



PROJECT TOPIC: VIOLENT AND NON-VIOLENT SITUATION DETECTION

- ► This project aims to detect violent or non-violent situations from the images using Computer Vision.
- The datasets are taken from various sources such as Kaggle and Roboflow.com. Some data are images and some are in video form. Different images will be taken from videos for the project.

Resources for dataset:

- ► 1.Kaggle.com(images and videos)
- 2.Universal roboflow.com(images)

PROJECT

The main goal of this project is to develop a computer vision system that could perform

an object (for example, coins, faces, animals, cars, plants) recognition (classification)

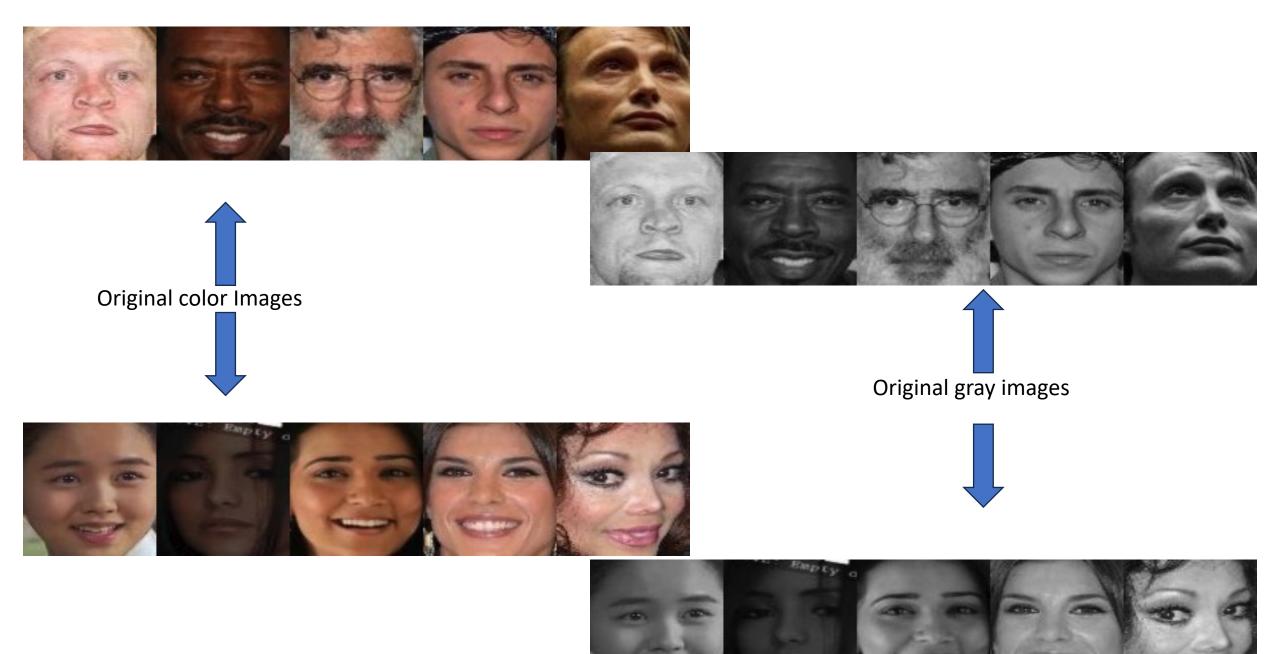
task by using at least 5 local and global features (Haralick Texture, Box Counting,

Chain Code, Area, and Local Binary Pattern)

Project Introduction

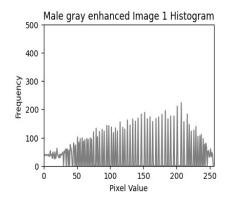
This project is about gender detection where 5 male and 5 female images are used for preprocessing and feature extraction. Finally, the correlation coefficient is used to measure the similarity between males and females respectively.

