

## Group-10:

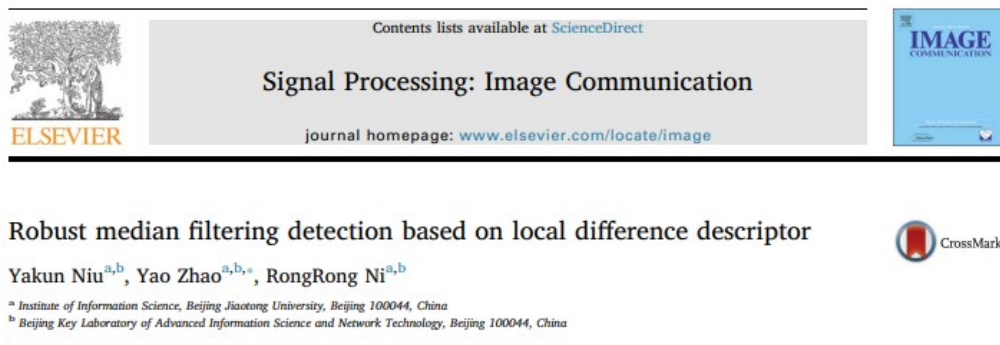
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### Research Paper:

We have selected the research paper with **Code Not Available**.

All the code has been written by us in Full.



## THEORY

We will use a **Local Difference Descriptor (LDD)** which is a set of features that we will use to discriminate between median filtered and non-median filtered images.

*LDD* will consist of:

1. **Joint Histogram of Rotation Invariant Uniform LBP**
2. **Correlation Coefficient of PDM**

### 1. Joint Histogram of Rotation Invariant Uniform LBP:

#### *LBP (Local Binary Pattern):*

LBP is a simple yet very efficient texture operator which labels the pixels of an image by thresholding the neighborhood of each pixel and considers the result as a binary number. Due to its discriminative power and computational simplicity, LBP texture operator has become a popular approach in various applications. It can be used for texture classification, segmentation, face recognition, and other applications.

#### **Formulation:**

The LBP operator can be written as:

$$LBP_{P,R} = \sum_{p=0}^{P-1} s(g_p - g_c) 2^p$$

where  $g_c$  is the intensity of the center pixel and  $g_p$  is the intensity of the neighboring pixels.  $s(x)$  is a function that returns 0 if  $x$  is negative and 1 otherwise.  $P$  is the number of pixels in the neighborhood and  $R$  is the radius of the neighborhood.

#### Rotation Invariant LBP:

In rotation invariant LBP, we cycle through all possible rotations of the binary pattern and use the smallest one as the result. For example, if the binary pattern is 10010000, we would get 01001000, 00100100, 00010010, 00001001, 00000100, 00000010, 00000001, and 10000010. The smallest one is 00000001, which is the result of rotation invariant LBP.

#### Formulation:

The rotation invariant LBP operator can be written as:

$$LBP_{P,R}^{ri} = \min\{ROR(x, i)\}$$

where  $ROR(x, i)$  is the result of rotating the binary pattern  $x$  by  $i$  bits to the right.

#### Uniform LBP:

A binary pattern is called uniform if it contains at most two bitwise transitions from 0 to 1 or vice versa when the bit pattern is traversed circularly. For example, 00000000 and 01110000 are uniform patterns, while 01010000 and 11000000 are not.

#### Rotation Invariant Uniform LBP:

In the paper, they have talked about rotation invariant uniform LBP because it is more discriminative than the original LBP.

#### Formulation:

The rotation invariant uniform LBP operator can be written as:

$$LBP_{P,R}^{riu2} = \begin{cases} \sum_{p=0}^{P-1} s(g_p - g_c) 2^p & \text{if } U(LBP_{P,R}) \leq 2 \\ P+1 & \text{otherwise} \end{cases}$$

where  $U(x)$  is the number of bitwise transitions in the binary pattern  $x$ .

#### CLBP (Completed Local Binary Pattern):

Because the conventional **LBP** descriptor just use the sign information of image local difference, and the magnitude information is discarded. Another paper (referenced by this paper) proposed a **completed LBP** descriptor by decomposing the image local difference into two complementary components, i.e., the *Sign*( $s_p$ ) and the *Magnitude*( $m_p$ ), respectively.

$$s_p = s(g_p - g_c)$$

$$m_p = |g_p - g_c|$$

where  $g_p$ ,  $g_c$  and  $s(x)$  are defined above.

Then they defined the  $CLBP - Sign (CLBPS)$  (i.e.,  $LBP$ ) and  $CLBP - Magnitude (CLBPM)$  as:

$$CLBPM_{p,R} = \sum_{p=0}^{P-1} s(m_p - c) 2^p$$

where  $c$  denotes the mean value of  $m_p$  in the whole image.

Hence, we get:

$$CLBPS_{p,R}^{riu2} = \begin{cases} \sum_{p=0}^{P-1} s(g_p - g_c) 2^p & \text{if } U(LBP_{p,R}) \leq 2 \\ P+1 & \text{otherwise} \end{cases}$$

$$CLBPM_{p,R}^{riu2} = \begin{cases} \sum_{p=0}^{P-1} s(m_p - c) 2^p & \text{if } U(LBP_{p,R}) \leq 2 \\ P+1 & \text{otherwise} \end{cases}$$

**Note: The Outputs will be converted to Rotation Invariant Form.**

The center gray level component of  $CLBP$  is discard, because it describes the global information of an image. The histogram of the  $CLBPC$  cannot distinguish the median filtered images from non-median filtered images. The mappings from  $LBP_{p,R}$  to  $CLBPS_{p,R}^{riu2}$  and  $CLBPM_{p,R}^{riu2}$ , which have  $P+2$  distinct output values, can be implemented with a lookup table of  $2^P$  elements.

Finally, we use the joint 2D histogram of the  $CLBPS_{p,R}^{riu2}$  and  $CLBPM_{p,R}^{riu2}$  codes as our first features set ( $JHLBP$ ) with  $(P+2)^2$  dimensions for median filtering detection.

## 2. Correlation Coefficients of PDM

The local pixel differences can better describe how pixel values change and implicitly encode important micro structures. To show the behavior of local pixel difference pair, we present the joint probability distribution of local pixel difference pair which is denoted as:

$$P_{i,j}^{p,q}(t_x, t_y) = Pr(g_{i,j}^p - g_{i,j} = t_x, g_{i,j}^q - g_{i,j} = t_y)$$

where  $g_{i,j}^p$  and  $g_{i,j}^q$  are the pixel values of the  $p_{th}$  and  $q_{th}$  neighbors of the center pixel  $(i, j)$ , respectively.  $t_x$  and  $t_y$  are the local pixel difference pair.

Essentially it is the Probability of  $p_{th}$  neighbour differing by a difference of  $t_x$  from the center pixel  $g_{i,j}$  and  $q_{th}$  neighbour differing by a difference of  $t_y$  from the center pixel  $g_{i,j}$ .

In order to take advantage of the correlation between the local pixel difference pair, we compute the correlation coefficients of the PDM.

#### **PDM (Pixel Difference Matrix):**

First, we obtain a PDV (Pixel Difference Vector) for each pixel in the image. The PDV is a vector of length  $P$  which contains the difference between the center pixel and its  $P$  neighbours. The PDV is denoted as:

$$d_i = [d_{i,1}, d_{i,2}, \dots, d_{i,P}]^T$$

where  $d_i$  is the PDV of  $i^{th}$  pixel in the image.  $d_{i,j}$  is the difference between the center pixel and its  $j^{th}$  neighbour.

Now, we eliminate the PDVs whose elements are all 0 values.

Then, we construct a PDM (Pixel Difference Matrix) for the remaining PDVs. The PDM is denoted as:

$$M = \begin{bmatrix} d_{1,1} & d_{2,1} & \dots & d_{N,1} \\ d_{1,2} & d_{2,2} & \dots & d_{N,2} \\ \vdots & \vdots & \ddots & \vdots \\ d_{1,P} & d_{2,P} & \dots & d_{N,P} \end{bmatrix}$$

where  $M$  is the PDM of the image.  $N$  is the number of PDVs after eliminating the PDVs whose elements are all 0 values.

The PDV measures the differences between the center point and neighboring pixels within a patch, thus it can better describe how pixel values change and can implicitly encode important visual patterns such as edges and lines in images.

Now, Joint probability is suitable to elaborate the behavior of local difference pairs. Therefore, the *Normalized Cross Correlation (NCC)* coefficients can be used as features to capture the joint probability of local difference pairs.

#### **Normalized Cross Correlation (NCC):**

The *NCC* coefficient of  $i^{th}$  and  $j^{th}$  rows in the PDM is denoted as:

$$NCC(i, j) = \frac{cov([d_{1,i}, d_{2,i}, \dots, d_{N,i}]^T, [d_{1,j}, d_{2,j}, \dots, d_{N,j}]^T)}{\sqrt{D([d_{1,i}, d_{2,i}, \dots, d_{N,i}]^T) D([d_{1,j}, d_{2,j}, \dots, d_{N,j}]^T)}}$$

where  $cov$  is the covariance of two vectors and  $D$  is defined as:

$$D([d_{1,i}, d_{2,i}, \dots, d_{N,i}]^T) = cov([d_{1,i}, d_{2,i}, \dots, d_{N,i}]^T, [d_{1,i}, d_{2,i}, \dots, d_{N,i}]^T)$$

The *NCC* coefficient is a measure of similarity between two vectors. It is equal to 1 when the two vectors are identical, and it is equal to 0 when the two vectors are orthogonal.

Our second set of features can be summarized as follows:

1. Group the  $P$   $DV$  s to form the  $P$   $D$   $M$  with  $P$  rows;
2. Consider an arbitrary row of  $P$   $D$   $M$  as a random variable and obtain the  $NCC$  coefficients of any different variables;
3. Concatenate all the  $NCC$  coefficients of  $P$   $D$   $M$  ( $C$   $P$   $D$   $M$ ) to yield a  $\square^P C_2 - Dimensional$  feature vector  $C$ .

### 3. Final Features Set

Combining  $JHLBP$  features and  $C$   $P$   $D$   $M$  features, we obtain the final  $LDD$  features with  $(P+2)^2 + \frac{P(P-1)}{2}$  elements for median filtering detection.

**NOTE:** As instructed in the research paper, we will use 8 neighbours i.e.  $P=8$  and  $R=1$  where  $P$  is the number of neighbours and  $R$  is the radius of the neighbourhood.

## IMPLEMENTATION

### Importing Libraries

```
import numpy as np
import cv2
import matplotlib.pyplot as plt
import threading
from sklearn.discriminant_analysis import LinearDiscriminantAnalysis
```

### Importing Dataset

```
DATASET_PATH = 'ucid_dataset/ucid_v2/'
```

```
# Read All Images from the UCID Dataset
```

```
def read_images():
```

```
    images = []
```

```
    for i in range(1338):
```

```
        file_name = str(i+1)
```

```
        while(len(file_name) < 5):
```

```
            file_name = '0' + file_name
```

```
        # print("Reading Image: " + file_name)
```

```
        img = cv2.imread(DATASET_PATH + "ucid" + file_name + '.tif')
```

```
        # Convert to Gray Scale
```

```
        img = cv2.cvtColor(img, cv2.COLOR_BGR2GRAY)
```

```
        images.append(img)
```

```
    return images
```

```
ucid_images = read_images()
```

```

len(ucid_images)

1338

ucid_images[0].shape

(384, 512)

# Checking Shapes of All UCID Images
shapes = {}
for img in ucid_images:
    shapes[img.shape] = shapes.get(img.shape, 0) + 1

shapes

{(384, 512): 885, (512, 384): 453}

# All Images which are of shape (512, 384), rotate them by 90 degrees
for i in range(len(ucid_images)):
    if ucid_images[i].shape == (512, 384):
        ucid_images[i] = np.rot90(ucid_images[i])

# Checking Shapes of All UCID Images
shapes = {}
for img in ucid_images:
    shapes[img.shape] = shapes.get(img.shape, 0) + 1

shapes

{(384, 512): 1338}

```

Now all Images are of same shape

### Checking a Sample Image

```

# Displaying a Sample Image
plt.imshow(ucid_images[0], cmap='gray')
# Remove Axis
plt.axis('off')
plt.show()

```



### Creating Median Filtered and Average Filtered Images

*# Creating Median Filtered Dataset and Average Filtered Dataset*

```
median_filtered_images_3 = []  
median_filtered_images_5 = []  
average_filtered_images_3 = []
```

```
for img in ucid_images:
```

```
    median_filtered_images_3.append(cv2.medianBlur(img, 3))  
    median_filtered_images_5.append(cv2.medianBlur(img, 5))  
    average_filtered_images_3.append(cv2.blur(img, (3, 3)))
```

*# Converting to Numpy Arrays*

```
ucid_images = np.array(ucid_images)  
median_filtered_images_3 = np.array(median_filtered_images_3)  
median_filtered_images_5 = np.array(median_filtered_images_5)  
average_filtered_images_3 = np.array(average_filtered_images_3)
```

*# Displaying Multiple Versions of a Sample Image*

```
fig, axs = plt.subplots(2, 2, figsize=(15, 15))
```

```
axs[0, 0].imshow(ucid_images[5], cmap='gray')  
axs[0, 0].set_title('Original Image')
```

```
axs[0, 1].imshow(median_filtered_images_3[5], cmap='gray')  
axs[0, 1].set_title('Median Filtered Image (3x3)')
```



```

axs[1, 0].imshow(median_filtered_images_5[5], cmap='gray')
axs[1, 0].set_title('Median Filtered Image (5x5)')

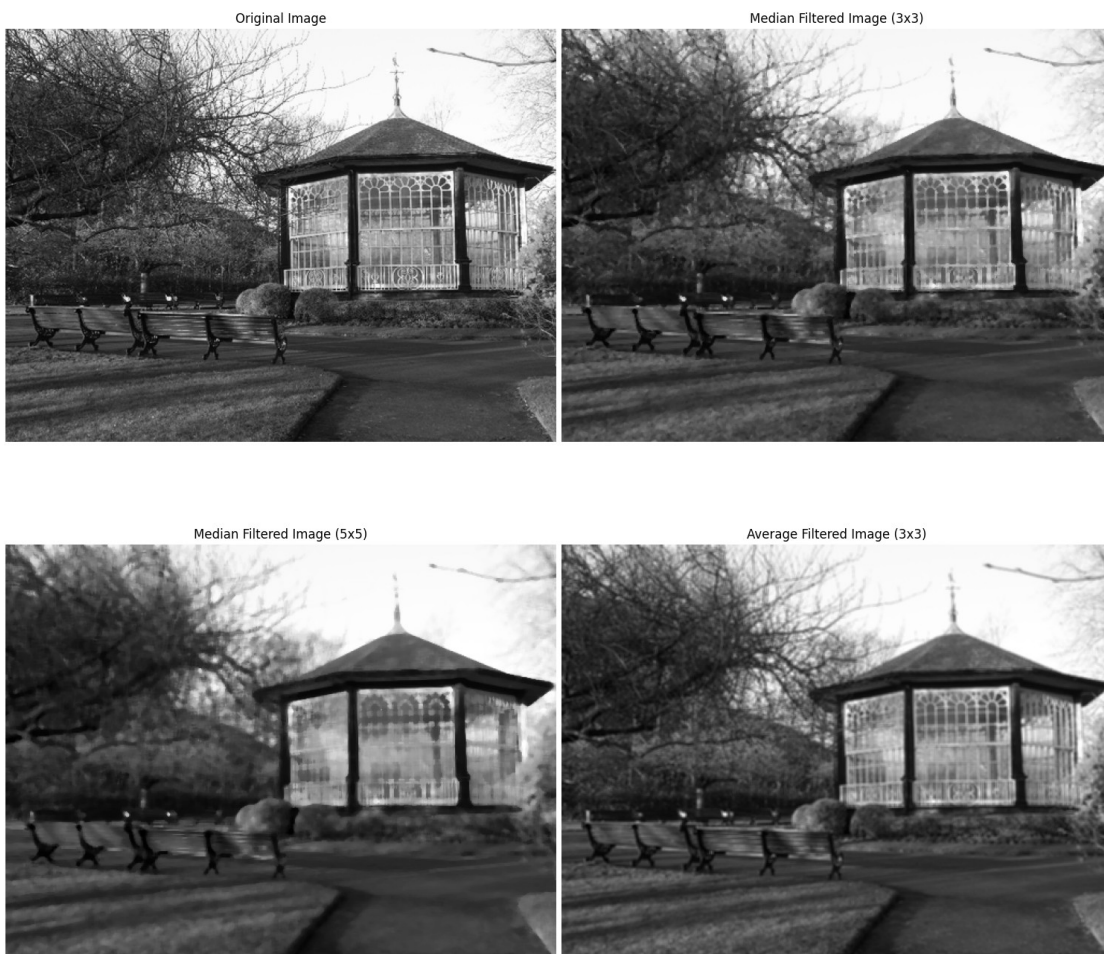
axs[1, 1].imshow(average_filtered_images_3[5], cmap='gray')
axs[1, 1].set_title('Average Filtered Image (3x3)')

# Remove Axis
for ax in axs.flat:
    ax.axis('off')

# Gap in first row and second row
plt.tight_layout(pad=0.5)

plt.show()

