

# CLASSIFICATION OF PULSES IN THE LUX-ZEPLIN DARK MATTER DETECTOR

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Elisa Ghetti

Kai Jenkins

# LUX-ZEPLIN DETECTOR

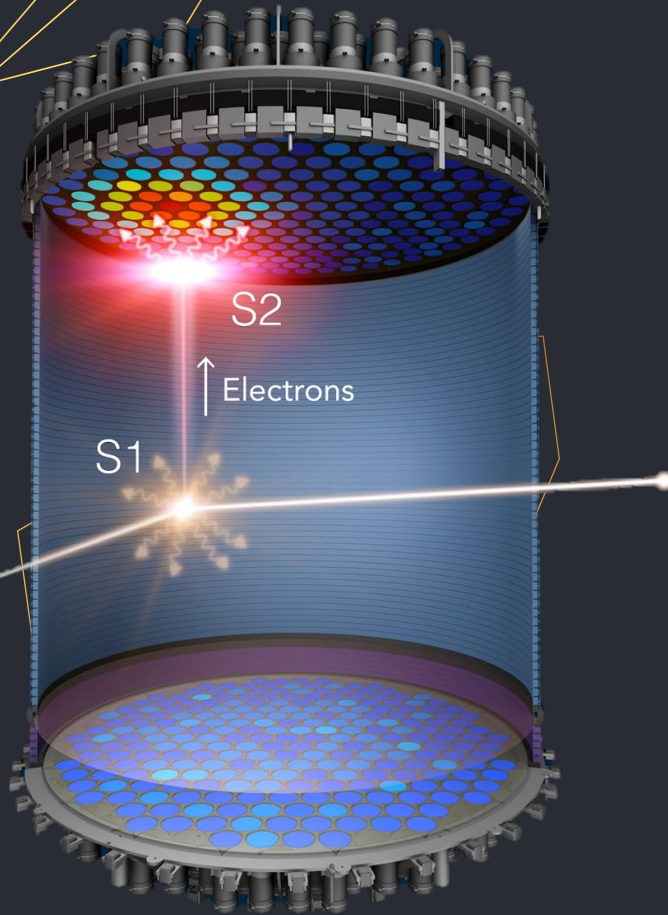
Direct detection of dark matter based on liquid xenon scintillator

The interaction of an incident particle produces two signals:

- **S1**: scintillation light
- **S2**: electroluminescence light

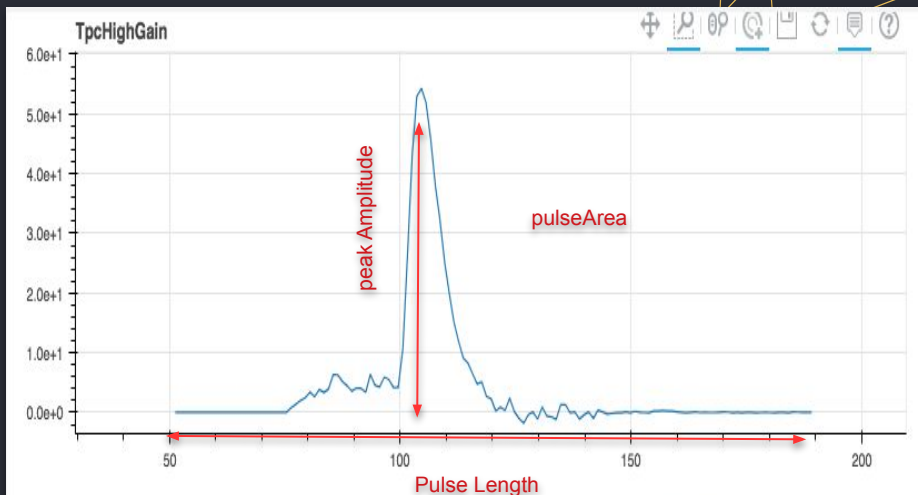
Machine learning algorithms can be used to discriminate between them.

**Goal:** reach **>99%** overall classification accuracy



## Classifier Input - 20 features (RQs):

- Pulse area (**pA**)
- Pulse amplitude (**pH**)
- Pulse length (**pL**, **pL90** - length at 90% area)
- Prompt fraction (**pF**) - fraction of area at start of pulse: 50, 100, 200, 500, 1k, 2k and 5k ns window
- Top-bottom asymmetry (**TBA**)
- Area fraction time (**aft**) time when pulse reaches X% of total area: 5%, 25%, 50%, 75%, 95%
- Peak Time (**pHT**) Time of maximum
- RMS Width (**pRMSW**)
- (**coincidence**) Number of channels that had non-zero contribution to pulse



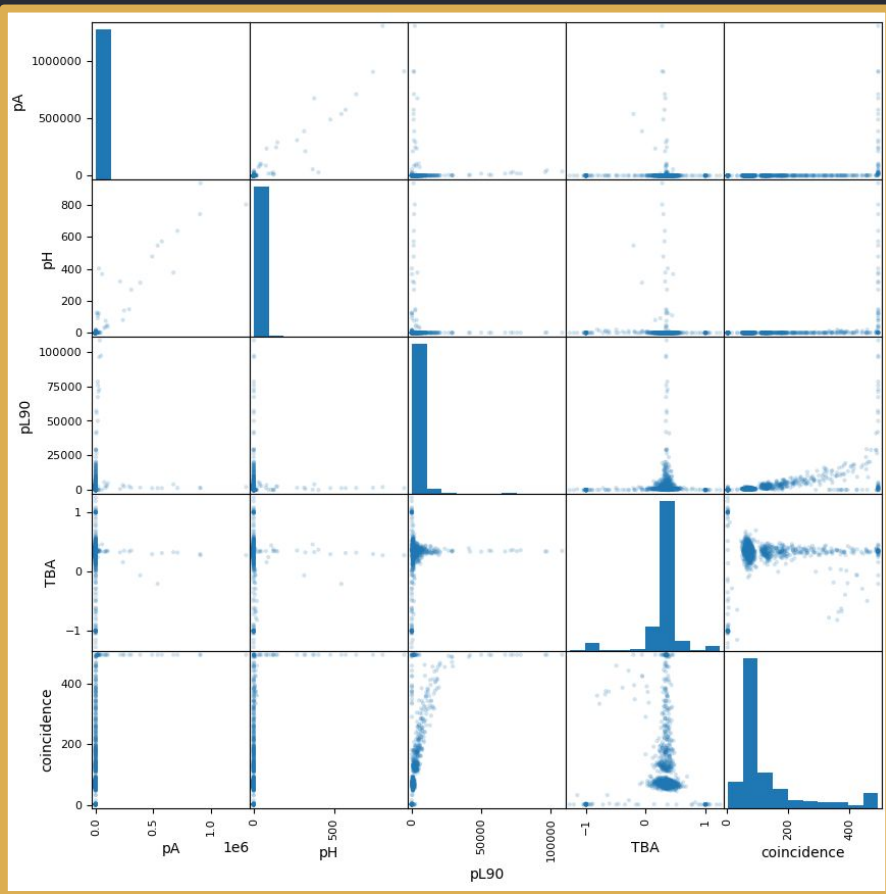
## Classifier Output - 4 classes:

[0] Other

[1] S1 (scintillation)

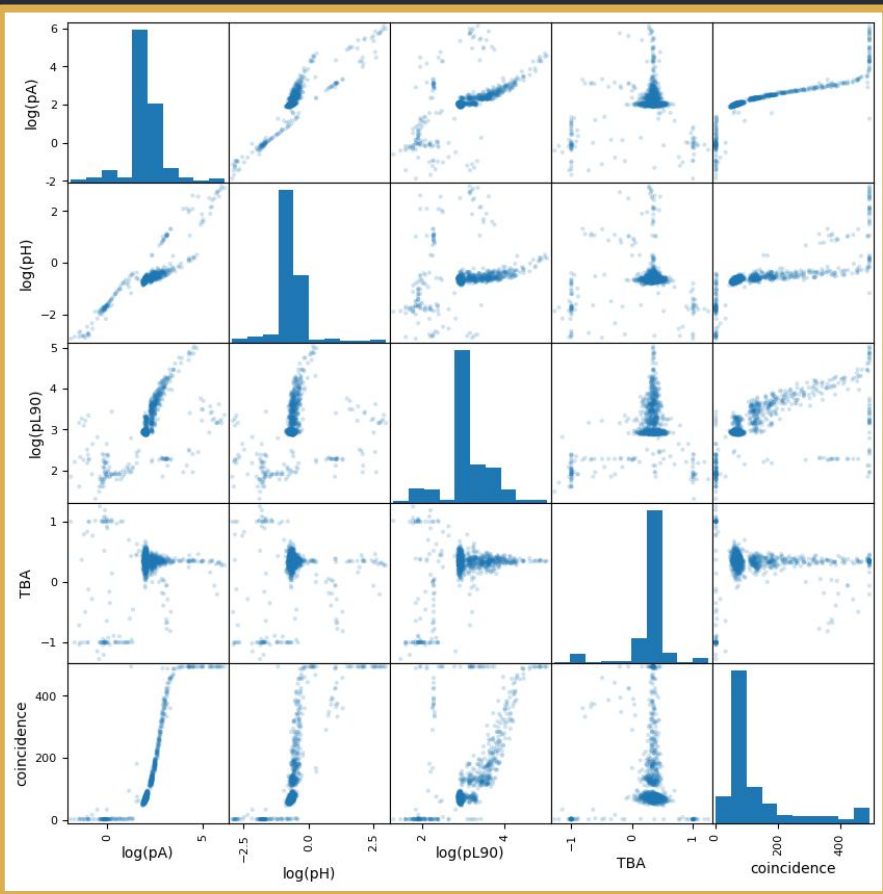
[2] S2 (electroluminescence)

