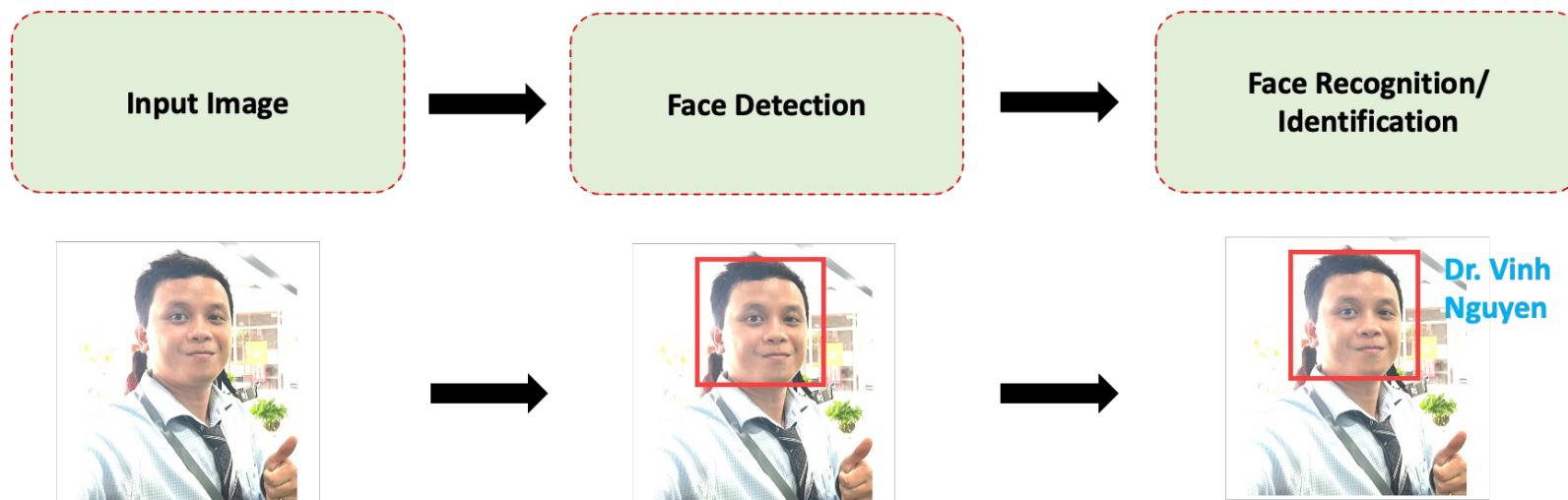


# Face Detection and Identification

*(Computer Vision Foundation)*



Vinh Dinh Nguyen  
PhD in Computer Science

# Outline

- Applications of Face Detection
- Face Detection in Computer Vision
- Haar Feature
- Integral Image
- Adaboost For Face Detection
- Cascade Classifier
- Face Recognition

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# What is Face Detection



Locate human faces in images

1. Application of face detection
2. Haar feature for face detection
3. Integral Image
4. Cascade Classifier

# Applications of Face Detection



Security Use Case of Face Recognition



A Face Attendance System



Healthcare Application



Face Recognition in Retail



Face Recognition in Finance



Automatic Camera Settings

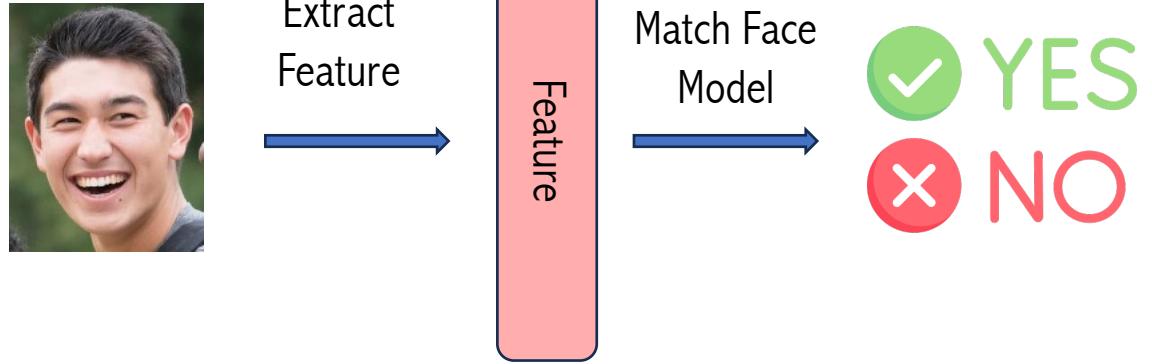
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# Face Detection in Computer Vision



Slides windows of different sizes across image.  
At each location match window to face model



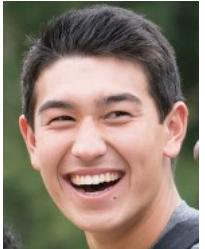
## Features:

- How to extract feature?
- Which features represent face well?

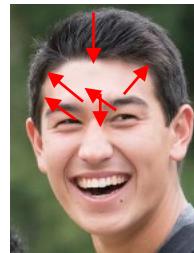
## Classifier:

- How to build a model and classify features as face or not?

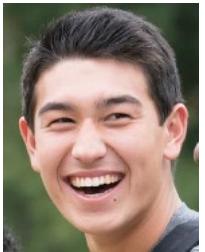
# What are good features?



Interest Point (Edge, Corners, SIFT,...)



- Extreme Fast to Compute
- Millions of windows in an image



Facial Components (Templates)?

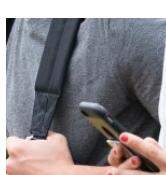


The key aspect in face recognition is detecting relevant features in human face like eyes, eyebrows, nose, lips. So how do we detect these features in real time/in an image ?

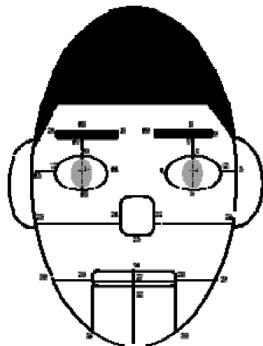


Discriminative Face / None Face

$\neq$

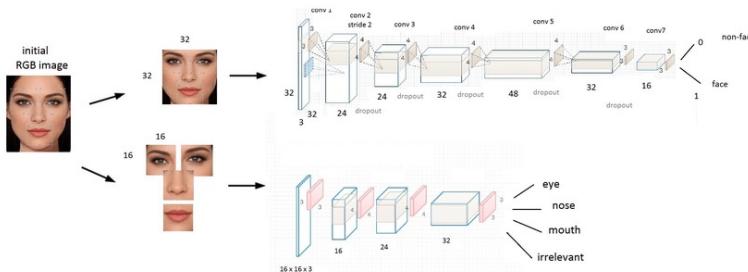


# Methods for Face Detection



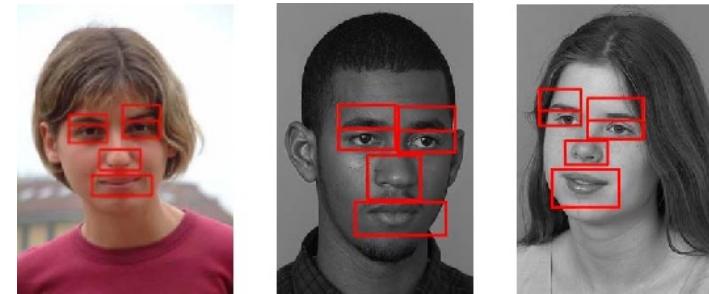
## Knowledge Based

- Rule based (Ex: X must have eyes, x must have a nose)
- Too many rules and variables with this method



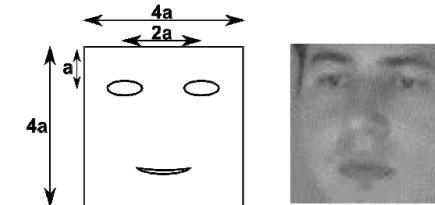
## Appearance Based

Learn the characteristics of a face. Example: CNN's  
Accuracy depends on training data (which can be scarce)



## Feature Based

Locate and extract structural features in the face  
Find a differential between facial and non facial regions in an image



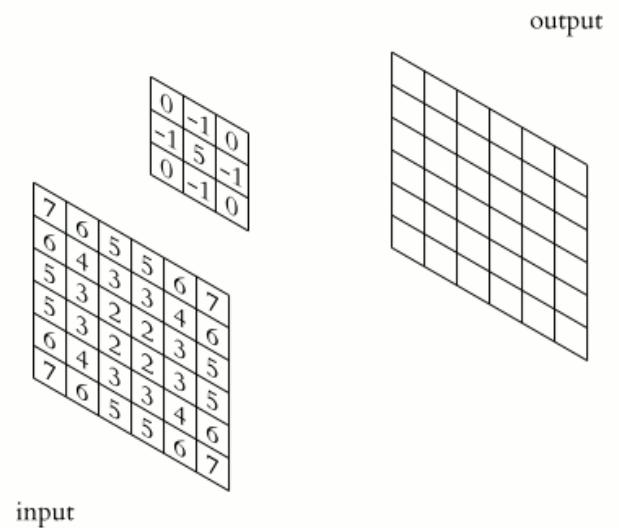
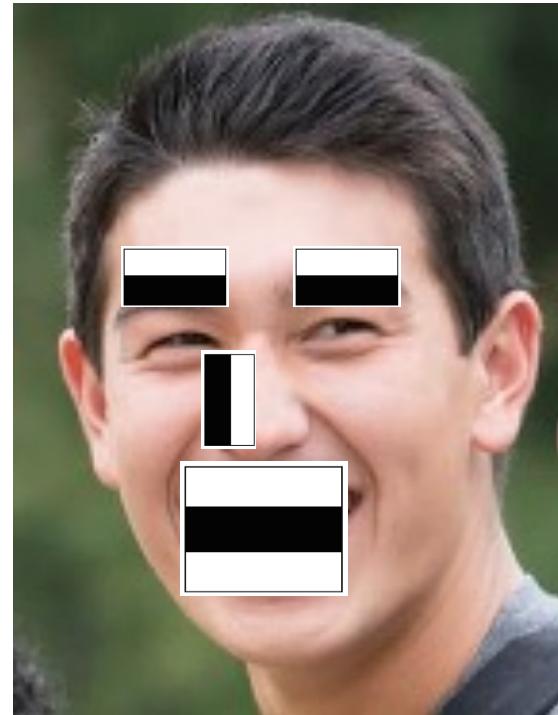
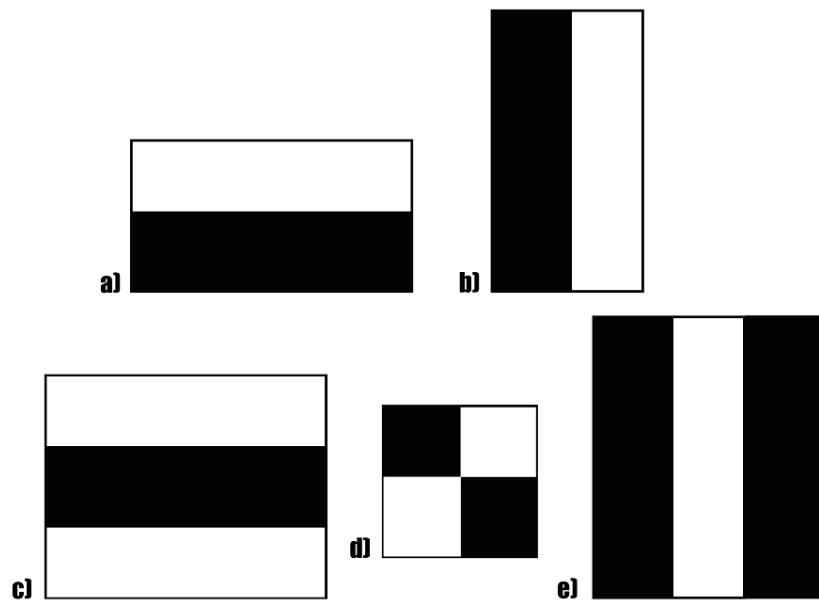
## Template

Using predefined templates for edge detection. Quick and easy  
A trade off for speed over accuracy

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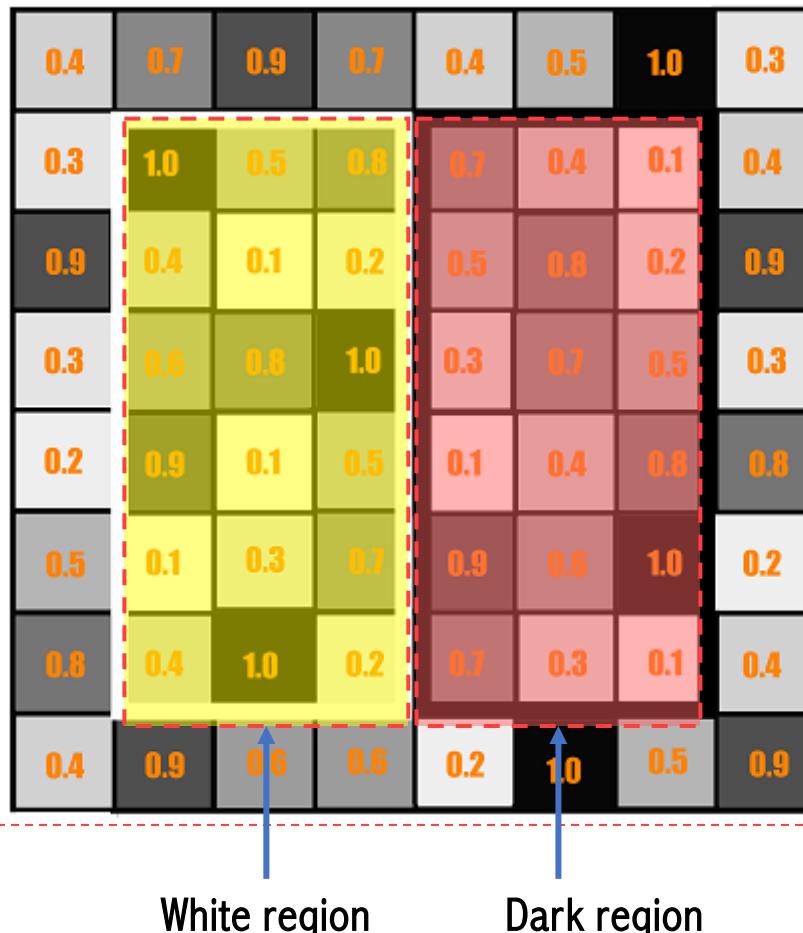
# Haar Feature



Viola-Jones uses 24\*24 as base window size and calculates the above features all over the image shifting by 1 PX (Template + Feature). There are over 160,000 possible feature combinations that can fit into a 24x24 pixel image, and over 250,000 for a 28x28 image

P. Viola and M. Jones, "Rapid object detection using a boosted cascade of simple features," Proceedings of the 2001 IEEE Computer Society Conference on Computer Vision and Pattern Recognition. CVPR 2001, Kauai, HI, USA, 2001, pp. I-I, doi: 10.1109/CVPR.2001.990517.

