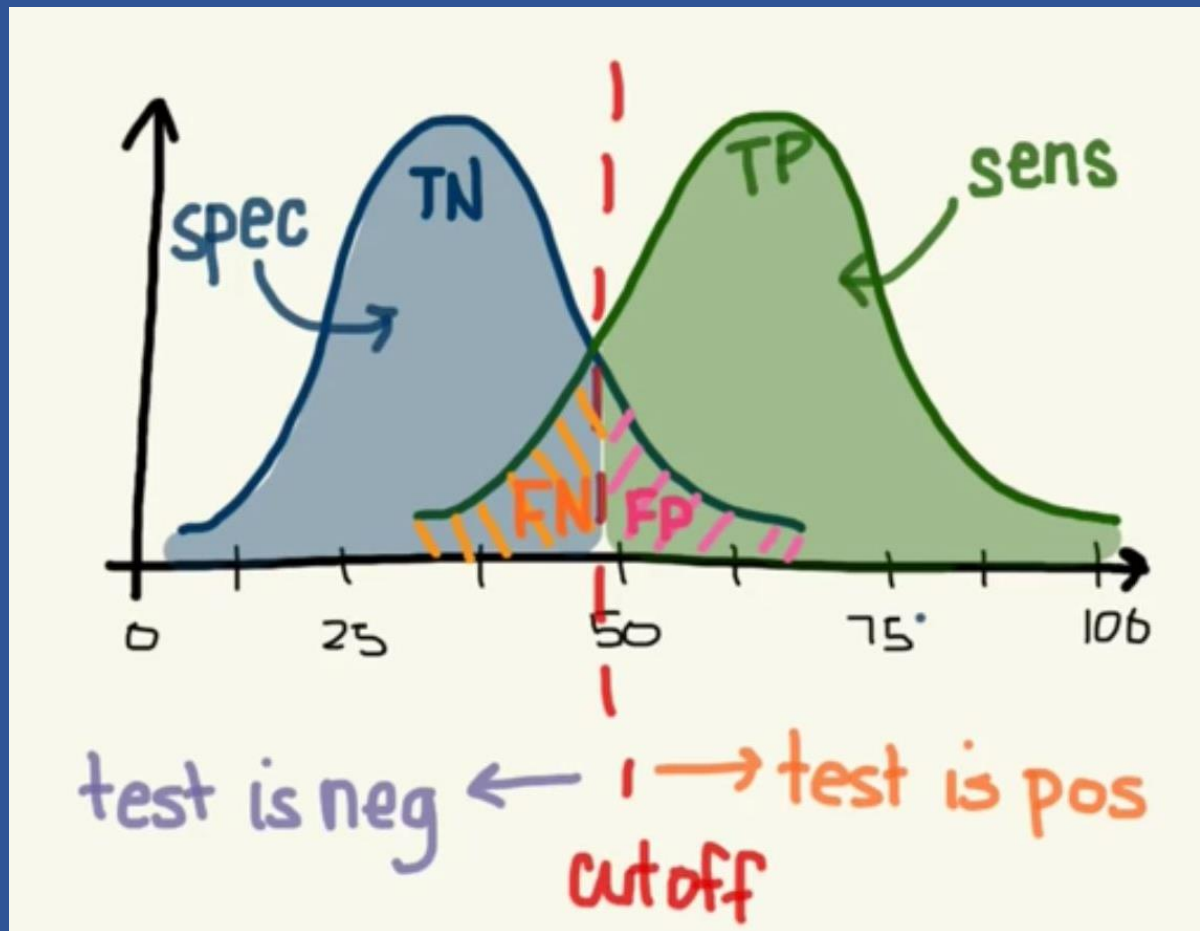


ML Evaluation

ROC Specificity / Sensitivity

