Classification

PHYS591000 2022.02.23

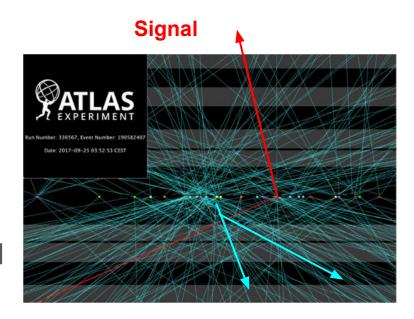
Announcement

• From this week (week 02) do *not* accept lab (homework) submission if you are absent from class without permission.

- As usual, take 3 mins to introduce yourself to your teammate for this week!
 - "How do you like the class last week?"
 - If your teammate just joined the class this week, tell your teammate what we did last week.

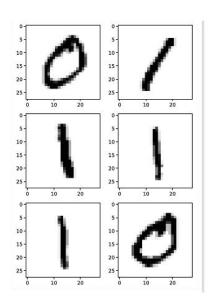
Classification

- Focus on binary classification
 - Example: Distinguish particles from hard collisions (signal) and particles from soft collisions (background) at the LHC

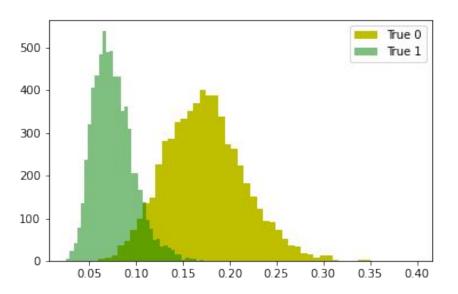


Background

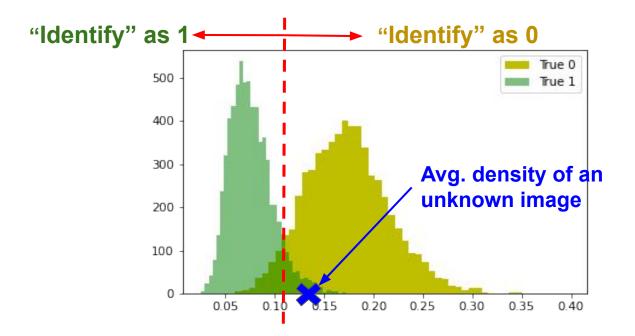
- In-class: Use the MNIST database
 - x_train: 28*28 images; each pixel
 is associated with a number from 0-255
 (~ the amount of 'ink')
- Task: Separate 0 (background) and 1 (signal)



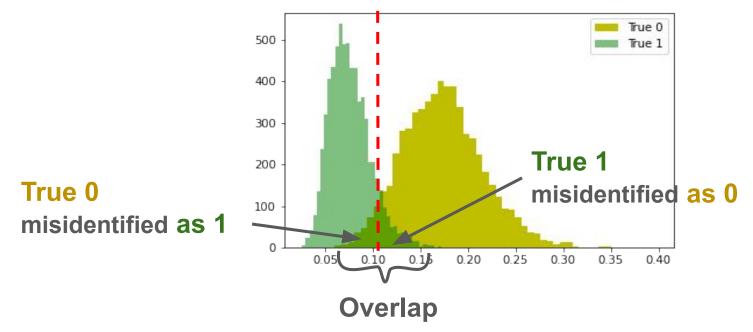
 Week 01: A useful **feature** is the average pixel density ('average amount of ink on each pixel')



Given an (unknown) image, calculate its average pixel density

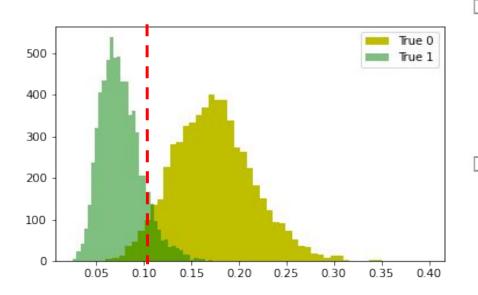


 There is an overlap of signal and background due to spreads of their distributions