```



## Implementing LDD (Local Difference Descriptor) Features

### 1. Joint Histogram of Rotation Invariant Uniform LBP

```
ucid_images.shape
```

```
(1338, 384, 512)
```



## Function to get Joint Histogram of Rotation Invariant Uniform LBP

*# Function to calculate Average value of  $|G_p - G_c|$  where  $c$  is the center pixel and  $p$  is the neighbour pixel*

```
def get_average_difference(img):  
  
    c = 0  
    cnt = 0  
  
    # Neighbours  
    dx = [-1, -1, -1, 0, 0, 1, 1, 1]  
    dy = [-1, 0, 1, -1, 1, -1, 0, 1]  
  
    for i in range(1, img.shape[0]-1):  
        for j in range(1, img.shape[1]-1):  
  
            p = img[i][j]  
  
            for k in range(8):  
                x = i + dx[k]  
                y = j + dy[k]  
                cnt += 1  
                c += abs(int(p) - int(img[x][y]))  
  
    return c/cnt
```

*# Function to get rotationally invariant string for a given binary string*

```
def rotational_invariant(binary_string):  
  
    # Get all possible rotations  
    rotations = []  
    for i in range(len(binary_string)):  
        rotations.append(binary_string[i:] + binary_string[:i])  
  
    # Sort the rotations  
    rotations.sort()  
  
    # Return the first rotation  
    return rotations[0]
```

*# Function to get joint histogram of rotationally invariant uniform lbp*

```
def get_joint_histogram(img):  
  
    dx = [-1, -1, -1, 0, 0, 1, 1, 1]  
    dy = [-1, 0, 1, -1, 1, -1, 0, 1]
```

```

# BINS
pattern_to_index = {
    "00000000": 0,
    "00000001": 1,
    "00000011": 2,
    "00000111": 3,
    "00001111": 4,
    "00011111": 5,
    "00111111": 6,
    "01111111": 7,
    "11111111": 8,
    "OTHER": 9
}

# HISTOGRAM
joint_histogram = np.zeros((10, 10))

# Calculate Average value of |G_p - G_c| where c is the center
pixel and p is the neighbour pixel
c = get_average_difference(img)

# Calculate CLBPS and CLBPM for each pixel
for i in range(1, img.shape[0]-1):
    for j in range(1, img.shape[1]-1):

        # Get CLBPS and CLBPM for this pixel
        clbps = ""
        clbpm = ""
        for k in range(8):
            x = i + dx[k]
            y = j + dy[k]

            if img[x][y] > img[i][j]:
                clbps += "1"
            else:
                clbps += "0"

            if abs(int(img[x][y]) - int(img[i][j])) > c:
                clbpm += "1"
            else:
                clbpm += "0"

        # Convert till rotationally invariant
        clbps = rotational_invariant(clbps)
        clbpm = rotational_invariant(clbpm)

        # Check Number of 0-1 and 1-0 transitions
        transitions_clbps = 0
        transitions_clbpm = 0

```

```

    for k in range(1, len(clbps)):
        if clbps[k] != clbps[k-1]:
            transitions_clbps += 1
        if clbpm[k] != clbpm[k-1]:
            transitions_clbpm += 1

    if transitions_clbps > 2:
        # P+1 ROTATION INVARIANT FORM
        clbps = "00001001"
    if transitions_clbpm > 2:
        # P+1 ROTATION INVARIANT FORM
        clbpm = "00001001"

    if clbps not in pattern_to_index:
        clbps = "OTHER"

    if clbpm not in pattern_to_index:
        clbpm = "OTHER"

    # Update Joint Histogram
    joint_histogram[pattern_to_index[clbps]]
[pattern_to_index[clbpm]] += 1

    # Convert to Numpy Array
    joint_histogram = np.array(joint_histogram)

    # Return Joint Histogram
    return joint_histogram

# Try for a sample image
joint_histogram = get_joint_histogram(ucid_images[5])

joint_histogram.shape

(10, 10)

# Function to calculate Joint Histogram for all images
def calculate_joint_histograms(images, TARGET_LIST,
DATASET_NAME="DEFAULT"):
    for i in range(len(images)):
        if i%100 == 0:
            print(f"{i}/1338 images done for {DATASET_NAME}")
            TARGET_LIST.append(get_joint_histogram(images[i]))

# Calculate Joint Histogram for all images
joint_histograms_ucid = []
joint_histograms_median_filtered_3 = []
joint_histograms_median_filtered_5 = []
joint_histograms_average_filtered_3 = []

```

```
# RUN FOR EVERY DATASET IN A SEPARATE THREAD
t1 = threading.Thread(target=lambda:
calculate_joint_histograms(ucid_images, joint_histograms_ucid,
"UCID"))
t2 = threading.Thread(target=lambda:
calculate_joint_histograms(median_filtered_images_3,
joint_histograms_median_filtered_3, "MEDIAN_FILTERED_3"))
t3 = threading.Thread(target=lambda:
calculate_joint_histograms(median_filtered_images_5,
joint_histograms_median_filtered_5, "MEDIAN_FILTERED_5"))
t4 = threading.Thread(target=lambda:
calculate_joint_histograms(average_filtered_images_3,
joint_histograms_average_filtered_3, "AVERAGE_FILTERED_3"))
```

```
# START ALL THREADS
```

```
t1.start()
t2.start()
t3.start()
t4.start()
```

```
# WAIT FOR ALL THREADS TO COMPLETE
```

```
t1.join()
t2.join()
t3.join()
t4.join()
```

```
0/1338 images done for UCID
0/1338 images done for MEDIAN_FILTERED_3
0/1338 images done for MEDIAN_FILTERED_5
0/1338 images done for AVERAGE_FILTERED_3
100/1338 images done for MEDIAN_FILTERED_3
100/1338 images done for UCID
100/1338 images done for MEDIAN_FILTERED_5
100/1338 images done for AVERAGE_FILTERED_3
200/1338 images done for MEDIAN_FILTERED_3
200/1338 images done for UCID
200/1338 images done for MEDIAN_FILTERED_5
200/1338 images done for AVERAGE_FILTERED_3
300/1338 images done for MEDIAN_FILTERED_3
300/1338 images done for UCID
300/1338 images done for MEDIAN_FILTERED_5
300/1338 images done for AVERAGE_FILTERED_3
400/1338 images done for UCID
400/1338 images done for MEDIAN_FILTERED_3
400/1338 images done for MEDIAN_FILTERED_5
400/1338 images done for AVERAGE_FILTERED_3
500/1338 images done for UCID
500/1338 images done for MEDIAN_FILTERED_3
500/1338 images done for MEDIAN_FILTERED_5
500/1338 images done for AVERAGE_FILTERED_3
```

```

600/1338 images done for UCID
600/1338 images done for MEDIAN_FILTERED_3
600/1338 images done for MEDIAN_FILTERED_5
700/1338 images done for UCID
600/1338 images done for AVERAGE_FILTERED_3
700/1338 images done for MEDIAN_FILTERED_3
700/1338 images done for MEDIAN_FILTERED_5
800/1338 images done for UCID
800/1338 images done for MEDIAN_FILTERED_3
700/1338 images done for AVERAGE_FILTERED_3
800/1338 images done for MEDIAN_FILTERED_5
900/1338 images done for UCID
900/1338 images done for MEDIAN_FILTERED_3
800/1338 images done for AVERAGE_FILTERED_3
1000/1338 images done for UCID
900/1338 images done for MEDIAN_FILTERED_5
1000/1338 images done for MEDIAN_FILTERED_3
900/1338 images done for AVERAGE_FILTERED_3
1100/1338 images done for UCID
1100/1338 images done for MEDIAN_FILTERED_3
1000/1338 images done for MEDIAN_FILTERED_5
1200/1338 images done for UCID
1000/1338 images done for AVERAGE_FILTERED_3
1200/1338 images done for MEDIAN_FILTERED_3
1100/1338 images done for MEDIAN_FILTERED_5
1300/1338 images done for UCID
1300/1338 images done for MEDIAN_FILTERED_3
1100/1338 images done for AVERAGE_FILTERED_3
1200/1338 images done for MEDIAN_FILTERED_5
1200/1338 images done for AVERAGE_FILTERED_3
1300/1338 images done for MEDIAN_FILTERED_5
1300/1338 images done for AVERAGE_FILTERED_3

```

```

# Convert to Numpy Arrays

```

```

joint_histograms_ucid = np.array(joint_histograms_ucid)
joint_histograms_median_filtered_3 =
np.array(joint_histograms_median_filtered_3)
joint_histograms_median_filtered_5 =
np.array(joint_histograms_median_filtered_5)
joint_histograms_average_filtered_3 =
np.array(joint_histograms_average_filtered_3)

```

```

joint_histograms_ucid.shape, joint_histograms_median_filtered_3.shape,
joint_histograms_median_filtered_5.shape,
joint_histograms_average_filtered_3.shape

```

```

((1338, 10, 10), (1338, 10, 10), (1338, 10, 10), (1338, 10, 10))

```

```

# Save Joint Histograms

```

```

# np.save("joint_histograms_ucid.npy", joint_histograms_ucid)
# np.save("joint_histograms_median_filtered_3.npy",

```

```

joint_histograms_median_filtered_3)
# np.save("joint_histograms_median_filtered_5.npy",
joint_histograms_median_filtered_5)
# np.save("joint_histograms_average_filtered_3.npy",
joint_histograms_average_filtered_3)

# Read Joint Histograms
# joint_histograms_ucid = np.load("joint_histograms_ucid.npy")
# joint_histograms_median_filtered_3 =
np.load("joint_histograms_median_filtered_3.npy")
# joint_histograms_median_filtered_5 =
np.load("joint_histograms_median_filtered_5.npy")
# joint_histograms_average_filtered_3 =
np.load("joint_histograms_average_filtered_3.npy")

joint_histograms_ucid.shape, joint_histograms_median_filtered_3.shape,
joint_histograms_median_filtered_5.shape,
joint_histograms_average_filtered_3.shape

((1338, 10, 10), (1338, 10, 10), (1338, 10, 10), (1338, 10, 10))

```

So now for every image, we have a feature vector of size 100.

## 2. Correlation Coefficients of PDM

```
ucid_images.shape
```

```
(1338, 384, 512)
```

### Function to get PDM (Pixel Difference Matrix) from an Image

```
# Function to calculate PDM of an Image
```

```
def calculate_pdm(img):
```

```
    # Calculate PDV's for all pixels
```

```
    pdm = []
```

```
    # Neighbours
```

```
    dx = [-1, -1, -1, 0, 0, 1, 1, 1]
```

```
    dy = [-1, 0, 1, -1, 1, -1, 0, 1]
```

```
    # Loop over all pixels
```

```
    for i in range(1, img.shape[0] - 1):
```

```
        for j in range(1, img.shape[1]-1):
```

```
            pdv = []
```

```
            # Flag to check if all elements are 0
```

```
            flag = False
```

```
            # Check for all 8 neighbours of the pixel
```

```
            for k in range(8):
```

```
                difference = int(img[i + dx[k]][j + dy[k]]) -
```

```

int(img[i][j])
        pdv.append(difference)
        if difference != 0:
            flag = True

        # If all elements are 0, then don't add it to the list
        if flag:
            pdm.append(pdv)

pdm = np.array(pdm)
# Take Transpose
pdm = pdm.T

return pdm

```

### Function to get Correlation Coefficients of PDM from an Image

```

# UTILS
def get_D(pdm_row):
    return np.cov(pdm_row, pdm_row)[0][1]

# Function to get Correlation Coefficients of PDM from an Image
def get_correlation_coefficients(img):

    # Calculate PDM
    pdm = calculate_pdm(img)

    # Calculate Correlation Coefficients
    corr_coefficients = []

    for i in range(pdm.shape[0]):
        for j in range(i+1, pdm.shape[0]):
            # Calculate Correlation Coefficient
            D1 = get_D(pdm[i])
            D2 = get_D(pdm[j])
            corr = np.cov(pdm[i], pdm[j])[0][1] / np.sqrt(D1 * D2)
            corr_coefficients.append(corr)

    return np.array(corr_coefficients)

# Calculating PDM of a Sample Image
calculate_pdm(ucid_images[0]).shape

(8, 186657)

# Calculating Correlation Coefficients of a Sample Image
get_correlation_coefficients(ucid_images[0]).shape

(28,)

```



```

# Calculating Correlation Coefficients of all Images
ucid_images_corr_coefficients = []
median_filtered_images_3_corr_coefficients = []
median_filtered_images_5_corr_coefficients = []
average_filtered_images_3_corr_coefficients = []

print("UCID IMAGES")
for img in ucid_images:

    ucid_images_corr_coefficients.append(get_correlation_coefficients(img)
    )

print("MEDIAN FILTERED IMAGES (3x3)")
for img in median_filtered_images_3:

    median_filtered_images_3_corr_coefficients.append(get_correlation_coef
    ficients(img))

print("MEDIAN FILTERED IMAGES (5x5)")
for img in median_filtered_images_5:

    median_filtered_images_5_corr_coefficients.append(get_correlation_coef
    ficients(img))

print("AVERAGE FILTERED IMAGES (3x3)")
for img in average_filtered_images_3:

    average_filtered_images_3_corr_coefficients.append(get_correlation_coe
    ficients(img))

UCID IMAGES
MEDIAN FILTERED IMAGES (3x3)
MEDIAN FILTERED IMAGES (5x5)
AVERAGE FILTERED IMAGES (3x3)

# Converting to Numpy Arrays
ucid_images_corr_coefficients =
np.array(ucid_images_corr_coefficients)
median_filtered_images_3_corr_coefficients =
np.array(median_filtered_images_3_corr_coefficients)
median_filtered_images_5_corr_coefficients =
np.array(median_filtered_images_5_corr_coefficients)
average_filtered_images_3_corr_coefficients =
np.array(average_filtered_images_3_corr_coefficients)

ucid_images_corr_coefficients.shape,
median_filtered_images_3_corr_coefficients.shape,
median_filtered_images_5_corr_coefficients.shape,
average_filtered_images_3_corr_coefficients.shape

((1338, 28), (1338, 28), (1338, 28), (1338, 28))

```

```

# import pickle

# # Save using Pickle
# with open('ucid_images_corr_coefficients.pkl', 'wb') as f:
#     pickle.dump(ucid_images_corr_coefficients, f)

# with open('median_filtered_images_3_corr_coefficients.pkl', 'wb') as
f:
#     pickle.dump(median_filtered_images_3_corr_coefficients, f)

# with open('median_filtered_images_5_corr_coefficients.pkl', 'wb') as
f:
#     pickle.dump(median_filtered_images_5_corr_coefficients, f)

# with open('average_filtered_images_3_corr_coefficients.pkl', 'wb')
as f:
#     pickle.dump(average_filtered_images_3_corr_coefficients, f)

# import pickle

# # Load using Pickle
# with open('ucid_images_corr_coefficients.pkl', 'rb') as f:
#     ucid_images_corr_coefficients = pickle.load(f)

# with open('median_filtered_images_3_corr_coefficients.pkl', 'rb') as
f:
#     median_filtered_images_3_corr_coefficients = pickle.load(f)

# with open('median_filtered_images_5_corr_coefficients.pkl', 'rb') as
f:
#     median_filtered_images_5_corr_coefficients = pickle.load(f)

# with open('average_filtered_images_3_corr_coefficients.pkl', 'rb')
as f:
#     average_filtered_images_3_corr_coefficients = pickle.load(f)

# Shapes
ucid_images_corr_coefficients.shape,
median_filtered_images_3_corr_coefficients.shape,
median_filtered_images_5_corr_coefficients.shape,
average_filtered_images_3_corr_coefficients.shape

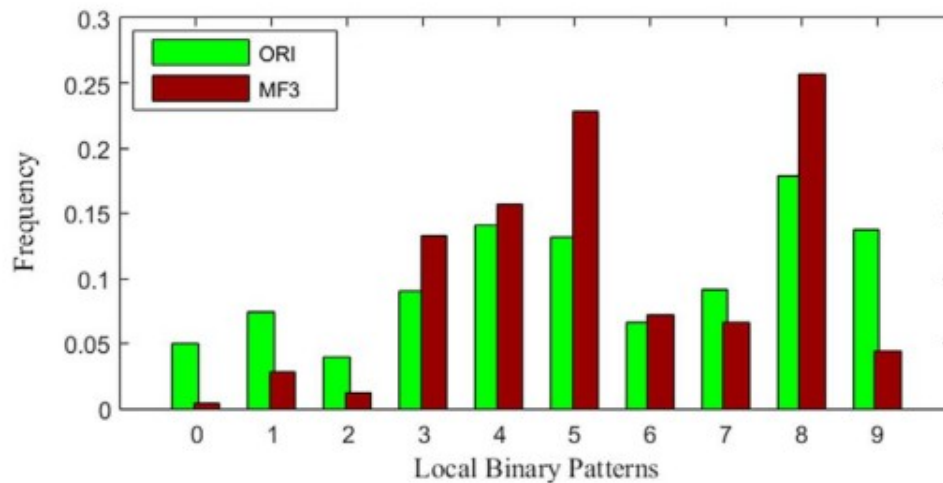
((1338, 28), (1338, 28), (1338, 28), (1338, 28))

```

So now for every image, we have a feature vector of size 28.