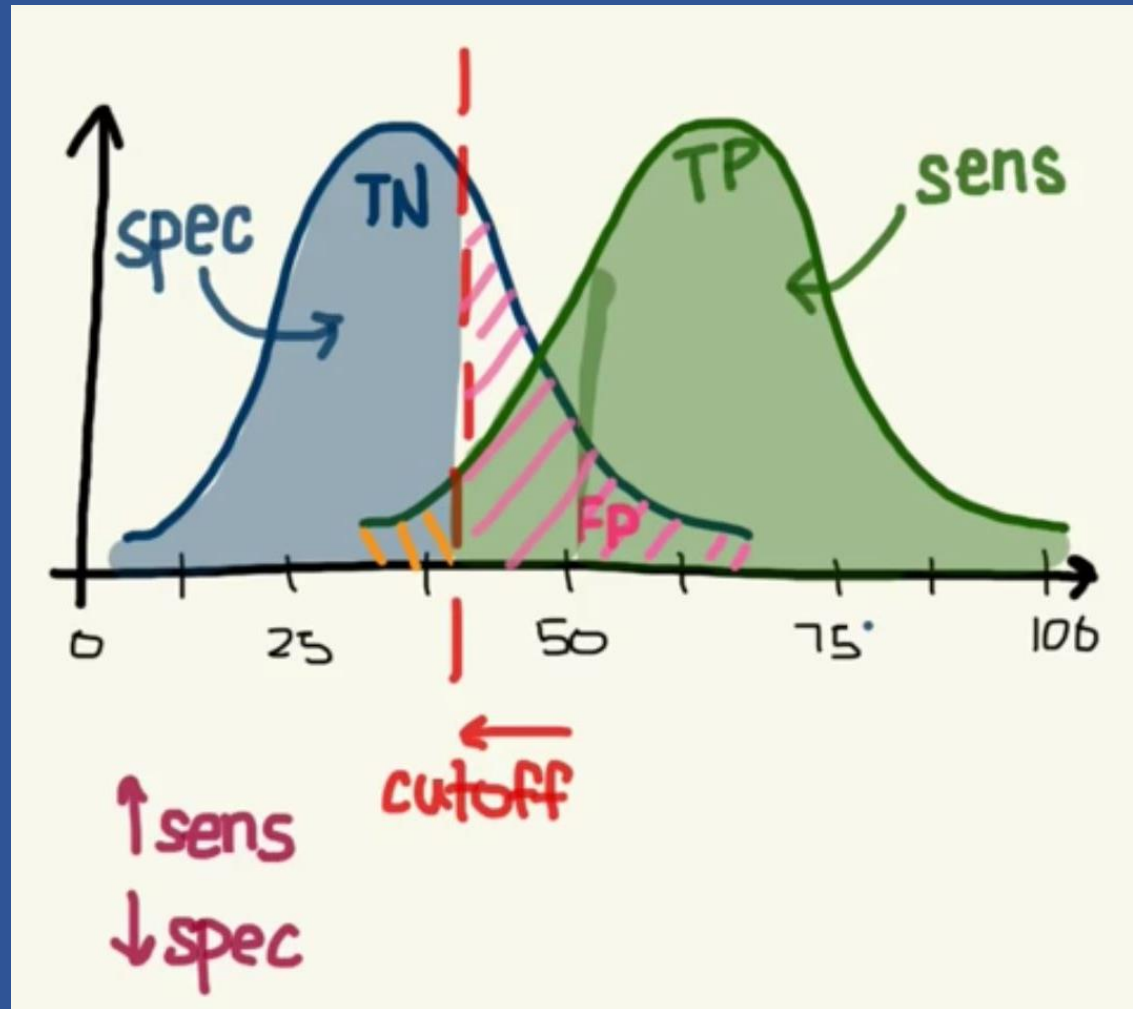
Specificity = True Negative Rate

Sensitivity (Recall) = True Positive Rate