[3] SE (single electron)



## FEATURE RESCALING

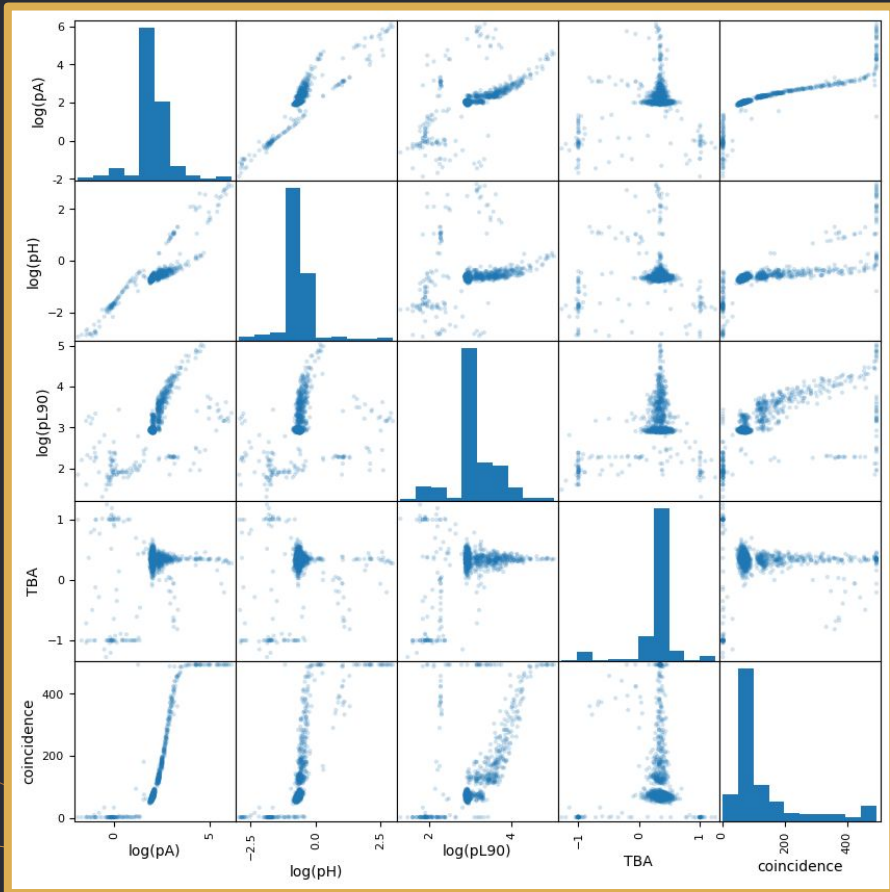
Data has to be in similar scale to avoid domination of features with larger values



## FEATURE RESCALING

Data has to be in similar scale to avoid domination of features with larger values.

$pA \rightarrow \log(pA)$   
 $pH \rightarrow \log(pH)$   
 $pL90 \rightarrow \log(pL90)$



## FEATURE RESCALING

Data has to be in similar scale to avoid domination of features with larger values.

pA  $\rightarrow$  log(pA)  
pH  $\rightarrow$  log(pH)  
pL90  $\rightarrow$  log(pL90)

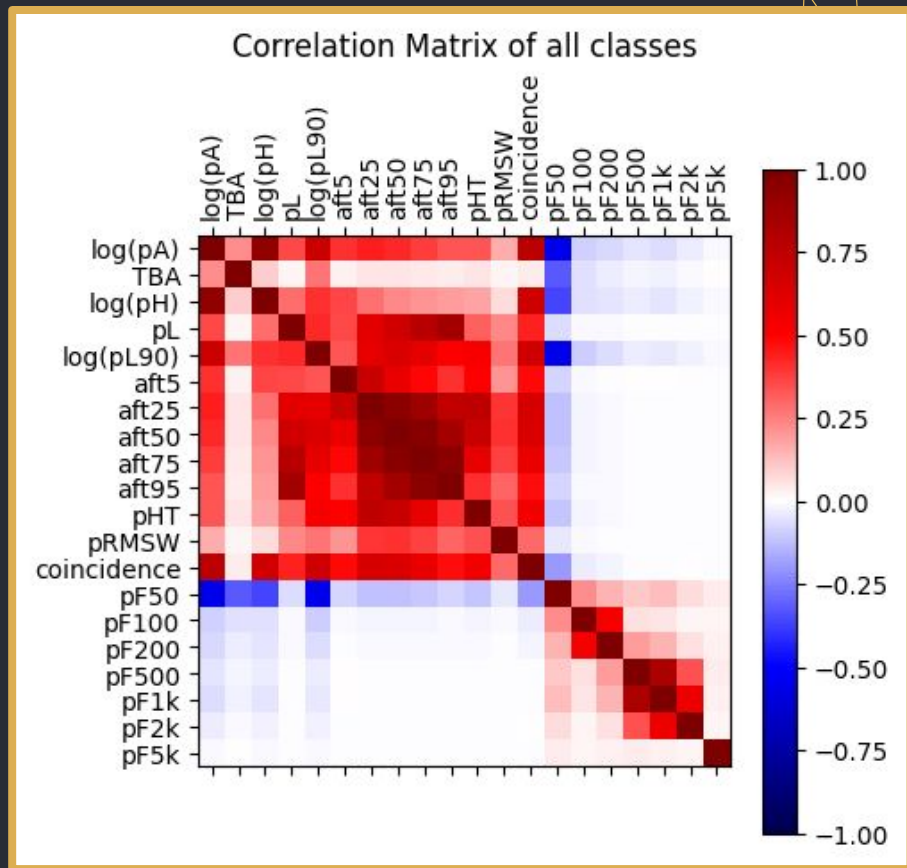
## NORMALISATION

*StandardScaler* normalisation:

- Mean = 0
- Standard deviation = 1

# CORRELATION MATRICES

Analysis of the **correlation** between features



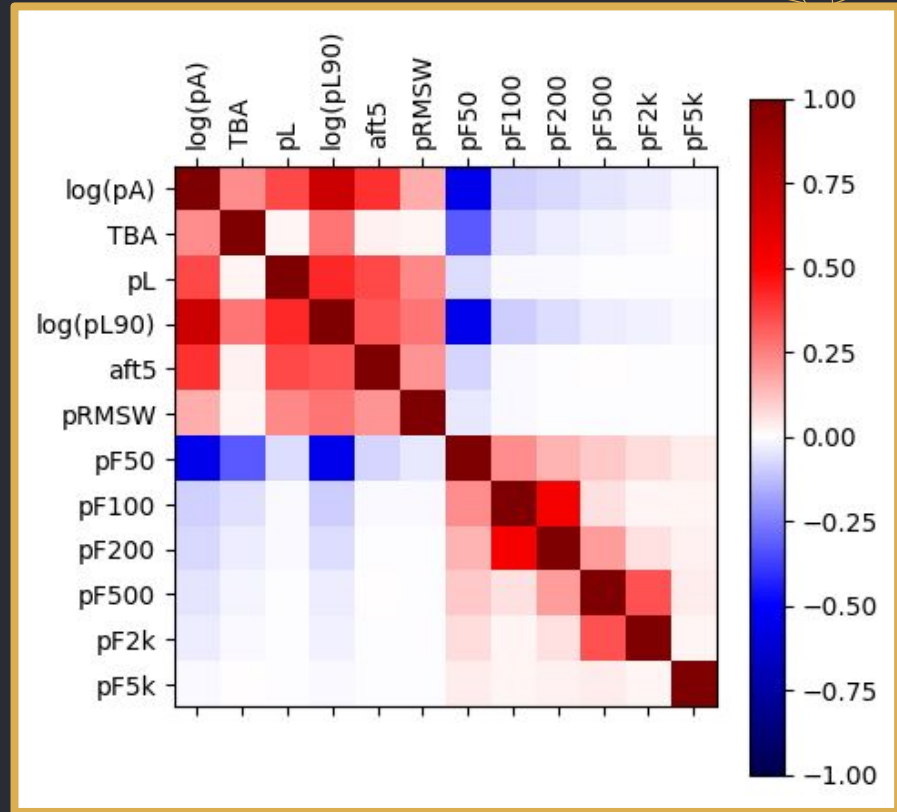
# CORRELATION MATRICES

Analysis of the **correlation** between features



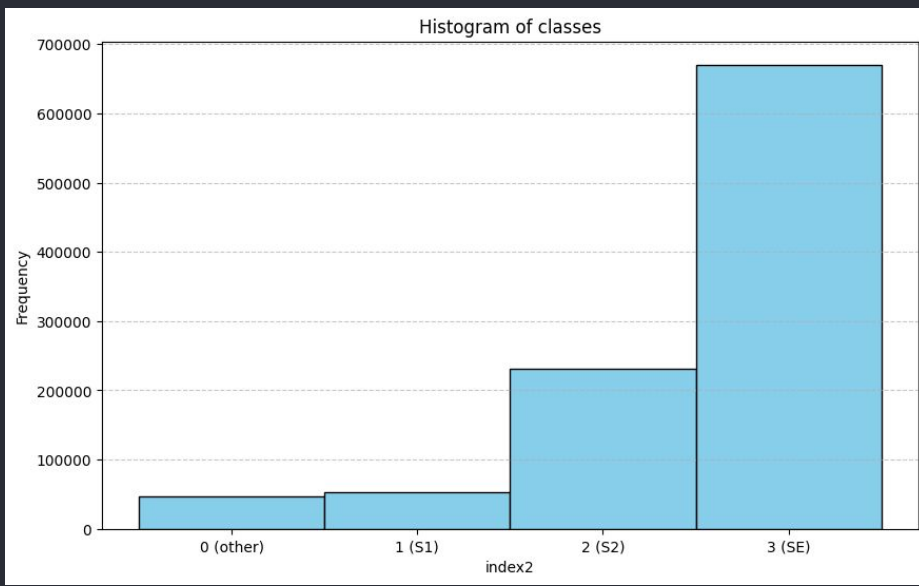
Highly correlated data can be rejected (adds no new information):

- log (pH)
- aft25, aft50, aft75, aft95
- pHT
- coincidence
- pF1k

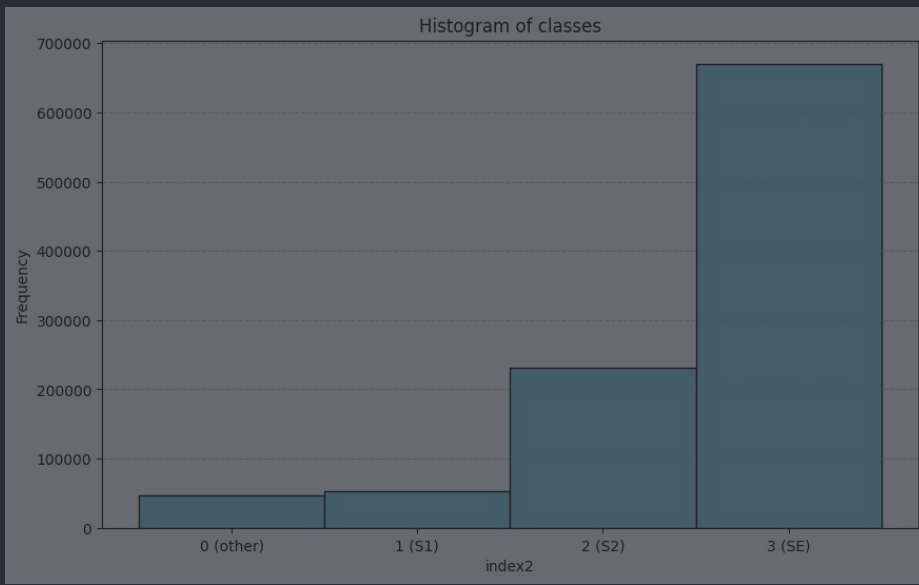




# PREPROCESSING OF THE LABELS DATASET

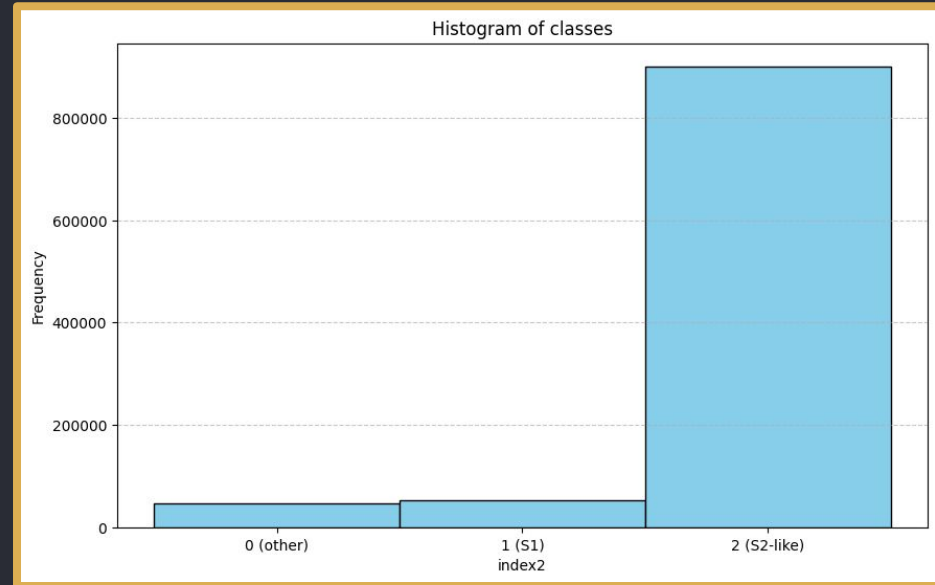


# PREPROCESSING OF THE LABELS DATASET



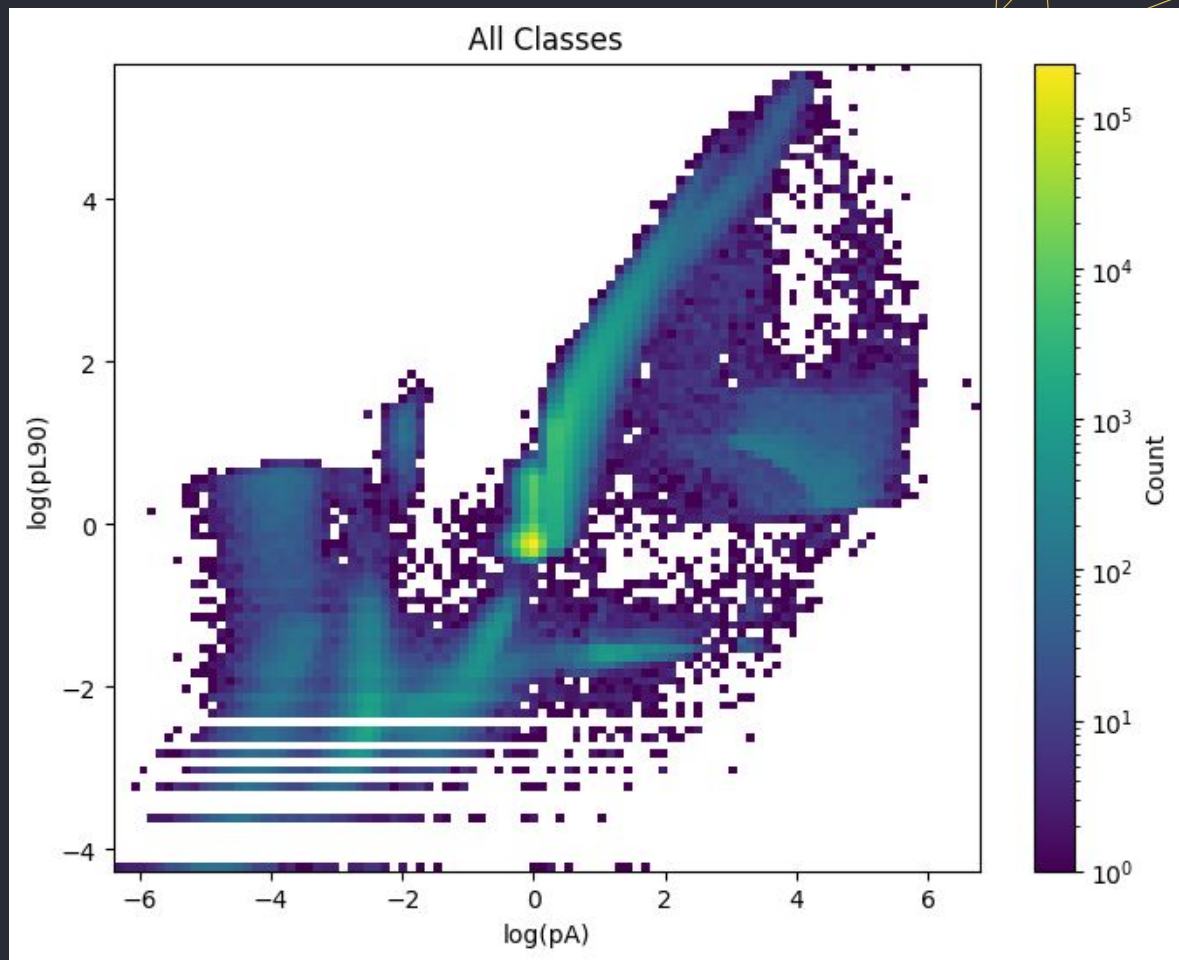
Labels S2 and SE can be combined into one S2-like label (both produced by electrons)