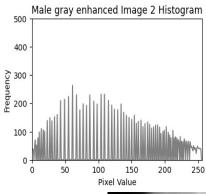
Images taken from Kaggle.com

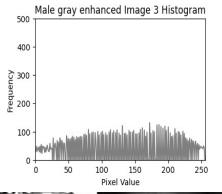


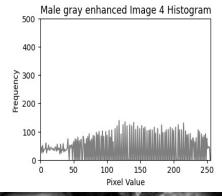
Enhanced Images with Histogram Equalization

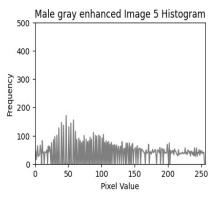


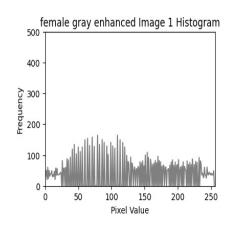


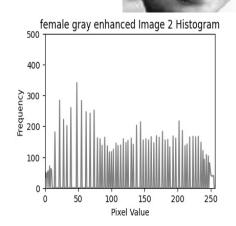


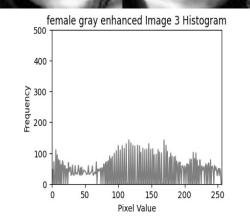


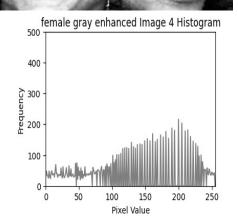


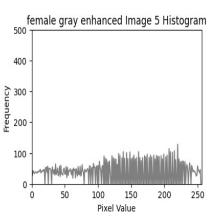






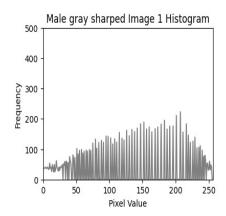




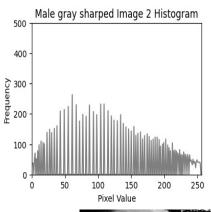


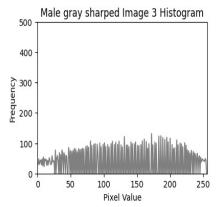
Sharpened Images Using

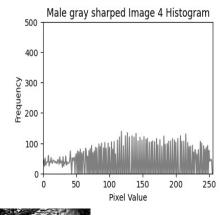
kernel = np.array([[-0.5,-.5,-.5], [-.5, 5,-.5], [-.5,-.5,-.5]])

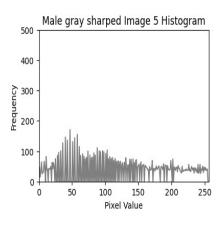


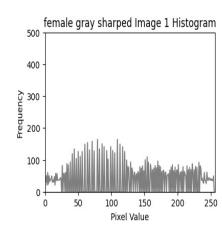


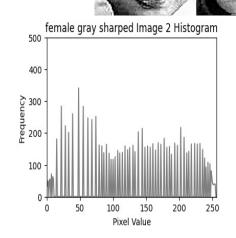


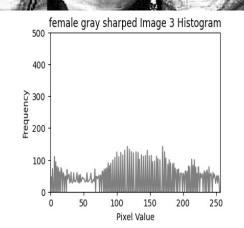


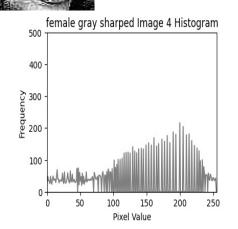


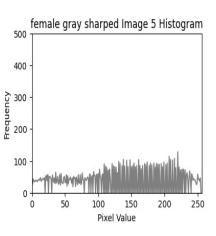












Images Preprocessing and feature generation

- 1. The images are of different sizes in the list, so resizing them to (100x100) size.
- 2. Applying Various feature extraction methods to extract features from images.

1. Haralick Texture features

Male Image 1 Haralick values: contrast:

4034.849494949494 homogeneity: 0.1968942557814769

energy: 0.12272569313528095 correlation:

0.7683814703584944

Male Image 2 Haralick values: contrast:

2312.089191919191 homogeneity: 0.18173083013656305

energy: 0.09992783846718796 correlation:

0.8508849152337469

Male Image 3 Haralick values: contrast:

3983.5212121212126 homogeneity: 0.17383596945223445

energy: 0.10829547736311224 correlation:

0.7590761552708296

Male Image 4 Haralick values: contrast:

2175.6860606060604 homogeneity: 0.16691790537028942

energy: 0.0916323362822626 correlation:

0.8595936550913024

Male Image 5 Haralick values: contrast:

1838.6469696969698 homogeneity: 0.16355958293158473

energy: 0.08306049612007312 correlation:

0.8697979204191079

L

2

3

4

5



Female Image 1 Haralick values: contrast: 1360.080505050505 homogeneity: 0.15385161438812073 energy: 0.05197197921778521 correlation: 0.8972944219136472 Female Image 2 Haralick values: contrast: 1572.30717171717 homogeneity: 0.13639498081009704 energy: 0.05412770254535551 correlation: 0.8805087357761103 Female Image 3 Haralick values: contrast: 1117.5352525252529 homogeneity: 0.147986796652242 energy: 0.05265247800878188 correlation: 0.9187720432715132 Female Image 4 Haralick values: contrast: 2481.29626262624 homogeneity: 0.14329502975479902 energy: 0.07394618973019998 correlation: 0.8328574672804369 Female Image 5 Haralick values: contrast: 2947.6292929294 homogeneity: 0.1332565737936554 energy: 0.07023587585643595 correlation:

0.8069268116709228

2.Box Counting

- Convert to grayscale to deal with binary operations and contour detection
- Setting threshold values to create a binary image from which contour can be detected
- _, thresh = cv2.threshold(gray, 128, 255, cv2.THRESH_BINARY)
- iii. Counter detection in the threshold image which are shapes or boundaries of the object in the image.

Image 1



Image 2



Image 3



Image 4



Image 5



Thresholded



Thresholded Thresholded





Thresholded



Thresholded



Box Counting Values for Male Images:

Male Image 1: 268

Male Image 2:

Male Image 3: 268

Male Image 4: 161

Male Image 5: 126

Box counting is a method used in image analysis to determine the complexity or granularity of an image. It involves counting the number of basic shapes (often squares or boxes) required to cover an image at different scales or resolutions.

The process typically involves thresholding the image to convert it into a binary image (black and white), where areas of interest become white (foreground) and the rest become black (background). Then, the number of contours or connected components in the binary image is counted.

Box Counting Values for Female Images:

Female Image 1: 113

Female Image 2: 145

Female Image 3: 79

Female Image 4: 169

Female Image 5: 208

Image 1



lmage 2



Image 3



Image 4







Thresholded Thresholded Thresholded Thresholded





3.Chain Code

- i. Convert to gray for threshold and contour detection
- ii. Apply binary threshold and convert to a binary image (0 or 255)
- iii. Detects contours in the threshold images

Features extracted

Male Chain Code



Male Chain Code



Male Chain Code



Female Chain Code



Female Chain Code



Female Chain Code



Male Chain Code



Male Chain Code



Female Chain Code



Female Chain Code



Sample values

```
Male Image 1 Chain
Code Values: [[], [],
[], [], [(0, 1)],
[(0, 1)], [(0, 1)],
[(1, 0)], [], [(-1,
1), (0, 1), (-1, 1),
(-1, 0)
```

There are 100's of values for (x,y) for each images

A chain code is a simple and efficient method used in image processing and computer vision for contour representation. It represents a contour by encoding the directions of consecutive boundary points of an object or shape in an image.

4. Extract Area

• Use Numpy to count the number of pixels with values greater than 0. Compute the sum of pixels and calculate the non-zero area regions









































Male 1:8274 Male 2:8303 Male 3:8118 Male 4:8141

Male 5:8046

Female 1:8992 Female 2:8303 Female 3:8830 Female 4:8708

Female 5:8678

The process of extracting an area from an image involves defining a specific region of interest (ROI) and then cropping the image to retain only that particular area, discarding the rest. This extraction can be based on various criteria such as pixel coordinates, shape, size, or features within the image.