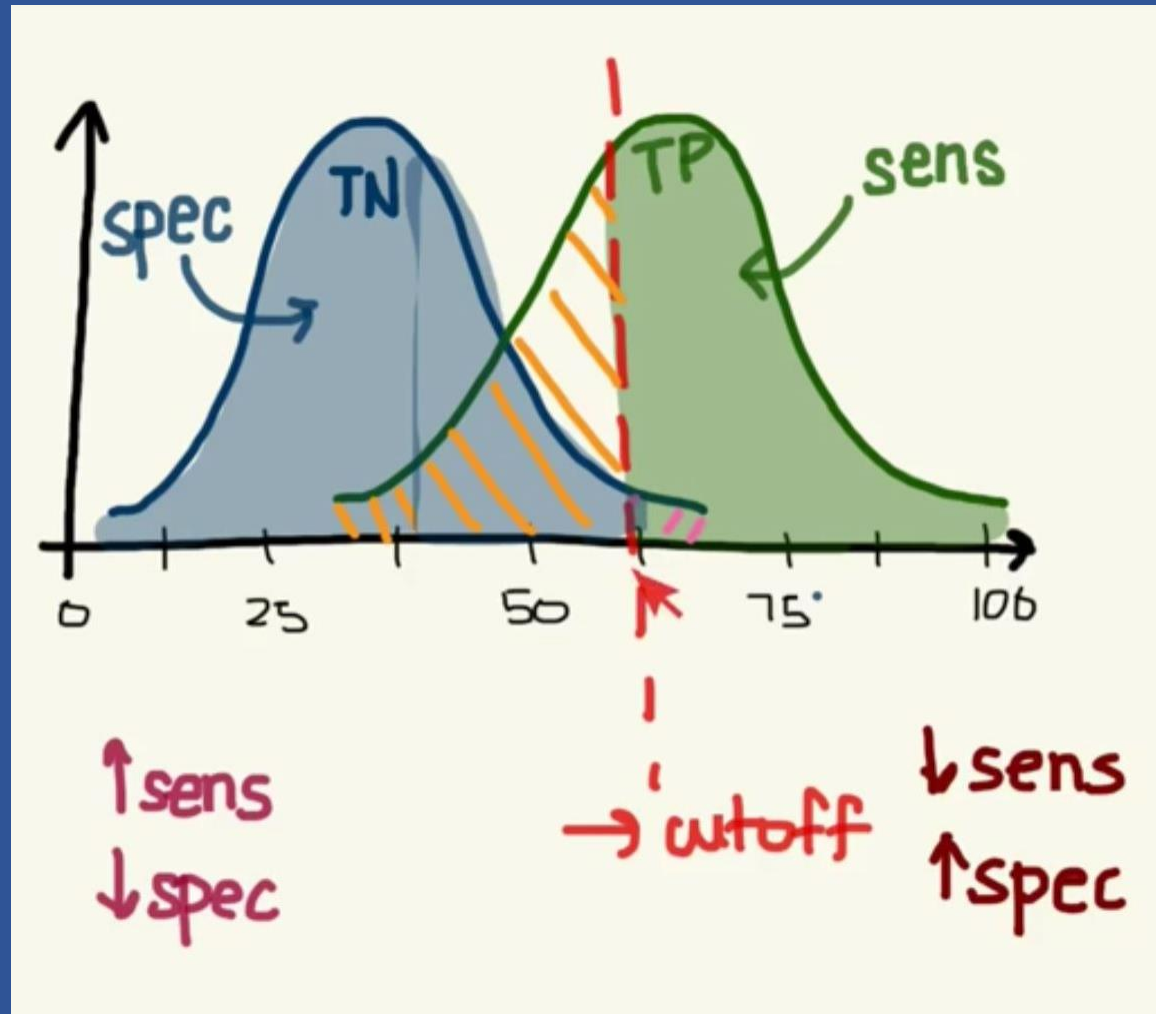
ML Evaluation

Move cutoff to the left Sens++ / Spec--