# Haar Feature



0	0	0	1	1	1
0	0	0	1	1	1
0	0	0	1	1	1
0	0	0	1	1	1
0	0	0	1	1	1
0	0	0	1	1	1

=

$$\frac{\text{SUM OF THE DARK PIXELS/NUMBER OF DARK PIXELS}}{\text{SUM OF THE LIGHT PIXELS/NUMBER OF THE LIGHT PIXELS}}$$

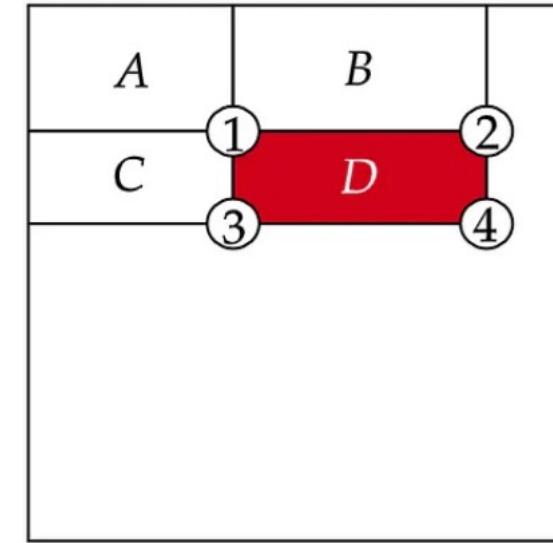
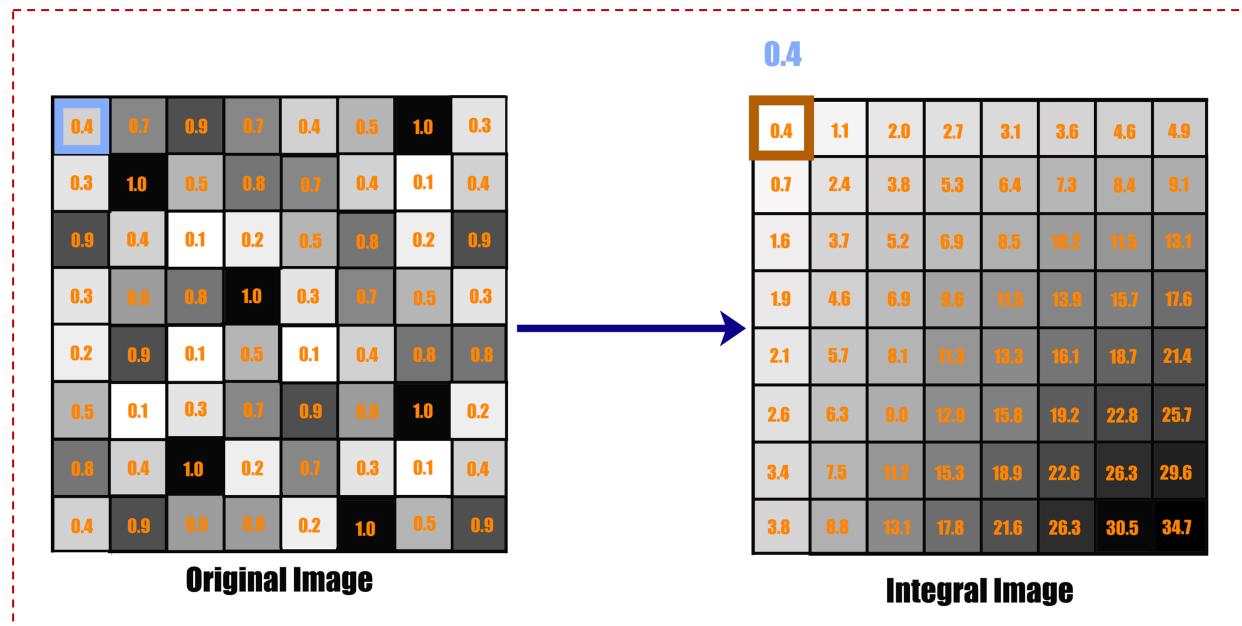
$$\frac{(0.7 + 0.4 + 0.1 + 0.5 + 0.8 + 0.2 + 0.3 + 0.7 + 0.5 + 0.1 + 0.4 + 0.8 + 0.9 + 0.6 + 1.0 + 0.7 + 0.3 + 0.1)/18}{(1.0 + 0.5 + 0.8 + 0.4 + 0.1 + 0.2 + 0.6 + 0.8 + 1.0 + 0.9 + 0.1 + 0.5 + 0.1 + 0.3 + 0.7 + 0.4 + 1.0 + 0.2)/18}$$

$$0.51 - 0.53 = -0.02$$

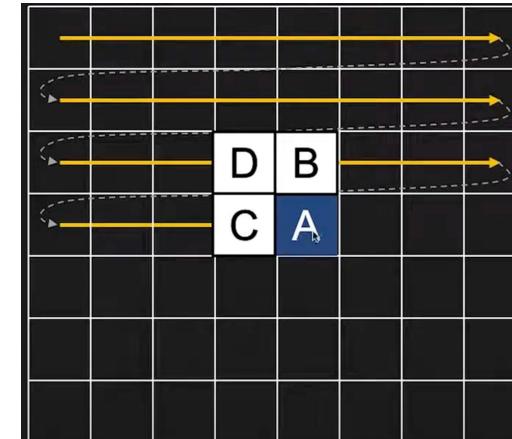
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# Integral Image

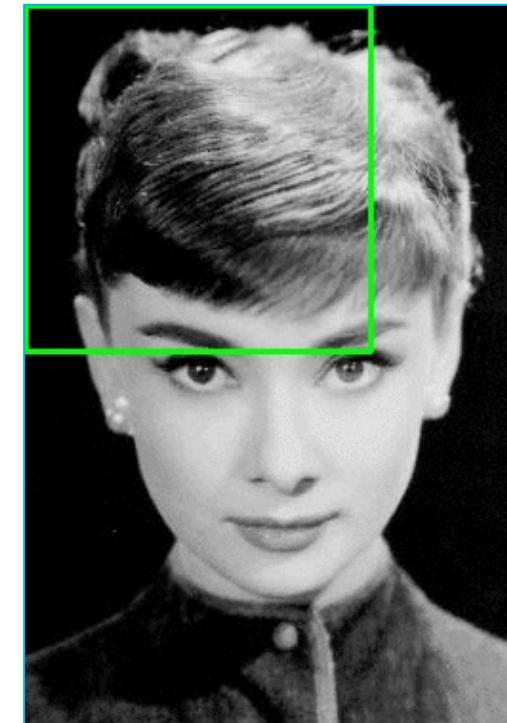
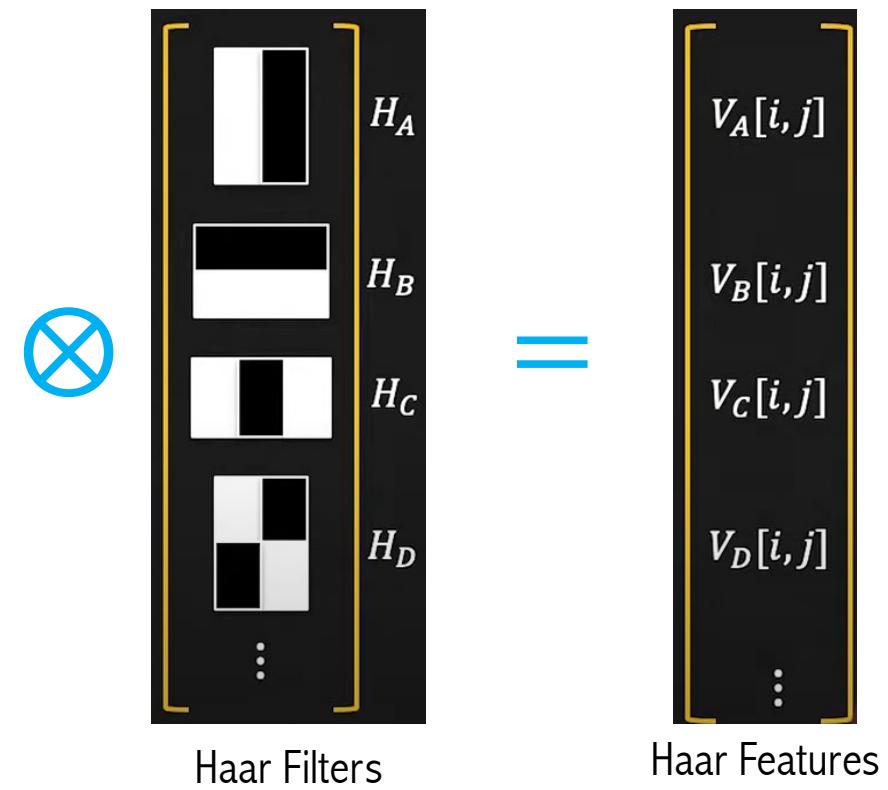


- The value of the integral image at point 1 is the sum of the pixels in rectangle A
- The value at point 2 is A + B
- The value at point 3 is A + C
- The value at point 4 is A + B + C + D.
- Therefore, the sum of pixels in region D can simply be computed as :  $4+1-(2+3)$ .

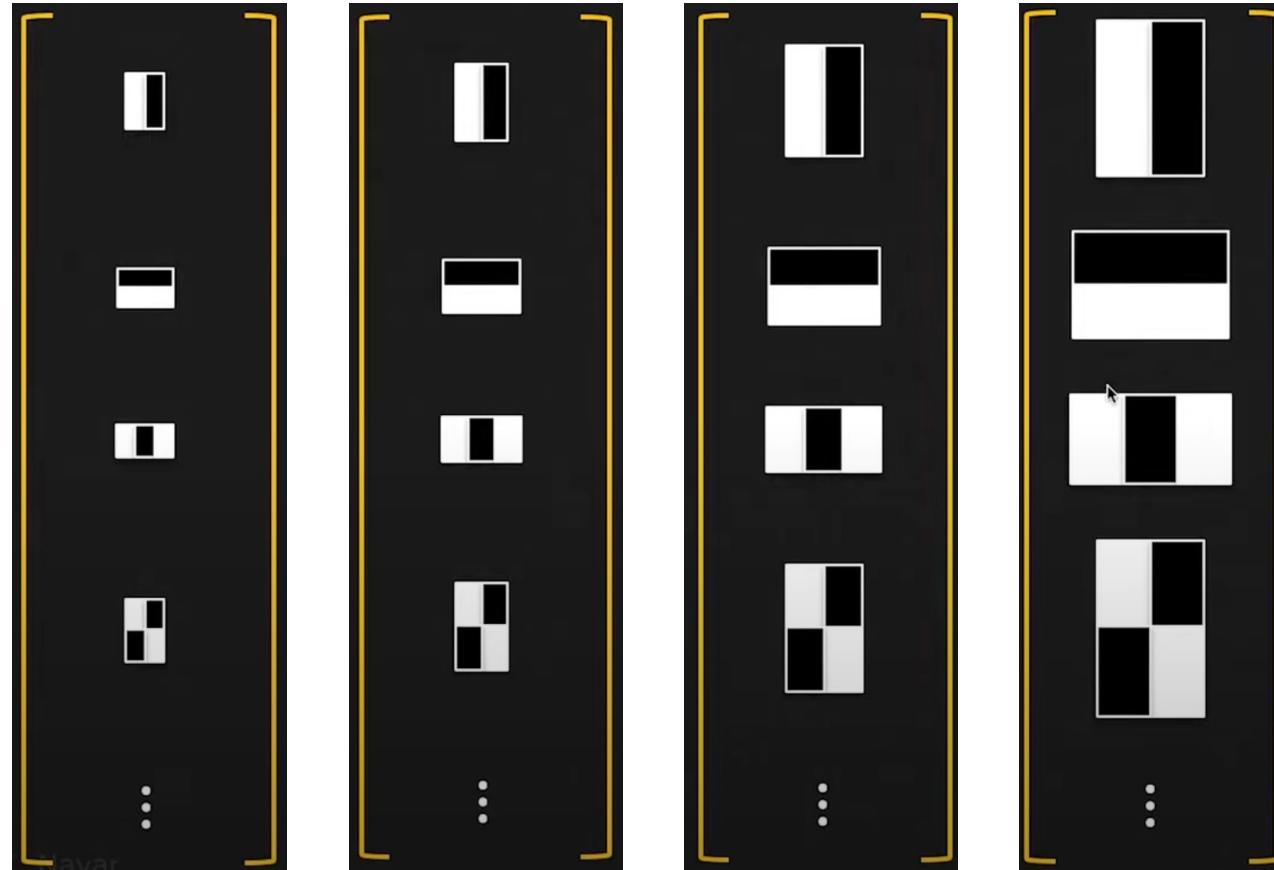


Raster Scanning

# Haar Feature



# Haar Features for Multiple Scales



Can we create a good classifier using just a small subset of all possible features?

$$V_A[i,j] = \sum_m \sum_n I[m - i, n - j] H_A[m, n]$$

Compute Haar Features at different scales to detect faces of different sizes

As stated previously there can be approximately 160,000 + feature values within a detector at 24\*24 base resolution which need to be calculated. But it is to be understood that only a few set of features will be useful among all these features to identify a face.



Relevant feature



Irrelevant feature

Feature detecting a vertical edge is useful detecting a nose but irrelevant detecting a lip.

# Outline

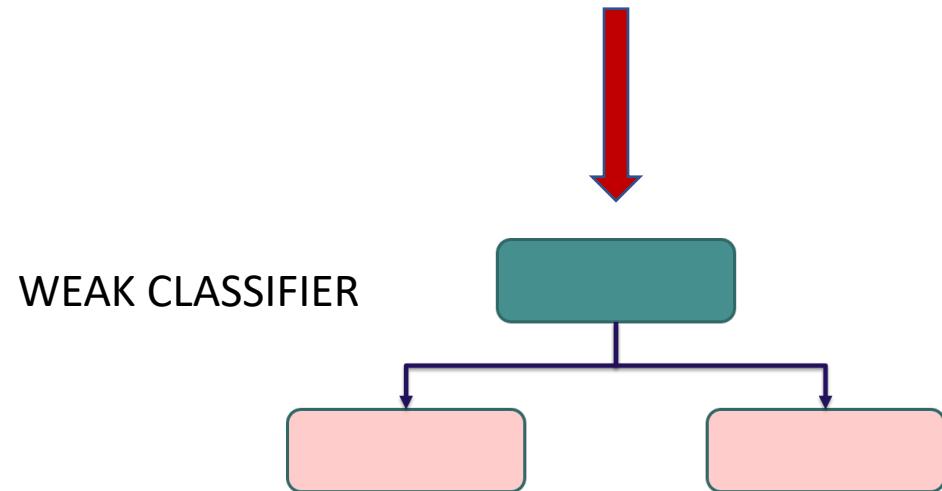
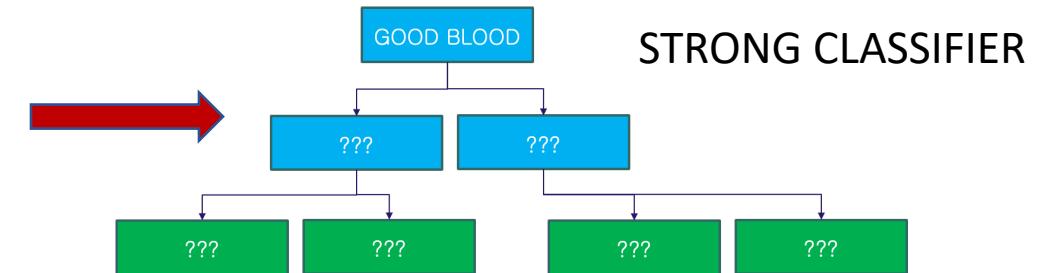
- Applications of Face Detection
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# Sample Dataset

CHEST PAIN	GOOD BLOOD CIRCULATION	BLOCKED ARTERIES	WEIGHT	HEART DISEASE
NO	NO	NO	125	NO
YES	YES	YES	180	YES
YES	YES	NO	210	NO
YES	NO	YES	167	YES

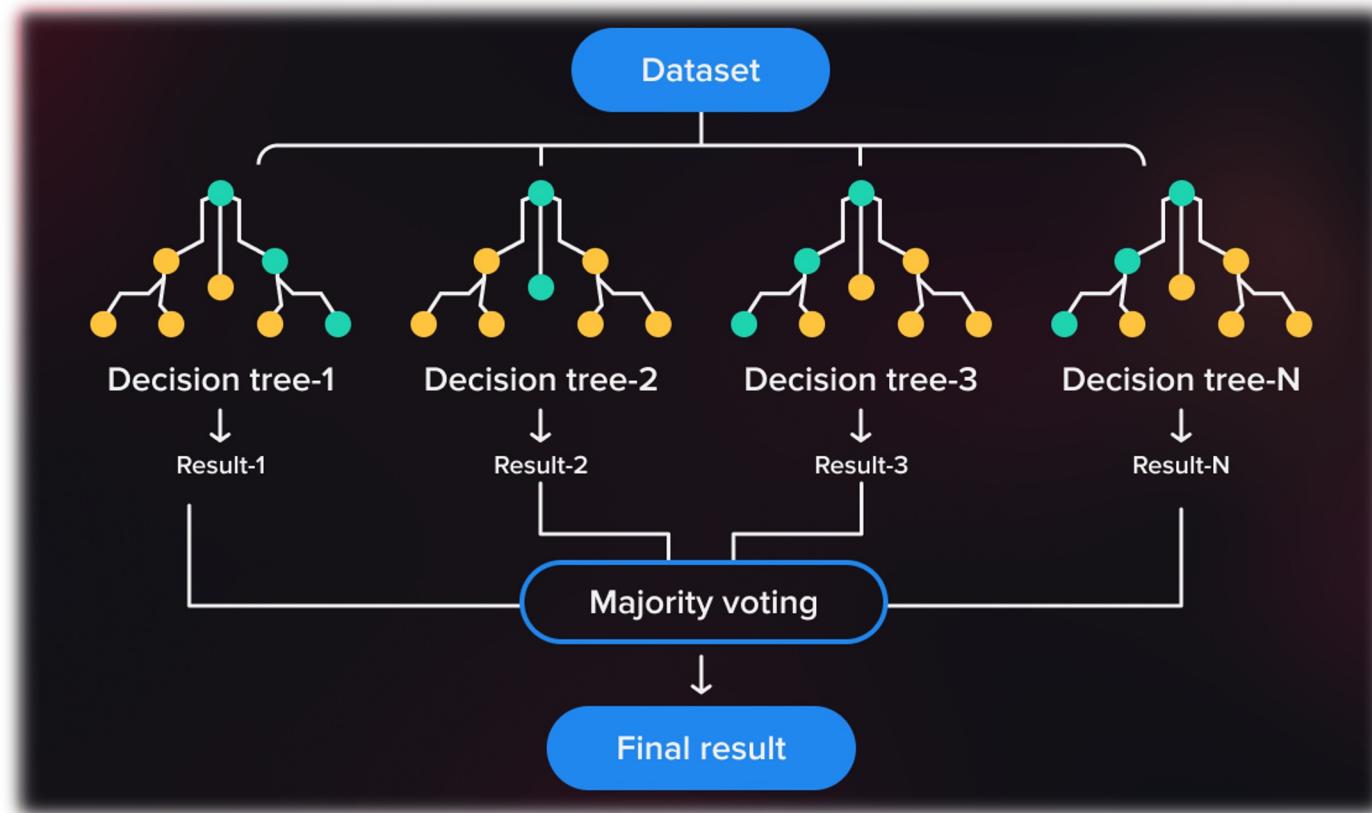
# Sample Dataset

CHEST PAIN	GOOD BLOOD CIRCULATION	BLOCKED ARTERIES	WEIGHT	HEART DISEASE
NO	NO	NO	125	NO
YES	YES	YES	180	YES
YES	YES	NO	210	NO
YES	NO	YES	167	YES