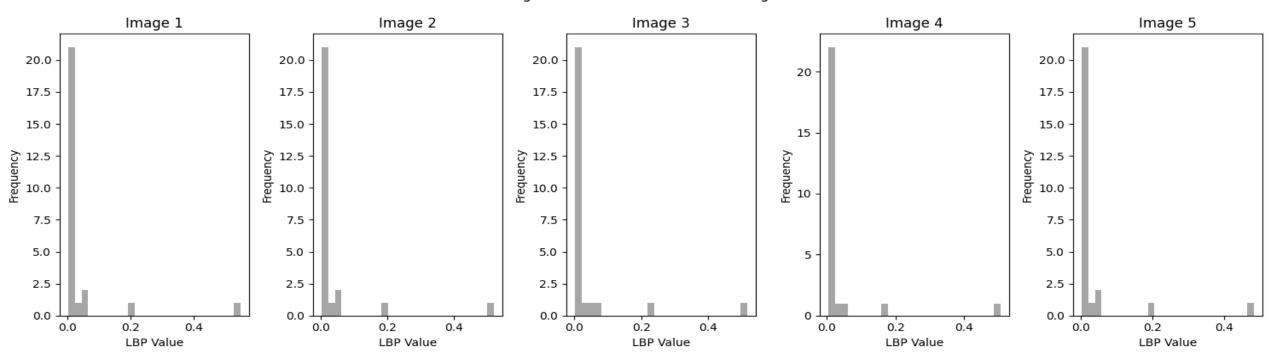
5.Extract lbp(local binary pattern)

- i. Convert to gray
- ii. Set the radius and calculate n_points(points in the LBP neighborhood
- iii. Computes the LBP image using the parameters
- iv. Compute the histogram of the LBP image considering a range of values from 0 to

Density =true normalize the hist to be the probability density

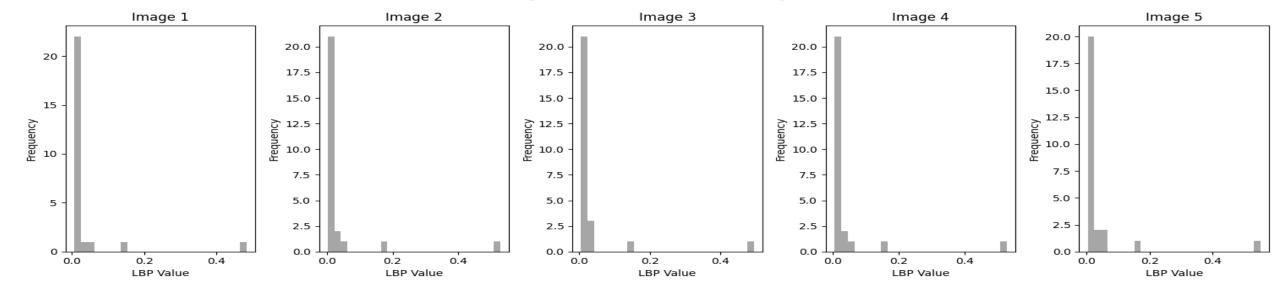


Histograms of LBP Values for Male Images





Histograms of LBP Values for female Images



Male Image LBP Values: Image 1 LBP values: [0.0608 0.0611 0.0229 0.0105 0.0049 0.0041 0.0019 0.0029 0.0026 0.0031 0.0028 0.0028 0.0024 0.0038 0.0025 0.003 0.0019 0.0034 0.0032 0.0032 0.0044 0.0065 0.0089 0.0183 0.211 0.5471] Image 2 LBP values: [0.0556 0.0437 0.0305 0.0146 0.009 0.0071 0.006 0.0055 0.0054 0.0057 0.0076 0.008 0.0085 0.01 0.0064 0.0059 0.0038 0.0037 0.004 0.005 0.0051 0.0083 0.0145 0.0164 0.1898 0.5199] Image 3 LBP values: [0.0652 0.0518 0.0264 0.0132 0.0088 0.0065 0.0029 0.0032 0.0036 0.0046 0.0035 0.0045 0.0046 0.0052 0.0035 0.0039 0.003 0.0032 0.0024 0.0033 0.0032 0.0066 0.0111 0.0154 0.2246 0.5164] Image 4 LBP values: [0.0528 0.0428 0.0205 0.0117 0.0063 0.0056 0.0059 0.0071 0.0066 0.0079 0.0124 0.0144 0.0203 0.0181 0.0099 0.0085 0.007 0.006 0.0043 0.0063 0.0066 0.0097 0.013 0.0187 0.1721 0.5055] Image 5 LBP values: [0.0525 0.0425 0.0237 0.0154 0.0116 0.0099 0.0071 0.0085 0.0083 0.0083 0.0137 0.0169 0.0189 0.0168 0.0088 0.0076 0.0057 0.0051 0.0041 0.0028 0.0028 0.0063 0.0085 0.0101 0.2008 0.4833]