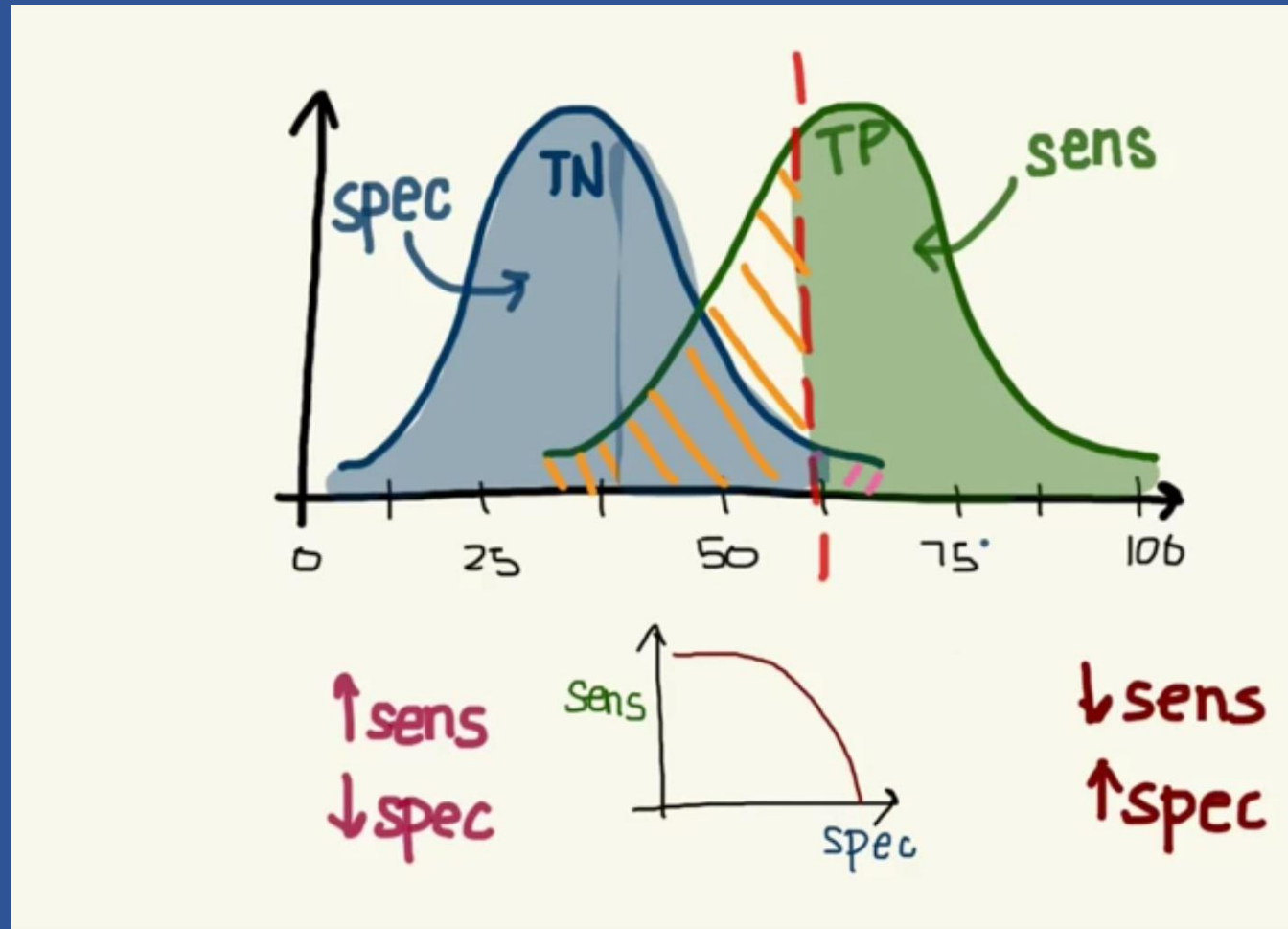
ML Evaluation

Move cutoff to right Sens-- / Spec++