Note: will be applying balancing in all our models to adjusting for S2 frequency

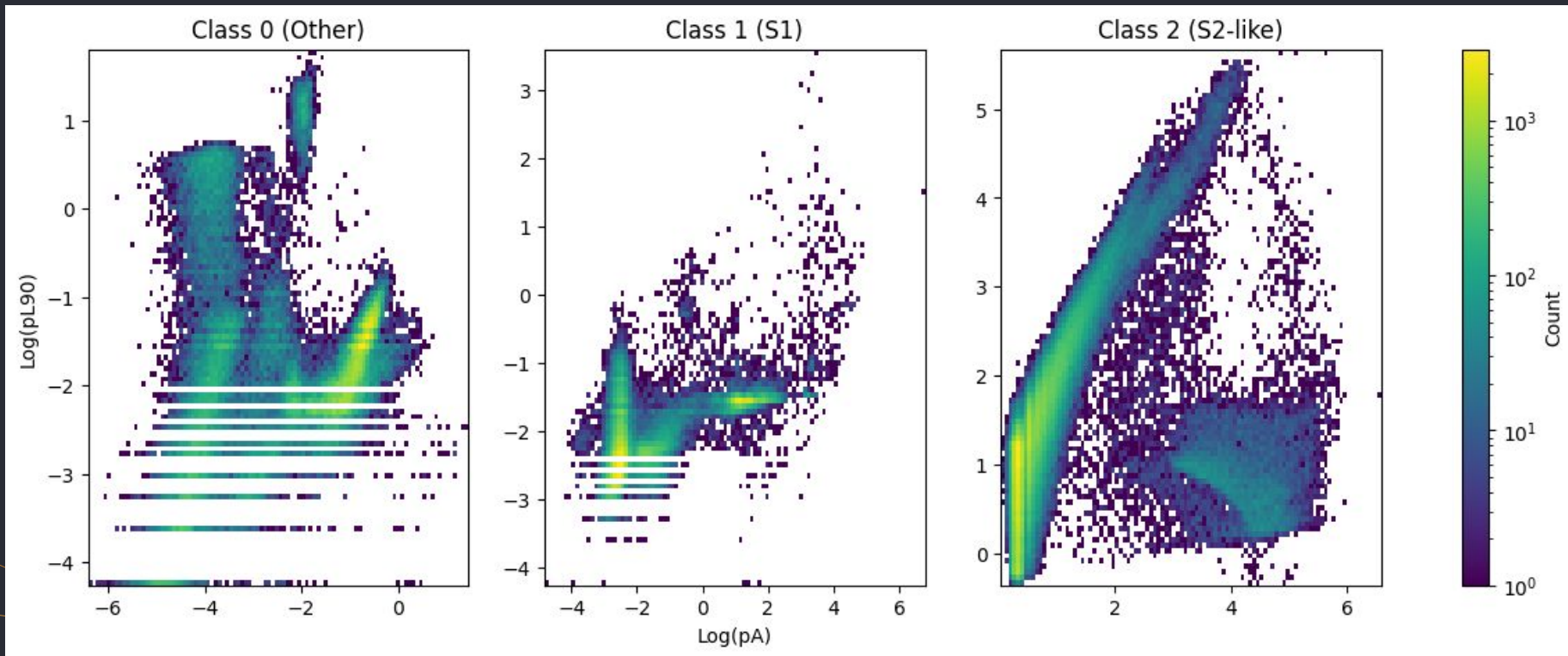


# DENSITY PLOTS

- Looked at density plots between different features to visualise their relationships
- Can already see some groups in this plot

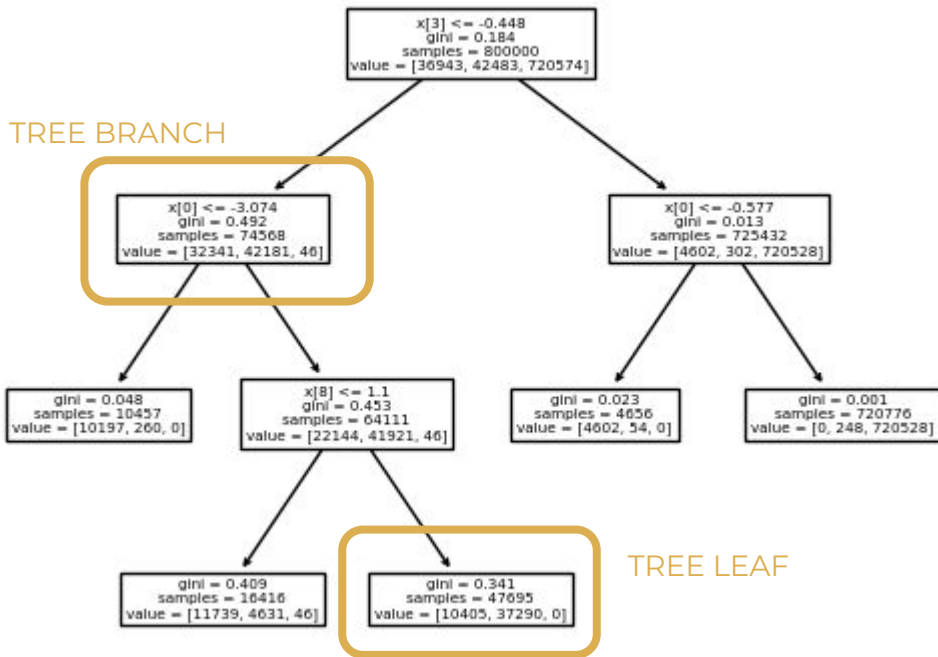


# DENSITY PLOTS (Individual classes)



# DecisionTreeClassifier MODEL

TREE BRANCH

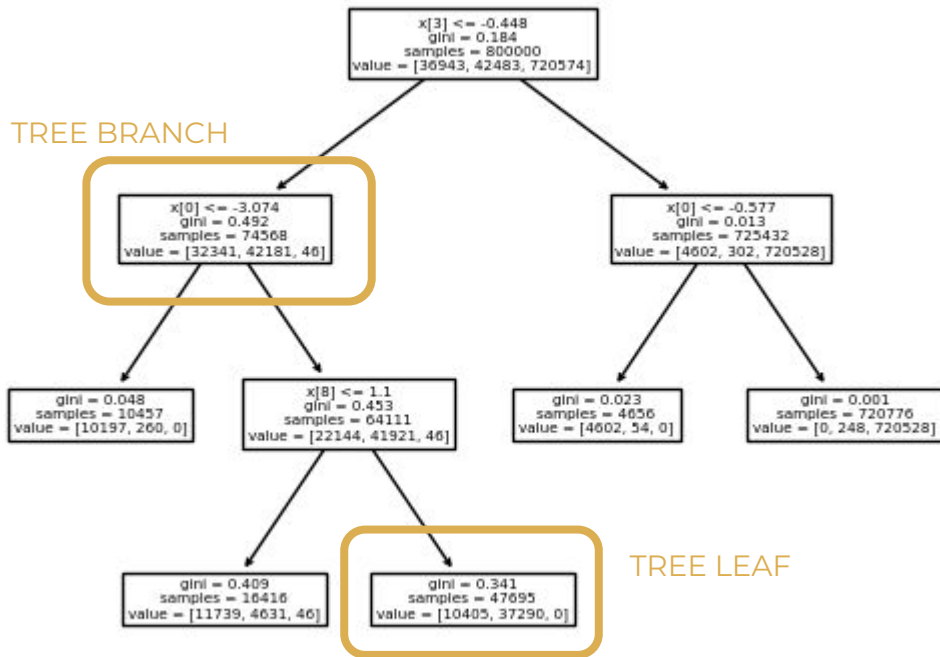


Recursive partitioning of the data based on the minimization of an impurity function

→ **GINI** impurity (likelihood of new data being misclassified if given a random class label.)

# DecisionTreeClassifier MODEL

TREE BRANCH



PARAMETERS:

- *random\_state*: set to 0 for reproducibility
- *max\_leaf\_nodes*
- *max\_depth*

NOTE:

all hyperparameters  
in this project where  
optimized with  
OPTUNA

## DecisionTreeClassifier MODEL

The model's performance can be tested by calculating the **score**:

→ Test set score: **98.80%**

A simple tree model is very simple yet powerful for a classification problem like this.