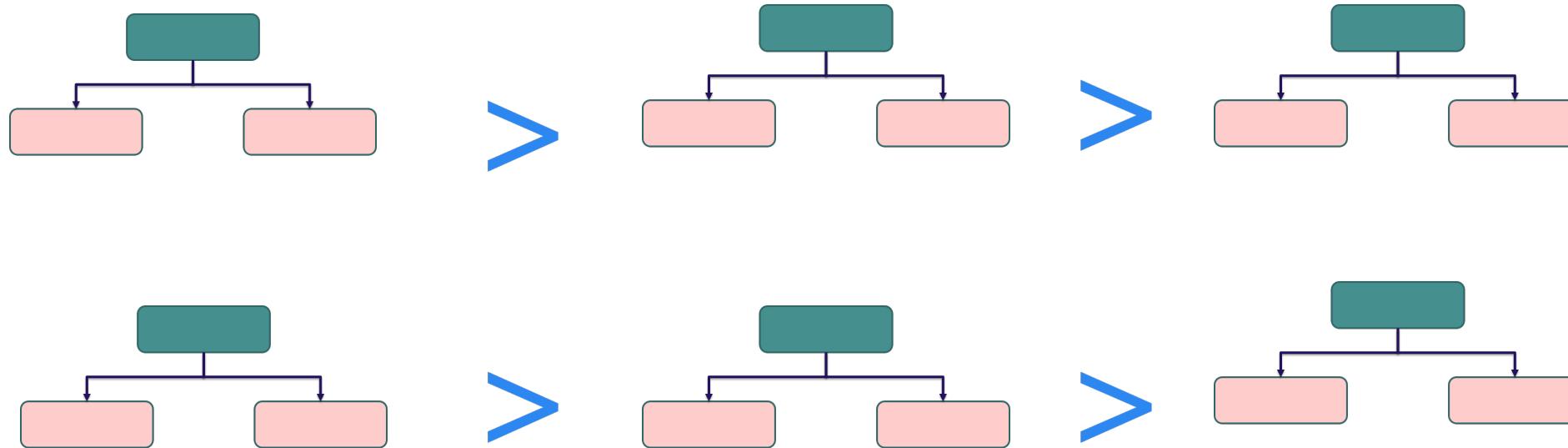
# RANDOM FOREST

- Each tree in the random forest has equal votes(weights) on the final decision

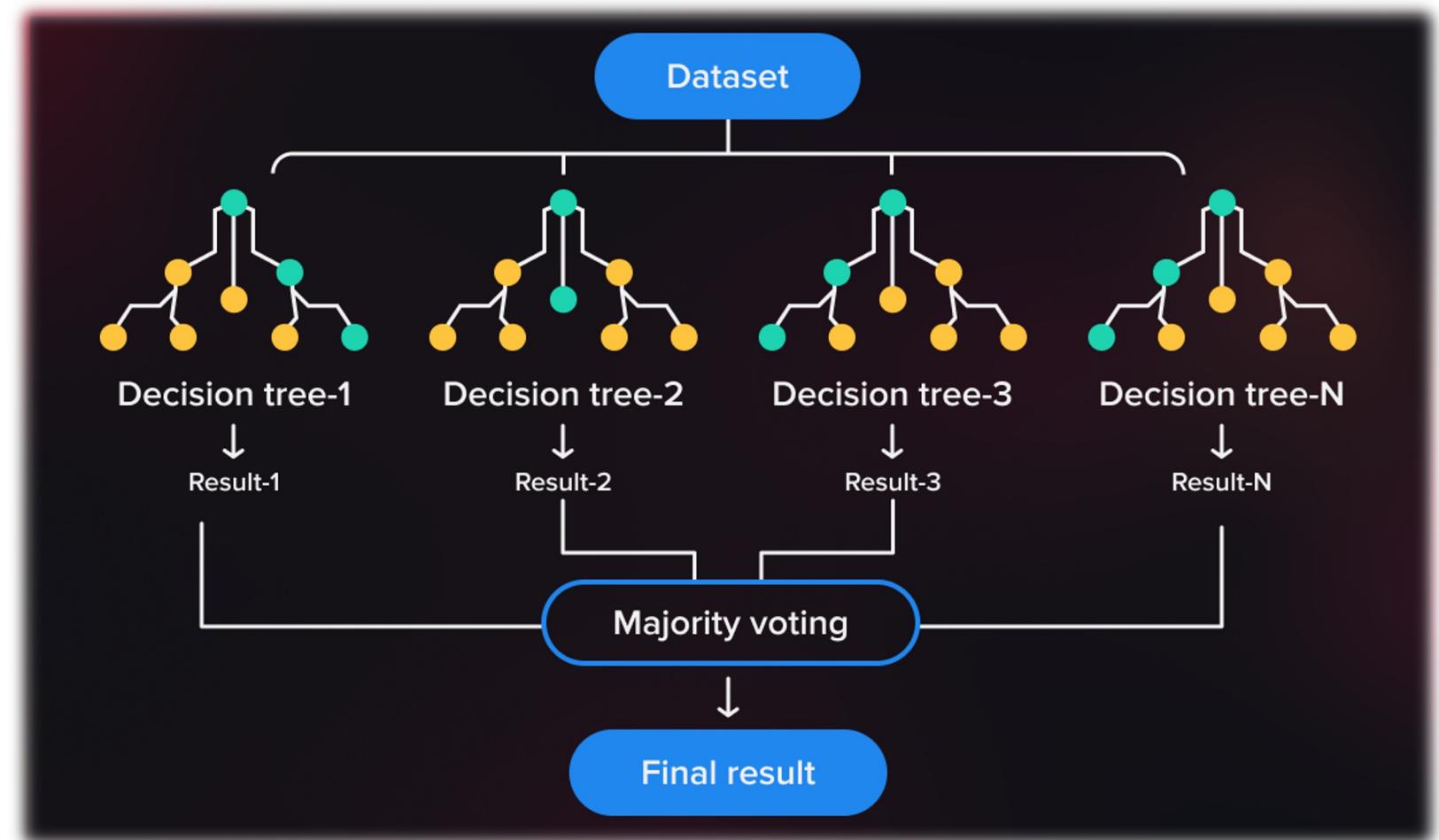


# AdaBoost: FOREST OF STUMP

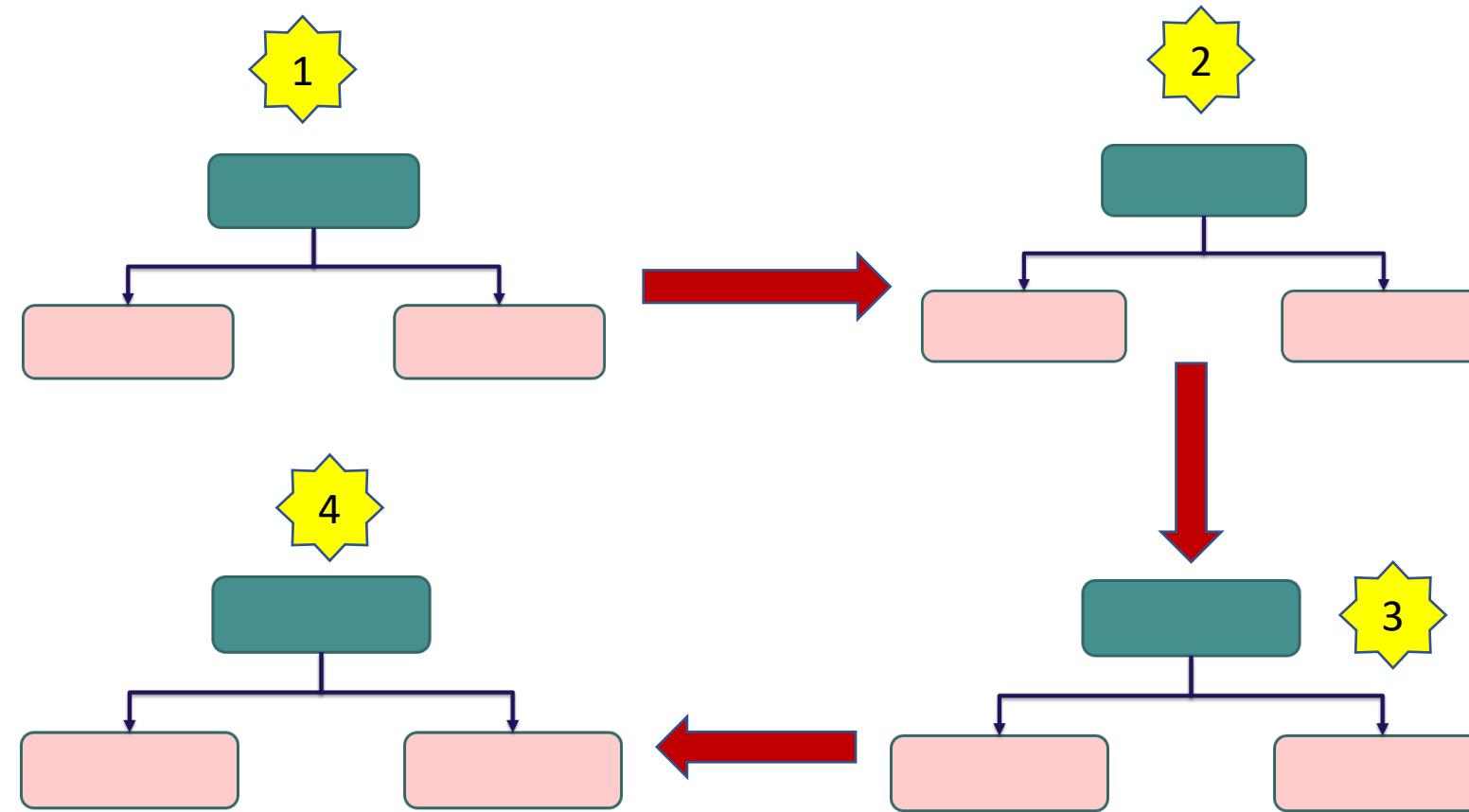
- Stump are not equally weighted in the final decision.
- Stump that create more error will have less contribution in the final decision



# Random Forest



# AdaBoost: FOREST OF STUMP



# Heart Disease Dataset

Chest Pain	Blocked Arteries	Patient Weight	Heart Disease
Yes	Yes	205	Yes
No	Yes	180	Yes
Yes	No	210	Yes
Yes	Yes	167	Yes
No	Yes	156	No
No	Yes	125	No
Yes	No	168	No
Yes	Yes	172	No

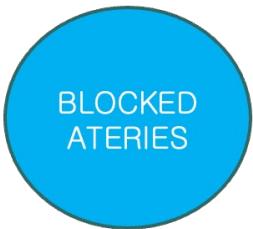
Important of sample = Sample weight =  $1 / \text{number of samples} = 1/8$

# 1st Stump in the Tree

Chest Pain	Blocked Arteries	Patient Weight	Heart Disease
Yes	Yes	205	Yes
No	Yes	180	Yes
Yes	No	210	Yes
Yes	Yes	167	Yes
No	Yes	156	No
No	Yes	125	No
Yes	No	168	No
Yes	Yes	172	No



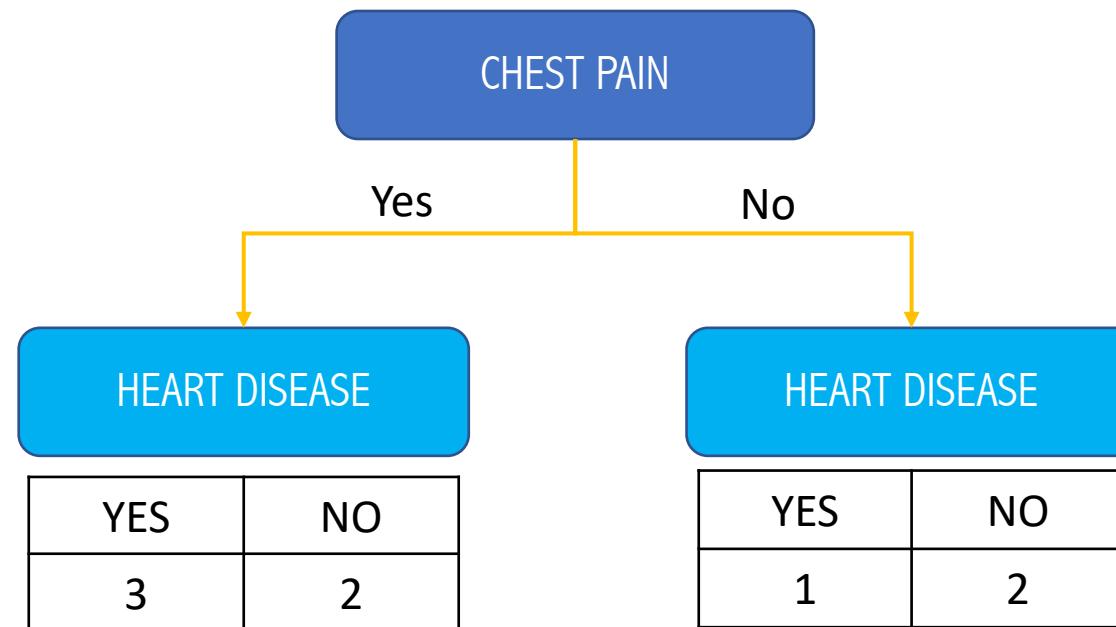
W H I C H  
o N E ?



# Compute Gini Index For Chest Pain

Chest Pain	Heart Disease	Sample Weight
Yes	Yes	1/8
No	Yes	1/8
Yes	Yes	1/8
Yes	Yes	1/8
No	No	1/8
No	No	1/8
Yes	No	1/8
Yes	No	1/8

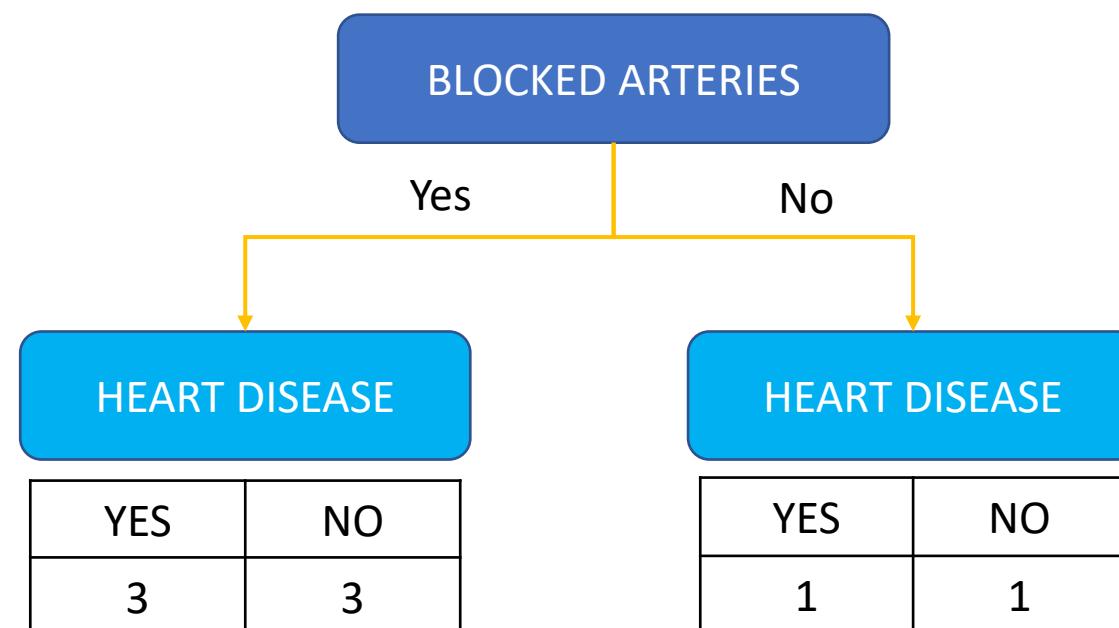
$$\text{Gini index} = 5/8 * (1 - (3/5)^2 - (2/5)^2) + 3/8 * (1 - (1/3)^2 - (2/3)^2) = 0.57$$



# GINI INDEX FOR BLOCKED ARTERIES

Blocked Arteries	Heart Disease	Sample Weight
Yes	Yes	1/8
Yes	Yes	1/8
No	Yes	1/8
Yes	Yes	1/8
Yes	No	1/8
Yes	No	1/8
No	No	1/8
Yes	No	1/8