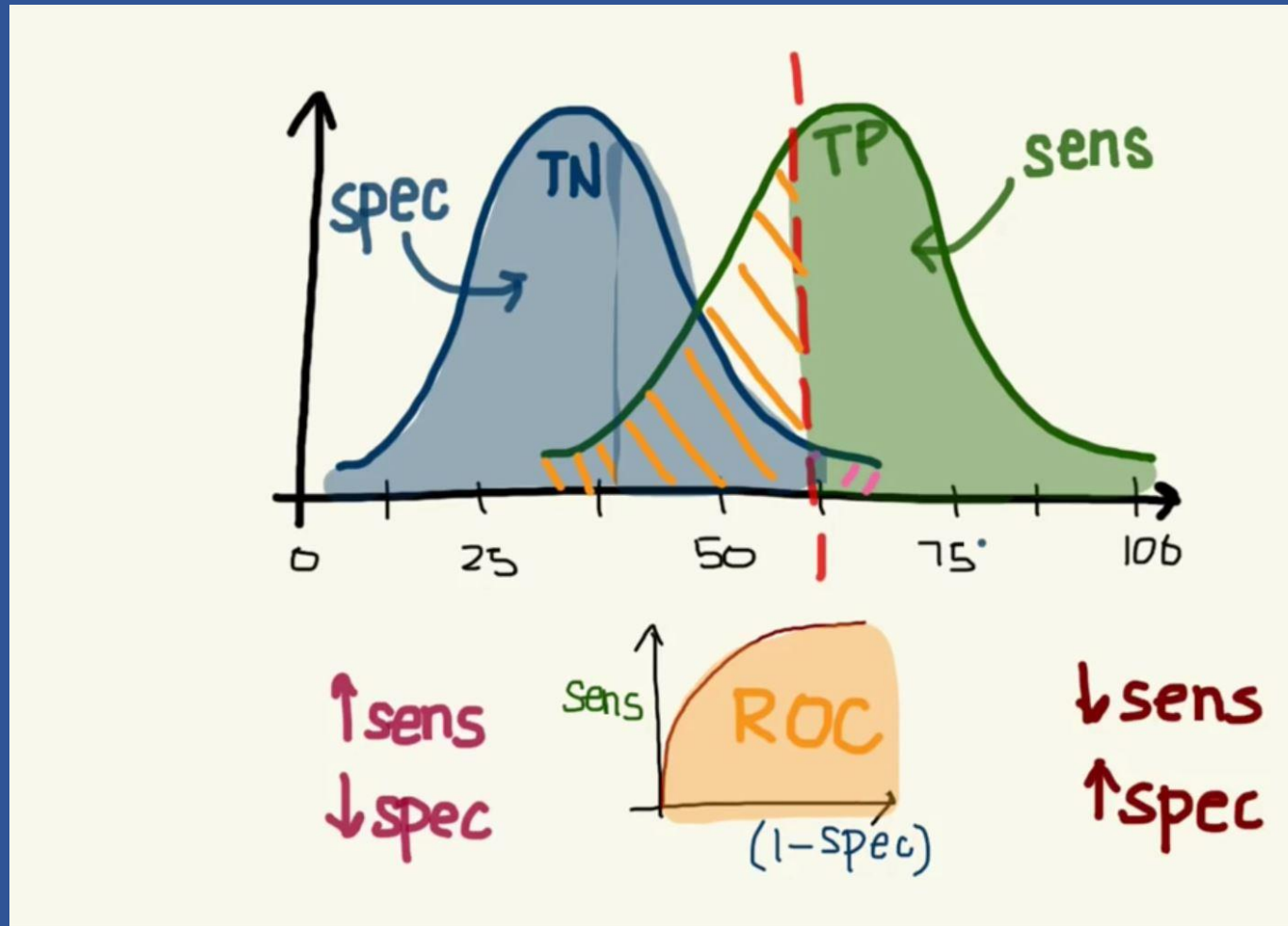
ML Evaluation

Chart proportion of Sens / Spec