### CONFUSION MATRIX

		PREDICTED CLASS		
		0	1	2
CLASS LABEL	0	8358	830	3
	1	909	9661	45
	2	21	595	179579

# RandomForest MODEL

- Ensemble of *DecisionTrees* where output is selected by **majority vote**
- **Bootstrapping:**
  - Reduces bias
  - More resistant to **overfitting**



# RandomForest MODEL

Test set score: **99.13 %**

## CONFUSION MATRIX

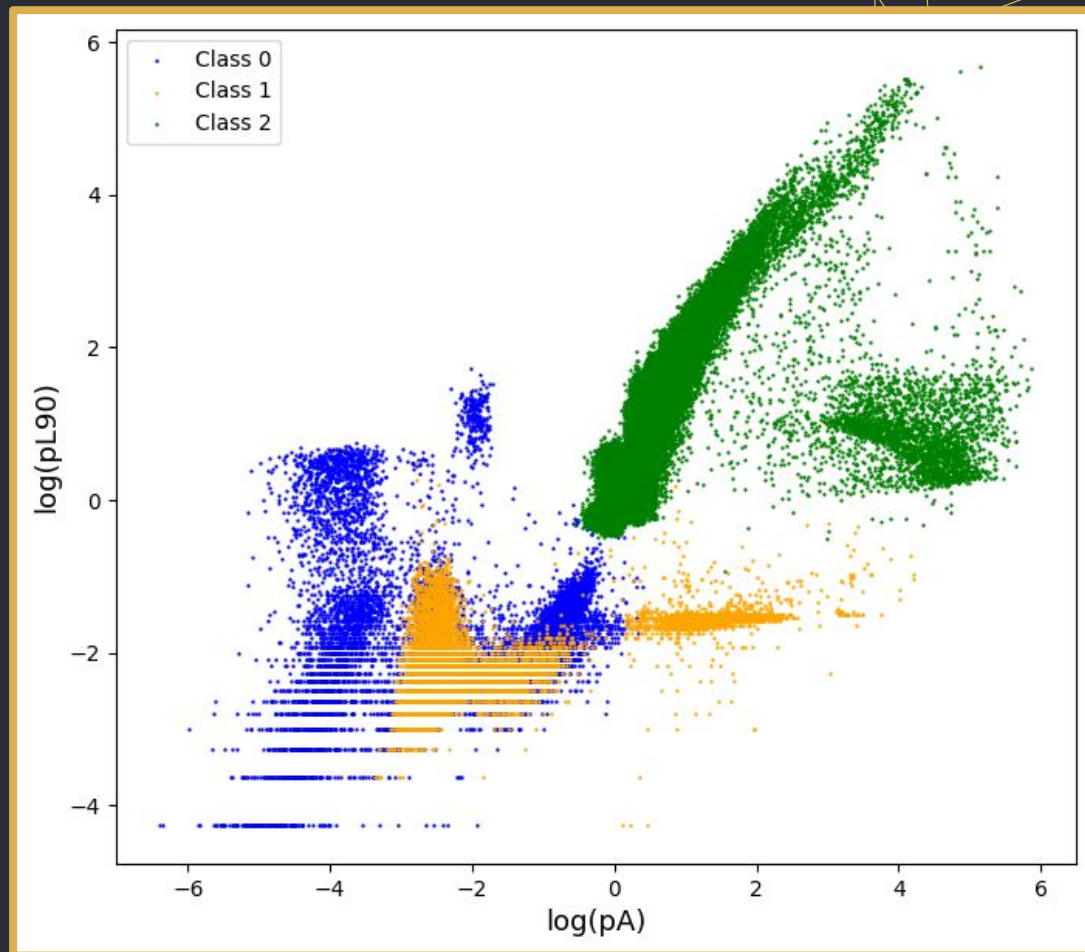
PREDICTED CLASS

0 1 2

0 8271 920 0

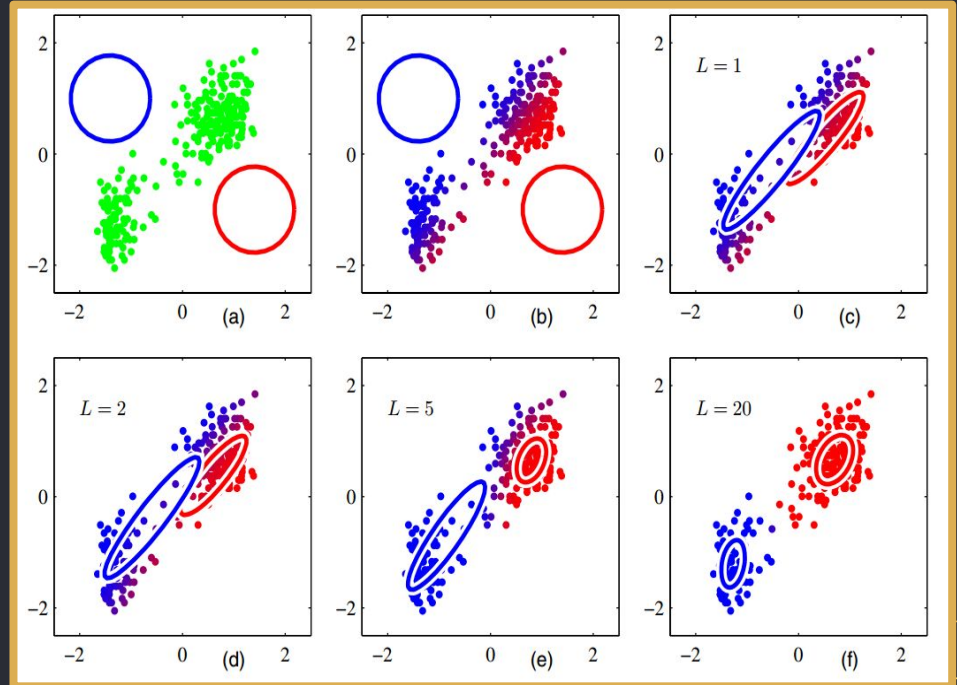
1 732 9844 39

2 44 0 180150

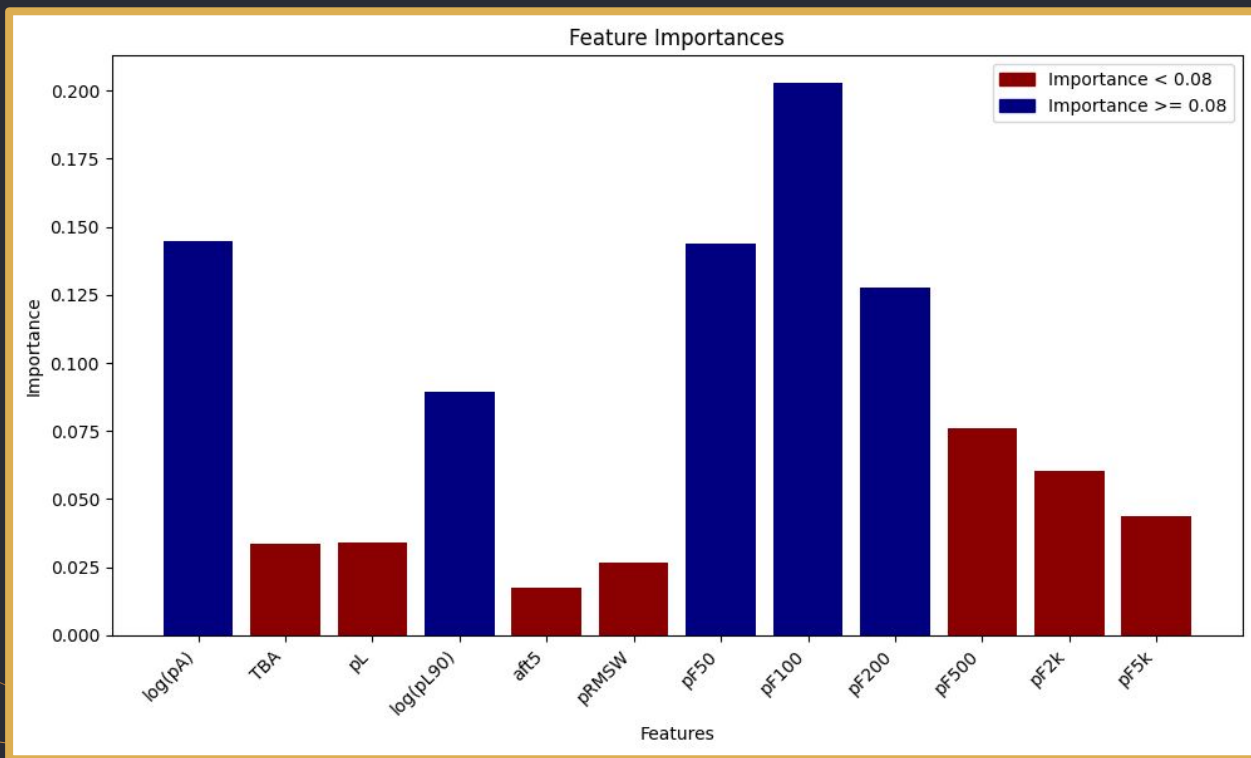


# GaussianMixture MODEL

- **Unsupervised** learning
- **Clustering analysis:** data is assumed to be distributed in a finite number of clusters
  - Linear superposition of **K gaussian distributions**



# FEATURE IMPORTANCE



**Relative importance**  
of each feature:

how much the tree  
nodes that use that  
feature reduce  
impurity on average

## *GaussianMixture* MODEL

GMM is a **density based** algorithm: a large number of components is necessary to fit less dense regions of the data