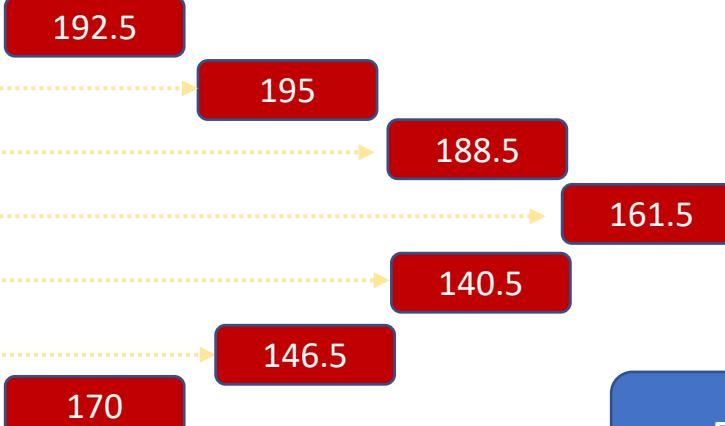
$$\text{Gini index} = 6/8 * (1 - (3/6)^2 - (3/6)^2) + 2/8 * (1 - (1/2)^2 - (1/2)^2) = 0.5$$



# Gini Index for Heart Disease

Patient Weight	Heart Disease	Sample Weight
205	Yes	1/8
180	Yes	1/8
210	Yes	1/8
167	Yes	1/8
156	No	1/8
125	No	1/8
168	No	1/8
172	No	1/8

$$\text{Gini index} = 4/8 * (1-(1/4)^2 - (3/4)^2) + 4/8 * (1-(1/4)^2 - (3/4)^2) = 0.375$$



PATIENT WEIGHT > 170

HEART DISEASE

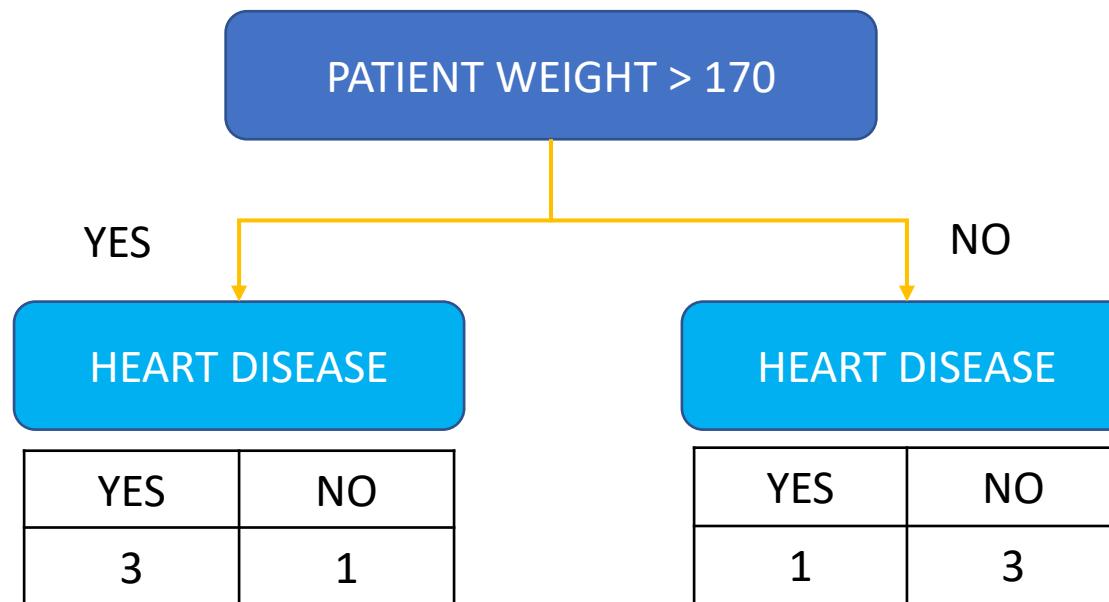
YES	NO
3	1

HEART DISEASE

YES	NO
1	3

# Amount of Say

How was this stump contribute to the final decision (classification)?



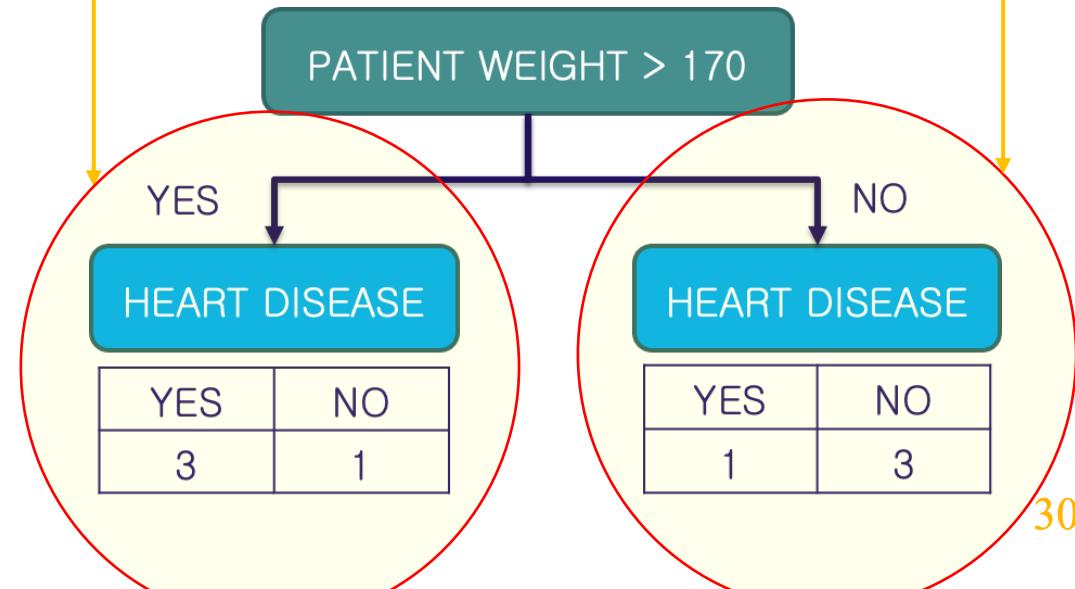
$$\text{Amount of Say} = \frac{1}{2} \log\left(\frac{1 - \text{Total Error}}{\text{Total Error}}\right)$$



# Amount of Say: Patient Weight

Chest Pain	Blocked Arteries	Patient Weight	Heart Disease	Sample Weight
Yes	Yes	205	Yes	1/8
No	Yes	180	Yes	1/8
Yes	No	210	Yes	1/8
Yes	Yes	167	Yes	1/8
No	Yes	156	No	1/8
No	Yes	125	No	1/8
Yes	No	168	No	1/8
Yes	Yes	172	No	1/8

- Total error is equal to the sum of the weights of the incorrect classified
- Amount of say =  $1/2 * \log((1-2/8) / (2/8)) = 0.55$

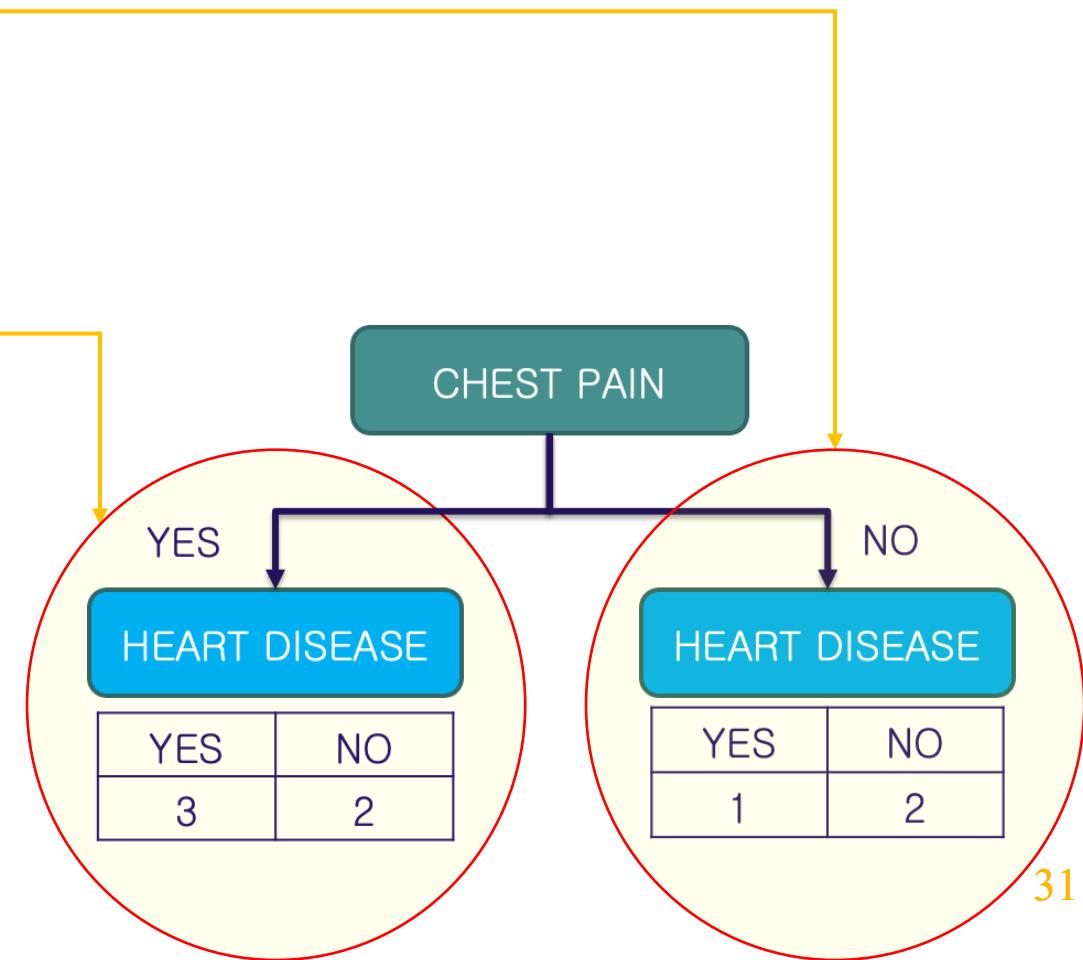


# Amount of say: Chest Pain

Chest Pain	Blocked Arteries	Patient Weight	Heart Disease	Sample Weight
Yes	Yes	205	Yes	1/8
No	Yes	180	Yes	1/8
Yes	No	210	Yes	1/8
Yes	Yes	167	Yes	1/8
No	Yes	156	No	1/8
No	Yes	125	No	1/8
Yes	No	168	No	1/8
Yes	Yes	172	No	1/8

- Total error is equal to the sum of the weights of the incorrect classified
- Amount of say =  $1/2 * \log((1-3/8) / (3/8)) = 0.25$

$$\log(\text{Odds}) = \log\left(\frac{1 - \text{Probability of incorrect prediction}}{\text{Probability of incorrect prediction}}\right)$$

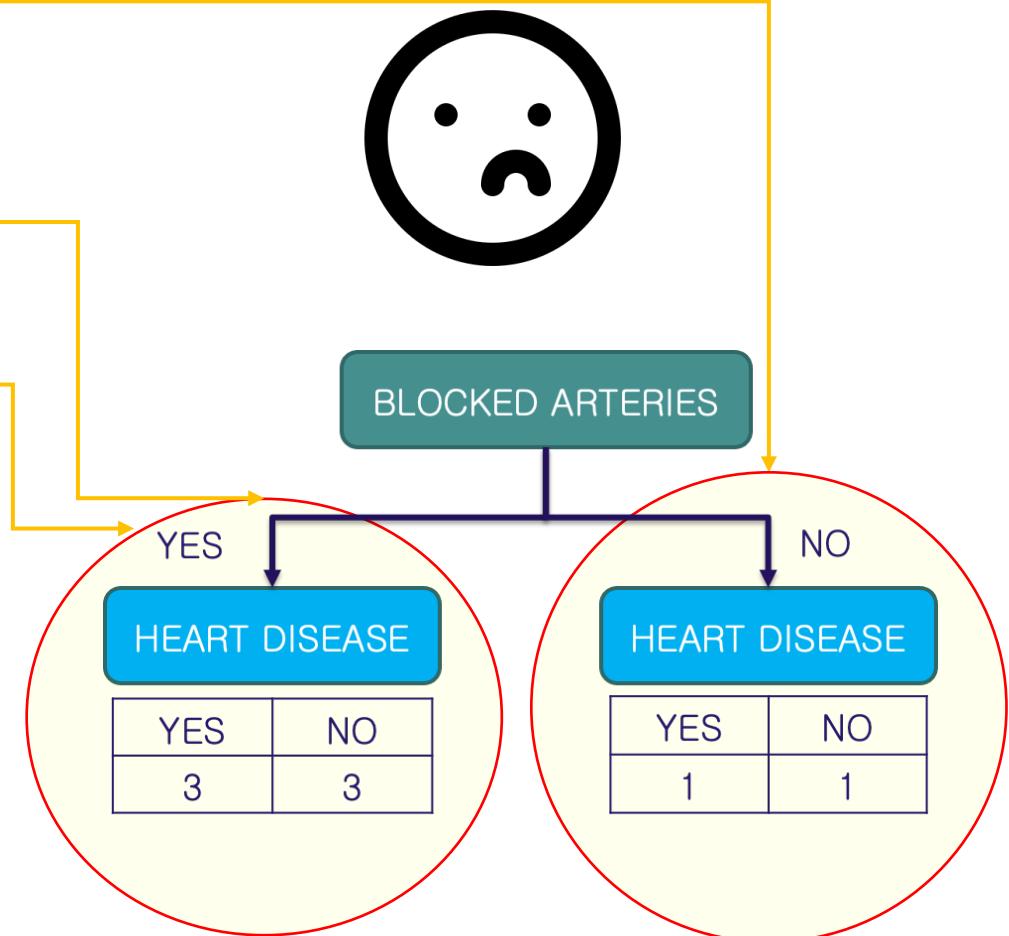


# Amout of Say: Blocked Arteries

Chest Pain	Blocked Arteries	Patient Weight	Heart Disease	Sample Weight
Yes	Yes	205	Yes	1/8
No	Yes	180	Yes	1/8
Yes	No	210	Yes	1/8
Yes	Yes	167	Yes	1/8
No	Yes	156	No	1/8
No	Yes	125	No	1/8
Yes	No	168	No	1/8
Yes	Yes	172	No	1/8

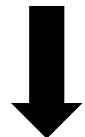
- Total error is equal to the sum of the weights of the incorrect classified
- Amount of say =  $1/2 * \log((1-4/8) / (4/8)) = 0$

WHY?



# Assumptions

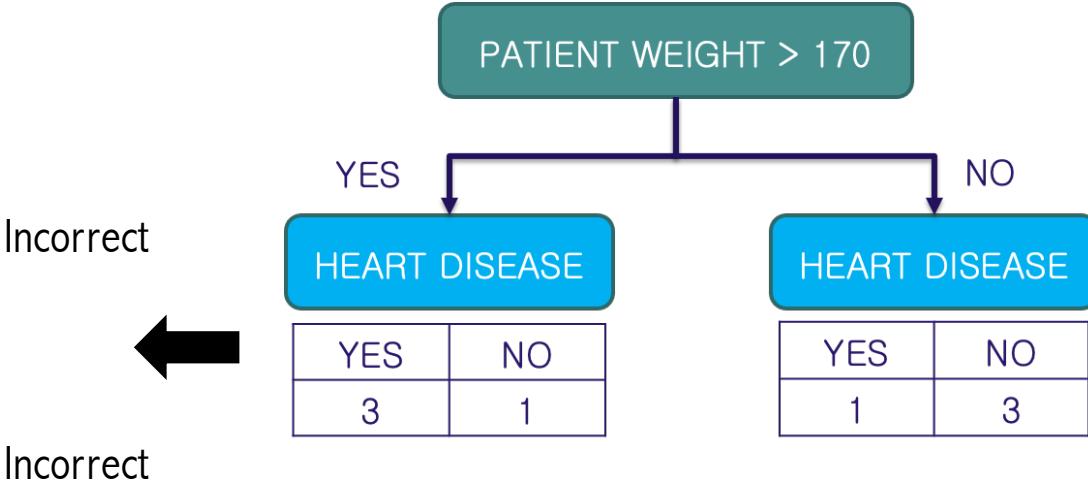
**Known:** weight cho các sample dự đoán sai được sử dụng để tính “Amount of Say” cho từng stump hiện tại.