ML Evaluation

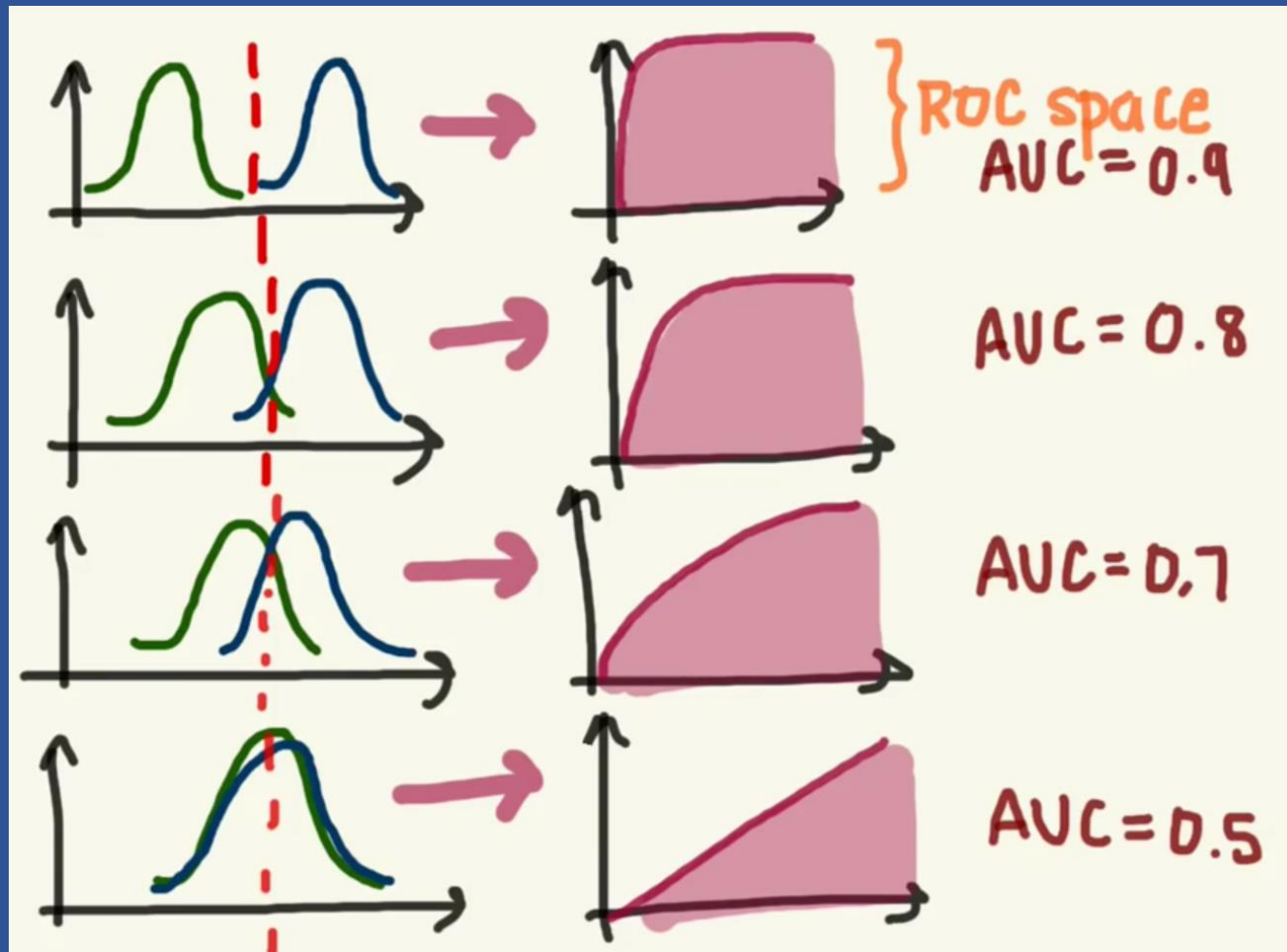
ROC curve = proportion of Sens / (1 – Spec)