### Recreating Plots....

a.) Fig 3.



**Fig. 3.** The uniform rotationally invariant LBP histogram from 10000  $3 \times 3$  median-filtered images (MF3) and 10000 original images (ORI). Bins 0–8 are the quantities of the uniform patterns; bin 9 is the quantity of the non-uniform patterns. All these images are from the BOWS2 database.

We have the UCID Dataset rather than BOWS2 Dataset. And we have 1338 images rather than 10,000 images.

Let us plot the same thing and see if the pattern is similar.

```
joint_histograms_ucid.shape
```

```
(1338, 10, 10)
```

```
joint_histograms_median_filtered_3.shape
```

```
(1338, 10, 10)
```

```
# Creating 1-D histograms for CLBPS and UCID Images
```

```
ucid_images_corr_coefficients_1d = []
```

```
median_filtered_images_3_corr_coefficients_1d = []
```

```
# Loop over all bins
```

```
for i in range(10):
```

```
    bin_value_ori = 0
```

```
    bin_value_med_3 = 0
```

```
# Loop over all images
```

```
    for j in range(1338):
```

```
        # Summation of the elements of ith row of jth histogram
```

```

        bin_value_ori += np.sum(joint_histograms_ucid[j][i])
        bin_value_med_3 +=
np.sum(joint_histograms_median_filtered_3[j][i])

    # Append to list
    ucid_images_corr_coefficients_1d.append(bin_value_ori)

median_filtered_images_3_corr_coefficients_1d.append(bin_value_med_3)

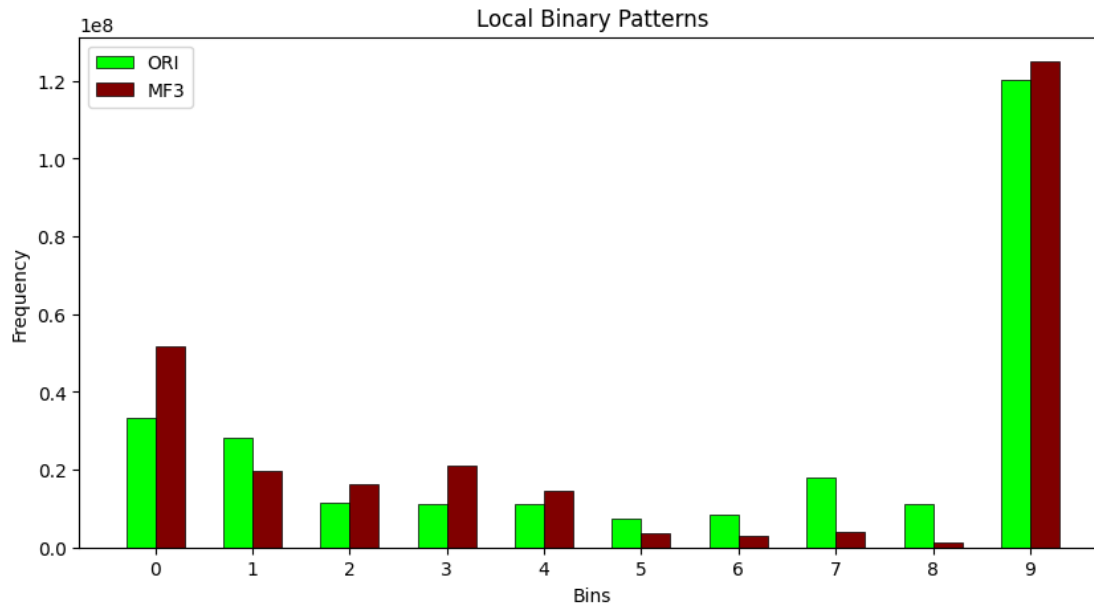
# Converting to Numpy Arrays
ucid_images_corr_coefficients_1d =
np.array(ucid_images_corr_coefficients_1d)
median_filtered_images_3_corr_coefficients_1d =
np.array(median_filtered_images_3_corr_coefficients_1d)

ucid_images_corr_coefficients_1d.shape,
median_filtered_images_3_corr_coefficients_1d.shape

((10,), (10,))

# Plotting 1-D Histograms
plt.figure(figsize=(10, 5))
plt.bar(np.arange(10), ucid_images_corr_coefficients_1d, width=0.3,
label="ORI", color="lime", edgecolor="black", linewidth=0.5)
plt.bar(np.arange(10)+0.3,
median_filtered_images_3_corr_coefficients_1d, width=0.3, label="MF3",
color="maroon", edgecolor="black", linewidth=0.5)
plt.xticks(np.arange(10)+0.15, np.arange(10))
plt.xlabel("Bins")
plt.ylabel("Frequency")
plt.title("Local Binary Patterns")
plt.legend()
plt.show()

```



Above we can see that we see the same pattern in the following bins:

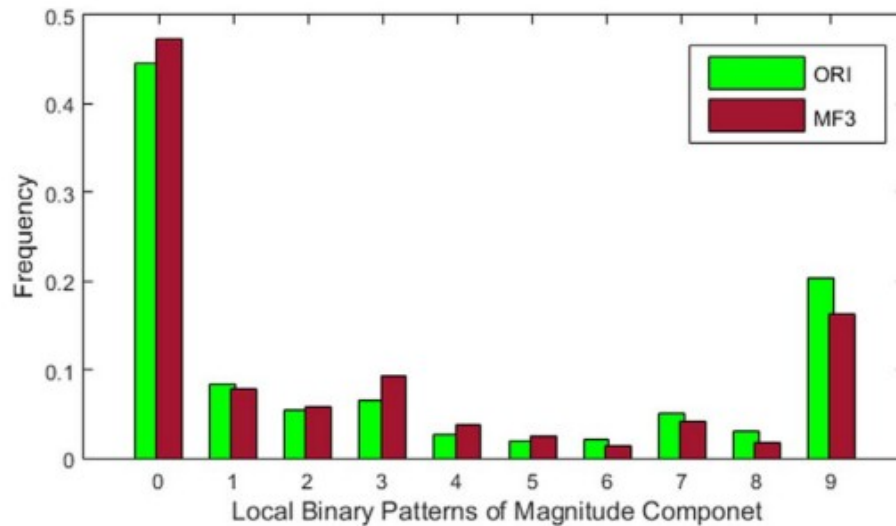
-- Bin #1

-- Bin #3

-- Bin #4

And different pattern in rest of the bins.

b.) Fig 4.



**Fig. 4.** The uniform rotationally invariant LBP histogram of magnitude component from 10000  $3 \times 3$  median-filtered images (MF3) and 10000 original images (ORI). Bins 0–8 are the quantities of the uniform patterns; bin 9 is the quantity of the non-uniform patterns. All these images are from the BOWS2 database.

*# Creating 1-D histograms for CLBPM and UCID Images*

```
ucid_images_corr_coefficients_1d = []
```

```
median_filtered_images_3_corr_coefficients_1d = []
```

*# Loop over all bins*

```
for i in range(10):
```

```
    bin_value_ori = 0
```

```
    bin_value_med_3 = 0
```

*# Loop over all images*

```
    for j in range(1338):
```

```
        # Summation of the elements of ith column of jth histogram
```

```
        bin_value_ori += np.sum(joint_histograms_ucid[j][:, i])
```

```
        bin_value_med_3 +=
```

```
        np.sum(joint_histograms_median_filtered_3[j][:, i])
```

*# Append to list*

```
        ucid_images_corr_coefficients_1d.append(bin_value_ori)
```

```
median_filtered_images_3_corr_coefficients_1d.append(bin_value_med_3)
```

*# Converting to Numpy Arrays*

```
ucid_images_corr_coefficients_1d =
```

```
np.array(ucid_images_corr_coefficients_1d)
```

```

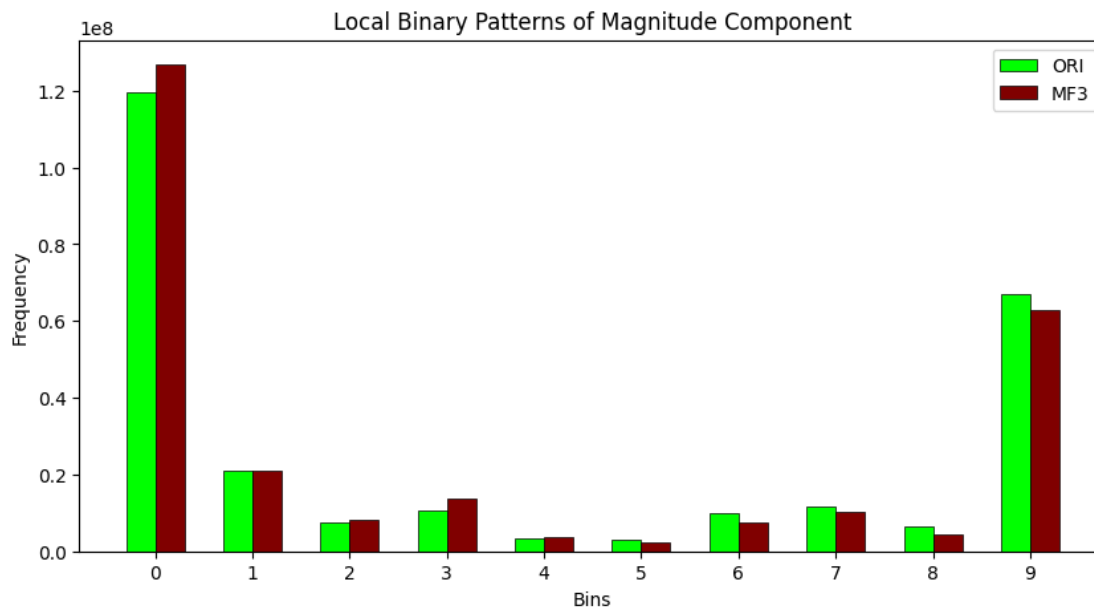
median_filtered_images_3_corr_coefficients_1d =
np.array(median_filtered_images_3_corr_coefficients_1d)

ucid_images_corr_coefficients_1d.shape,
median_filtered_images_3_corr_coefficients_1d.shape

((10,), (10,))

# Plotting 1-D Histograms
plt.figure(figsize=(10, 5))
plt.bar(np.arange(10), ucid_images_corr_coefficients_1d, width=0.3,
label="ORI", color="lime", edgecolor="black", linewidth=0.5)
plt.bar(np.arange(10)+0.3,
median_filtered_images_3_corr_coefficients_1d, width=0.3, label="MF3",
color="maroon", edgecolor="black", linewidth=0.5)
plt.xticks(np.arange(10)+0.15, np.arange(10))
plt.xlabel("Bins")
plt.ylabel("Frequency")
plt.title("Local Binary Patterns of Magnitude Component")
plt.legend()
plt.show()

```



Above we see the same pattern in all the bins.

### 3. Creating the LDD Feature Set

# Creating the LDD Features

```

joint_histograms_ucid.shape, joint_histograms_median_filtered_3.shape,
joint_histograms_median_filtered_5.shape,
joint_histograms_average_filtered_3.shape

```

```

((1338, 10, 10), (1338, 10, 10), (1338, 10, 10), (1338, 10, 10))

```



```

ucid_images_corr_coefficients.shape,
median_filtered_images_3_corr_coefficients.shape,
median_filtered_images_5_corr_coefficients.shape,
average_filtered_images_3_corr_coefficients.shape

((1338, 28), (1338, 28), (1338, 28), (1338, 28))

# Flattening the 2-D histogram features
histogram_features_ucid = []
histogram_features_median_filtered_3 = []
histogram_features_median_filtered_5 = []
histogram_features_average_filtered_3 = []

for i in range(1338):
    histogram_features_ucid.append(joint_histograms_ucid[i].flatten())

histogram_features_median_filtered_3.append(joint_histograms_median_filtered_3[i].flatten())

histogram_features_median_filtered_5.append(joint_histograms_median_filtered_5[i].flatten())

histogram_features_average_filtered_3.append(joint_histograms_average_filtered_3[i].flatten())

# Converting to Numpy Arrays
histogram_features_ucid = np.array(histogram_features_ucid)
histogram_features_median_filtered_3 =
np.array(histogram_features_median_filtered_3)
histogram_features_median_filtered_5 =
np.array(histogram_features_median_filtered_5)
histogram_features_average_filtered_3 =
np.array(histogram_features_average_filtered_3)

histogram_features_ucid.shape,
histogram_features_median_filtered_3.shape,
histogram_features_median_filtered_5.shape,
histogram_features_average_filtered_3.shape

((1338, 100), (1338, 100), (1338, 100), (1338, 100))

# # Save the Histogram Features
# np.save("histogram_features_ucid.npy", histogram_features_ucid)
# np.save("histogram_features_median_filtered_3.npy",
histogram_features_median_filtered_3)
# np.save("histogram_features_median_filtered_5.npy",
histogram_features_median_filtered_5)
# np.save("histogram_features_average_filtered_3.npy",
histogram_features_average_filtered_3)

```

```

# # Load the Histogram Features
# histogram_features_ucid = np.load("histogram_features_ucid.npy")
# histogram_features_median_filtered_3 =
np.load("histogram_features_median_filtered_3.npy")
# histogram_features_median_filtered_5 =
np.load("histogram_features_median_filtered_5.npy")
# histogram_features_average_filtered_3 =
np.load("histogram_features_average_filtered_3.npy")

```

```

histogram_features_ucid.shape,
histogram_features_median_filtered_3.shape,
histogram_features_median_filtered_5.shape,
histogram_features_average_filtered_3.shape

((1338, 100), (1338, 100), (1338, 100), (1338, 100))

```

Merging both sets of features into a single feature set.

```

# Creating the LDD Features

```

```

ldd_features_ucid = []
ldd_features_median_filtered_3 = []
ldd_features_median_filtered_5 = []
ldd_features_average_filtered_3 = []

```

```

for i in range(1338):

```

```

    ldd_features_ucid.append(np.concatenate((histogram_features_ucid[i],
    ucid_images_corr_coefficients[i]), axis=0))

```

```

    ldd_features_median_filtered_3.append(np.concatenate((histogram_features_median_filtered_3[i],
    median_filtered_images_3_corr_coefficients[i]), axis=0))

```

```

    ldd_features_median_filtered_5.append(np.concatenate((histogram_features_median_filtered_5[i],
    median_filtered_images_5_corr_coefficients[i]), axis=0))

```

```

    ldd_features_average_filtered_3.append(np.concatenate((histogram_features_average_filtered_3[i],
    average_filtered_images_3_corr_coefficients[i]), axis=0))

```

```

# Converting to Numpy Arrays

```

```

ldd_features_ucid = np.array(ldd_features_ucid)
ldd_features_median_filtered_3 =
np.array(ldd_features_median_filtered_3)
ldd_features_median_filtered_5 =
np.array(ldd_features_median_filtered_5)
ldd_features_average_filtered_3 =
np.array(ldd_features_average_filtered_3)

```

```

ldd_features_ucid.shape, ldd_features_median_filtered_3.shape,
ldd_features_median_filtered_5.shape,
ldd_features_average_filtered_3.shape

((1338, 128), (1338, 128), (1338, 128), (1338, 128))

# Saving the LDD Features
# np.save("ucid_ldd_features.npy", ldd_features_ucid)
# np.save("median_filtered_3_ldd_features.npy",
ldd_features_median_filtered_3)
# np.save("median_filtered_5_ldd_features.npy",
ldd_features_median_filtered_5)
# np.save("average_filtered_3_ldd_features.npy",
ldd_features_average_filtered_3)

# Loading the LDD Features
# ldd_features_ucid = np.load("ucid_ldd_features.npy")
# ldd_features_median_filtered_3 =
np.load("median_filtered_3_ldd_features.npy")
# ldd_features_median_filtered_5 =
np.load("median_filtered_5_ldd_features.npy")
# ldd_features_average_filtered_3 =
np.load("average_filtered_3_ldd_features.npy")

ldd_features_ucid.shape, ldd_features_median_filtered_3.shape,
ldd_features_median_filtered_5.shape,
ldd_features_average_filtered_3.shape

((1338, 128), (1338, 128), (1338, 128), (1338, 128))

```

### Recreating Plots....

c.) Fig 7.