Female Image LBP Values: Image 1 LBP values:

[0.0542 0.0396 0.0213 0.0123 0.01 0.0096 0.0095 0.0114 0.0139 0.016 0.0183 0.0211 0.023 0.0226 0.0128 0.0103 0.0075 0.0073 0.0063 0.007 0.0075 0.0079 0.0107 0.0166 0.1385 0.4848] Image 2 LBP values: [0.0519 0.0418 0.0266 0.0134 0.0104 0.0078 0.0081 0.0073 0.0073 0.0077 0.0099 0.0116 0.0134 0.0118 0.0081 0.0069 0.0063 0.0046 0.0023 0.0048 0.0057 0.0052 0.0111 0.0166 0.1713 0.5281] Image 3 LBP values: [0.0422 0.0385 0.0213 0.0152 0.0127 0.0082 0.0083 0.0105 0.0102 0.0127 0.0173 0.0234 0.0248 0.0207 0.0123 0.0106 0.0072 0.0068 0.0049 0.0064 0.0057 0.0075 0.0106 0.0122 0.1546 0.4952] Image 4 LBP values: [0.0523 0.0424 0.0256 0.011 0.0087 0.0063 0.0048 0.0059 0.0063 0.0059 0.0101 0.0133 0.0161 0.0159 0.01 0.0083 0.0071 0.0064 0.0043 0.0048 0.0058 0.0105 0.0138 0.0167 0.158 0.5297] Image 5 LBP values: [0.062 0.0507 0.0259 0.0125 0.0065 0.0045 0.0032 0.0032 0.0037 0.0067 0.0048 0.0062 0.0084 0.008 0.0055 0.0055 0.0043 0.004 0.0047 0.0031 0.0053 0.0087 0.0126 0.025 0.1622 0.5528]

Now, combining features

| | | Gender I | Haralick_Contrast Hara | alick_Homogeneity H | laralick_Energy Har | alick_Correlation Box_Coun | nting_Values Area_Va | Lues LBP_Values | Chain_Code_Values |
|------------------------|---|-----------|------------------------|---------------------|---------------------|----------------------------|----------------------|---|---|
| | 0 | Female | 1360.080505 | 0.153852 | 0.051972 | 0.897294 | 113 | 9291 [0.0542, 0.0396, 0.0213, 0.0123, 0.01, 0.0096, | [[(-1, 1), (0, 1), (-1, 1), (-1, 0), (1, 0), (|
| Null Values check | 1 | Female | 1572.307172 | 0.136395 | 0.054128 | 0.880509 | 145 | 9276 [0.0519, 0.0418, 0.0266, 0.0134, 0.0104, 0.007 | [[(1, 1), (1, 0), (1, 1), (1, 0), (1, 0), (1, |
| Gender 0 | 2 | Female | 1117.535253 | 0.147987 | 0.052652 | 0.918772 | 79 | 9382 [0.0422, 0.0385, 0.0213, 0.0152, 0.0127, 0.008 | [[], [], [], [(1, 0), (1, 0), (1, 0), (1, 1), |
| Haralick_Contrast 0 | 3 | Female | 2481.296263 | 0.143295 | 0.073946 | 0.832857 | 169 | 9055 [0.0523, 0.0424, 0.0256, 0.011, 0.0087, 0.0063 | [[(0, 1), (0, 1), (0, 1), (0, -1), (0, -1)], [|
| Haralick_Homogeneity (| 4 | Female | 2947.629293 | 0.133257 | 0.070236 | 0.806927 | 208 | 8994 [0.062, 0.0507, 0.0259, 0.0125, 0.0065, 0.0045 | [[], [(0, 1), (0, 1), (0, 1), (0, -1), (0, -1) |
| Haralick_Energy 0 | G | ender Har | alick_Contrast Haral: | ick_Homogeneity Ha | ralick_Energy Hara | lick_Correlation Box_Coun | ting_Values Area_Va | ues LBP_Values | Chain_Code_Value |
| Haralick_Correlation 0 | 0 | Male | 4034.849495 | 0.196894 | 0.122726 | 0.768381 | 268 | 3606 [0.0608, 0.0611, 0.0229, 0.0105, 0.0049, 0.004 | [[(1, 0), (1, 0), (-1, 0)], [(0, 1), (1, 1), (. |
| Box_Counting_Values 0 | 1 | Male | 2312.089192 | 0.181731 | 0.099928 | 0.850885 | 169 | 8827 [0.0556, 0.0437, 0.0305, 0.0146, 0.009, 0.0071 | [[], [], [], [(1, 0), (1, 0), (-1, 0)], [], [(. |
| Area_Values 0 | 2 | Male | 3983.521212 | 0.173836 | 0.108295 | 0.759076 | 268 | 8605 [0.0652, 0.0518, 0.0264, 0.0132, 0.0088, 0.006 | [[], [], [(0, 1)], [], [(1, 0)], [(0, 1), (0,]) |
| LBP_Values 0 | 3 | Male | 2175.686061 | 0.166918 | 0.091632 | 0.859594 | 161 | 3760 [0.0528, 0.0428, 0.0205, 0.0117, 0.0063, 0.005 | [[(0, 1), (0, 1), (0, 1), (0, 1), (0, 1), (0, . |
| Chain_Code_Values 0 | 4 | Male | 1838.646970 | 0.163560 | 0.083060 | 0.869798 | 126 | 3957 [0.0525, 0.0425, 0.0237, 0.0154, 0.0116, 0.009 | [[], [(-1, 1), (0, 1), (1, 1), (0, 1), (0, 1), |
| | | | | | | | | | |