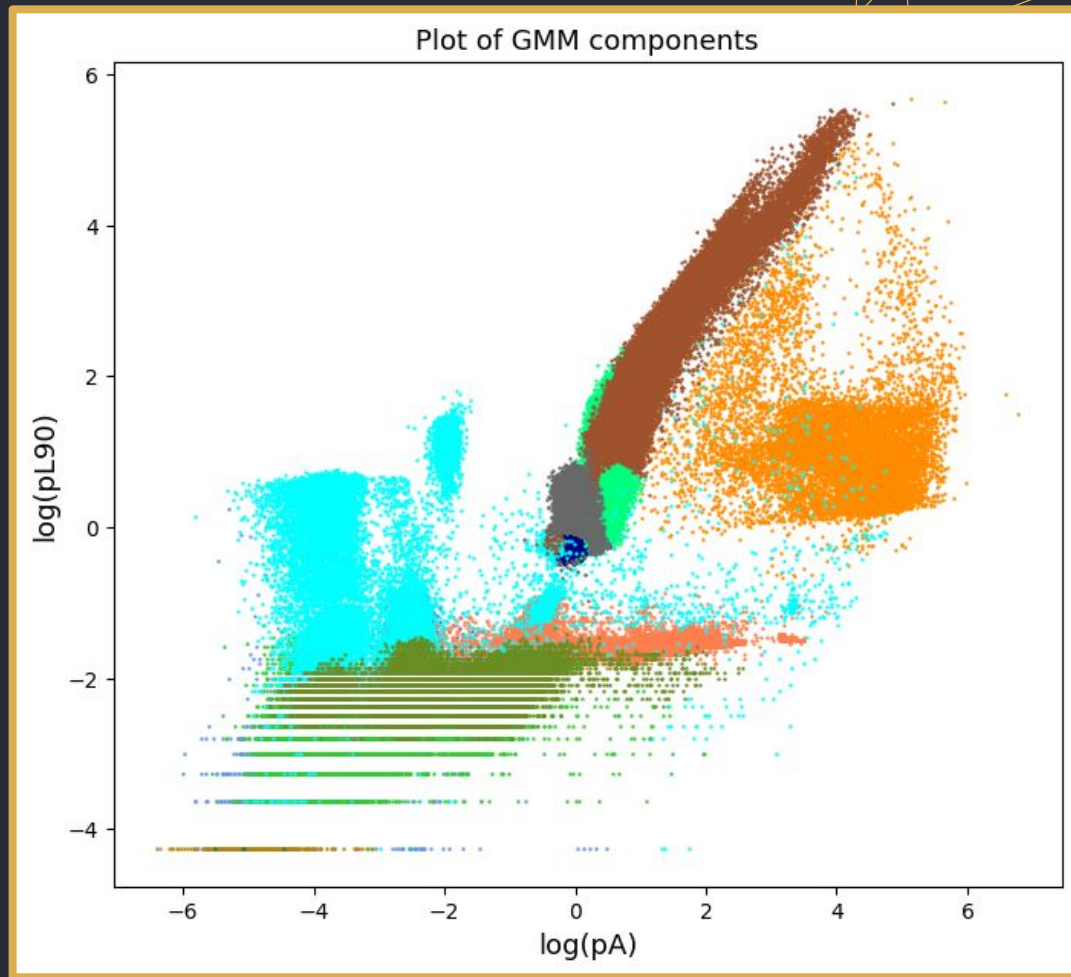
→ K has to be much larger than the number of classes

## *GaussianMixture* MODEL

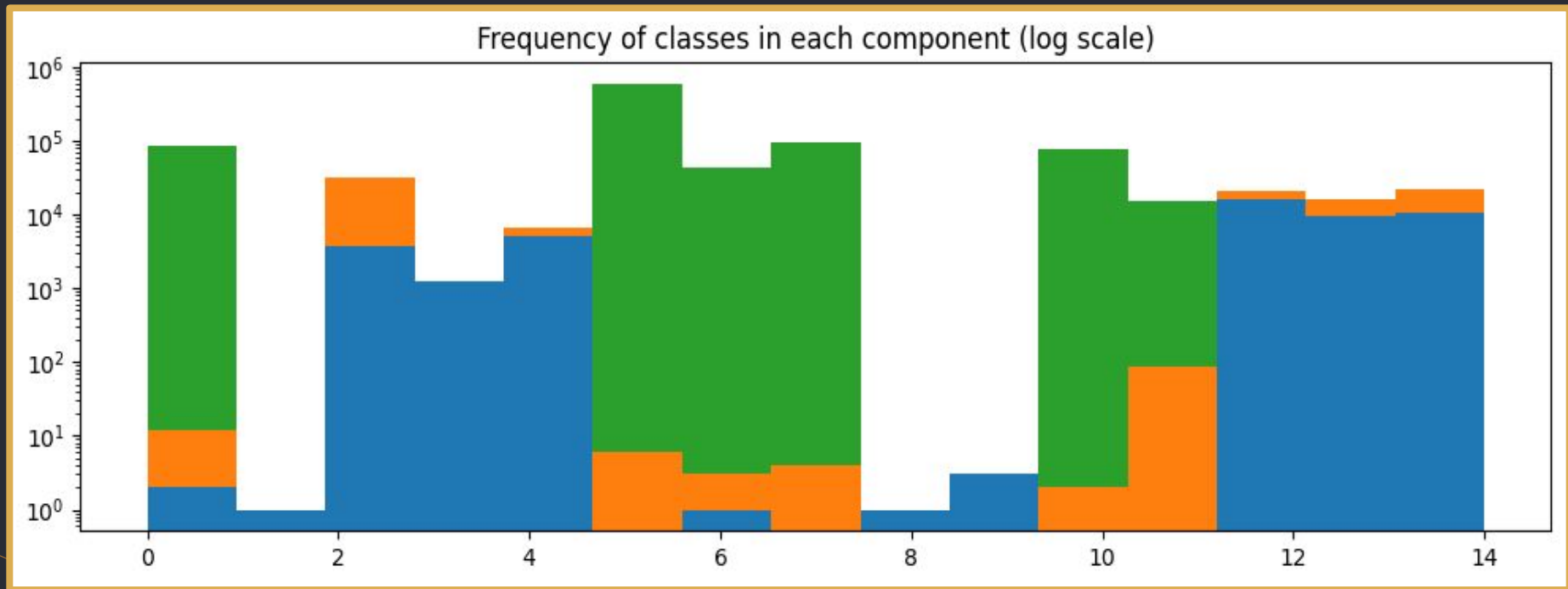
GMM is a **density based** algorithm: a large number of components is necessary to fit less dense regions of the data

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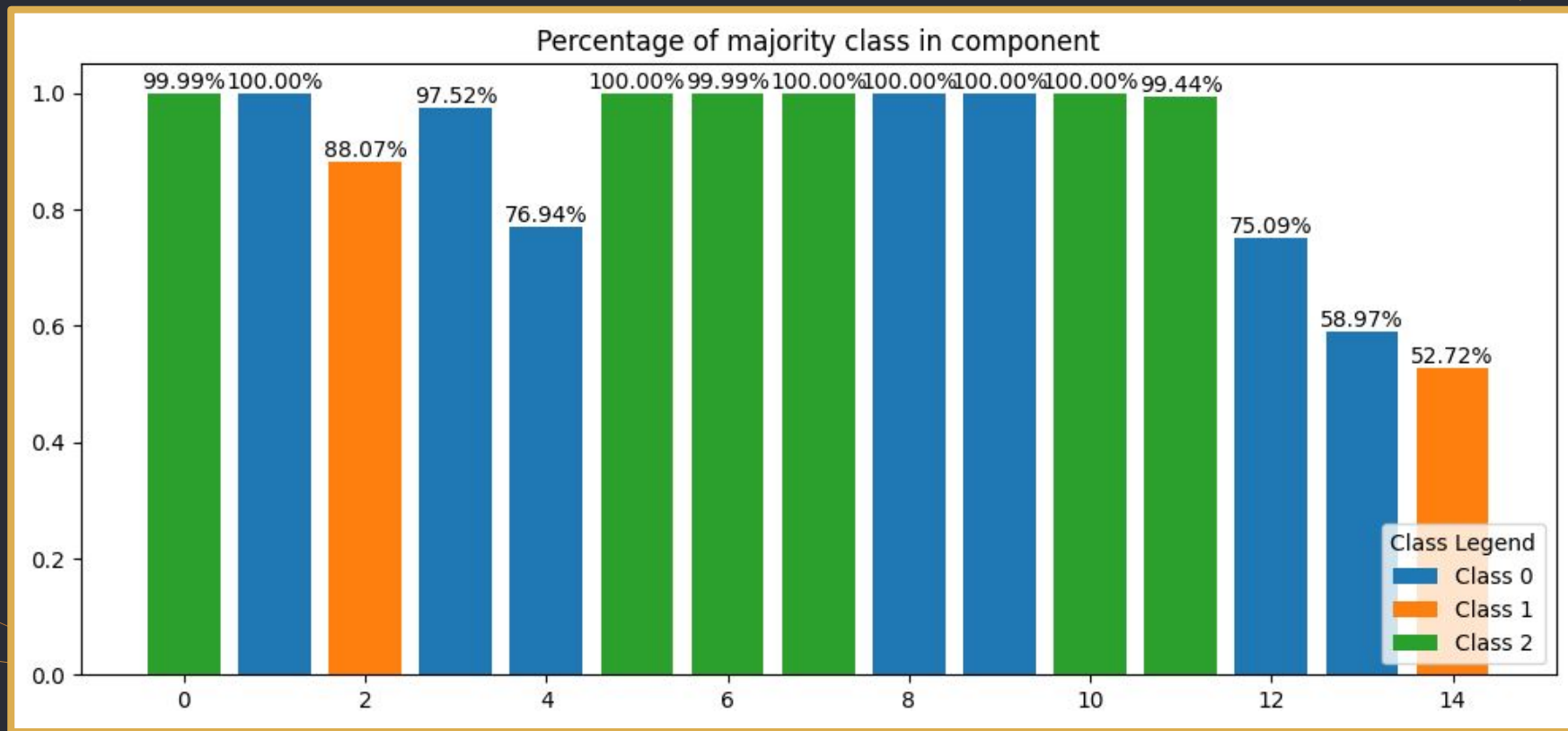
Fit with **K=15** gaussians