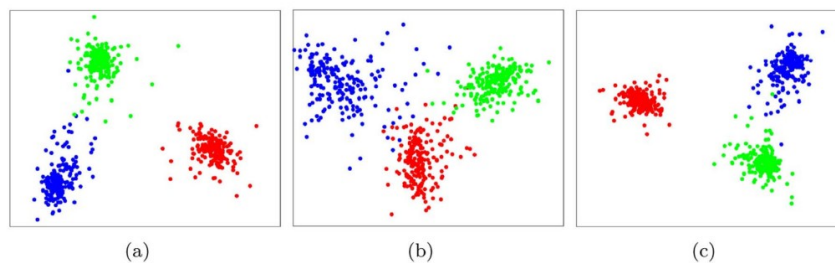


Fig. 7. Distribution of 2D projection results from the proposed features by using linear discriminant analysis (LDA). (a), (b), and (c) correspond to the JHLBP, CPDM and LDD features. Markers with the blue, red, green color denote the unaltered, median filtered and average filtered images source, respectively. Discrimination capability is shown via the clustering effects.

### # JHLBP Features Dataset

```

X_JHLBP = []
y_JHLBP = []

```

### # CPDM Features Dataset

```

X_CPDM = []

```

```

y_CPDM = []

# LDD Features Dataset
X_LDD = []
y_LDD = []

histogram_features_ucid.shape,
histogram_features_median_filtered_3.shape,
histogram_features_median_filtered_5.shape,
histogram_features_average_filtered_3.shape

((1338, 100), (1338, 100), (1338, 100), (1338, 100))

ucid_images_corr_coefficients.shape,
median_filtered_images_3_corr_coefficients.shape,
median_filtered_images_5_corr_coefficients.shape,
average_filtered_images_3_corr_coefficients.shape

((1338, 28), (1338, 28), (1338, 28), (1338, 28))

ldd_features_ucid.shape, ldd_features_median_filtered_3.shape,
ldd_features_median_filtered_5.shape,
ldd_features_average_filtered_3.shape

((1338, 128), (1338, 128), (1338, 128), (1338, 128))

# Creating Datasets
for i in range(1338):

    # JHLBP Features Dataset
    X_JHLBP.append(histogram_features_ucid[i])
    X_JHLBP.append(histogram_features_median_filtered_3[i])
    X_JHLBP.append(histogram_features_median_filtered_5[i])
    X_JHLBP.append(histogram_features_average_filtered_3[i])
    y_JHLBP.append("blue")
    y_JHLBP.append("red")
    y_JHLBP.append("yellow")
    y_JHLBP.append("green")

    # CPDM Features Dataset
    X_CPDM.append(ucid_images_corr_coefficients[i])
    X_CPDM.append(median_filtered_images_3_corr_coefficients[i])
    X_CPDM.append(median_filtered_images_5_corr_coefficients[i])
    X_CPDM.append(average_filtered_images_3_corr_coefficients[i])
    y_CPDM.append("blue")
    y_CPDM.append("red")
    y_CPDM.append("yellow")
    y_CPDM.append("green")

    # LDD Features Dataset
    X_LDD.append(ldd_features_ucid[i])
    X_LDD.append(ldd_features_median_filtered_3[i])

```

```

X_LDD.append(ldd_features_median_filtered_5[i])
X_LDD.append(ldd_features_average_filtered_3[i])
y_LDD.append("blue")
y_LDD.append("red")
y_LDD.append("yellow")
y_LDD.append("green")

```

*# Converting to Numpy Arrays*

```

X_JHLBP = np.array(X_JHLBP)
X_CPDM = np.array(X_CPDM)
X_LDD = np.array(X_LDD)

```

```

y_JHLBP = np.array(y_JHLBP)
y_CPDM = np.array(y_CPDM)
y_LDD = np.array(y_LDD)

```

```

X_JHLBP.shape, y_JHLBP.shape, X_CPDM.shape, y_CPDM.shape, X_LDD.shape,
y_LDD.shape

```

```

((5352, 100), (5352,), (5352, 28), (5352,), (5352, 128), (5352,))

```

*# Save the Datasets*

```

# np.save("X_JHLBP.npy", X_JHLBP)
# np.save("y_JHLBP.npy", y_JHLBP)
# np.save("X_CPDM.npy", X_CPDM)
# np.save("y_CPDM.npy", y_CPDM)
# np.save("X_LDD.npy", X_LDD)
# np.save("y_LDD.npy", y_LDD)

```

*# # Load the Datasets*

```

X_JHLBP = np.load("X_JHLBP.npy")
y_JHLBP = np.load("y_JHLBP.npy")
X_CPDM = np.load("X_CPDM.npy")
y_CPDM = np.load("y_CPDM.npy")
X_LDD = np.load("X_LDD.npy")
y_LDD = np.load("y_LDD.npy")

```

```

X_JHLBP.shape, y_JHLBP.shape, X_CPDM.shape, y_CPDM.shape, X_LDD.shape,
y_LDD.shape

```

```

((5352, 100), (5352,), (5352, 28), (5352,), (5352, 128), (5352,))

```

**from** sklearn.preprocessing **import** MinMaxScaler

*# Normalizing the Datasets*

```

scaler = MinMaxScaler()

```

```

X_JHLBP_Scaled = scaler.fit_transform(X_JHLBP)
X_CPDM_Scaled = scaler.fit_transform(X_CPDM)
X_LDD_Scaled = scaler.fit_transform(X_LDD)

```

```

X_JHLBP_Scaled.shape, X_CPDM_Scaled.shape, X_LDD_Scaled.shape
((5352, 100), (5352, 28), (5352, 128))

# LDA using JHLBP Features
lda = LinearDiscriminantAnalysis(n_components=2)

# Fitting the LDA
new_X_JHLBP = lda.fit_transform(X_JHLBP_Scaled, y_JHLBP)
new_X_CPDM = lda.fit_transform(X_CPDM_Scaled, y_CPDM)
new_X_LDD = lda.fit_transform(X_LDD_Scaled, y_LDD)

new_X_JHLBP.shape, new_X_CPDM.shape, new_X_LDD.shape
((5352, 2), (5352, 2), (5352, 2))

# Plotting the LDAs in 3 Subplots
fig, (ax1, ax2, ax3) = plt.subplots(1, 3, figsize=(20, 5))

# Plotting the JHLBP LDA
ax1.scatter(new_X_JHLBP[:, 0], new_X_JHLBP[:, 1], c=y_JHLBP)
ax1.set_title("JHLBP Features")
# axis off
ax1.axis('off')
# Border around the plot
ax1.set_frame_on(True)

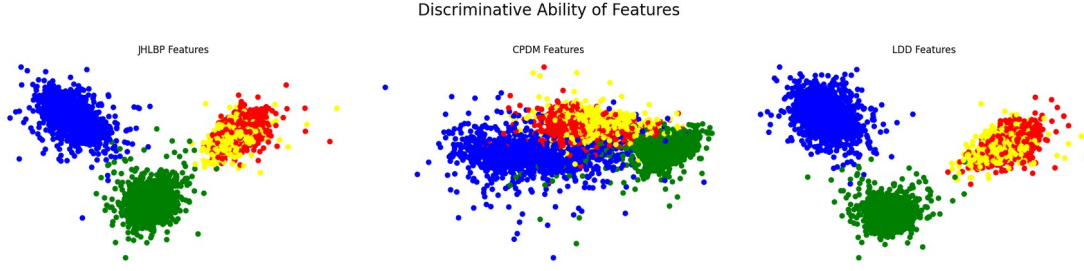
# Plotting the CPDM LDA
ax2.scatter(new_X_CPDM[:, 0], new_X_CPDM[:, 1], c=y_CPDM)
ax2.set_title("CPDM Features")
# axis off
ax2.axis('off')
# Box around the plot
ax2.set_frame_on(True)

# Plotting the LDD LDA
ax3.scatter(new_X_LDD[:, 0], new_X_LDD[:, 1], c=y_LDD)
ax3.set_title("LDD Features")
# axis off
ax3.axis('off')
# Box around the plot
ax3.set_frame_on(True)

# Title of Figure
fig.suptitle("Discriminative Ability of Features", fontsize=20)
# MArgin below the title
fig.tight_layout(pad=2.0)

plt.show()

```



*Blue* refers to Unaltered Images

*Red* refers to Images with Median filtering applied with a kernel size 3x3

*Yellow* refers to Images with Median filtering applied with a kernel size 5x5

*Green* refers to Images with Average filtering applied with a kernel size 3x3

The Discriminative ability is shown via Clustering Effect.

We see the same pattern as shown in the research paper i.e. LDD provides better discriminative ability than the other two methods.

JHLBP provides better discriminative ability than CPDM which has very low discriminative ability.

**NOTE:** We see MF3 and MF5 are a bit merged, but the median filtering as a whole is separated than the original images and the average filtered images.

## Experiments and Results

The  $C-SVM$  with *Gaussian Kernel* is used as the classifier in our experiments. Using the **five-fold cross-validation** in conjunction with a grid search, we obtain the best kernel parameters for the  $SVM$ . The grid search for the optimal parameters are performed on the multiplicative grid  $(C, \gamma) \in \{2^i, 2^j\} \forall i, j \in Z$ .

We use those optimal parameters to get the classifier model on the entire training set, and the trained classifier model is used to perform a classification on the testing set.

Specifically, the images in the  $UCID$  database are randomly divided into four folds of nearly equal size. The training set is composed of three folds, while the remaining fold is used for evaluation.

The performance of each detection method is summarized by the minimal average decision error of each technique under the assumption of equal priors and equal costs,

$$P_e = \min \left( \frac{P_{fp} + 1 - P_{tp}}{2} \right)$$

where  $P_{fp}$  and  $P_{tp}$  denote the false positive and true positive rates, respectively.



```
X_JHLBP.shape, y_JHLBP.shape, X_CPDM.shape, y_CPDM.shape, X_LDD.shape,
y_LDD.shape
```

```
((5352, 100), (5352,)), (5352, 28), (5352,)), (5352, 128), (5352,))
```

### **C-SVM with Gaussian Kernel**

```
# Importing the Libraries
```

```
from sklearn.model_selection import GridSearchCV
```

```
from sklearn.svm import SVC
```

```
# Creating the C-SVM Classifier
```

```
svc_jhlbp = SVC(kernel='rbf')
```

```
svc_cpdm = SVC(kernel='rbf')
```

```
svc_ldd = SVC(kernel='rbf')
```

```
# Creating the Grid Search Parameters
```

```
parameters = {'C': [0.1, 10, 100, 1000], 'gamma': [1, 0.1, 0.01,
0.001, 0.0001]}
```

```
# Used in Paper
```

```
parameters_2 = {'C': [2**i for i in range(-5, 16)], 'gamma': [2**i for
i in range(-15, 4)]}
```

```
# Creating the Grid Search
```

```
clf_jhlbp = GridSearchCV(svc_jhlbp, parameters, cv=5, n_jobs=-1,
refit=True)
```

```
clf_cpdm = GridSearchCV(svc_cpdm, parameters, cv=5, n_jobs=-1,
refit=True)
```

```
clf_ldd = GridSearchCV(svc_ldd, parameters, cv=5, n_jobs=-1,
refit=True)
```

```
# Using original parameters
```

```
clf_jhlbp_2 = GridSearchCV(svc_jhlbp, parameters_2, cv=5, n_jobs=-1,
refit=True)
```

```
clf_cpdm_2 = GridSearchCV(svc_cpdm, parameters_2, cv=5, n_jobs=-1,
refit=True)
```

```
clf_ldd_2 = GridSearchCV(svc_ldd, parameters_2, cv=5, n_jobs=-1,
refit=True)
```

### **JHLBP Features**

```
# Fitting the Grid Search on the JHLBP Features and print the best
parameters
```

```
clf_jhlbp.fit(X_JHLBP_Scaled, y_JHLBP)
```

```
GridSearchCV(cv=5, estimator=SVC(), n_jobs=-1,
              param_grid={'C': [0.1, 10, 100, 1000],
                           'gamma': [1, 0.1, 0.01, 0.001, 0.0001]})
```

```
# Fitting the Grid Search using Original Parameters in Paper
```

```
clf_jhlbp_2.fit(X_JHLBP_Scaled, y_JHLBP)
```

```

GridSearchCV(cv=5, estimator=SVC(), n_jobs=-1,
             param_grid={'C': [0.03125, 0.0625, 0.125, 0.25, 0.5, 1,
                                2, 4, 8,
                                16, 32, 64, 128, 256, 512, 1024, 2048,
                                4096,
                                8192, 16384, 32768],
                         'gamma': [3.0517578125e-05, 6.103515625e-05,
                                    0.0001220703125, 0.000244140625,
                                    0.00048828125, 0.0009765625,
                                    0.001953125,
                                    0.00390625, 0.0078125, 0.015625,
                                    0.03125,
                                    0.0625, 0.125, 0.25, 0.5, 1, 2, 4,
                                    8]})

```

```

# Printing the Best Parameters

```

```

print("Best Parameters for JHLBP Features using our own Params: ",
      clf_jhlbp.best_params_)
print("Best Parameters for JHLBP Features using Paper Params: ",
      clf_jhlbp_2.best_params_)

```

```

Best Parameters for JHLBP Features using our own Params: {'C': 1000,
'gamma': 0.1}

```

```

Best Parameters for JHLBP Features using Paper Params: {'C': 512,
'gamma': 0.125}

```

```

jhlbp_best_params = clf_jhlbp.best_params_
jhlbp_best_params_2 = clf_jhlbp_2.best_params_

```

```

# import pickle

```

```

# # Saving Best Parameters with Pickle

```

```

# with open("jhlbp_best_params.pickle", "wb") as f:
#     pickle.dump(jhlbp_best_params, f)

```

```

# with open("jhlbp_best_params_2.pickle", "wb") as f:
#     pickle.dump(jhlbp_best_params_2, f)