**Unknown:** Tiếp theo, chúng ta cần làm thế nào để sử dụng thông tin các weight của sample dự đoán sai này để **xây dựng stump tiếp và khắc phục các dự đoán sai này**

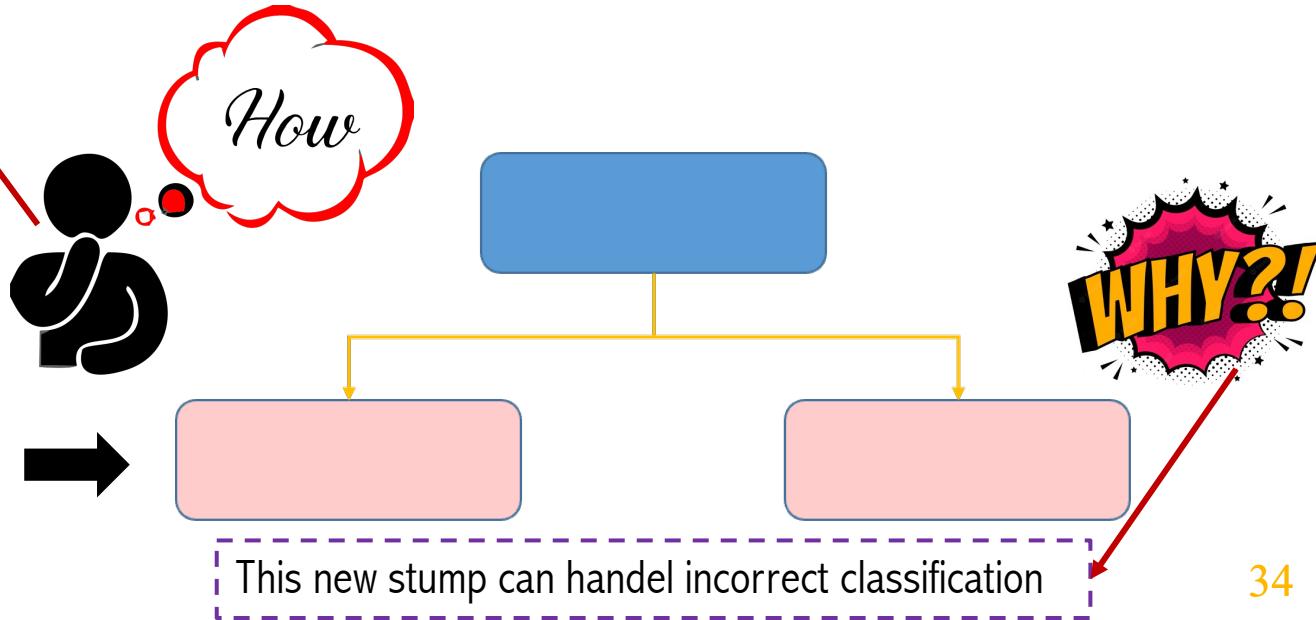
# Idea: Improved Bootstrapped Dataset

Chest Pain	Blocked Arteries	Patient Weight	Heart Disease
Yes	Yes	205	Yes
No	Yes	180	Yes
Yes	No	210	Yes
Yes	Yes	167	Yes
No	Yes	156	No
No	Yes	125	No
Yes	No	168	No
Yes	Yes	172	No



Chest Pain	Blocked Arteries	Patient Weight	Heart Disease
Yes	Yes	205	Yes
No	Yes	180	Yes
Yes	No	210	Yes
Yes	Yes	167	Yes
Yes	Yes	167	Yes
Yes	Yes	167	Yes
Yes	Yes	172	No
Yes	Yes	172	No

*Create new dataset*

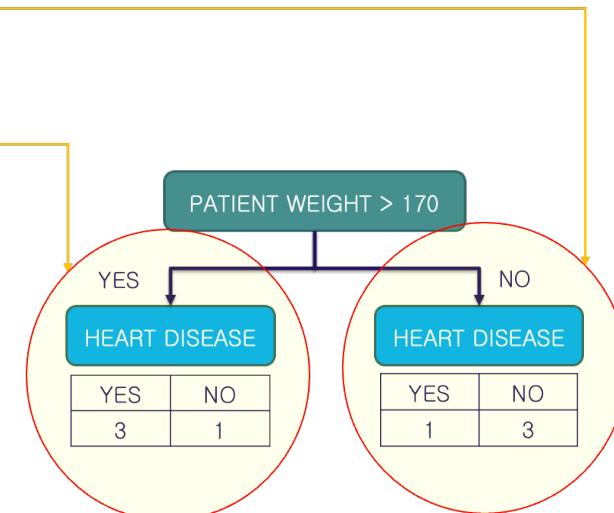


# How to build next Stump

Increase the sample weights of samples that were incorrectly classified and decrease sample weights of samples that were correctly classified. Label {-1, 1}

Chest Pain	Blocked Arteries	Patient Weight	Heart Disease	Sample Weight
Yes	Yes	205	Yes	1/8
No	Yes	180	Yes	1/8
Yes	No	210	Yes	1/8
Yes	Yes	167	Yes	1/8
No	Yes	156	No	1/8
No	Yes	125	No	1/8
Yes	No	168	No	1/8
Yes	Yes	172	No	1/8

- Total error is equal to the sum of the weights of the incorrect classified
- Amount of say =  $1/2 * \log((1-2/8) / (2/8)) = 0.55$



New Sample = sample weight  $\times e^{\text{amount of say}}$   
Weight

New sample weight =  $1/8 * e^{0.55} = 0.22$

```
def update_weights_formular1(w_i, alpha, y, y_pred):
    result = w_i * np.exp(-alpha * y * y_pred)
    w_norm = result / np.sum(result)
    return w_norm
```

# How to build next Stump

Increase the sample weights of samples that were incorrectly classified and decrease the sample weights of samples that were correctly classified. Label {-1, 1}

Chest Pain	Blocked Arteries	Patient Weight	Heart Disease
Yes	Yes	205	Yes
No	Yes	180	Yes
Yes	No	210	Yes
Yes	Yes	167	Yes
No	Yes	156	No
No	Yes	125	No
Yes	No	168	No
Yes	Yes	172	No

New Sample Weight = sample weight  $\times e^{-\text{amount of say}}$

New sample weight =  $1/8 * e^{-0.55} = 0.07$

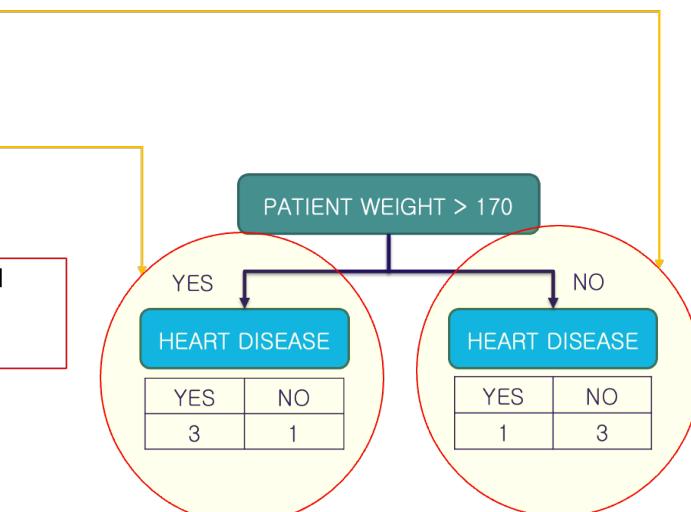
```
def update_weights_formular1(w_i, alpha, y, y_pred):
    result = w_i * np.exp(-alpha * y * y_pred)
    w_norm = result / np.sum(result)
    return w_norm
```

# How to build next Stump

Increase the sample weights of samples that were incorrectly classified and keep the sample weights of samples that were correctly classified. Label {0, 1}

Chest Pain	Blocked Arteries	Patient Weight	Heart Disease	Sample Weight
Yes	Yes	205	Yes	1/8
No	Yes	180	Yes	1/8
Yes	No	210	Yes	1/8
Yes	Yes	167	Yes	1/8
No	Yes	156	No	1/8
No	Yes	125	No	1/8
Yes	No	168	No	1/8
Yes	Yes	172	No	1/8

- Total error is equal to the sum of the weights of the incorrect classified
- Amount of say =  $1/2 * \log((1-2/8) / (2/8)) = 0.55$



$$w_i \leftarrow w_i \cdot \exp[\alpha_m \cdot I(y_i \neq G_m(x_i))],$$

New sample weight =  $1/8 * e^{0.55*1} = 0.22$

```

def update_weights_formular2(w_i, alpha, y, y_pred):
    result = w_i * np.exp(alpha * (
        np.not_equal(y, y_pred)).astype(int))
    w_norm = result / np.sum(result)
    return w_norm
  
```

# How to build next Stump

Increase the sample weights of samples that were incorrectly classified and keep the sample weights of samples that were correctly classified. Label {0, 1}

Chest Pain	Blocked Arteries	Patient Weight	Heart Disease
Yes	Yes	205	Yes
No	Yes	180	Yes
Yes	No	210	Yes
Yes	Yes	167	Yes
No	Yes	156	No
No	Yes	125	No
Yes	No	168	No
Yes	Yes	172	No

$$w_i \leftarrow w_i \cdot \exp[\alpha_m \cdot I(y_i \neq G_m(x_i))]$$

New sample weight =  $1/8 * e^{0.55*0} = 0.125$

```
def update_weights_formular2(w_i, alpha, y, y_pred):
    result = w_i * np.exp(alpha * (
        np.not_equal(y, y_pred)).astype(int))
    w_norm = result / np.sum(result)
    return w_norm
```

# New Sample Weight

Chest Pain	Blocked Arteries	Patient Weight	Heart Disease	Sample Weight	New Weight	Normal Weight
Yes	Yes	205	Yes	1/8	0.07	0.08
No	Yes	180	Yes	1/8	0.07	0.08
Yes	No	210	Yes	1/8	0.07	0.08
Yes	Yes	167	Yes	1/8	0.22	0.25
No	Yes	156	No	1/8	0.07	0.08
No	Yes	125	No	1/8	0.07	0.08
Yes	No	168	No	1/8	0.07	0.08
Yes	Yes	172	No	1/8	0.22	0.25
Sum				~1.0	0.86	~1.0

New Sample Weight = sample weight  $\times e^{\text{amount of say}}$

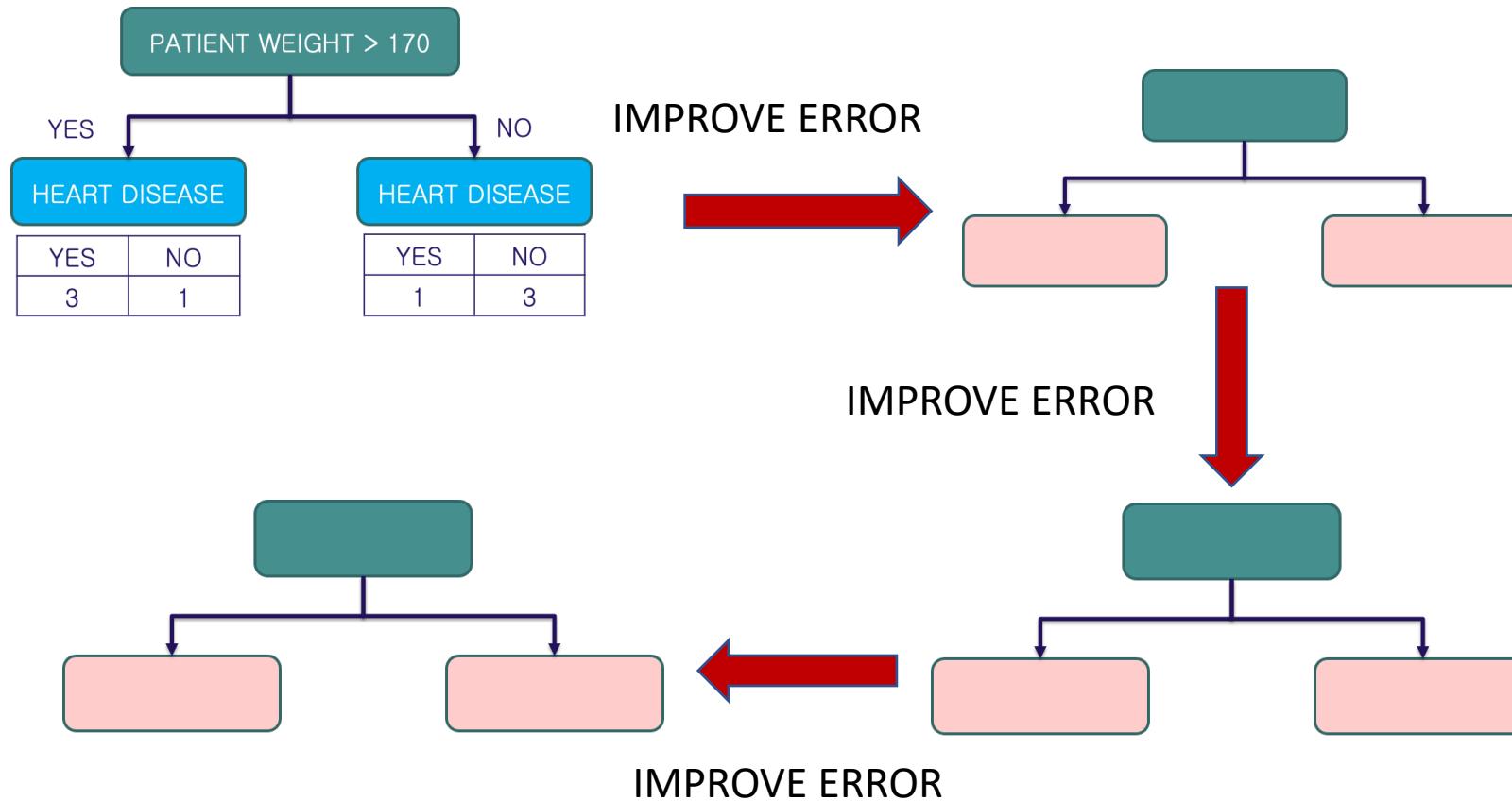


Update

# New Sample Weight

Chest Pain	Blocked Arteries	Patient Weight	Heart Disease	New Weight
Yes	Yes	205	Yes	0.08
No	Yes	180	Yes	0.08
Yes	No	210	Yes	0.08
Yes	Yes	167	Yes	0.25
No	Yes	156	No	0.08
No	Yes	125	No	0.08
Yes	No	168	No	0.08
Yes	Yes	172	No	0.25
Sum				~1.0

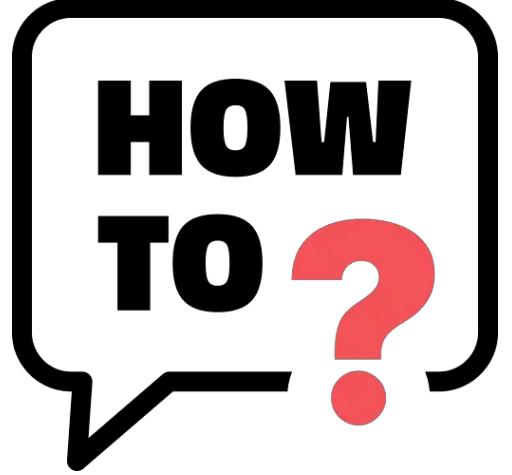
# AdaBoost: FOREST OF STUMPS



HOW

# New Dataset

Chest Pain	Blocked Arteries	Patient Weight	Heart Disease	Normal Weight
Yes	Yes	205	Yes	0.08
No	Yes	180	Yes	0.08
Yes	No	210	Yes	0.08
Yes	Yes	167	Yes	0.25
No	Yes	156	No	0.08
No	Yes	125	No	0.08
Yes	No	168	No	0.08
Yes	Yes	172	No	0.25
Sum			$\sim 1.0$	

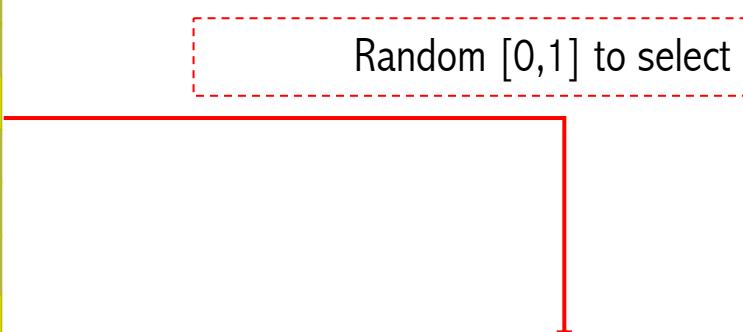



Chest Pain	Blocked Arteries	Patient Weight	Heart Disease	Normal Weight
Yes	Yes	205	Yes	0.08
No	Yes	180	Yes	0.08
Yes	No	210	Yes	0.08
Yes	Yes	167	Yes	0.25
No	Yes	156	No	0.08
No	Yes	125	No	0.08
Yes	No	168	No	0.08
Yes	Yes	172	No	0.25
Sum			$\sim 1.0$	

# New Dataset

Chest Pain	Blocked Arteries	Patient Weight	Heart Disease	Normal Weight
Yes	Yes	205	Yes	0.08
No	Yes	180	Yes	0.08
Yes	No	210	Yes	0.08
Yes	Yes	167	Yes	0.25
No	Yes	156	No	0.08
No	Yes	125	No	0.08
Yes	No	168	No	0.08
Yes	Yes	172	No	0.25
Sum			$\sim 1.0$	

Random [0,1] to select samples



Chest Pain	Blocked Arteries	Patient Weight	Heart Disease	Normal Weight
Yes	Yes	205	Yes	0.08
No	Yes	180	Yes	0.08
Yes	No	210	Yes	0.08
Yes	Yes	167	Yes	0.25
No	Yes	156	No	0.08
No	Yes	125	No	0.08
Yes	No	168	No	0.08
Yes	Yes	172	No	0.25
Sum			$\sim 1.0$	

# New Dataset

Chest Pain	Blocked Arteries	Patient Weight	Heart Disease	Normal Weight	Range
Yes	Yes	205	Yes	0.08	[0-0.08]
No	Yes	180	Yes	0.08	(0.08-0.16]
Yes	No	210	Yes	0.08	(0.16-0.24]
Yes	Yes	167	Yes	0.25	(0.24-0.495]
No	Yes	156	No	0.08	(0.495-0.575]
No	Yes	125	No	0.08	(0.575-0.655]
Yes	No	168	No	0.08	(0.655-0.735]
Yes	Yes	172	No	0.25	(0.735-1.0]
Sum				~1.0	

# New Dataset

Chest Pain	Blocked Arteries	Patient Weight	Heart Disease	Normal Weight
Yes	Yes	205	Yes	0.08
No	Yes	180	Yes	0.08
Yes	No	210	Yes	0.08
Yes	Yes	167	Yes	0.25
No	Yes	156	No	0.08
No	Yes	125	No	0.08
Yes	No	168	No	0.08
Yes	Yes	172	No	0.25
Sum				~1.0

Old dataset

Ý tưởng: các sample bị phân loại sai, sẽ được thêm vào dataset mới

Random [0,1]  
to select  
samples

CONTINUE TO BUILD  
THE NEXT STUMP

Chest Pain	Blocked Arteries	Patient Weight	Heart Disease
Yes	Yes	167	Yes
No	Yes	180	Yes
Yes	No	210	Yes
Yes	Yes	167	Yes
No	Yes	156	No
No	Yes	125	No
Yes	Yes	172	No
Yes	Yes	172	No

New dataset

# How to Classify The Final Result

**Algorithm 1** AdaBoost (*Freund & Schapire 1997*)

1. Initialize the observation weights  $w_i = 1/n$ ,  $i = 1, 2, \dots, n$ .

2. For  $m = 1$  to  $M$ :

(a) Fit a classifier  $T^{(m)}(\mathbf{x})$  to the training data using weights  $w_i$ .