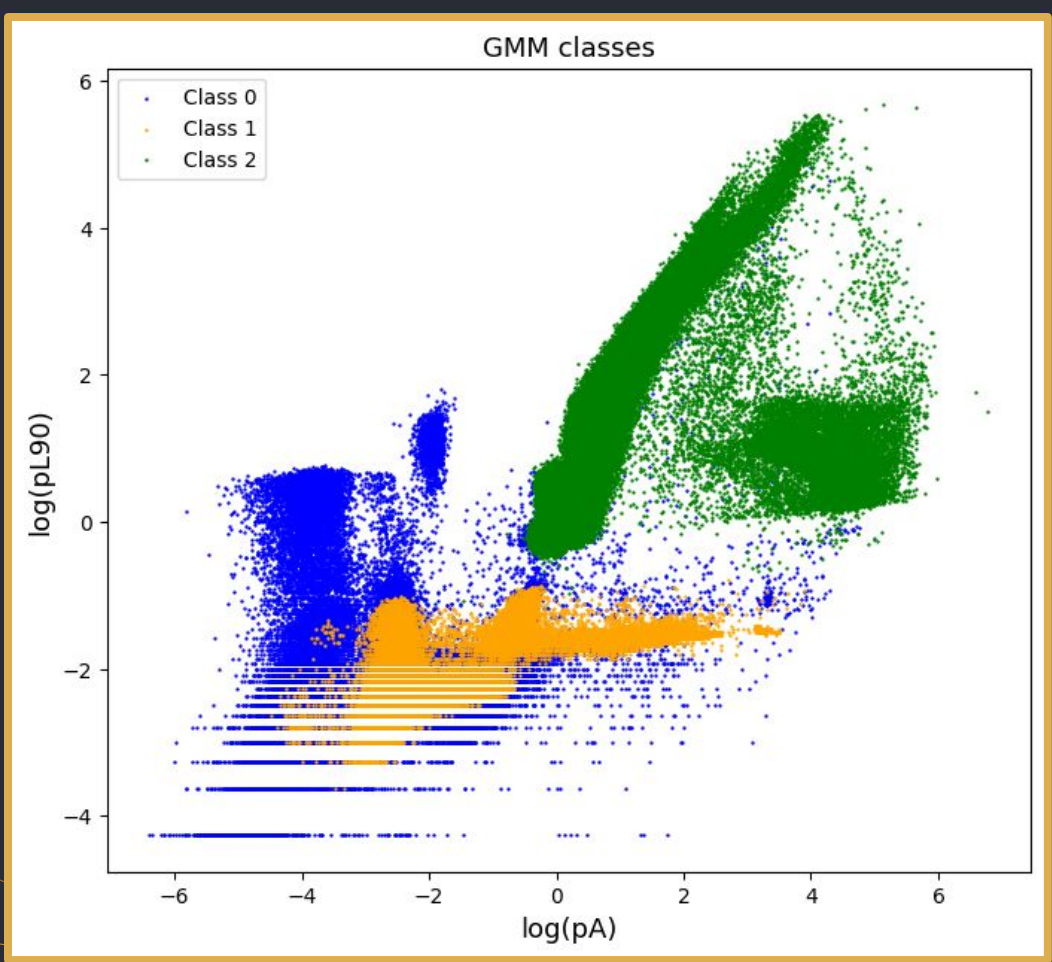


# GaussianMixture MODEL



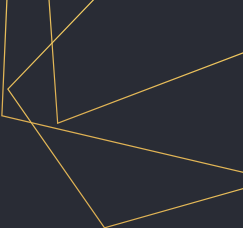
# GaussianMixture MODEL





Each gaussian component is associated to its majority class

New labels dataset can be used to train a more accurate Forest model





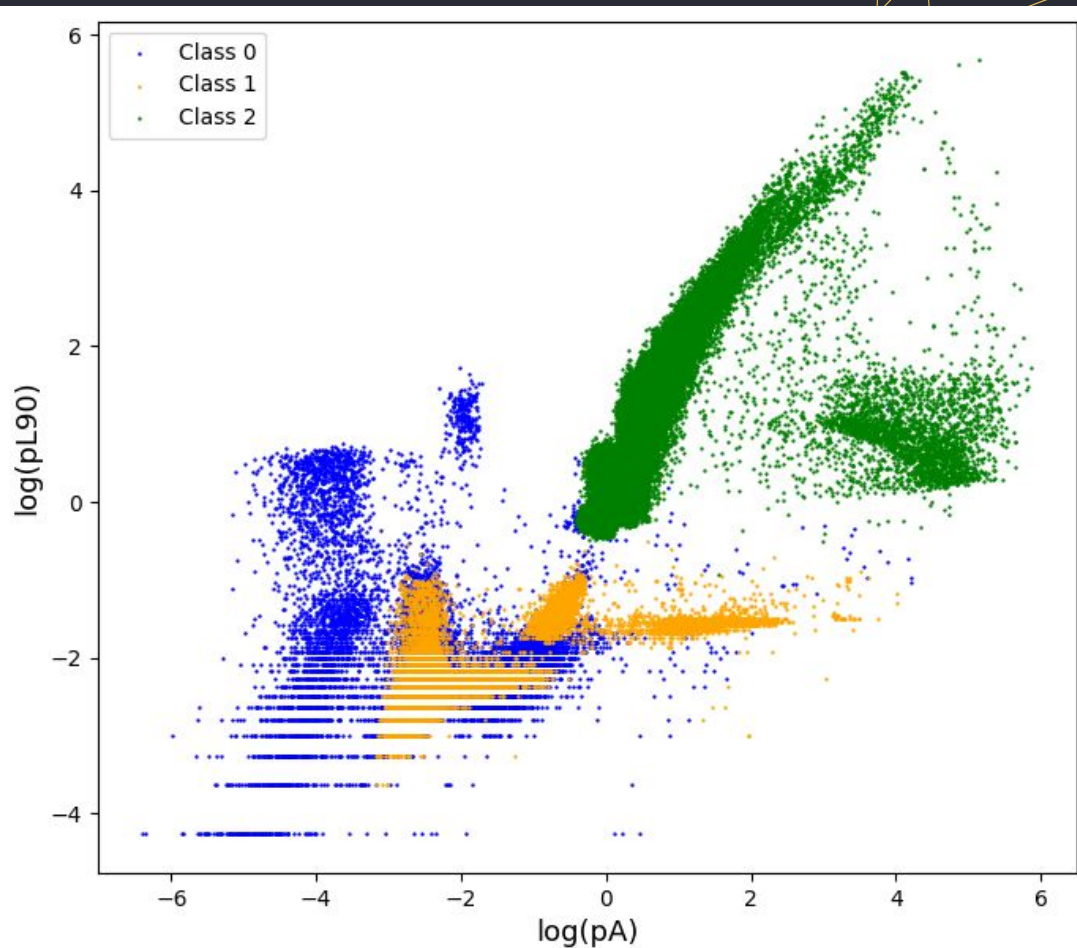
# GMM *RandomForest*

Testing score: **99.40 %**

## CONFUSION MATRIX

PREDICTED CLASS

	0	1	2
0	8377	555	39
1	492	10370	0
2	122	0	180045



# CONCLUSIONS

## DECISION TREE

Score:  
**98.80 %**

CLASS LABEL	PREDICTED CLASS		
	0	1	2
0	89.9%	7.49%	0.002%
1	9.7%	87.2%	0.03%
2	0.2%	5.4%	99.97%

## RANDOM FOREST

Score:  
**99.13 %**

CLASS LABEL	PREDICTED CLASS		
	0	1	2
0	88.8%	9.38%	0%
1	9.99%	90.6%	0.02%
2	1.23%	0%	99.98%

## RANDOM FOREST WITH GMM DATA

Score:  
**99.40 %**

CLASS LABEL	PREDICTED CLASS		
	0	1	2
0	93.2%	5.08%	0.022%
1	5.47%	94.9%	0%
2	1.13%	0%	99.98%

\* percentage of actual class label over total predictions of one class label

# FUTURE DEVELOPMENTS

## PERMUTATION IMPORTANCE

Randomly permuting variables in a tree and comparing its accuracy with the one of the original tree

→ accounts for **highly correlated** features

## INCREASING K

Better fit of less dense regions and decrease in relevance of singularities

## NEURAL NETWORK (TriNeT)

Ensemble of Neural Networks which focus on separating one feature from the others