```

```

import pickle

```

```

# Loading the Best Parameters

```

```

with open("jhlbp_best_params.pickle", "rb") as f:
    jhlbp_best_params = pickle.load(f)

```

```

with open("jhlbp_best_params_2.pickle", "rb") as f:
    jhlbp_best_params_2 = pickle.load(f)

```

```

jhlbp_best_params, jhlbp_best_params_2

```

```

({'C': 1000, 'gamma': 0.1}, {'C': 512, 'gamma': 0.125})

```

## CPDM Features

```
# Fitting the Grid Search on the CPDM Features and print the best
parameters
clf_cpdm.fit(X_CPDM_Scaled, y_CPDM)

GridSearchCV(cv=5, estimator=SVC(), n_jobs=-1,
             param_grid={'C': [0.1, 10, 100, 1000],
                         'gamma': [1, 0.1, 0.01, 0.001, 0.0001]})

# Fitting the Grid Search using Original Parameters in Paper
clf_cpdm_2.fit(X_CPDM_Scaled, y_CPDM)

GridSearchCV(cv=5, estimator=SVC(), n_jobs=-1,
             param_grid={'C': [0.03125, 0.0625, 0.125, 0.25, 0.5, 1,
                               2, 4, 8,
                               16, 32, 64, 128, 256, 512, 1024, 2048,
                               4096,
                               8192, 16384, 32768],
                         'gamma': [3.0517578125e-05, 6.103515625e-05,
                                   0.0001220703125, 0.000244140625,
                                   0.00048828125, 0.0009765625,
                                   0.001953125,
                                   0.00390625, 0.0078125, 0.015625,
                                   0.03125,
                                   0.0625, 0.125, 0.25, 0.5, 1, 2, 4,
                                   8]})

# Printing the Best Parameters
print("Best Parameters for CPDM Features: ", clf_cpdm.best_params_)
print("Best Parameters for CPDM Features using Paper Params: ",
      clf_cpdm_2.best_params_)

Best Parameters for CPDM Features:  {'C': 1000, 'gamma': 1}
Best Parameters for CPDM Features using Paper Params:  {'C': 16384,
'gamma': 0.125}

cpdm_best_params = clf_cpdm.best_params_
cpdm_best_params_2 = clf_cpdm_2.best_params_

# import pickle

# # Saving Best Parameters with Pickle
# with open("cpdm_best_params.pickle", "wb") as f:
#     pickle.dump(cpdm_best_params, f)

# with open("cpdm_best_params_2.pickle", "wb") as f:
#     pickle.dump(cpdm_best_params_2, f)

import pickle

# Loading the Best Parameters
```

```

with open("cpdm_best_params.pickle", "rb") as f:
    cpdm_best_params = pickle.load(f)

with open("cpdm_best_params_2.pickle", "rb") as f:
    cpdm_best_params_2 = pickle.load(f)

cpdm_best_params, cpdm_best_params_2
({'C': 1000, 'gamma': 1}, {'C': 16384, 'gamma': 0.125})

```

## LDD Features

```

# Fitting the Grid Search on the LDD Features and print the best parameters
clf_ldd.fit(X_LDD_Scaled, y_LDD)

GridSearchCV(cv=5, estimator=SVC(), n_jobs=-1,
             param_grid={'C': [0.1, 10, 100, 1000],
                         'gamma': [1, 0.1, 0.01, 0.001, 0.0001]})

# Fitting the Grid Search using Original Parameters in Paper
clf_ldd_2.fit(X_LDD_Scaled, y_LDD)

GridSearchCV(cv=5, estimator=SVC(), n_jobs=-1,
             param_grid={'C': [0.03125, 0.0625, 0.125, 0.25, 0.5, 1,
                                2, 4, 8,
                                16, 32, 64, 128, 256, 512, 1024, 2048,
                                4096,
                                8192, 16384, 32768],
                         'gamma': [3.0517578125e-05, 6.103515625e-05,
                                   0.0001220703125, 0.000244140625,
                                   0.00048828125, 0.0009765625,
                                   0.001953125,
                                   0.00390625, 0.0078125, 0.015625,
                                   0.03125,
                                   0.0625, 0.125, 0.25, 0.5, 1, 2, 4,
                                   8]})

# Printing the Best Parameters
print("Best Parameters for LDD Features: ", clf_ldd.best_params_)
print("Best Parameters for LDD Features using Paper Params: ",
      clf_ldd_2.best_params_)

Best Parameters for LDD Features: {'C': 1000, 'gamma': 0.1}
Best Parameters for LDD Features using Paper Params: {'C': 1024,
                                                       'gamma': 0.125}

ldd_best_params = clf_ldd.best_params_
ldd_best_params_2 = clf_ldd_2.best_params_

# import pickle

```

```

# # Saving Best Parameters with Pickle
# with open("ldd_best_params.pickle", "wb") as f:
#     pickle.dump(ldd_best_params, f)

# with open("ldd_best_params_2.pickle", "wb") as f:
#     pickle.dump(ldd_best_params_2, f)

import pickle

# Loading the Best Parameters
with open("ldd_best_params.pickle", "rb") as f:
    ldd_best_params = pickle.load(f)

with open("ldd_best_params_2.pickle", "rb") as f:
    ldd_best_params_2 = pickle.load(f)

ldd_best_params, ldd_best_params_2

({'C': 1000, 'gamma': 0.1}, {'C': 1024, 'gamma': 0.125})

```

## Training and Testing

### Splitting the Data in Training and Testing

```

from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score, confusion_matrix,
classification_report
from sklearn.metrics import roc_curve, auc

# Split into Train and Test
X_train_JHLBP, X_test_JHLBP, y_train_JHLBP, y_test_JHLBP =
train_test_split(X_JHLBP, y_JHLBP, test_size=0.25, random_state=42,
stratify=y_JHLBP)
X_train_CPDM, X_test_CPDM, y_train_CPDM, y_test_CPDM =
train_test_split(X_CPDM, y_CPDM, test_size=0.25, random_state=42,
stratify=y_CPDM)
X_train_LDD, X_test_LDD, y_train_LDD, y_test_LDD =
train_test_split(X_LDD, y_LDD, test_size=0.25, random_state=42,
stratify=y_LDD)

X_train_JHLBP.shape, X_test_JHLBP.shape, y_train_JHLBP.shape,
y_test_JHLBP.shape, X_train_CPDM.shape, X_test_CPDM.shape,
y_train_CPDM.shape, y_test_CPDM.shape, X_train_LDD.shape,
X_test_LDD.shape, y_train_LDD.shape, y_test_LDD.shape

((4014, 100),
 (1338, 100),
 (4014, ),
 (1338, ),
 (4014, 28),
 (1338, 28),
 (4014, ),

```

```
(1338,),  
(4014, 128),  
(1338, 128),  
(4014,),  
(1338,))
```

```
# Scaling the Data using MinMaxScaler
```

```
X_train_JHLBP = scaler.fit_transform(X_train_JHLBP)  
X_test_JHLBP = scaler.transform(X_test_JHLBP)
```

```
X_train_CPDM = scaler.fit_transform(X_train_CPDM)  
X_test_CPDM = scaler.transform(X_test_CPDM)
```

```
X_train_LDD = scaler.fit_transform(X_train_LDD)  
X_test_LDD = scaler.transform(X_test_LDD)
```

### **Train Using Optimal Parameters and plot the ROC Curve**

```
# Train on Features using the best parameters
```

```
svc_jhlbp = SVC(kernel='rbf', C=jhlbp_best_params['C'],  
gamma=jhlbp_best_params['gamma'])  
svc_cpdm = SVC(kernel='rbf', C=cpdm_best_params['C'],  
gamma=cpdm_best_params['gamma'])  
svc_ldd = SVC(kernel='rbf', C=ldd_best_params['C'],  
gamma=ldd_best_params['gamma'])
```

```
svc_jhlbp_2 = SVC(kernel='rbf', C=jhlbp_best_params_2['C'],  
gamma=jhlbp_best_params_2['gamma'])  
svc_cpdm_2 = SVC(kernel='rbf', C=cpdm_best_params_2['C'],  
gamma=cpdm_best_params_2['gamma'])  
svc_ldd_2 = SVC(kernel='rbf', C=ldd_best_params_2['C'],  
gamma=ldd_best_params_2['gamma'])
```

```
# Fitting the JHLBP Features
```

```
svc_jhlbp.fit(X_train_JHLBP, y_train_JHLBP)
```

```
SVC(C=1000, gamma=0.1)
```

```
# Fitting the JHLBP Features using Original Parameters in Paper
```

```
svc_jhlbp_2.fit(X_train_JHLBP, y_train_JHLBP)
```

```
SVC(C=512, gamma=0.125)
```

```
# Fitting the CPDM Features
```

```
svc_cpdm.fit(X_train_CPDM, y_train_CPDM)
```

```
SVC(C=1000, gamma=1)
```

```
# Fitting the CPDM Features using Original Parameters in Paper
```

```
svc_cpdm_2.fit(X_train_CPDM, y_train_CPDM)
```

```
SVC(C=16384, gamma=0.125)
```

```

# Fitting the LDD Features
svc_ldd.fit(X_train_LDD, y_train_LDD)

SVC(C=1000, gamma=0.1)

# Fitting the LDD Features using Original Parameters in Paper
svc_ldd_2.fit(X_train_LDD, y_train_LDD)

SVC(C=1024, gamma=0.125)

# Predicting on the basis of JHLBP Features
y_pred_jhlbp = svc_jhlbp.predict(X_test_JHLBP)

# Predicting on the basis of CPDM Features
y_pred_cpdm = svc_cpdm.predict(X_test_CPDM)

# Predicting on the basis of LDD Features
y_pred_ldd = svc_ldd.predict(X_test_LDD)

# Predicting on the basis of JHLBP Features using Original Parameters
in Paper
y_pred_jhlbp_2 = svc_jhlbp_2.predict(X_test_JHLBP)

# Predicting on the basis of CPDM Features using Original Parameters
in Paper
y_pred_cpdm_2 = svc_cpdm_2.predict(X_test_CPDM)

# Predicting on the basis of LDD Features using Original Parameters in
Paper
y_pred_ldd_2 = svc_ldd_2.predict(X_test_LDD)

# Classification Report for JHLBP Features
print("Classification Report for JHLBP Features: \n",
classification_report(y_test_JHLBP, y_pred_jhlbp))

Classification Report for JHLBP Features:

```

	precision	recall	f1-score	support
blue	1.00	0.99	1.00	335
green	0.99	0.99	0.99	334
red	0.99	0.99	0.99	335
yellow	0.99	0.99	0.99	334
accuracy			0.99	1338
macro avg	0.99	0.99	0.99	1338
weighted avg	0.99	0.99	0.99	1338

```

# Classification Report for JHLBP Features using Original Parameters
in Paper
print("Classification Report for JHLBP Features using Original

```



```
Parameters in Paper: \n", classification_report(y_test_JHLBP,
y_pred_jhlbp_2))
```

Classification Report for JHLBP Features using Original Parameters in Paper:

	precision	recall	f1-score	support
blue	1.00	0.99	1.00	335
green	0.99	0.99	0.99	334
red	0.99	0.99	0.99	335
yellow	0.98	0.99	0.99	334
accuracy			0.99	1338
macro avg	0.99	0.99	0.99	1338
weighted avg	0.99	0.99	0.99	1338

```
# Classification Report for CPDM Features
```

```
print("Classification Report for CPDM Features: \n",
classification_report(y_test_CPDM, y_pred_cpd))
```

Classification Report for CPDM Features:

	precision	recall	f1-score	support
blue	0.92	0.94	0.93	335
green	0.99	0.98	0.99	334
red	0.88	0.81	0.84	335
yellow	0.83	0.89	0.86	334
accuracy			0.91	1338
macro avg	0.91	0.91	0.91	1338
weighted avg	0.91	0.91	0.91	1338

```
# Classification Report for CPDM Features using Original Parameters in
Paper
```

```
print("Classification Report for CPDM Features using Original
Parameters in Paper: \n", classification_report(y_test_CPDM,
y_pred_cpd_2))
```

Classification Report for CPDM Features using Original Parameters in Paper:

	precision	recall	f1-score	support
blue	0.96	0.94	0.95	335
green	0.99	1.00	0.99	334
red	0.89	0.82	0.85	335
yellow	0.84	0.90	0.87	334
accuracy			0.92	1338

macro avg	0.92	0.92	0.92	1338
weighted avg	0.92	0.92	0.92	1338

*# Classification Report for LDD Features*

```
print("Classification Report for LDD Features: \n",
classification_report(y_test_LDD, y_pred_ldd))
```

Classification Report for LDD Features:

	precision	recall	f1-score	support
blue	1.00	1.00	1.00	335
green	1.00	0.99	1.00	334
red	0.99	0.98	0.99	335
yellow	0.98	0.99	0.99	334
accuracy			0.99	1338
macro avg	0.99	0.99	0.99	1338
weighted avg	0.99	0.99	0.99	1338

*# Classification Report for LDD Features using Original Parameters in Paper*

```
print("Classification Report for LDD Features using Original
Parameters in Paper: \n", classification_report(y_test_LDD,
y_pred_ldd_2))
```

Classification Report for LDD Features using Original Parameters in Paper:

	precision	recall	f1-score	support
blue	1.00	1.00	1.00	335
green	1.00	0.99	1.00	334
red	0.99	0.99	0.99	335
yellow	0.98	0.99	0.99	334
accuracy			0.99	1338
macro avg	0.99	0.99	0.99	1338
weighted avg	0.99	0.99	0.99	1338

## RESULT:

Voila! From the above Classification Reports we can see that LDD features are the best, followed by JHLBP and then CPDM.

## Recreating Plots....

### d.) Fig 8.

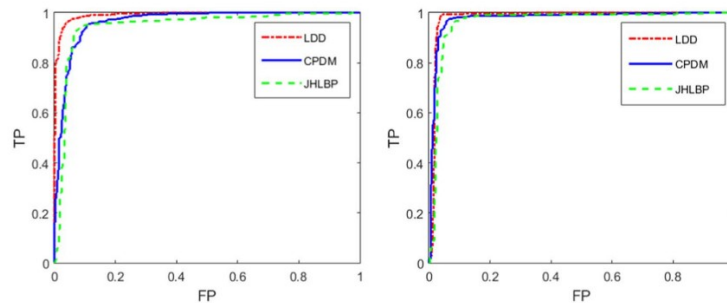


Fig. 8. Roc cures show 3×3 (left) and 5×5 (right) median filtering detection performance on JPEG compressed images. The proposed JHLBP, CPDM and LDD features are tested.