(b) Compute

$$err^{(m)} = \sum_{i=1}^n w_i \mathbb{I}(c_i \neq T^{(m)}(\mathbf{x}_i)) / \sum_{i=1}^n w_i.$$

(c) Compute

$$\alpha^{(m)} = \log \frac{1 - err^{(m)}}{err^{(m)}}.$$

(d) Set

$$w_i \leftarrow w_i \cdot \exp \left( \alpha^{(m)} \cdot \mathbb{I}(c_i \neq T^{(m)}(\mathbf{x}_i)) \right), \quad i = 1, 2, \dots, n.$$

(e) Re-normalize  $w_i$ .

3. Output

$$C(\mathbf{x}) = \arg \max_k \sum_{m=1}^M \alpha^{(m)} \cdot \mathbb{I}(T^{(m)}(\mathbf{x}) = k).$$

# How to Classify The Final Result

SAMME — Stagewise Additive  
Modeling using a Multi-class  
Exponential loss function

## Algorithm 2 SAMME

1. Initialize the observation weights  $w_i = 1/n$ ,  $i = 1, 2, \dots, n$ .

2. For  $m = 1$  to  $M$ :

(a) Fit a classifier  $T^{(m)}(\mathbf{x})$  to the training data using weights  $w_i$ .

(b) Compute

$$err^{(m)} = \sum_{i=1}^n w_i \mathbb{I}(c_i \neq T^{(m)}(\mathbf{x}_i)) / \sum_{i=1}^n w_i.$$

(c) Compute

$$\alpha^{(m)} = \log \frac{1 - err^{(m)}}{err^{(m)}} + \log(K - 1). \quad (1)$$

(d) Set

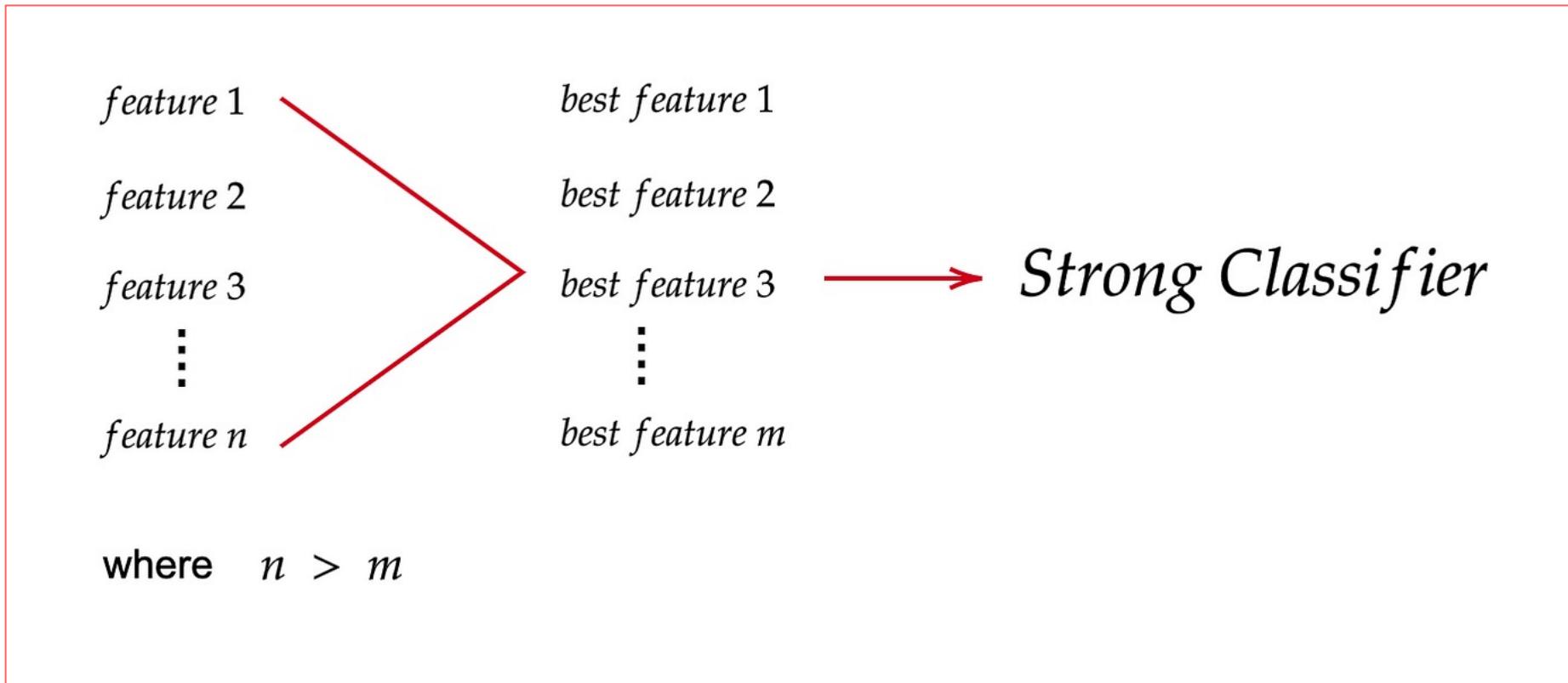
$$w_i \leftarrow w_i \cdot \exp \left( \alpha^{(m)} \cdot \mathbb{I}(c_i \neq T^{(m)}(\mathbf{x}_i)) \right), \quad i = 1, \dots, n.$$

(e) Re-normalize  $w_i$ .

3. Output

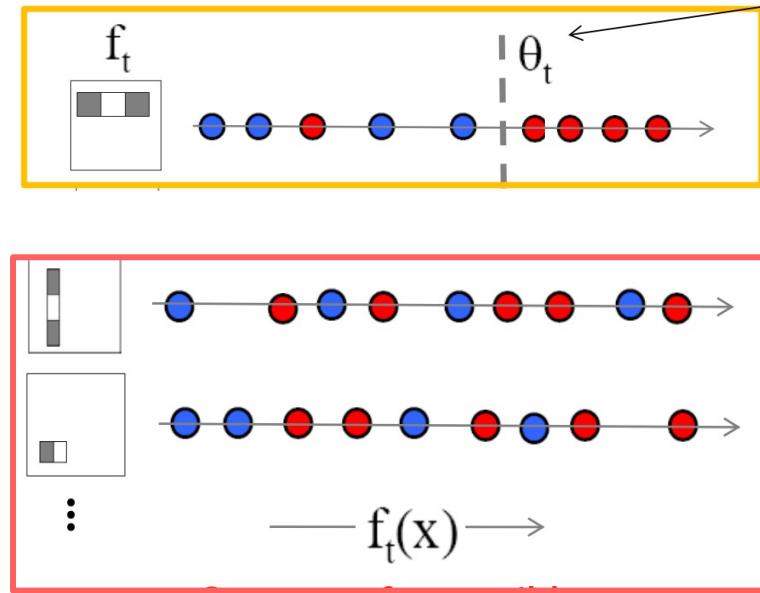
$$C(\mathbf{x}) = \arg \max_k \sum_{m=1}^M \alpha^{(m)} \cdot \mathbb{I}(T^{(m)}(\mathbf{x}) = k).$$

# Adaboost for Face Detection



The goal of using the AdaBoost algorithm is to extract the best features from n features

# AdaBoost for feature+classifier selection



$\theta_t$  is a threshold for classifier  $h^t$

**Resulting weak classifier:**

$$h_t(x) = \begin{cases} +1 & \text{if } f_t(x) > \theta_t \\ 0 & \text{otherwise} \end{cases}$$

For next round, reweight the examples according to errors, choose another filter/threshold combo.

- Each weak classifier works on exactly one rectangle feature.
- Each weak classifier has 3 associated variables
  - its threshold  $\theta$
  - its polarity  $p$
  - its weight  $\alpha$
- The polarity can be 0 or 1
- The weak classifier computes its one feature  $f$ 
  - When the polarity is 1, we want  $f > \theta$  for face
  - When the polarity is 0, we want  $f < \theta$  for face
- The weight will be used in the final classification by AdaBoost.

$$h(x) = 1 \text{ if } p \cdot f(x) < p\theta, \text{ else } 0$$

The code does not actually compute  $h$ .

Want to select the single rectangle feature and threshold that best separates positive (faces) and negative (non-faces) training examples, in terms of weighted error

- Initialized weights for each training example
- Normalized all the weights
- Then we selected the best weak classifier based on the weighted error of the training examples
- Then updated the weights according to the error of the chosen best weak classifier
- Repeated steps 2-4 N times, where N is the desired number of weak classifiers

# AdaBoost for feature+classifier selection

- Given example images  $(x_1, y_1), \dots, (x_n, y_n)$  where  $y_i = 0, 1$  for negative and positive examples respectively.
- Initialize weights  $w_{1,i} = \frac{1}{2m}, \frac{1}{2l}$  for  $y_i = 0, 1$  respectively, where  $m$  and  $l$  are the number of negatives and positives respectively.
- For  $t = 1, \dots, T$ :

- Normalize the weights,

$$w_{t,i} \leftarrow \frac{w_{t,i}}{\sum_{j=1}^n w_{t,j}}$$

so that  $w_t$  is a probability distribution.

- For each feature,  $j$ , train a classifier  $h_j$  which is restricted to using a single feature. The error is evaluated with respect to  $w_t$ ,  $\epsilon_j = \sum_i w_i |h_j(x_i) - y_i|$ .
- Choose the classifier,  $h_t$ , with the lowest error  $\epsilon_t$ .
- Update the weights:

$$w_{t+1,i} = w_{t,i} \beta_t^{1-\epsilon_t}$$

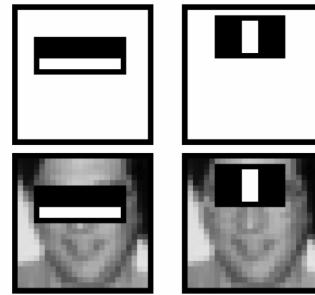
where  $e_i = 0$  if example  $x_i$  is classified correctly,  $e_i = 1$  otherwise, and  $\beta_t = \frac{\epsilon_t}{1-\epsilon_t}$ .

- The final strong classifier is:

$$h(x) = \begin{cases} 1 & \sum_{t=1}^T \alpha_t h_t(x) \geq \frac{1}{2} \sum_{t=1}^T \alpha_t \\ 0 & \text{otherwise} \end{cases}$$

where  $\alpha_t = \log \frac{1}{\beta_t}$

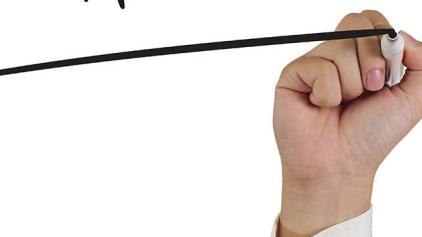
First two features selected by boosting:



This feature combination can yield 100% detection rate and 50% false positive rate



LIMITATION

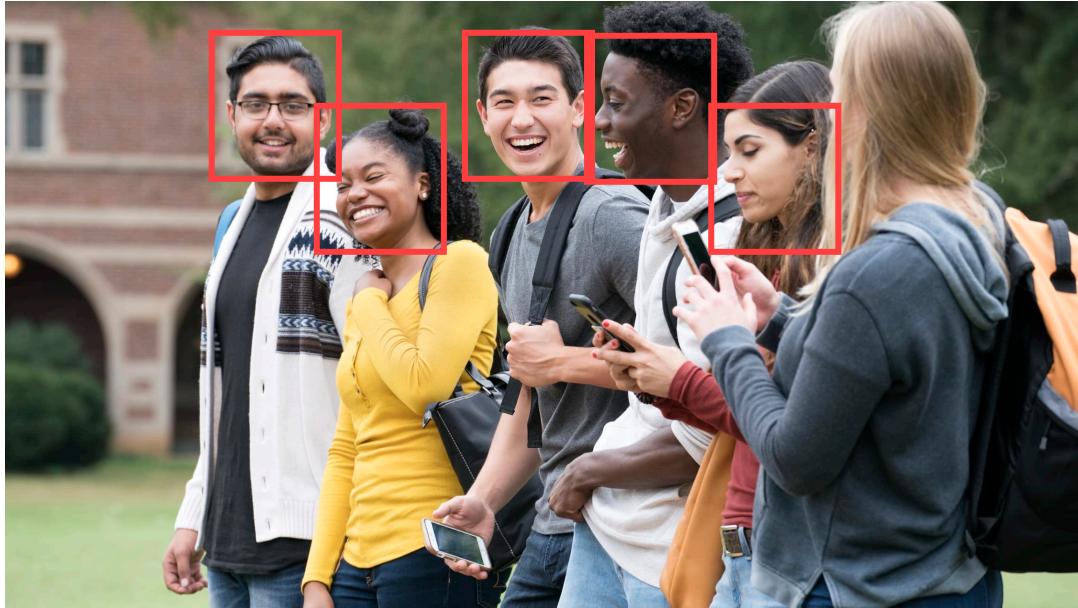


A 200-feature classifier can yield 95% detection rate and a false positive rate of 1 in 14084 50

# Outline

- Applications of Face Detection
- Face Detection in Computer Vision
- Haar Feature
- Integral Image
- Adaboost For Face Detection
- Cascade Classifier
- Face Recognition

# The Cascade Classifier

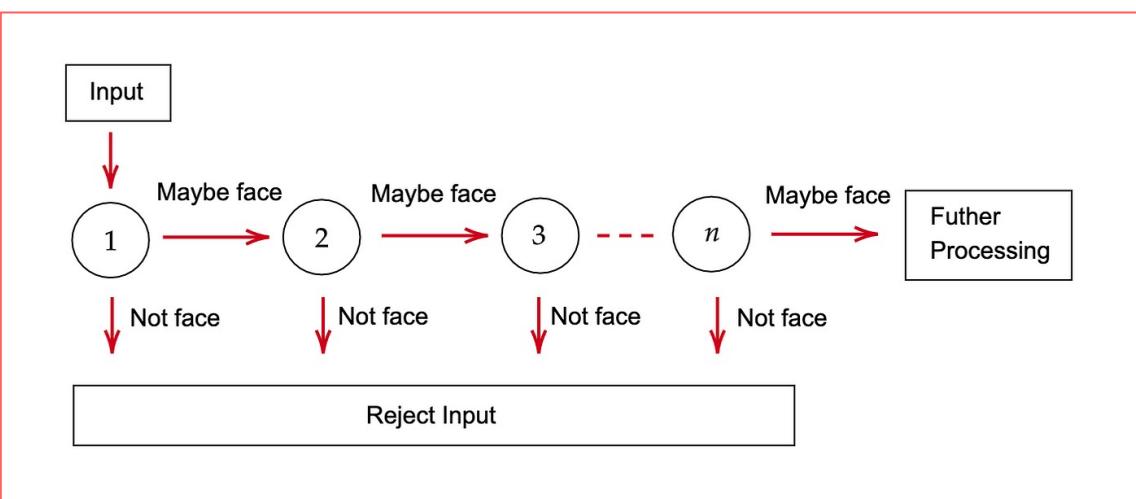


A series of classifiers are applied to every sub-window. These classifiers are simple decision trees :