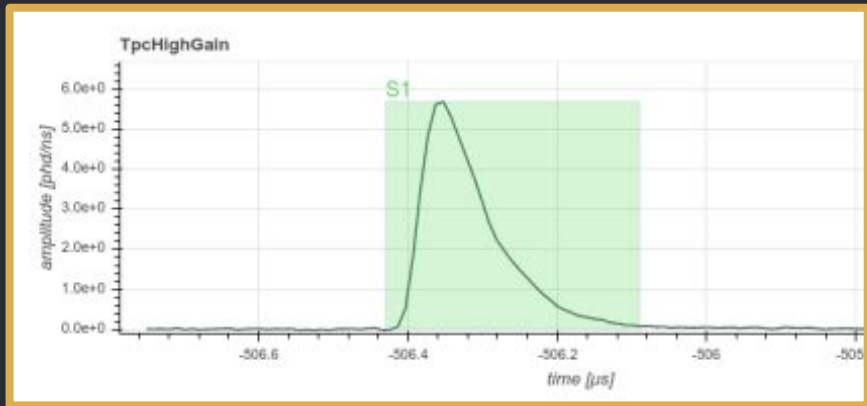
THANK YOU FOR  
YOUR ATTENTION



BACKUP

# PULSES IN LZ

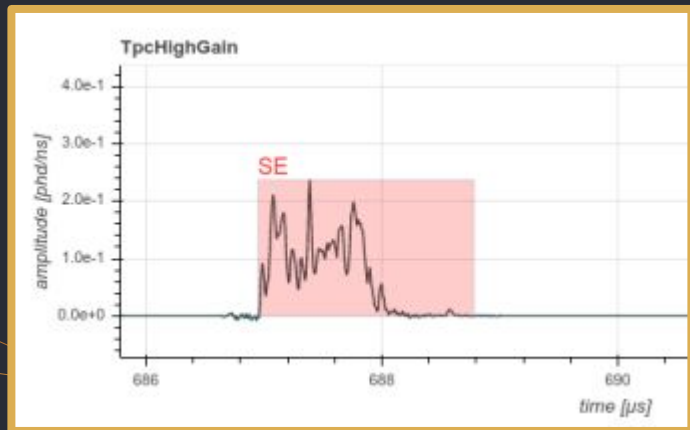
S1



S2



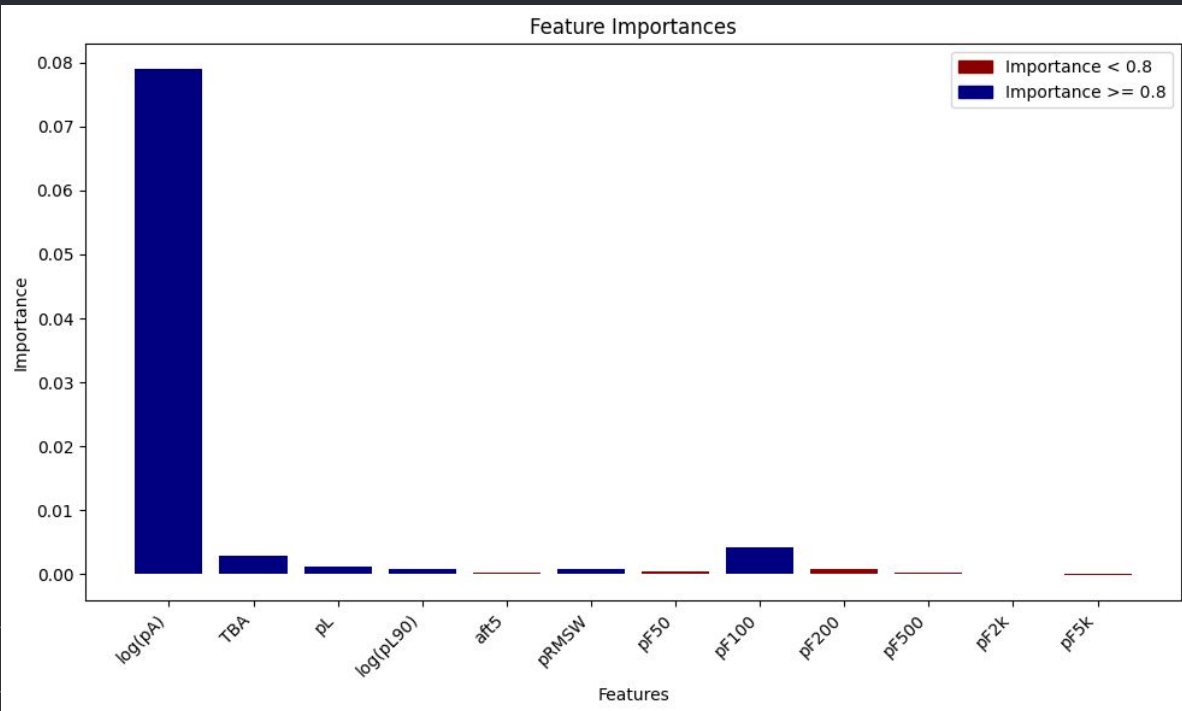
SE



OTHER



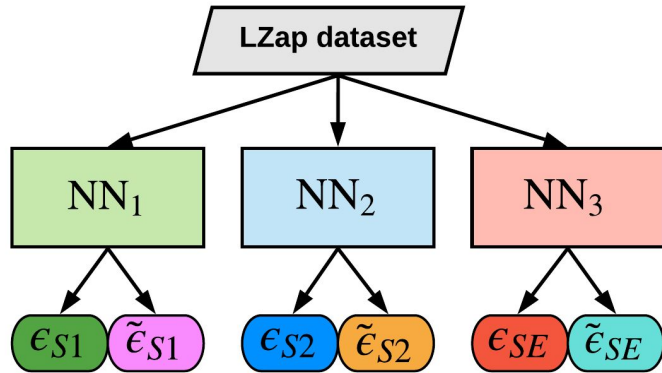
# PERMUTATION IMPORTANCE



Randomly permuting variables in a tree reduces its efficiency  
→ Comparing its accuracy with the one of the original tree you can get the variable's importance

Permutation importance accounts for **highly correlated** features

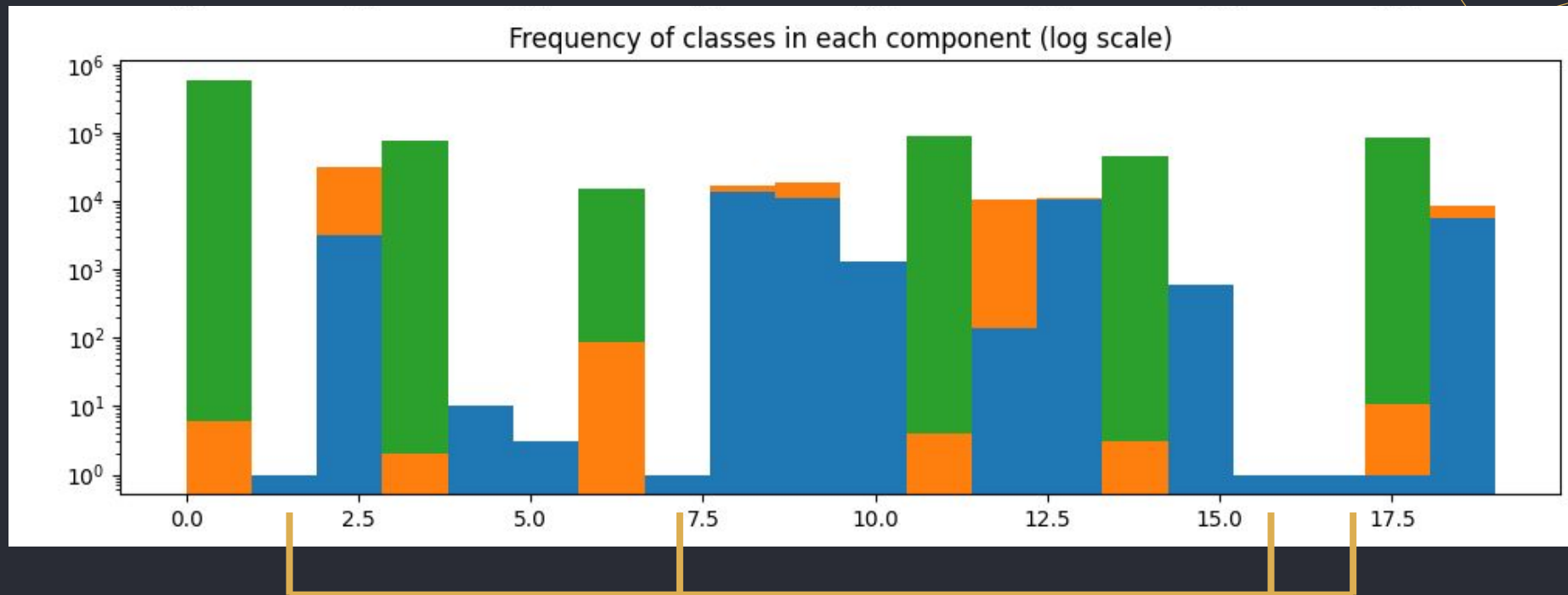
# TriNet CLASSIFIER



Ensemble of Neural Networks trained ad **One-VS-All**:

- Each NN only learns one designated class, the rest of the pulses are labelled as “not of that class”
- Trained using pre-existing labels dataset



GMM WITH  $K=20$ 

GMM with 15 or 20 doesn't change much, as the number of singularities increases: to see improvement we would need a much larger  $K$ , which requires too much computational power