## Plotting the ROC Curves

*# Plot ROC Curve for all classes (following OvR Strategy)*

### *# JHLBP Features*

```
red_y_test_JHLBP = np.where(y_test_JHLBP == 'red', 1, 0)
blue_y_test_JHLBP = np.where(y_test_JHLBP == 'blue', 1, 0)
green_y_test_JHLBP = np.where(y_test_JHLBP == 'green', 1, 0)
yellow_y_test_JHLBP = np.where(y_test_JHLBP == 'yellow', 1, 0)
```

```
red_y_pred_jhlbp = np.where(y_pred_jhlbp == 'red', 1, 0)
blue_y_pred_jhlbp = np.where(y_pred_jhlbp == 'blue', 1, 0)
green_y_pred_jhlbp = np.where(y_pred_jhlbp == 'green', 1, 0)
yellow_y_pred_jhlbp = np.where(y_pred_jhlbp == 'yellow', 1, 0)
```

```
red_y_pred_jhlbp_2 = np.where(y_pred_jhlbp_2 == 'red', 1, 0)
blue_y_pred_jhlbp_2 = np.where(y_pred_jhlbp_2 == 'blue', 1, 0)
green_y_pred_jhlbp_2 = np.where(y_pred_jhlbp_2 == 'green', 1, 0)
yellow_y_pred_jhlbp_2 = np.where(y_pred_jhlbp_2 == 'yellow', 1, 0)
```

### *# CPDM Features*

```
red_y_test_CPDM = np.where(y_test_CPDM == 'red', 1, 0)
blue_y_test_CPDM = np.where(y_test_CPDM == 'blue', 1, 0)
green_y_test_CPDM = np.where(y_test_CPDM == 'green', 1, 0)
yellow_y_test_CPDM = np.where(y_test_CPDM == 'yellow', 1, 0)
```

```
red_y_pred_cpdm = np.where(y_pred_cpdm == 'red', 1, 0)
blue_y_pred_cpdm = np.where(y_pred_cpdm == 'blue', 1, 0)
green_y_pred_cpdm = np.where(y_pred_cpdm == 'green', 1, 0)
yellow_y_pred_cpdm = np.where(y_pred_cpdm == 'yellow', 1, 0)
```

```
red_y_pred_cpdm_2 = np.where(y_pred_cpdm_2 == 'red', 1, 0)
blue_y_pred_cpdm_2 = np.where(y_pred_cpdm_2 == 'blue', 1, 0)
green_y_pred_cpdm_2 = np.where(y_pred_cpdm_2 == 'green', 1, 0)
yellow_y_pred_cpdm_2 = np.where(y_pred_cpdm_2 == 'yellow', 1, 0)
```

#### *# LDD Features*

```
red_y_test_LDD = np.where(y_test_LDD == 'red', 1, 0)
blue_y_test_LDD = np.where(y_test_LDD == 'blue', 1, 0)
green_y_test_LDD = np.where(y_test_LDD == 'green', 1, 0)
yellow_y_test_LDD = np.where(y_test_LDD == 'yellow', 1, 0)
```

```
red_y_pred_ldd = np.where(y_pred_ldd == 'red', 1, 0)
blue_y_pred_ldd = np.where(y_pred_ldd == 'blue', 1, 0)
green_y_pred_ldd = np.where(y_pred_ldd == 'green', 1, 0)
yellow_y_pred_ldd = np.where(y_pred_ldd == 'yellow', 1, 0)
```

```
red_y_pred_ldd_2 = np.where(y_pred_ldd_2 == 'red', 1, 0)
blue_y_pred_ldd_2 = np.where(y_pred_ldd_2 == 'blue', 1, 0)
green_y_pred_ldd_2 = np.where(y_pred_ldd_2 == 'green', 1, 0)
yellow_y_pred_ldd_2 = np.where(y_pred_ldd_2 == 'yellow', 1, 0)
```

#### *# Plotting the ROC Curve for JHLBP Features*

```
fpr_red_jhlbp, tpr_red_jhlbp, thresholds_red_jhlbp =
roc_curve(red_y_test_JHLBP, red_y_pred_jhlbp)
fpr_blue_jhlbp, tpr_blue_jhlbp, thresholds_blue_jhlbp =
roc_curve(blue_y_test_JHLBP, blue_y_pred_jhlbp)
fpr_green_jhlbp, tpr_green_jhlbp, thresholds_green_jhlbp =
roc_curve(green_y_test_JHLBP, green_y_pred_jhlbp)
fpr_yellow_jhlbp, tpr_yellow_jhlbp, thresholds_yellow_jhlbp =
roc_curve(yellow_y_test_JHLBP, yellow_y_pred_jhlbp)
```

```
fpr_red_jhlbp_2, tpr_red_jhlbp_2, thresholdss_red_jhlbp_2 =
roc_curve(red_y_test_JHLBP, red_y_pred_jhlbp_2)
fpr_blue_jhlbp_2, tpr_blue_jhlbp_2, thresholdss_blue_jhlbp_2 =
roc_curve(blue_y_test_JHLBP, blue_y_pred_jhlbp_2)
fpr_green_jhlbp_2, tpr_green_jhlbp_2, thresholdss_green_jhlbp_2 =
roc_curve(green_y_test_JHLBP, green_y_pred_jhlbp_2)
fpr_yellow_jhlbp_2, tpr_yellow_jhlbp_2, thresholdss_yellow_jhlbp_2 =
roc_curve(yellow_y_test_JHLBP, yellow_y_pred_jhlbp_2)
```

#### *# Plotting the ROC Curve for CPDM Features*

```
fpr_red_cpdm, tpr_red_cpdm, thresholds_red_cpdm =
roc_curve(red_y_test_CPDM, red_y_pred_cpdm)
fpr_blue_cpdm, tpr_blue_cpdm, thresholds_blue_cpdm =
roc_curve(blue_y_test_CPDM, blue_y_pred_cpdm)
fpr_green_cpdm, tpr_green_cpdm, thresholds_green_cpdm =
roc_curve(green_y_test_CPDM, green_y_pred_cpdm)
fpr_yellow_cpdm, tpr_yellow_cpdm, thresholds_yellow_cpdm =
roc_curve(yellow_y_test_CPDM, yellow_y_pred_cpdm)
```

```
fpr_red_cpdm_2, tpr_red_cpdm_2, thresholds_red_cpdm_2 =
roc_curve(red_y_test_CPDM, red_y_pred_cpdm_2)
fpr_blue_cpdm_2, tpr_blue_cpdm_2, thresholds_blue_cpdm_2 =
roc_curve(blue_y_test_CPDM, blue_y_pred_cpdm_2)
```

```
fpr_green_cpdm_2, tpr_green_cpdm_2, thresholds_green_cpdm_2 =
roc_curve(green_y_test_CPDM, green_y_pred_cpdm_2)
fpr_yellow_cpdm_2, tpr_yellow_cpdm_2, thresholds_yellow_cpdm_2 =
roc_curve(yellow_y_test_CPDM, yellow_y_pred_cpdm_2)
```

#### *# Plotting the ROC Curve for LDD Features*

```
fpr_red_ldd, tpr_red_ldd, thresholds_red_ldd =
roc_curve(red_y_test_LDD, red_y_pred_ldd)
fpr_blue_ldd, tpr_blue_ldd, thresholds_blue_ldd =
roc_curve(blue_y_test_LDD, blue_y_pred_ldd)
fpr_green_ldd, tpr_green_ldd, thresholds_green_ldd =
roc_curve(green_y_test_LDD, green_y_pred_ldd)
fpr_yellow_ldd, tpr_yellow_ldd, thresholds_yellow_ldd =
roc_curve(yellow_y_test_LDD, yellow_y_pred_ldd)
```

```
fpr_red_ldd_2, tpr_red_ldd_2, thresholds_red_ldd_2 =
roc_curve(red_y_test_LDD, red_y_pred_ldd_2)
fpr_blue_ldd_2, tpr_blue_ldd_2, thresholds_blue_ldd_2 =
roc_curve(blue_y_test_LDD, blue_y_pred_ldd_2)
fpr_green_ldd_2, tpr_green_ldd_2, thresholds_green_ldd_2 =
roc_curve(green_y_test_LDD, green_y_pred_ldd_2)
fpr_yellow_ldd_2, tpr_yellow_ldd_2, thresholds_yellow_ldd_2 =
roc_curve(yellow_y_test_LDD, yellow_y_pred_ldd_2)
```

### **Median Filtered 3x3 Images ROC Curve**

Below Curve is based on Model Trained on Custom Parameters

#### *# Plotting the ROC Curve for JHLBP Features Red Class*

```
plt.plot(fpr_red_jhlbp, tpr_red_jhlbp, label='JHLBP (area = %0.2f)' %
auc(fpr_red_jhlbp, tpr_red_jhlbp), color='lime', linestyle='--')
```

#### *# Plotting the ROC Curve for CPDM Features Red Class*

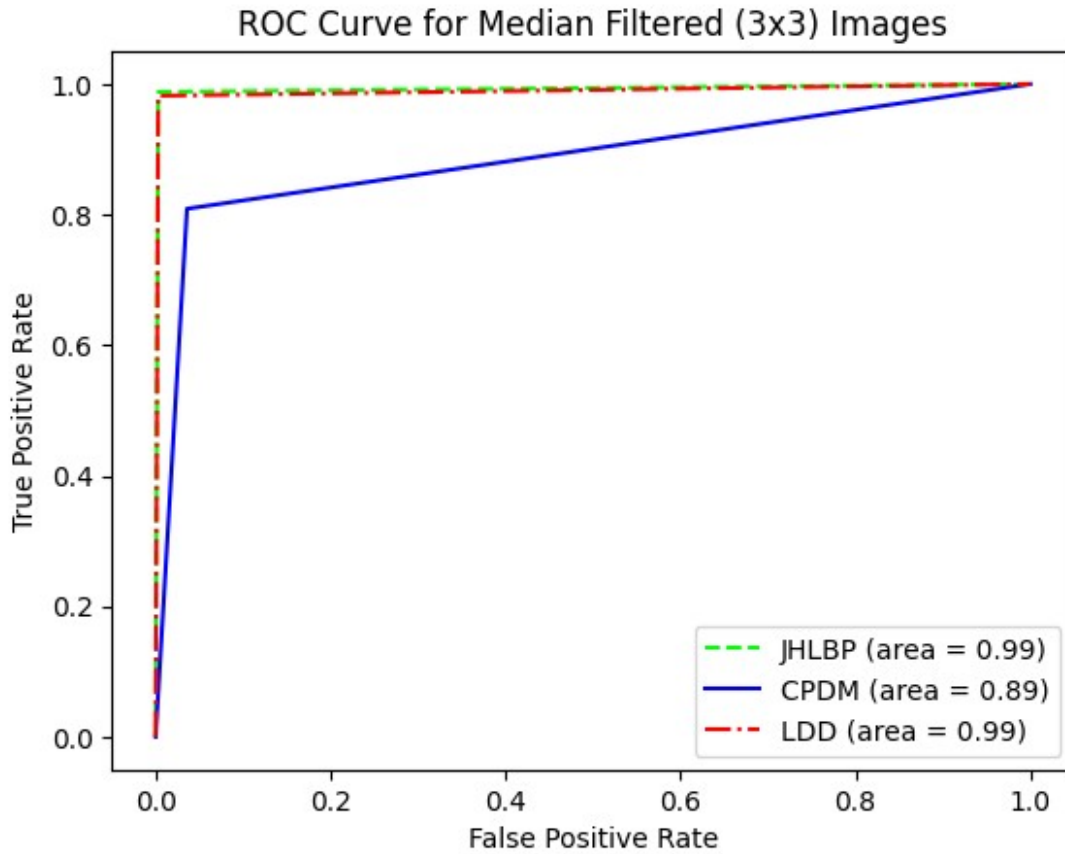
```
plt.plot(fpr_red_cpdm, tpr_red_cpdm, label='CPDM (area = %0.2f)' %
auc(fpr_red_cpdm, tpr_red_cpdm), color='blue')
```

#### *# Plotting the ROC Curve for LDD Features Red Class*

```
plt.plot(fpr_red_ldd, tpr_red_ldd, label='LDD (area = %0.2f)' %
auc(fpr_red_ldd, tpr_red_ldd), color='red', linestyle='-.')
```

#### *# Title and Labels*

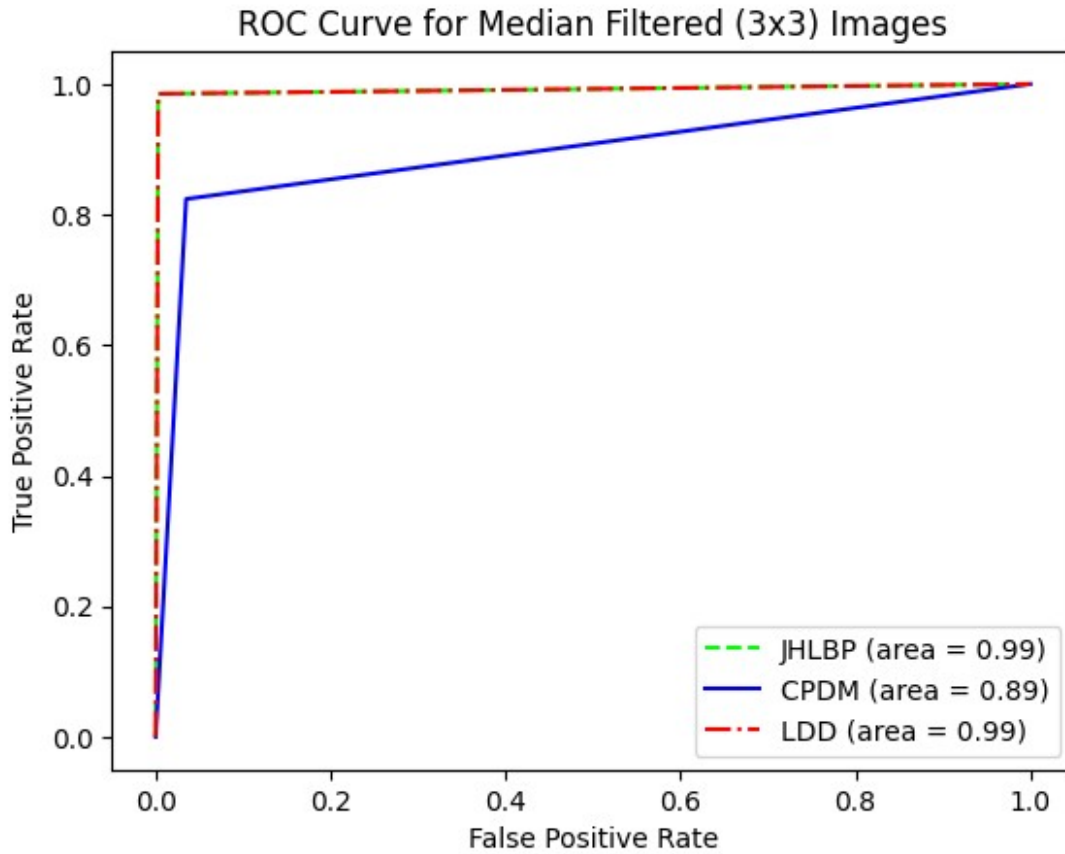
```
plt.title('ROC Curve for Median Filtered (3x3) Images')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.legend(loc="lower right")
plt.show()
```



Below Curve is based on Model Trained on Original Parameters

```
# Plotting the ROC Curve for JHLBP Features Red Class using Original
Parameters
plt.plot(fpr_red_jhlbp_2, tpr_red_jhlbp_2, label='JHLBP (area =
%0.2f)' % auc(fpr_red_jhlbp_2, tpr_red_jhlbp_2), color='lime',
linestyle='--')
# Plotting the ROC Curve for CPDM Features Red Class using Original
Parameters
plt.plot(fpr_red_cpdm_2, tpr_red_cpdm_2, label='CPDM (area = %0.2f)' %
auc(fpr_red_cpdm_2, tpr_red_cpdm_2), color='blue')
# Plotting the ROC Curve for LDD Features Red Class using Original
Parameters
plt.plot(fpr_red_ldd_2, tpr_red_ldd_2, label='LDD (area = %0.2f)' %
auc(fpr_red_ldd_2, tpr_red_ldd_2), color='red', linestyle='-.')

# Title and Labels
plt.title('ROC Curve for Median Filtered (3x3) Images')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.legend(loc="lower right")
plt.show()
```