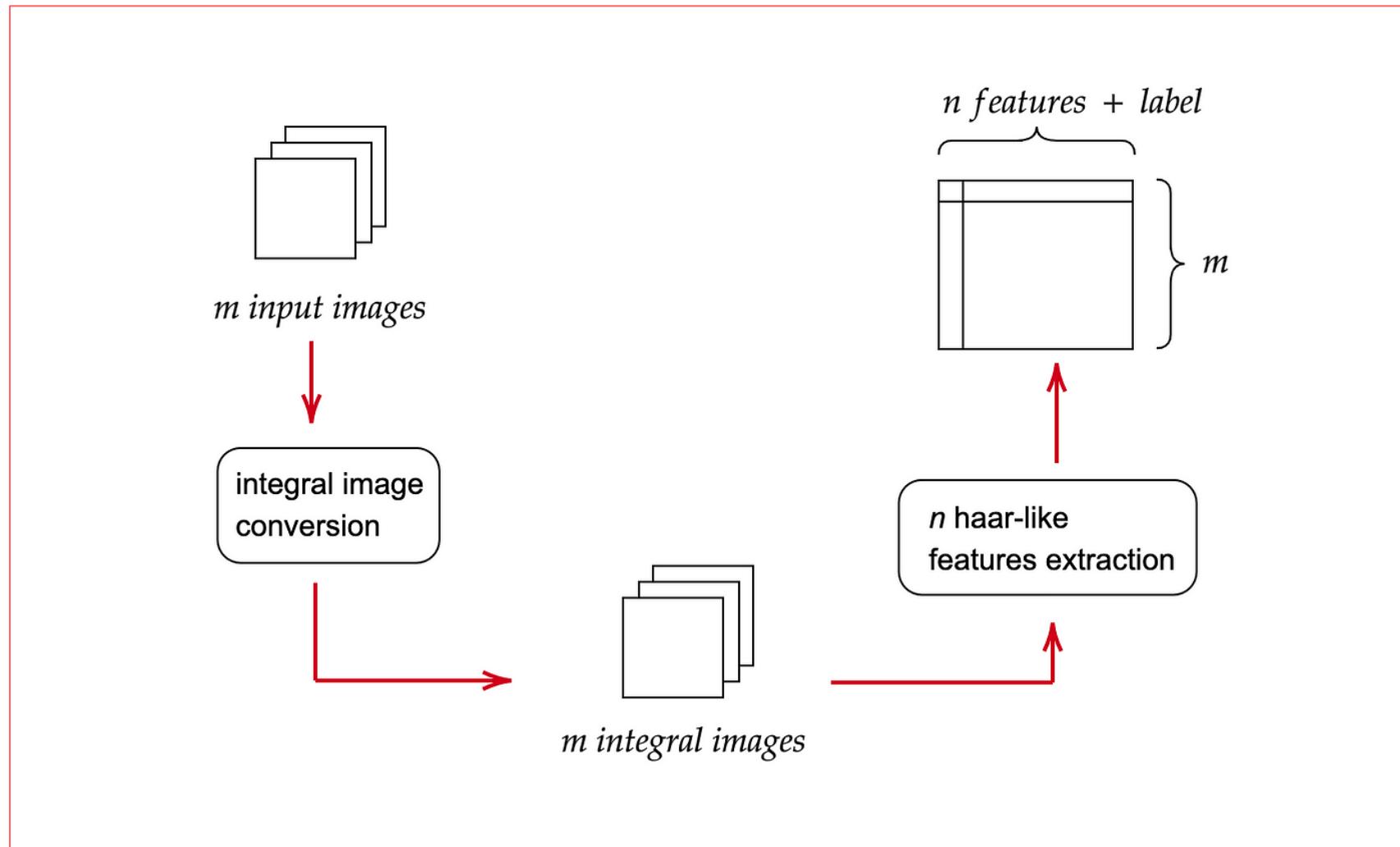
- if the first classifier is positive, we move on to the second
- if the second classifier is positive, we move on to the third.
- Any negative result at some point leads to a rejection of the sub-window as potentially containing a face

In an image, most of the image is a non-face region

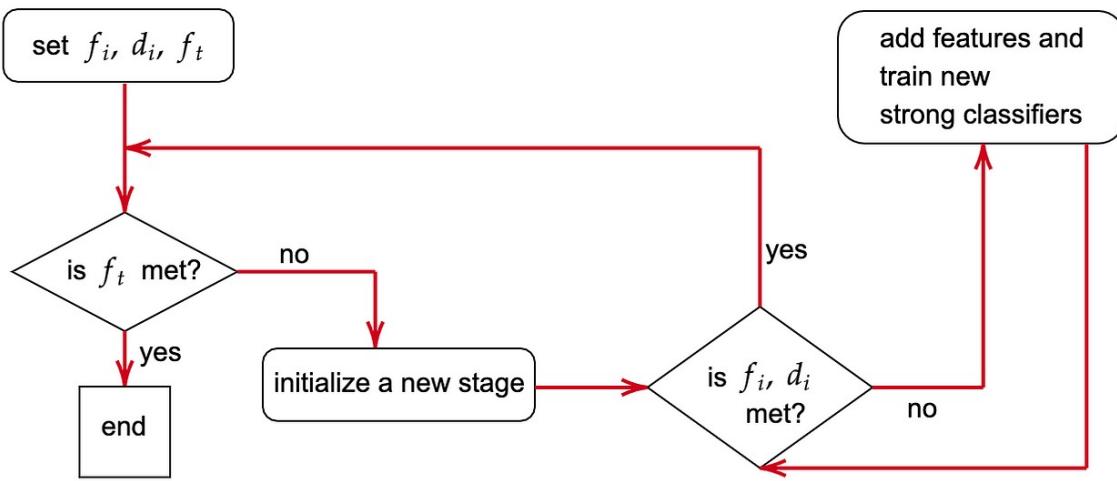
The key idea is to reject sub-windows that do not contain faces while identifying regions that do. Since the task is to identify properly the face, we want to minimize the false negative rate, i.e the sub-windows that contain a face and have not been identified as such.



# Data Preparation



# AdaBoost Algorithm + the Cascade Classifier



$f_i$  = maximum acceptable false positive rate per stage

$d_i$  = minimum acceptable true positive rate per stage

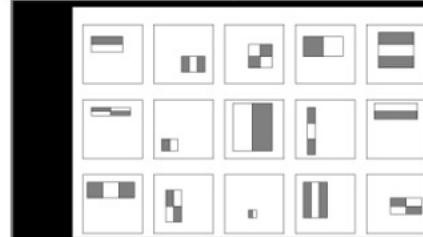
$f_t$  = target overall false positive rate

*In their paper, Viola & Jones (2001) mentioned that their Cascade Classifier has 38 stages (38 strong classifiers) that are made up of over 6000 features*

# Dicussion



Train cascade of classifiers with AdaBoost



Train with 5K positives, 350M negatives

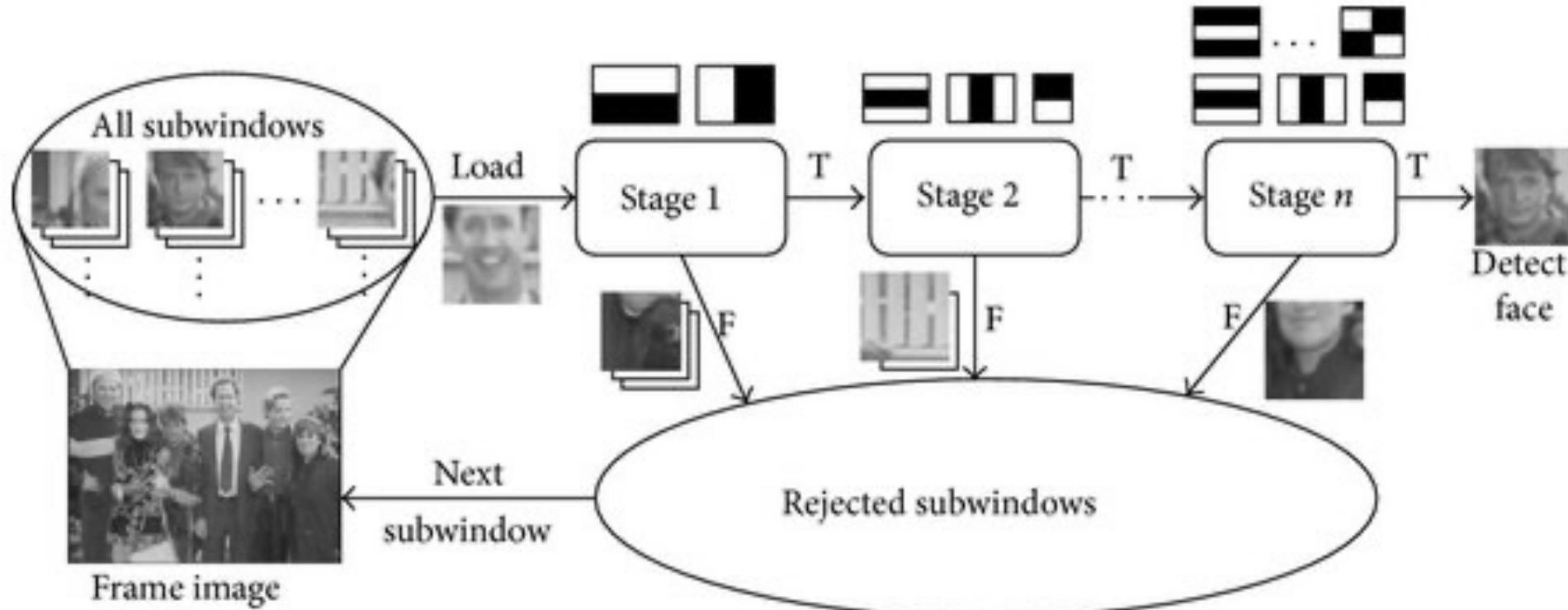
Real-time detector using 38 layer cascade

6061 features in final layer

[Implementation available in OpenCV:

<http://www.intel.com/technology/computing/opencv/>

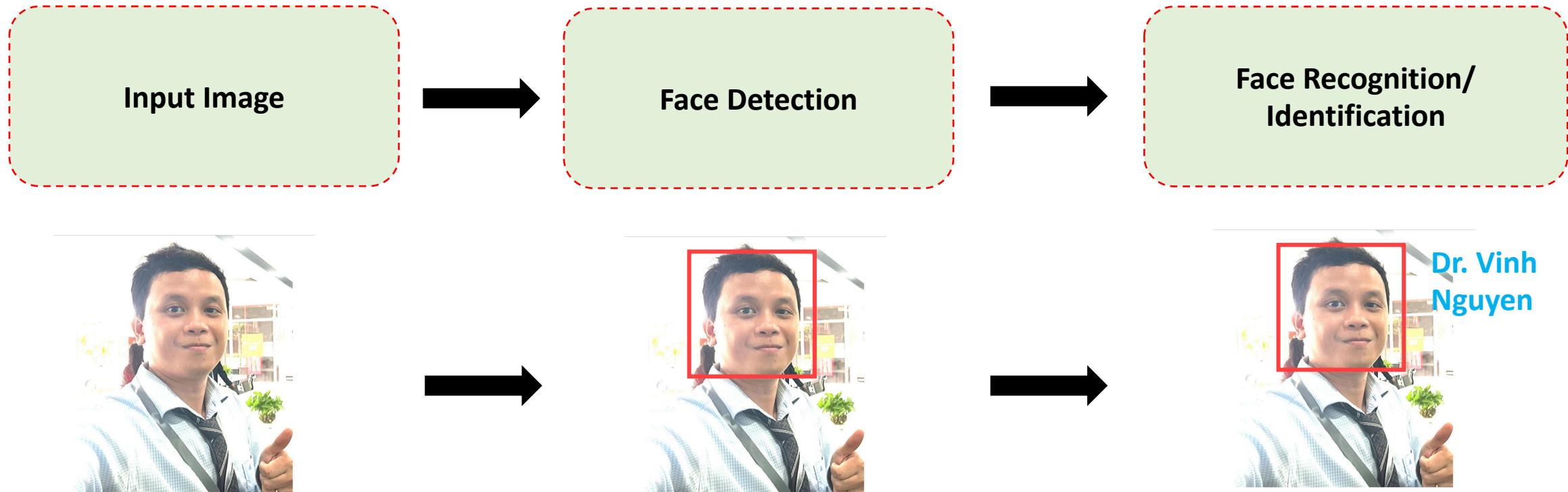
# Dicussion



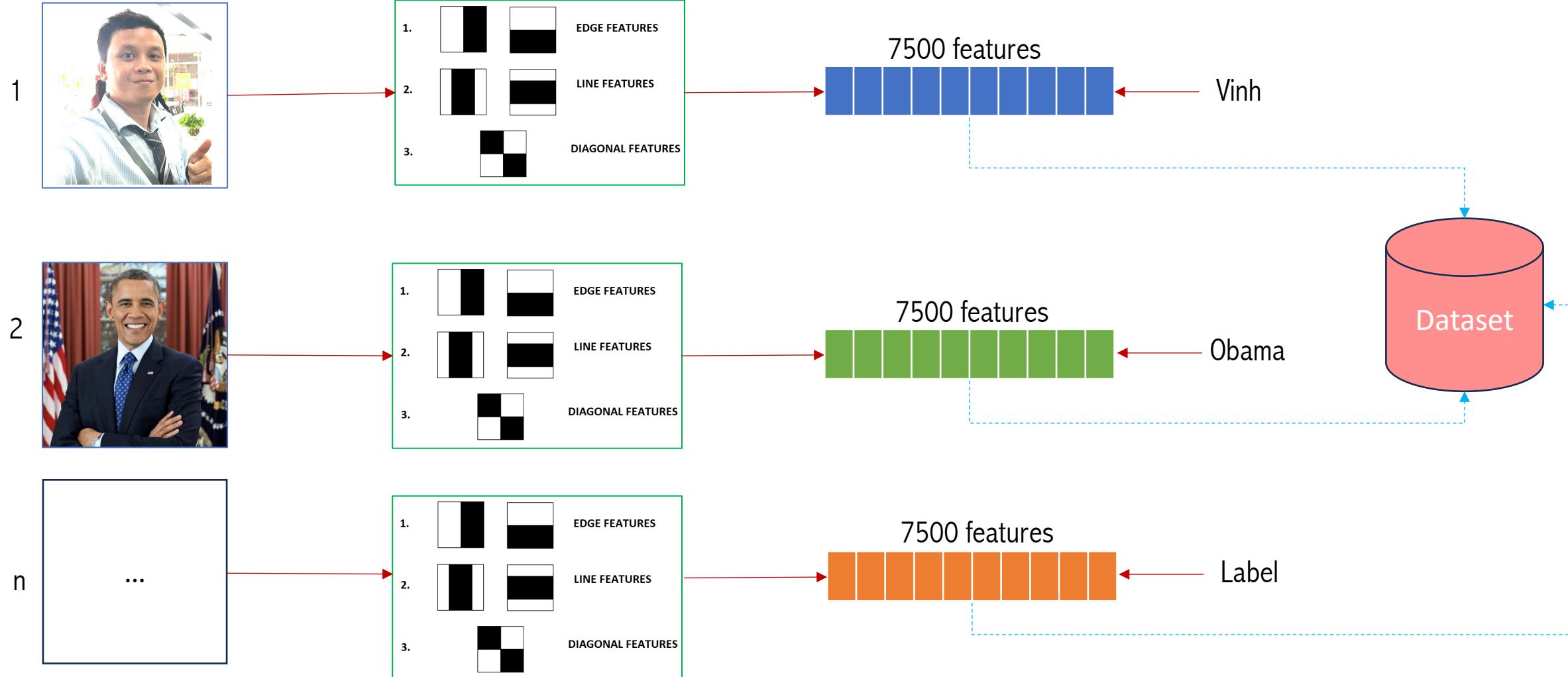
# Outline

- Applications of Face Detection
- Face Detection in Computer Vision
- Haar Feature
- Integral Image
- Adaboost For Face Detection
- Cascade Classifier
- Face Recognition