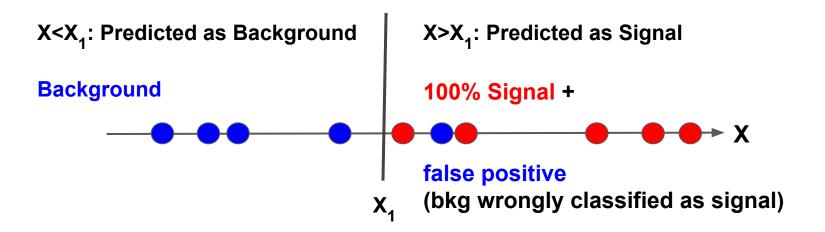
 The performance of the cut (the 'classifier') thus depends on the cut value (threshold) we choose

Courtesy of Prof. Kai-Feng Chen (NTU)

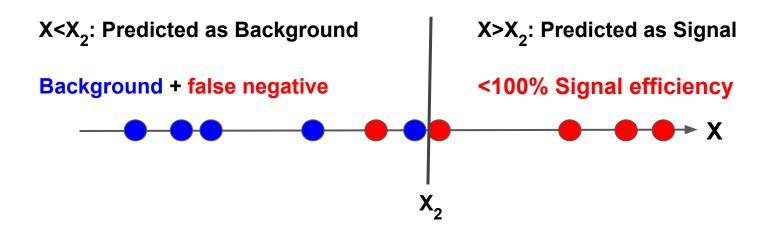


- If a threshold of 0.11 is set:
 93.0% of the "ones" are selected;
 94.5% of the "zeros" are rejected.
 (or 5.5% of the zeros are misidentified)
- If a threshold of 0.16 is set:
 99.8% of the "ones" are selected;
 61.2% of the "zeros" are rejected.
 (or 38.8% of the zeros are misidentified)

- The cut is a linear classifier which maps data into a 'score' X
- The performance depends on the cut value we choose, e.g.



- The cut is a linear classifier which maps data into a 'score' X
- The performance depends on the cut value we choose, e.g.



 One way to quantify the performance of the chosen cut is to construct the corresponding confusion matrix

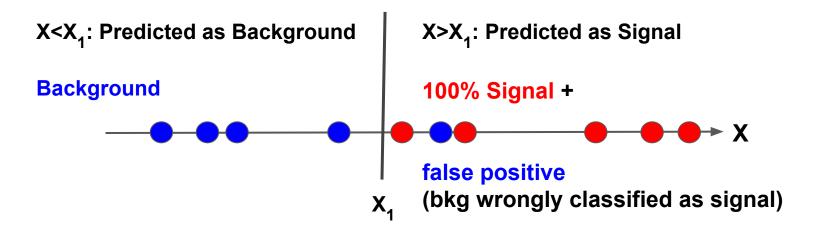
	Actual Signal	Actual Background	
Predicted as Signal	True Positives	False Positives True Negatives	
Predicted as Background	False Negatives		

One confusion matrix for each cut value

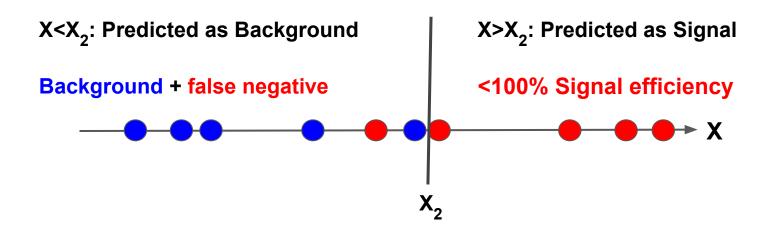
X ₁	Actual Signal	Actual Background	
Predicted as Signal	5	1	
Predicted as Background	0	4	

X_2	Actual Signal	Actual Background
Predicted as Signal	4	0
Predicted as Background	1	5

 Which cut to choose depends on what is more important to your goal: E.g. Searching for rare signals → Identify as many signals as possible → Choose X₁



 Which cut to choose depends on what is more important to your goal: E.g. Perform a precise measurement → Reject as many background as possible → Choose X₂



 There are many metrics for comparing the performances of the models based on the confusion matrix.

	Actual Signal	Actual Background	
Predicted as Signal	True Positives	False Positives	
Predicted as Background	False Negatives	True Negatives	

$$sensitivity = \frac{number\ of\ true\ positives}{number\ of\ true\ positives + number\ of\ false\ negatives}$$

 There are many metrics for comparing the performances of the models based on the confusion matrix.