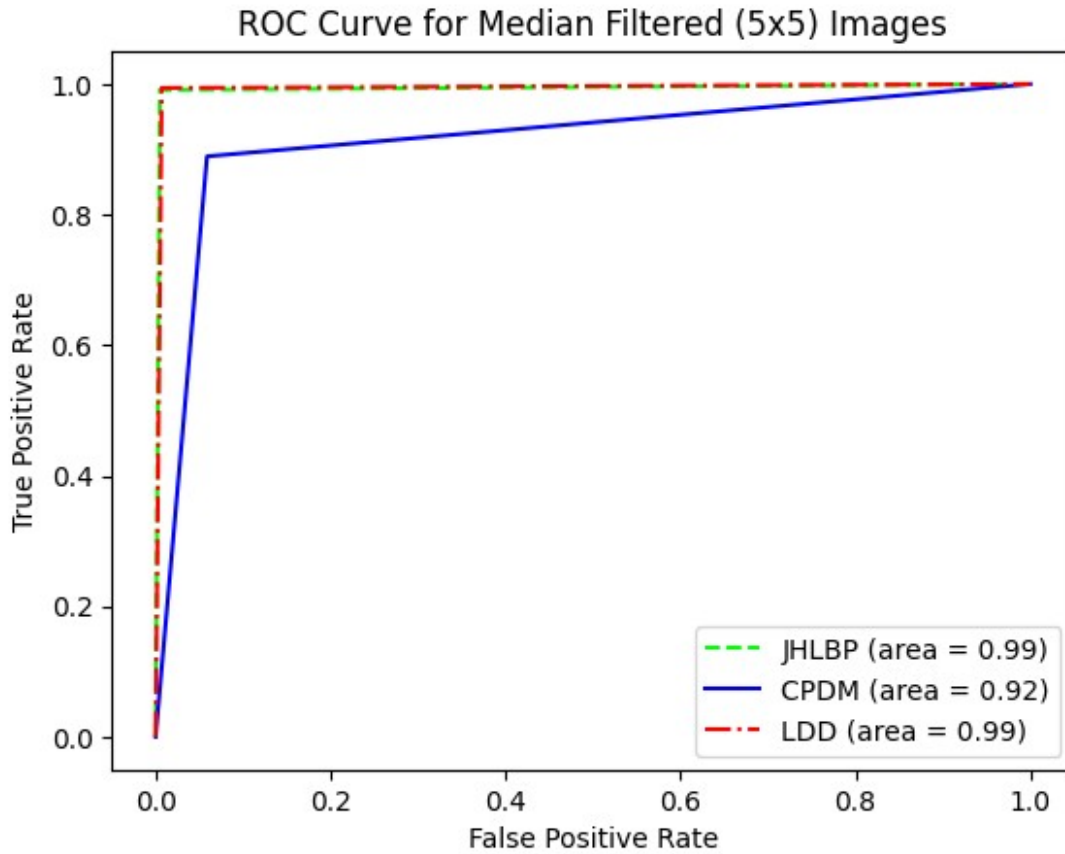


### Median Filtered 5x5 Images ROC Curve

Below Curve is based on Model Trained on Custom Parameters

```
# Plotting the ROC Curve for JHLBP Features Yellow Class
plt.plot(fpr_yellow_jhlbp, tpr_yellow_jhlbp, label='JHLBP (area = %0.2f)' % auc(fpr_yellow_jhlbp, tpr_yellow_jhlbp), color='lime', linestyle='--')
# Plotting the ROC Curve for CPDM Features Yellow Class
plt.plot(fpr_yellow_cpdm, tpr_yellow_cpdm, label='CPDM (area = %0.2f)' % auc(fpr_yellow_cpdm, tpr_yellow_cpdm), color='blue')
# Plotting the ROC Curve for LDD Features Yellow Class
plt.plot(fpr_yellow_ldd, tpr_yellow_ldd, label='LDD (area = %0.2f)' % auc(fpr_yellow_ldd, tpr_yellow_ldd), color='red', linestyle='-.')

# Title and Labels
plt.title('ROC Curve for Median Filtered (5x5) Images')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.legend(loc="lower right")
plt.show()
```

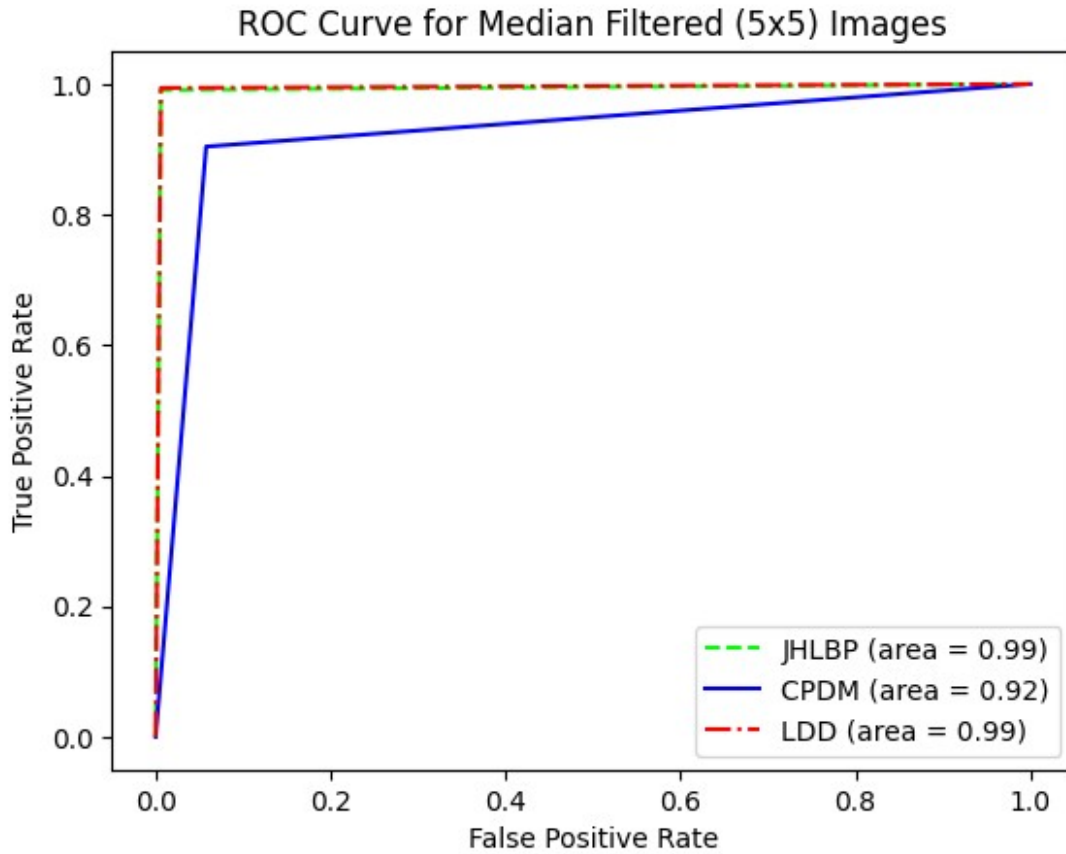


Below Curve is based on Model Trained on Original Parameters

```
# Plotting the ROC Curve for JHLBP Features Yellow Class using
Original Parameters
plt.plot(fpr_yellow_jhlbp_2, tpr_yellow_jhlbp_2, label='JHLBP (area =
%0.2f)' % auc(fpr_yellow_jhlbp_2, tpr_yellow_jhlbp_2), color='lime',
linestyle='--')
# Plotting the ROC Curve for CPDM Features Yellow Class using Original
Parameters
plt.plot(fpr_yellow_cpdm_2, tpr_yellow_cpdm_2, label='CPDM (area =
%0.2f)' % auc(fpr_yellow_cpdm_2, tpr_yellow_cpdm_2), color='blue')
# Plotting the ROC Curve for LDD Features Yellow Class using Original
Parameters
plt.plot(fpr_yellow_ldd_2, tpr_yellow_ldd_2, label='LDD (area =
%0.2f)' % auc(fpr_yellow_ldd_2, tpr_yellow_ldd_2), color='red',
linestyle='-.')

# Title and Labels
plt.title('ROC Curve for Median Filtered (5x5) Images')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.legend(loc="lower right")
plt.show()
```



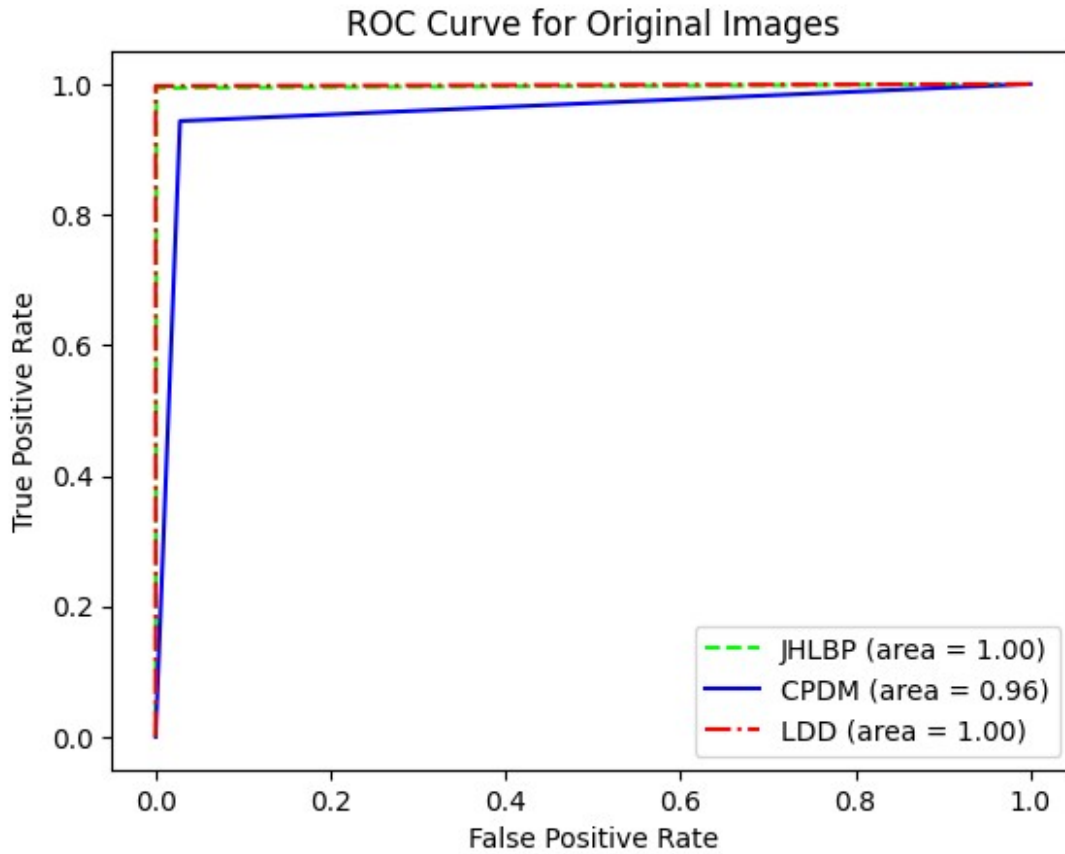


### Original Images ROC Curve

Below Curve is based on Model Trained on Custom Parameters

```
# Plotting the ROC Curve for JHLBP Features blue Class
plt.plot(fpr_blue_jhlbp, tpr_blue_jhlbp, label='JHLBP (area = %0.2f)'
% auc(fpr_blue_jhlbp, tpr_blue_jhlbp), color='lime', linestyle='--')
# Plotting the ROC Curve for CPDM Features blue Class
plt.plot(fpr_blue_cpdm, tpr_blue_cpdm, label='CPDM (area = %0.2f)' %
auc(fpr_blue_cpdm, tpr_blue_cpdm), color='blue')
# Plotting the ROC Curve for LDD Features blue Class
plt.plot(fpr_blue_ldd, tpr_blue_ldd, label='LDD (area = %0.2f)' %
auc(fpr_blue_ldd, tpr_blue_ldd), color='red', linestyle='-.')

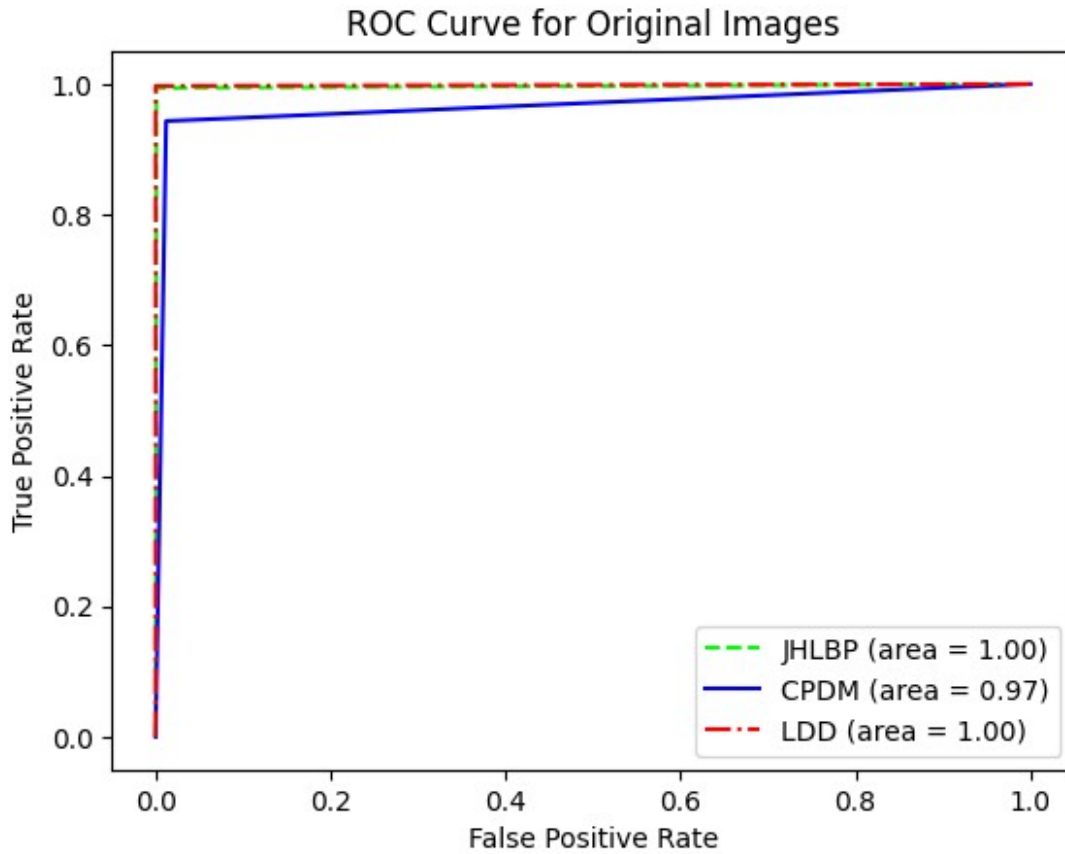
# Title and Labels
plt.title('ROC Curve for Original Images')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.legend(loc="lower right")
plt.show()
```



Below Curve is based on Model Trained on Original Parameters

```
# Plotting the ROC Curve for JHLBP Features blue Class using Original
Parameters
plt.plot(fpr_blue_jhlbp_2, tpr_blue_jhlbp_2, label='JHLBP (area =
%0.2f)' % auc(fpr_blue_jhlbp_2, tpr_blue_jhlbp_2), color='lime',
linestyle='--')\
# Plotting the ROC Curve for CPDM Features blue Class using Original
Parameters
plt.plot(fpr_blue_cpdm_2, tpr_blue_cpdm_2, label='CPDM (area = %0.2f)'
% auc(fpr_blue_cpdm_2, tpr_blue_cpdm_2), color='blue')
# Plotting the ROC Curve for LDD Features blue Class using Original
Parameters
plt.plot(fpr_blue_ldd_2, tpr_blue_ldd_2, label='LDD (area = %0.2f)' %
auc(fpr_blue_ldd_2, tpr_blue_ldd_2), color='red', linestyle='-.')

# Title and Labels
plt.title('ROC Curve for Original Images')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.legend(loc="lower right")
plt.show()
```