Similarity Check for features using the Correlation Coefficient

Using LBP and Chain Code features complicated the calculations as they have 1000 of (x,y) values in array form, also they reduced the Similarity measures, so dropping these two features from the data frame and also tried distance metrics which doesn't seem to be working as threshold needed to be about 5000.

We are taking row 0 to compare with other rows to check the similarity measures

| G | ender | Haralick_Contrast | Haralick_Homogeneity | Haralick_Energy | Haralick_Correlation | Box_Counting_Values | Area_Values | LBP_Values | Chain_Code_Values |
|---|-------|-------------------|----------------------|-----------------|----------------------|---------------------|-------------|--|--|
| 0 | Male | 4034.849495 | 0.196894 | 0.122726 | 0.768381 | 268 | 8606 | [0.0608, 0.0611, 0.0229, 0.0105, 0.0049, 0.004 | [[(1, 0), (1, 0), (-1, 0)], [(0, 1), (1, 1), (|
| 1 | Male | 2312.089192 | 0.181731 | 0.099928 | 0.850885 | 169 | 8827 | [0.0556, 0.0437, 0.0305, 0.0146, 0.009, 0.0071 | [[], [], [], [(1, 0), (1, 0), (-1, 0)], [], [(|
| 2 | Male | 3983.521212 | 0.173836 | 0.108295 | 0.759076 | 268 | 8605 | [0.0652, 0.0518, 0.0264, 0.0132, 0.0088, 0.006 | [[], [], [(0, 1)], [], [(1, 0)], [(0, 1), (0, |
| 3 | Male | 2175.686061 | 0.166918 | 0.091632 | 0.859594 | 161 | 8760 | [0.0528, 0.0428, 0.0205, 0.0117, 0.0063, 0.005 | {(0, 1), (0, 1), (0, 1), (0, 1), (0, 1), (0, |
| 4 | Male | 1838.646970 | 0.163560 | 0.083060 | 0.869798 | 126 | 8957 | [0.0525, 0.0425, 0.0237, 0.0154, 0.0116, 0.009 | [[], [(-1, 1), (0, 1), (1, 1), (0, 1), (0, 1), |

Male Images features

| Row 0 Threshold | Similarity | Dissimilarity |
|--------------------|------------------------|---------------------|
| 0.91 | row1, row2, row3, row4 | None |
| 0.95 | row1,row2,row3,row4 | None |
| 0.99 | row2 | row1, row3, row4 |
| 1 | None | row1,row2,row3,row4 |