	Actual Signal	Actual Background	
Predicted as Signal	True Positives	False Positives	
Predicted as Background	False Negatives	True Negatives	

$$specificity = \frac{number\ of\ true\ negatives}{number\ of\ true\ negatives + number\ of\ false\ positives}$$

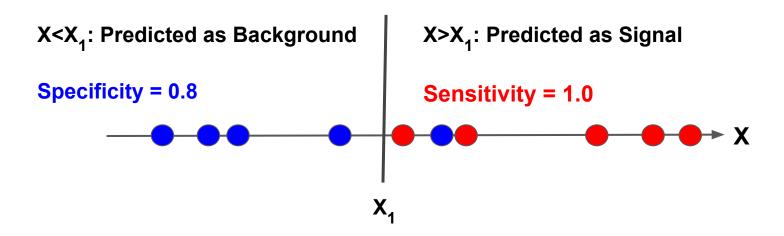
X ₁ Better Sensitivity	Actual Signal	Actual Background	
Predicted as Signal	5	1	
Predicted as Background	0	4	

X_2	Actual Signal	Actual Background
Predicted as Signal	4	0
Predicted as Background	1	5

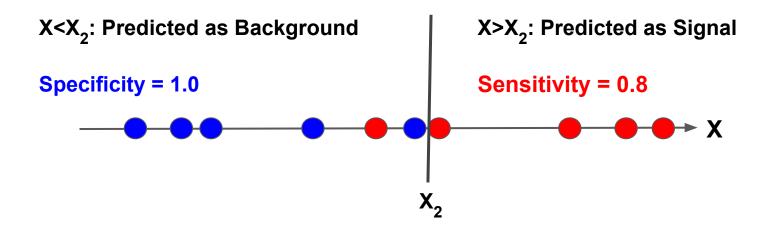
X ₁	Actual Signal	Actual Background
Predicted as Signal	5	1
Predicted as Background	0	4

X ₂ Better Specificity	Actual Signal	Actual Background	
Predicted as Signal	4	0	
Predicted as Background	1	5	

 E.g. Searching for rare signals → Correctly identify positives is (Sensitivity) more important → Choose X₁

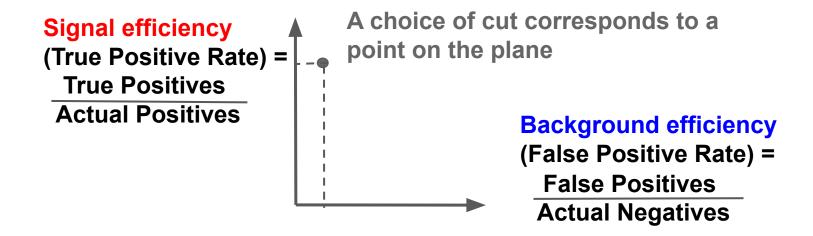


 E.g. Perform a precise measurement → Correctly identify negatives (Specificity) is more important → Choose X₂