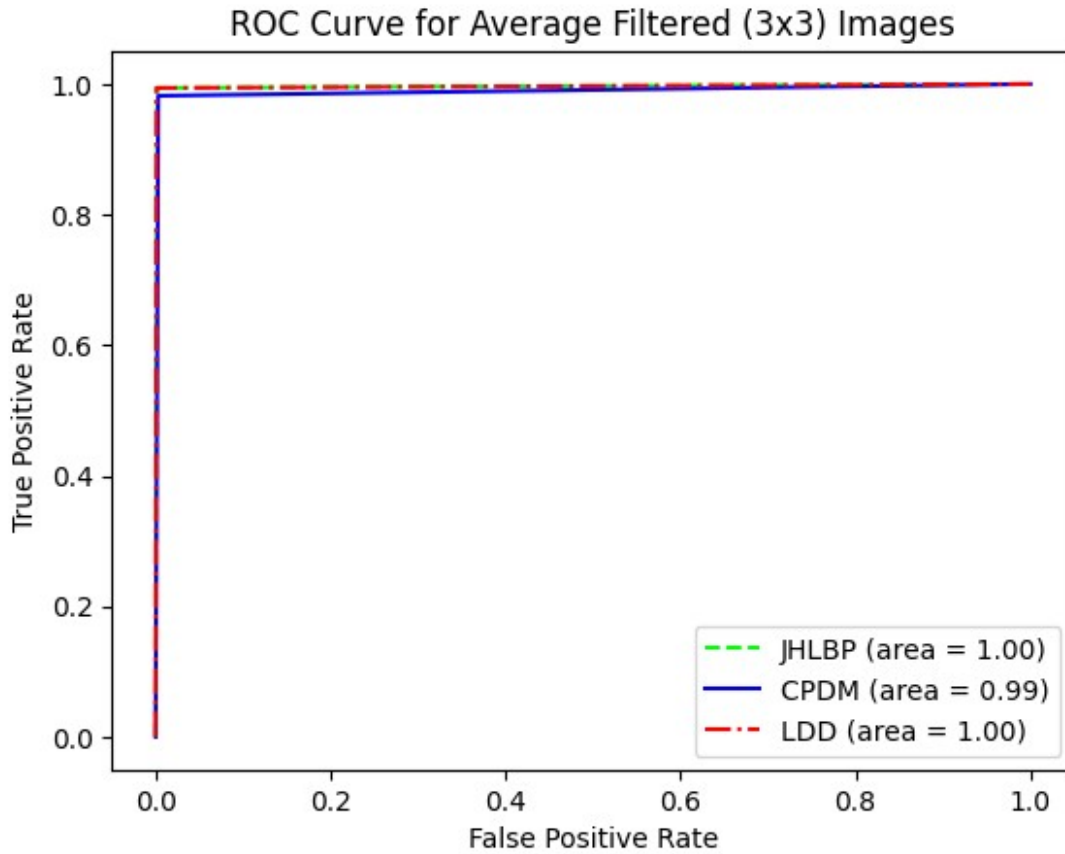


### Average Filtered 3x3 Images ROC Curve

Below Curve is based on Model Trained on Custom Parameters

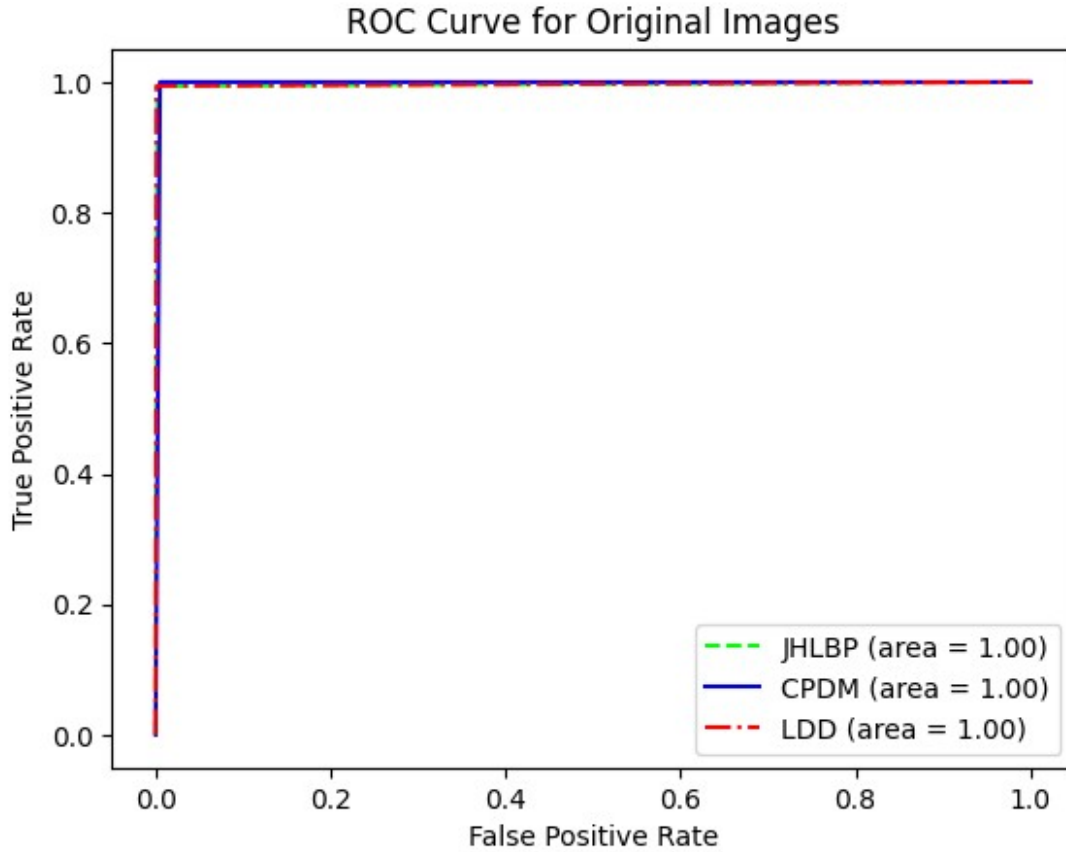
```
# Plotting the ROC Curve for JHLBP Features green Class
plt.plot(fpr_green_jhlbp, tpr_green_jhlbp, label='JHLBP (area = %0.2f)' % auc(fpr_green_jhlbp, tpr_green_jhlbp), color='lime',
linestyle='--')
# Plotting the ROC Curve for CPDM Features green Class
plt.plot(fpr_green_cpdm, tpr_green_cpdm, label='CPDM (area = %0.2f)' % auc(fpr_green_cpdm, tpr_green_cpdm), color='blue')
# Plotting the ROC Curve for LDD Features green Class
plt.plot(fpr_green_ldd, tpr_green_ldd, label='LDD (area = %0.2f)' % auc(fpr_green_ldd, tpr_green_ldd), color='red', linestyle='-.')

# Title and Labels
plt.title('ROC Curve for Average Filtered (3x3) Images')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.legend(loc="lower right")
plt.show()
```



Below Curve is based on Model Trained on Original Parameters

```
# Plotting the ROC Curve for JHLBP Features green Class using Original
Parameters
plt.plot(fpr_green_jhlbp_2, tpr_green_jhlbp_2, label='JHLBP (area =
%0.2f)' % auc(fpr_green_jhlbp_2, tpr_green_jhlbp_2), color='lime',
linestyle='--')
# Plotting the ROC Curve for CPDM Features green Class using Original
Parameters
plt.plot(fpr_green_cpdm_2, tpr_green_cpdm_2, label='CPDM (area =
%0.2f)' % auc(fpr_green_cpdm_2, tpr_green_cpdm_2), color='blue')
# Plotting the ROC Curve for LDD Features green Class using Original
Parameters
plt.plot(fpr_green_ldd_2, tpr_green_ldd_2, label='LDD (area = %0.2f)'
% auc(fpr_green_ldd_2, tpr_green_ldd_2), color='red', linestyle='-.')
# Title and Labels
plt.title('ROC Curve for Original Images')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.legend(loc="lower right")
plt.show()
```



**Finding  $P_e$ :**  $P_e = \min \left( \frac{P_{fp} + 1 - P_{tp}}{2} \right)$

*# Calculate the  $P_e$  for JHLBP Features*

```
Pe_red_jhlbp = min((fpr_red_jhlbp + 1 - tpr_red_jhlbp)/2)
Pe_blue_jhlbp = min((fpr_blue_jhlbp + 1 - tpr_blue_jhlbp)/2)
Pe_green_jhlbp = min((fpr_green_jhlbp + 1 - tpr_green_jhlbp)/2)
Pe_yellow_jhlbp = min((fpr_yellow_jhlbp + 1 - tpr_yellow_jhlbp)/2)
```

*# Calculate the  $P_e$  for JHLBP Features using Original Parameters*

```
Pe_red_jhlbp_2 = min((fpr_red_jhlbp_2 + 1 - tpr_red_jhlbp_2)/2)
Pe_blue_jhlbp_2 = min((fpr_blue_jhlbp_2 + 1 - tpr_blue_jhlbp_2)/2)
Pe_green_jhlbp_2 = min((fpr_green_jhlbp_2 + 1 - tpr_green_jhlbp_2)/2)
Pe_yellow_jhlbp_2 = min((fpr_yellow_jhlbp_2 + 1 - tpr_yellow_jhlbp_2)/2)
```

*# Calculate the  $P_e$  for CPDM Features*

```
Pe_red_cpdm = min((fpr_red_cpdm + 1 - tpr_red_cpdm)/2)
Pe_blue_cpdm = min((fpr_blue_cpdm + 1 - tpr_blue_cpdm)/2)
Pe_green_cpdm = min((fpr_green_cpdm + 1 - tpr_green_cpdm)/2)
Pe_yellow_cpdm = min((fpr_yellow_cpdm + 1 - tpr_yellow_cpdm)/2)
```

*# Calculate the  $P_e$  for CPDM Features using Original Parameters*

```
Pe_red_cpdm_2 = min((fpr_red_cpdm_2 + 1 - tpr_red_cpdm_2)/2)
```

```
Pe_blue_cpdm_2 = min((fpr_blue_cpdm_2 + 1 - tpr_blue_cpdm_2)/2)
Pe_green_cpdm_2 = min((fpr_green_cpdm_2 + 1 - tpr_green_cpdm_2)/2)
Pe_yellow_cpdm_2 = min((fpr_yellow_cpdm_2 + 1 - tpr_yellow_cpdm_2)/2)
```

```
# Calculate the Pe for LDD Features
```

```
Pe_red_ldd = min((fpr_red_ldd + 1 - tpr_red_ldd)/2)
Pe_blue_ldd = min((fpr_blue_ldd + 1 - tpr_blue_ldd)/2)
Pe_green_ldd = min((fpr_green_ldd + 1 - tpr_green_ldd)/2)
Pe_yellow_ldd = min((fpr_yellow_ldd + 1 - tpr_yellow_ldd)/2)
```

```
# Calculate the Pe for LDD Features using Original Parameters
```

```
Pe_red_ldd_2 = min((fpr_red_ldd_2 + 1 - tpr_red_ldd_2)/2)
Pe_blue_ldd_2 = min((fpr_blue_ldd_2 + 1 - tpr_blue_ldd_2)/2)
Pe_green_ldd_2 = min((fpr_green_ldd_2 + 1 - tpr_green_ldd_2)/2)
Pe_yellow_ldd_2 = min((fpr_yellow_ldd_2 + 1 - tpr_yellow_ldd_2)/2)
```

```
# Print the Pe for JHLBP Features
```

```
print("Pe (%) for JHLBP Features with Params as powers of 10:")
print(f"MF3 Class: {Pe_red_jhlbp*100:.4f}%")
print(f"MF5 Class: {Pe_yellow_jhlbp*100:.4f}%")
print(f"ORI Class: {Pe_blue_jhlbp*100:.4f}%")
print(f"AVE Class: {Pe_green_jhlbp*100:.4f}%")
```

```
Pe (%) for JHLBP Features with Params as powers of 10:
```

```
MF3 Class: 0.7466%
MF5 Class: 0.6981%
ORI Class: 0.3484%
AVE Class: 0.3990%
```

```
# Print the Pe for JHLBP Features using Original Parameters
```

```
print("Pe (%) for JHLBP Features with Original Params:")
print(f"MF3 Class: {Pe_red_jhlbp_2*100:.4f}%")
print(f"MF5 Class: {Pe_yellow_jhlbp_2*100:.4f}%")
print(f"ORI Class: {Pe_blue_jhlbp_2*100:.4f}%")
print(f"AVE Class: {Pe_green_jhlbp_2*100:.4f}%")
```

```
Pe (%) for JHLBP Features with Original Params:
```

```
MF3 Class: 0.8958%
MF5 Class: 0.7479%
ORI Class: 0.3484%
AVE Class: 0.3990%
```

```
# Print the Pe for CPDM Features
```

```
print("Pe (%) for CPDM Features")
print(f"MF3 Class: {Pe_red_cpdm*100:.4f}%")
print(f"MF5 Class: {Pe_yellow_cpdm*100:.4f}%")
print(f"ORI Class: {Pe_blue_cpdm*100:.4f}%")
print(f"AVE Class: {Pe_green_cpdm*100:.4f}%")
```

```
Pe (%) for CPDM Features
```

```
MF3 Class: 11.3469%
MF5 Class: 8.4772%
```

ORI Class: 4.2316%  
AVE Class: 1.0476%

*# Print the Pe for CPDM Features using Original Parameters*

```
print("Pe (%) for CPDM Features with Original Params:")
print(f"MF3 Class: {Pe_red_cpd_2*100:.4f}%")
print(f"MF5 Class: {Pe_yellow_cpd_2*100:.4f}%")
print(f"ORI Class: {Pe_blue_cpd_2*100:.4f}%")
print(f"AVE Class: {Pe_green_cpd_2*100:.4f}%")
```

Pe (%) for CPDM Features with Original Params:

MF3 Class: 10.5507%  
MF5 Class: 7.6789%  
ORI Class: 3.4340%  
AVE Class: 0.2490%

*# Print the Pe for LDD Features*

```
print("Pe (%) for LDD Features")
print(f"MF3 Class: {Pe_red_ldd*100:.4f}%")
print(f"MF5 Class: {Pe_yellow_ldd*100:.4f}%")
print(f"ORI Class: {Pe_blue_ldd*100:.4f}%")
print(f"AVE Class: {Pe_green_ldd*100:.4f}%")
```

Pe (%) for LDD Features

MF3 Class: 1.0451%  
MF5 Class: 0.6480%  
ORI Class: 0.1493%  
AVE Class: 0.3492%

*# Print the Pe for LDD Features using Original Parameters*

```
print("Pe (%) for LDD Features with Original Params:")
print(f"MF3 Class: {Pe_red_ldd_2*100:.4f}%")
print(f"MF5 Class: {Pe_yellow_ldd_2*100:.4f}%")
print(f"ORI Class: {Pe_blue_ldd_2*100:.4f}%")
print(f"AVE Class: {Pe_green_ldd_2*100:.4f}%")
```

Pe (%) for LDD Features with Original Params:

MF3 Class: 0.8958%  
MF5 Class: 0.5982%  
ORI Class: 0.1493%  
AVE Class: 0.3492%

**Displaying Results of  $P_e$  in a Tabular Form**

```
import pandas as pd
```

*# Creating Dataframe for Pe*

```
Pe_df = pd.DataFrame({'JHLBP_10': [Pe_red_jhlbp*100,
Pe_yellow_jhlbp*100, Pe_blue_jhlbp*100, Pe_green_jhlbp*100],
                      'JHLBP_2': [Pe_red_jhlbp_2*100,
Pe_yellow_jhlbp_2*100, Pe_blue_jhlbp_2*100, Pe_green_jhlbp_2*100],
                      'CPDM_10': [Pe_red_cpd_2*100, Pe_yellow_cpd_2*100,
```

```

Pe_blue_cpdm*100, Pe_green_cpdm*100],
    'CPDM_2': [Pe_red_cpdm_2*100,
Pe_yellow_cpdm_2*100, Pe_blue_cpdm_2*100, Pe_green_cpdm_2*100],
    'LDD_10': [Pe_red_ldd*100, Pe_yellow_ldd*100,
Pe_blue_ldd*100, Pe_green_ldd*100],
    'LDD_2': [Pe_red_ldd_2*100, Pe_yellow_ldd_2*100,
Pe_blue_ldd_2*100, Pe_green_ldd_2*100]}
    ,index=['MF3 - Pe (%)', 'MF5 - Pe (%)', 'ORI -
Pe (%)', 'AVE - Pe (%)'])

```

Pe\_df

	JHLBP_10	JHLBP_2	CPDM_10	CPDM_2	LDD_10
LDD_2					
MF3 - Pe (%)	0.746566	0.895820	11.346855	10.550736	1.045074
0.895820					
MF5 - Pe (%)	0.698106	0.747907	8.477169	7.678865	0.648007
0.598206					
ORI - Pe (%)	0.348358	0.348358	4.231633	3.434026	0.149254
0.149254					
AVE - Pe (%)	0.399003	0.399003	1.047606	0.249004	0.349202
0.349202					

## Conclusion

As a conclusion, we see that *CPDM* in its own are not the best features but when combined with *LBP* features, they compliment them beautifully and create a new feature set i.e. *LDD* feature set which has the best discriminative ability amongst all the 3 feature sets discussed in this report.