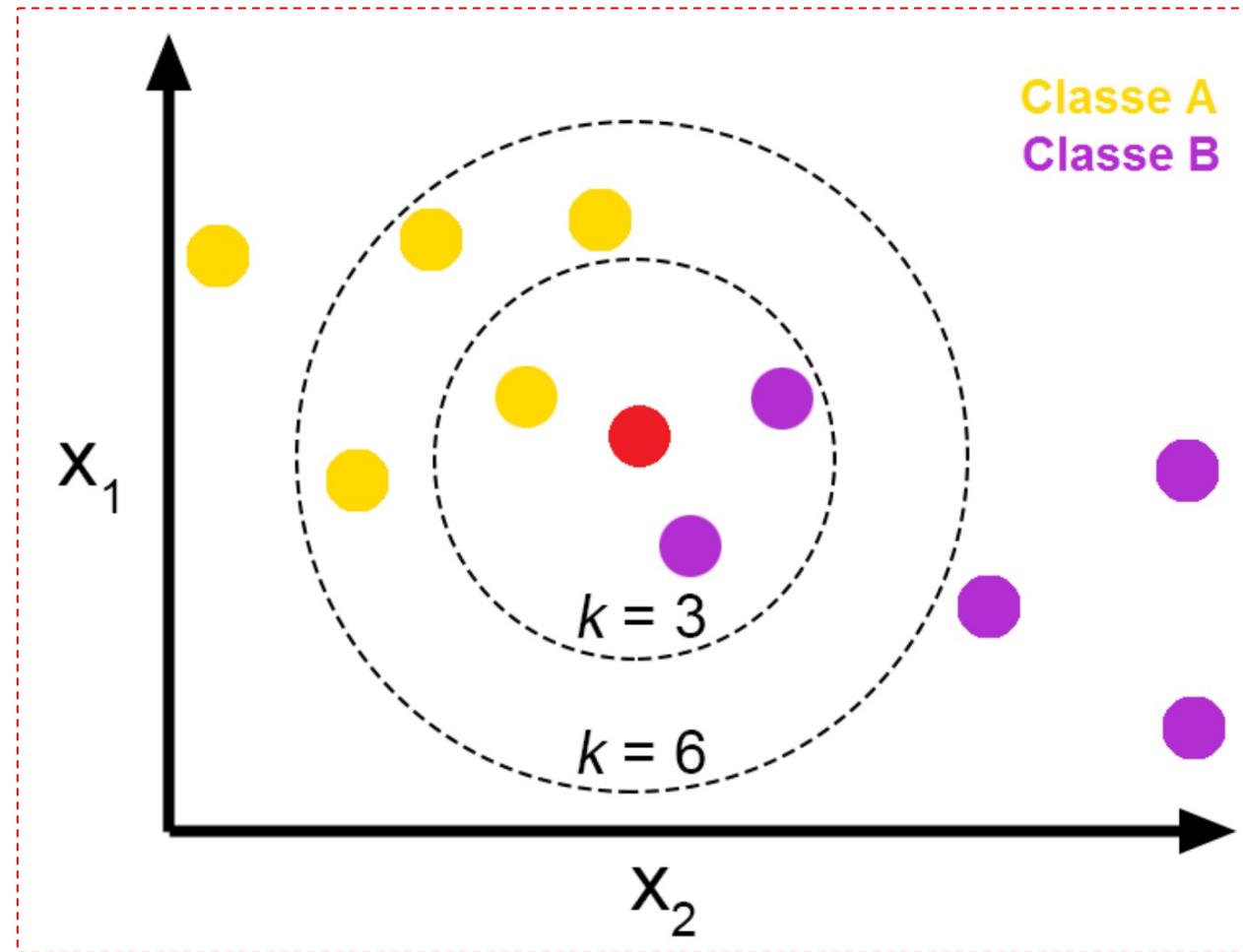
# Face Recognition/Identification



# Data Preparation



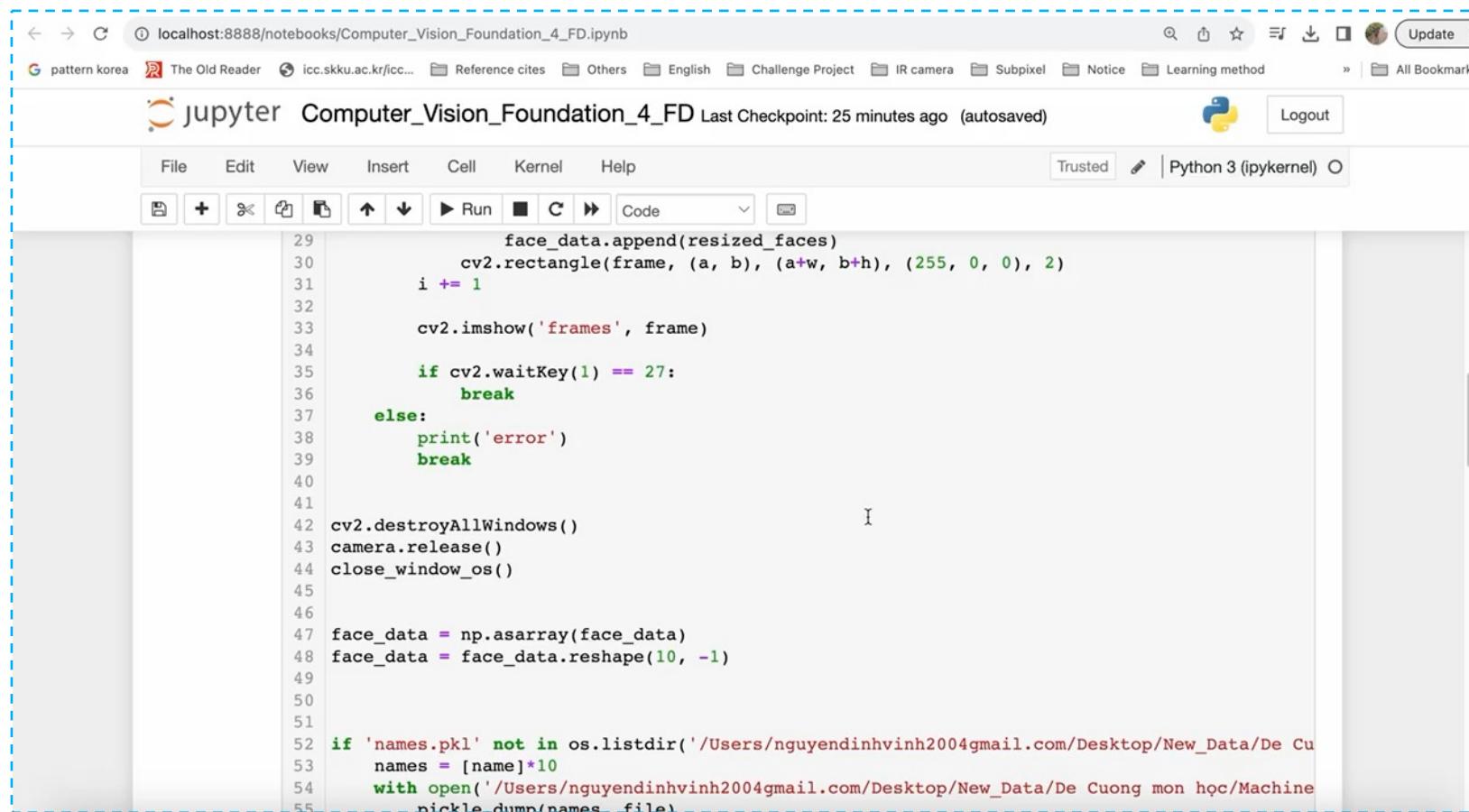
# KNN Review



# KNN for Face Recognition



# Face Recognition with KNN



The screenshot shows a Jupyter Notebook interface with a blue dashed border. The title bar says "localhost:8888/notebooks/Computer\_Vision\_Foundation\_4\_FD.ipynb". The toolbar includes File, Edit, View, Insert, Cell, Kernel, Help, Trusted, Python 3 (ipykernel), and Logout. Below the toolbar is a toolbar with icons for file operations like Open, Save, Run, and Cell. The main area contains the following Python code:

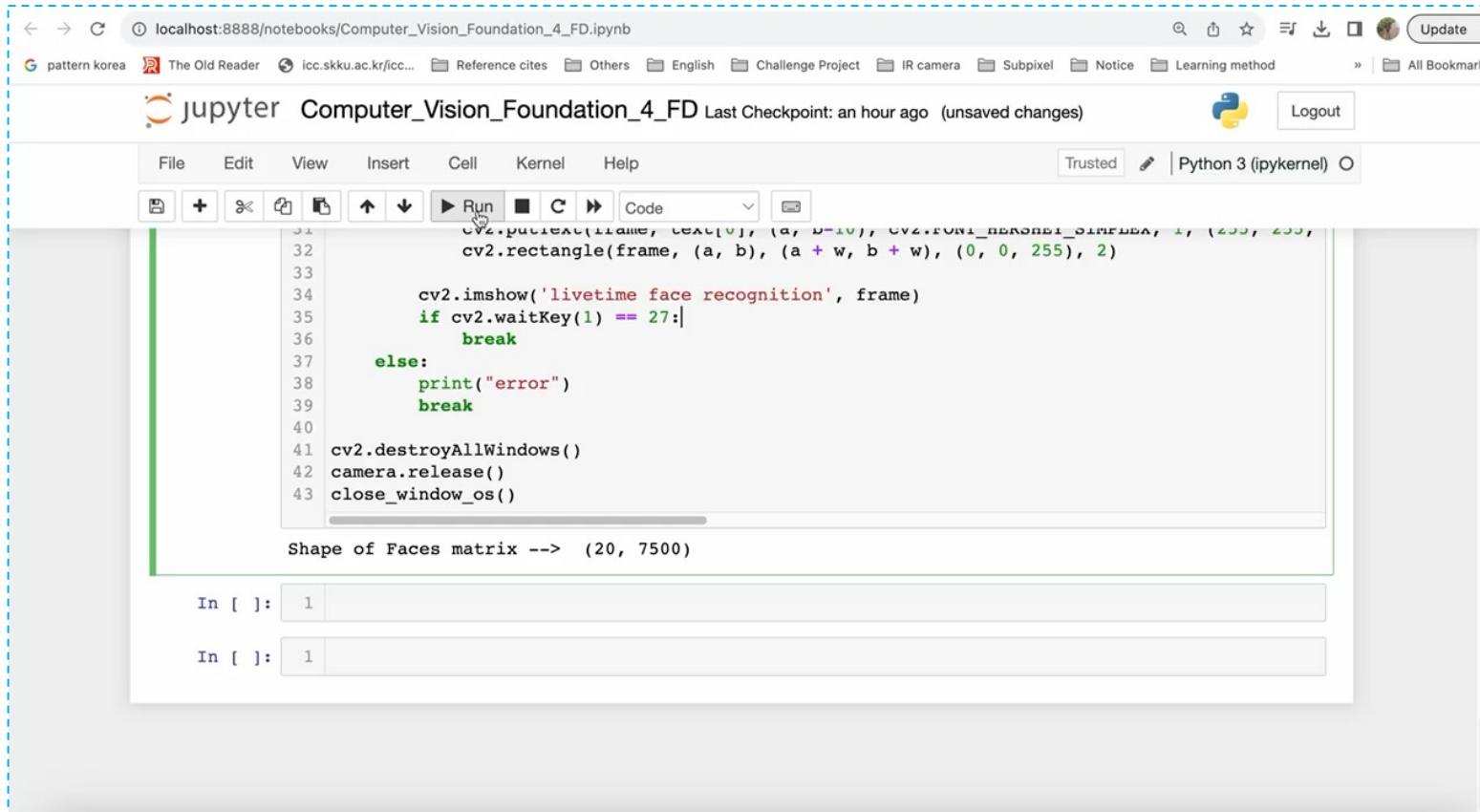
```
29     face_data.append(resized_faces)
30     cv2.rectangle(frame, (a, b), (a+w, b+h), (255, 0, 0), 2)
31     i += 1
32
33     cv2.imshow('frames', frame)
34
35     if cv2.waitKey(1) == 27:
36         break
37     else:
38         print('error')
39         break
40
41 cv2.destroyAllWindows()
42 camera.release()
43 close_window_os()
44
45
46 face_data = np.asarray(face_data)
47 face_data = face_data.reshape(10, -1)
48
49
50
51
52 if 'names.pkl' not in os.listdir('/Users/nguyendinhvinh2004@gmail.com/Desktop/New_Data/De Cuong mon hoc/Machine')
53     names = [name]*10
54     with open('/Users/nguyendinhvinh2004@gmail.com/Desktop/New_Data/De Cuong mon hoc/Machine/nickle_duan/names_file1.pkl', 'wb') as f:
55         pickle.dump(names, f)
```

Data preparation in 10 seconds (Demo)

Collect new faces and store to the database

Play Video

# Face Recognition with KNN



```
cv2.putText(frame, text, (x, y), cv2.FONT_HERSHEY_SIMPLEX, 1, (255, 255, 255), 2)
cv2.rectangle(frame, (a, b), (a + w, b + h), (0, 0, 255), 2)

cv2.imshow('livetime face recognition', frame)
if cv2.waitKey(1) == 27:
    break
else:
    print("error")

cv2.destroyAllWindows()
camera.release()
close_window_os()

Shape of Faces matrix --> (20, 7500)
```

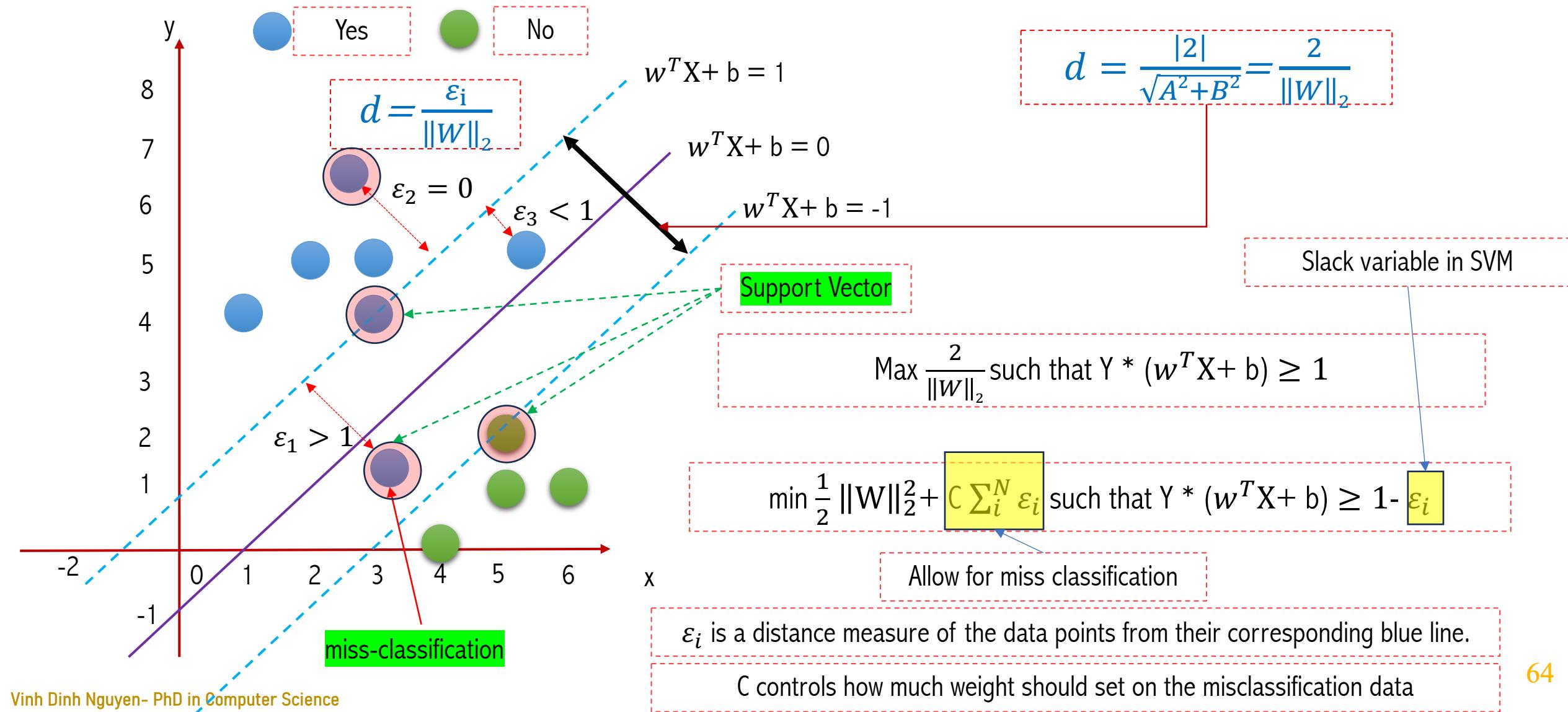
In [ ]: 1

In [ ]: 1

Provide instant detection and identification  
after 10 seconds

Play Video

# SVM Review



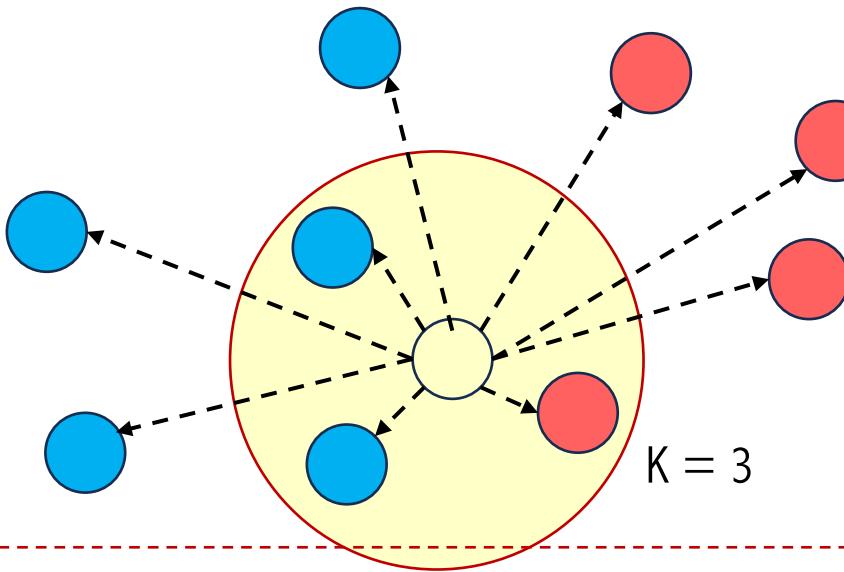
# SVM for Face Recognition



# Decision Tree Review

## KNN Limitations

Full dataset



Data cần phân loại

Đúng

Sub-dataset

Điều kiện

Sub-dataset

## Decision Tree Idea

Sai

Sub-dataset

Điều kiện

Sub-dataset

Sub-dataset

Dự đoán class A

Dự đoán class B

Dự đoán class A

Dự đoán class B

KNN has some drawbacks and challenges, such as computational expense, slow speed, for large datasets

# Decision Tree for Face Recognition