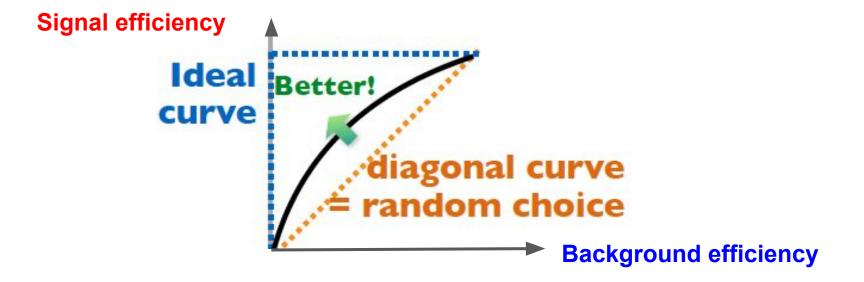


		Predicted condition		Sources	s: [1][2][3][4][5][6][7][8] view ·talk·edit
	Total population = P + N	Positive (PP)	Negative (PN)	Informedness, bookmaker informedness (BM) = TPR + TNR - 1	Prevalence threshold (PT) = √TPR×FPR – FPR TPR – FPR
Actual condition	Positive (P)	True positive (TP),	False negative (FN), type II error, miss, underestimation	True positive rate (TPR), recall, sensitivity (SEN), probability of detection, hit rate, power $= \frac{TP}{P} = 1 - FNR$	False negative rate (FNR), miss rate $= \frac{FN}{P} = 1 - TPR$
Actual	Negative (N)	False positive (FP), type I error, false alarm, overestimation	True negative (TN), correct rejection	False positive rate (FPR), probability of false alarm, fall-out $= \frac{FP}{N} = 1 - TNR$	True negative rate (TNR), specificity (SPC), selectivity $= \frac{TN}{N} = 1 - FPR$
	Prevalence = P P+N	Positive predictive value (PPV), precision = TP PP = 1 - FDR	False omission rate (FOR) = FN = 1 - NPV	Positive likelihood ratio (LR+) = TPR FPR	Negative likelihood ratio (LR-) = FNR TNR
	Accuracy (ACC) $= \frac{TP + TN}{P + N}$	False discovery rate (FDR) $= \frac{FP}{PP} = 1 - PPV$	Negative predictive value (NPV) = TN PN = 1 - FOR	Markedness (MK), deltaP (Δp) = PPV + NPV - 1	Diagnostic odds ratio (DOR) = LR+ LR-
	Balanced accuracy (BA) $= \frac{\text{TPR} + \text{TNR}}{2}$	$F_{1} \text{ score}$ $= \frac{2PPV \times TPR}{PPV + TPR} = \frac{2TP}{2TP + FP + FN}$	Fowlkes–Mallows index (FM) = √PPV×TPR	Matthews correlation coefficient (MCC) = √TPR×TNR×PPV×NPV - √FNR×FPR×FOR×FDR	Threat score (TS), critical success index (CSI), Jaccard index = TP TP + FN + FP

 A good way to summarize all confusion matrices is the ROC curve (receiver operating characteristic curve)

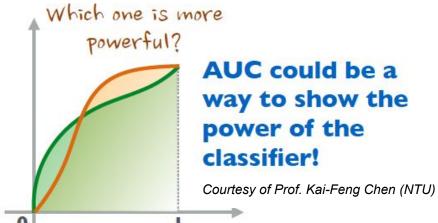


 The ROC curve illustrates the ability of the binary classifier when the discrimination threshold is varied.

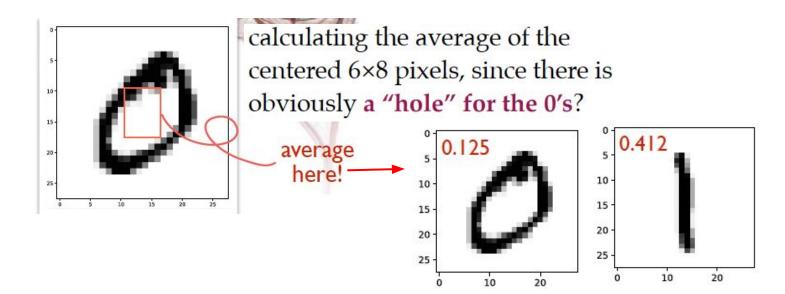


ROC curves can be used to compare two or more classifiers:
 The more it bends away from the diagonal line, the better its performance is.

 Another way to estimate the performance is the AUC (area under the curve), which varies from 0.5 (diagonal line) to 1.0 (ideal case)



 Can we do better by using more information, e.g. take the average density in the center of the image?



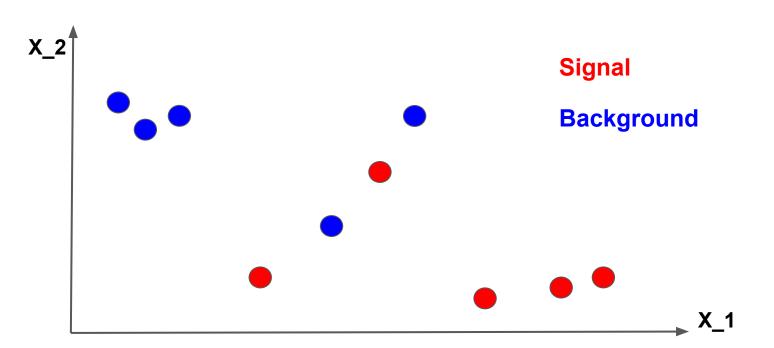
- Let's turn to this week's <u>in-class exercise</u> (Join Competition → code → inClass-exercise-02) and
 - Compute and plot the average center density
 - Compare the ROC curves of the two classifier (one with full average and another with center average)
 - 'Combine' the two distributions (full and center averages) and construct a more powerful classifier.