ML Evaluation

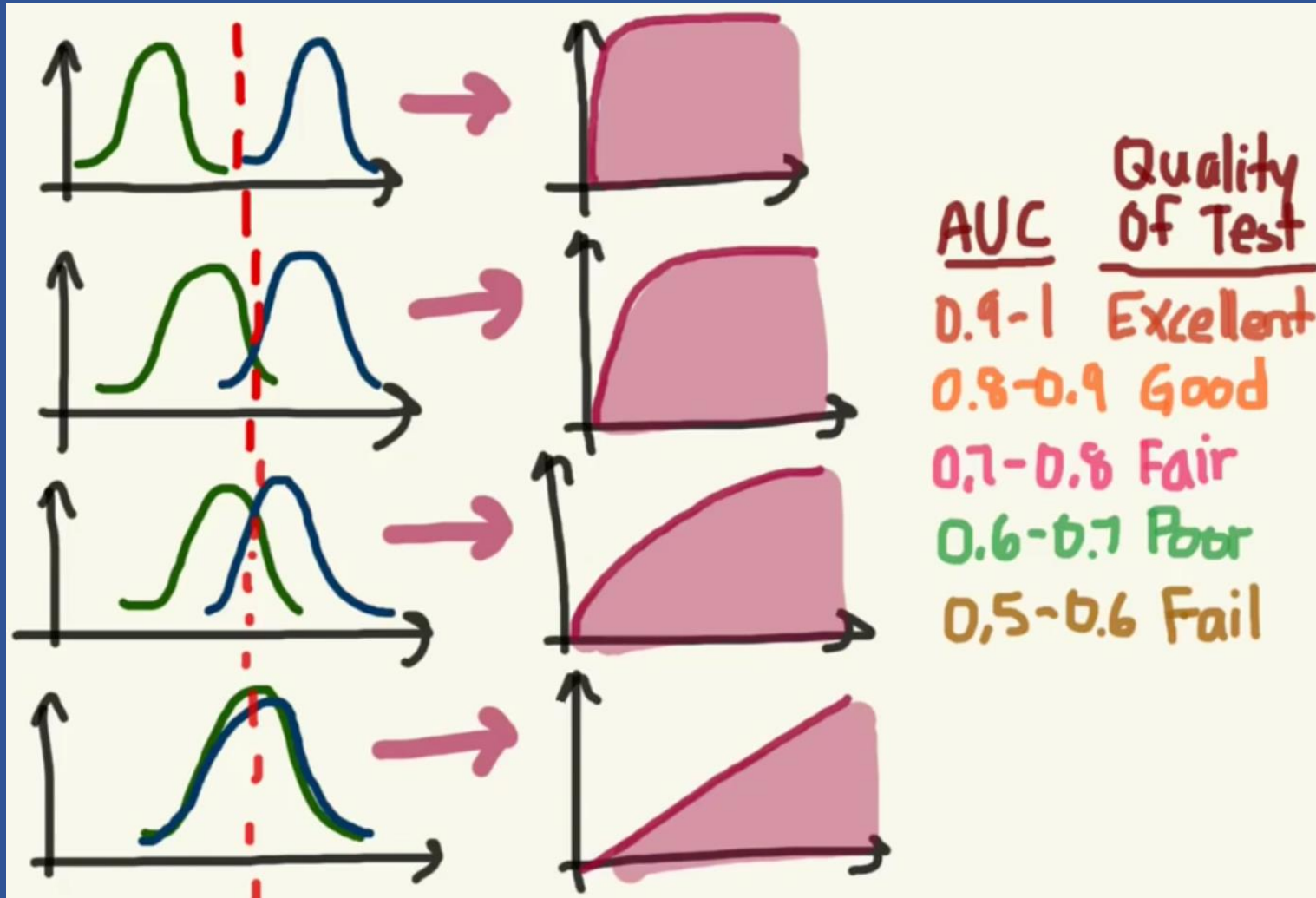
Area Under the Curve (AUC)

AUC is used to determine which of the used models predicts the classes best.