 We will use the Scikit-Learn, which is a machine learning library with Python: http://scikit-learn.org/

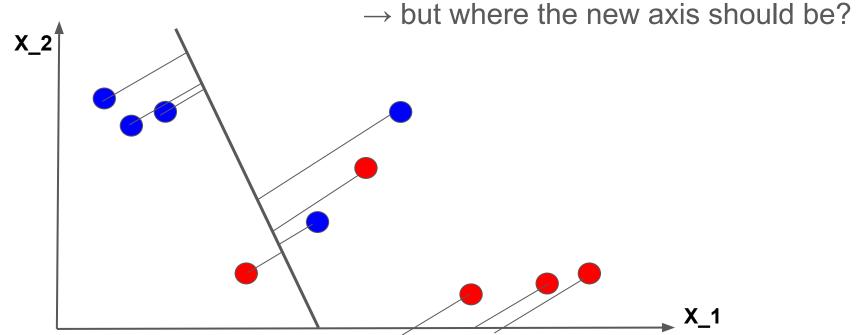
How to combine the two distributions (full and center averages)
 and construct a new, more powerful linear classifier?

→ Linear Discriminant Analysis (LDA)

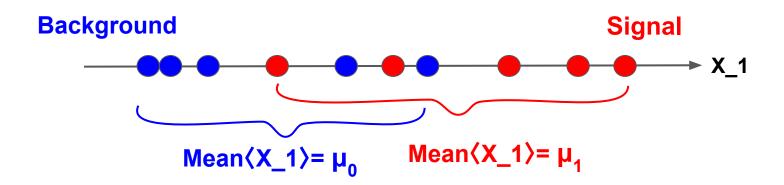
Want to transform the 2D information into a number (1D)



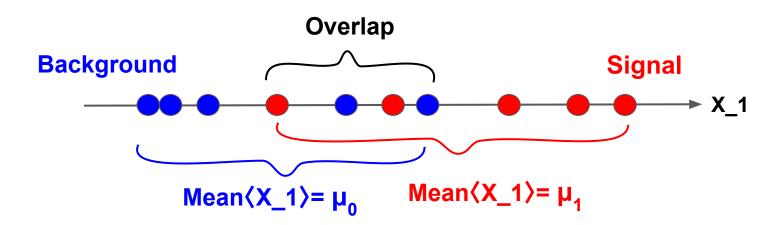
LDA: find a new axis and projects all data onto the new axis



 Recall that a variable X_1 can be used to separate signals and backgrounds when they have different means of X_1 distributions

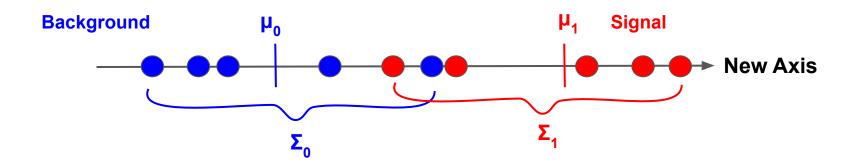


 There is an overlap of signals and backgrounds due to their spreads of X_1 distributions



- Better separation power means
 - Larger difference of the means between the signal and the background
 - Smaller spreads of the distributions of the signal and the background

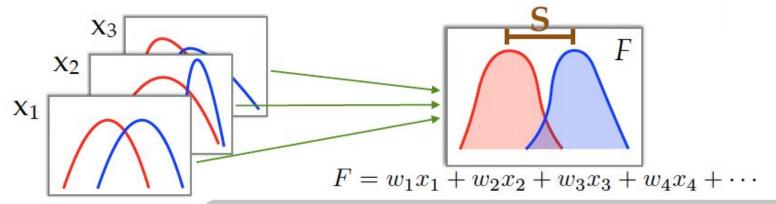
- LDA creates the new axis by combining all distributions with optimized weights which
 - Maximize the distance between the means (µ)
 - Minimize the spreads ("covariance" Σ) within each category



Performing LDA 'by hand': Fisher's discriminant

Courtesy of Prof. Kai-Feng Chen (NTU)

Now let's practice the easiest/simplest algorithm: Linear discriminant analysis (LDA), or even simpler, the Fisher's discriminant, by combining the multiple features into one variable:



Calculate the weights (\mathbf{w}_i) to maximize the separation \mathbf{S} .

LDA with Fisher's discriminant

Courtesy of Prof. Kai-Feng Chen (NTU)

- Consider a set of observables: $\overrightarrow{x} = (x_1, x_2, x_3, \cdots)$
- For 2 different event classes, the **mean** and **covariance** of the observables are: $\overrightarrow{\mu}_0$, $\overrightarrow{\mu}_1$, Σ_0 , Σ_1

$$\overrightarrow{\mu} = \langle \overrightarrow{x} \rangle$$

$$\Sigma = \langle (\overrightarrow{x} - \overrightarrow{\mu}) \cdot (\overrightarrow{x} - \overrightarrow{\mu})^T \rangle$$

 $\begin{array}{c} \overrightarrow{\mu} = \langle \overrightarrow{x} \rangle & \Sigma = \langle (\overrightarrow{x} - \overrightarrow{\mu}) \cdot (\overrightarrow{x} - \overrightarrow{\mu})^T \rangle \\ \hline \blacksquare \text{ The separation } S \text{ is given by} & \text{distance of } \mu \text{ (large)} \\ S = \frac{(\overrightarrow{w} \cdot \overrightarrow{\mu}_1 - \overrightarrow{w} \cdot \overrightarrow{\mu}_0)^2}{\overrightarrow{w}^T \Sigma_1 \overrightarrow{w} + \overrightarrow{w}^T \Sigma_0 \overrightarrow{w}} & \text{covariance } \Sigma \text{ (small)} \end{array}$

■ The optimal weights can be determined by maximizing the *S*:

$$\overrightarrow{w} \propto (\Sigma_0 + \Sigma_1)^{-1} (\overrightarrow{\mu}_1 - \overrightarrow{\mu}_0)$$

With Scikit-learn:

```
from sklearn.discriminant_analysis import LinearDiscriminantAnalysis
```

Courtesy of Prof. Kai-Feng Chen (NTU)

LDA in Scikit-learn

 In Scikit-learn LDA makes predictions by estimating the probability of an event belongs to each class, assuming the probability of each class is Gaussian and shares the same covariance.

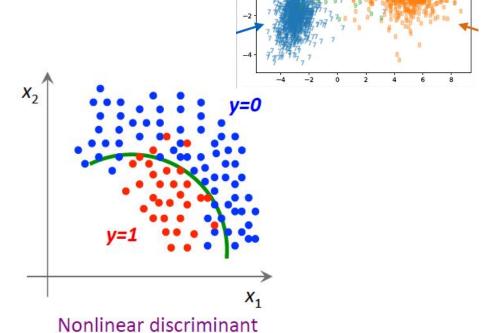
The predicted class is the one with the highest probability.

 Let's go back to the in-class exercise and play with LDA using Scikit-learn!

Outlook

What if we want to do multiclass classification?

Or if we need a Non-linear Discriminant?



Outlook

What if we want to do multiclass classification?

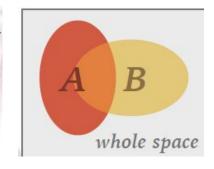
Or if we need a Non-linear Discriminant?

 We will make use of more sophisticated algorithms such as support vector machines (SVM), decision trees, neural networks... next time!

LDA in Scikit-learn

- The probability is calculated using Bayes' Theorem
- Then the conditional probability, *P*(*A* | *B*), the probability that an elementary event, known to belong to the set *B*, and is also a member of set *A*:

$$P(A \text{ and } B) = P(A|B)P(B) = P(B|A)P(A)$$



Bayes theorem
$$P(A|B) = P(B|A) \cdot P(A)/P(B)$$

Courtesy of Prof. Kai-Feng Chen (NTU)