ML Evaluation

AUC score



ML Evaluation

PRECISION/RECALL

Precision: the number of items correctly predicted as belonging to that class divided by the total number of items predicted as belonging to the class. $TP / (TP + FP)$

Recall: the number of items correctly predicted as belonging to that class divided by the total number of items that actually belong to the class. $TP / (TP + FN)$

ML Evaluation

Evaluation metrics

Score Bin	Positive Examples	Negative Examples	Fraction Above Threshold
(0.900,1.000]	59	8	0.251
(0.800,0.900]	3	4	0.277
(0.700,0.800]	0	1	0.281
(0.600,0.700]	0	1	0.285

Accuracy	F1 Score	Precision	Recall	Negative Precision	Negative Recall	Cumulative AUC
0.824	0.715	0.881	0.602	0.805	0.953	0.023
0.820	0.721	0.838	0.633	0.813	0.929	0.038
0.816	0.717	0.827	0.633	0.813	0.923	0.041
0.813	0.713	0.816	0.633	0.812	0.917	0.045

ML Evaluation

Evaluation metrics variable

- **True Positive (TP)**: Correctly identified e.g. Sick people correctly diagnosed as sick
- **False Positive (FP)**: Incorrectly identified e.g. healthy people incorrectly identified as sick
- **True Negative (TN)**: Correctly rejected e.g. healthy people correctly identified as healthy
- **False Negative (FN)**: Incorrectly rejected e.g. Sick people incorrectly identified as healthy
- **Accuracy** : The proportion of the total number of predictions that is correct. $(TP + TN) / (TP + TN + FP + FN)$

ML Evaluation

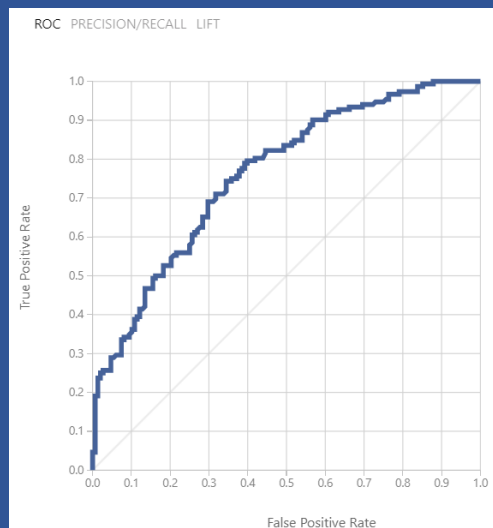
- **Precision:** is the proportion of positive cases that were correctly identified. $TP / (TP + FP)$
- **Recall:** Sensitivity or Recall is the proportion of actual positive cases which are correctly identified. $TP / (TP + FN)$
- **F1 Score:** is the harmonic mean of precision and Recall. $2TP / (2TP + FP + FN)$
- **Threshold:** Threshold is the value above which it belongs to first class and all other values to the second class. E.g. if the threshold is 0.5 then any patient scored more than or equal to 0.5 is identified as sick else healthy.

ML Evaluation

Sentiment evaluation results

- Positive Label: 1 = **Good Text (GT)**
- Negative Label: 0 = **Bad Text (BT)**
- True Positive (TP): correctly predict **GT**
- True Negative (TN): correctly predict **BT**
- False Positive (FP): incorrectly predict **GT**
- False Negative (FN): incorrectly predict **BT**

AUC 0.761



True Positive	False Negative	Accuracy	Precision	Threshold
106	46	0.690	0.693	0.5
False Positive	True Negative	Recall	F1 Score	
47	101	0.697	0.695	
Positive Label	Negative Label			
1	0			

ML Evaluation

Metrics for Binary Classification

METRICS	DESCRIPTION	LOOK FOR
Accuracy	proportion of correct predictions with a test data set	The closer to 1.00, the better
AUC	Area under the curve: This is measuring the area under the curve created by sweeping the true positive rate vs. the false positive rate.	The closer to 1.00, the better
AUCPR	Area under the curve of a Precision-Recall curve: Useful measure of success of prediction when the classes are very imbalanced (highly skewed datasets).	The closer to 1.00, the better
F1-score	the harmonic mean of the precision and recall. F1 Score is helpful when you want to seek a balance between Precision and Recall.	The closer to 1.00, the better

Next Step

Create Sentiment model using AutoML