| Gender | Haralick_Contrast | Haralick_Homogeneity | Haralick_Energy | ${\tt Haralick_Correlation}$ | Box_Counting_Values | Area_Values | LBP_Values | Chain_Code_Values |
|----------|-------------------|----------------------|-----------------|-------------------------------|---------------------|-------------|--|--|
| 0 Female | 1360.080505 | 0.153852 | 0.051972 | 0.897294 | 113 | 9291 | [0.0542, 0.0896, 0.0213, 0.0123, 0.01, 0.0096, | [[(-1, 1), (0, 1), (-1, 1), (-1, 0), (1, 0), (|
| 1 Female | 1572.307172 | 0.136395 | 0.054128 | 0.880509 | 145 | 9276 | [0.0519, 0.0418, 0.0266, 0.0134, 0.0104, 0.007 | [[(1, 1), (1, 0), (1, 1), (1, 0), (1, 0), (1, |
| 2 Female | 1117.535253 | 0.147987 | 0.052652 | 0.918772 | 79 | 9382 | [0.0422,0.0385,0.0213,0.0152,0.0127,0.008 | [[], [], [], [(1, 0), (1, 0), (1, 0), (1, 1), |
| 3 Female | 2481.296263 | 0.143295 | 0.073946 | 0.832857 | 169 | 9055 | [0.0523,0.0424,0.0256,0.011,0.0087,0.0063 | [[(0, 1), (0, 1), (0, 1), (0, -1), (0, -1)], [|
| 4 Female | 2947.629293 | 0.133257 | 0.070236 | 0.806927 | 208 | 8994 | [0.062, 0.0507, 0.0259, 0.0125, 0.0065, 0.0045 | [[], [(0, 1), (0, 1), (0, 1), (0, -1), (0, -1) |

Female Images features

| Threshold | |
|-----------|--|
| 0.91 | |
| 0.95 | |
| 0.99 | |
| 1 | |

| Similarity | |
|------------------------|--|
| row1, row2, row3, row4 | |
| row1, row2, row3, row4 | |
| row1, row2, row3 | |
| None | |
| | |

None None row4 row1, row2, row3, row4



Applied chi-square

| | Gender | ${\tt Haralick_Contrast}$ | Haralick_Homogeneity | Haralick_Energy | ${\tt Haralick_Correlation}$ | Box_Counting_Values | Area_Values | LBP_Values | Chain_Code_Values |
|---|--------|----------------------------|----------------------|-----------------|-------------------------------|---------------------|-------------|--|--|
| 0 | Male | 4034.849495 | 0.196894 | 0.122726 | 0.768381 | 268 | 8606 | [0.0608, 0.0611, 0.0229, 0.0105, 0.0049, 0.004 | [[(1, 0), (1, 0), (-1, 0)], [(0, 1), (1, 1), (|
| 1 | Male | 2312.089192 | 0.181731 | 0.099928 | 0.850885 | 169 | 8827 | [0.0556, 0.0437, 0.0395, 0.0146, 0.009, 0.0071 | [[], [], [], [(1, 0), (1, 0), (-1, 0)], [], [(|
| 2 | Male | 3983.521212 | 0.173836 | 0.108295 | 0.759076 | 268 | 8605 | [0.0652, 0.0518, 0.0264, 0.0132, 0.0688, 0.006 | [[],[],[(0,1)],[],[(1,0)],[(0,1),(0, |
| 3 | Male | 2175.686061 | 0.166918 | 0.091632 | 0.859594 | 161 | 8760 | [0.0528, 0.0428, 0.0205, 0.0117, 0.0063, 0.005 | [[(0, 1), (0, 1), (0, 1), (0, 1), (0, 1), (0, |
| 4 | Male | 1838.646970 | 0.163560 | 0.083060 | 0.869798 | 126 | 8957 | [0.0525, 0.0425, 0.0237, 0.0154, 0.0116, 0.009 | [[], [(-1, 1), (0, 1), (1, 1), (0, 1), (0, 1), |

<mark>Male</mark>

Threshold =0.05 No similar rows Threshold = 0.5 Row 2 similar to row 0 Threshold= 1000 Row1, Row2, Row3, Row4 similar to Row

| Gender | Haralick_Contrast | Haralick_Homogeneity | Haralick_Energy | Haralick_Correlation | Box_Counting_Values | Area_Values | LBP_Values | Chain_Code_Values |
|----------|-------------------|----------------------|-----------------|----------------------|---------------------|-------------|--|--|
| 0 Female | 1360.080505 | 0.153852 | 0.051972 | 0.897294 | 113 | 9291 | [0.0542, 0.0396, 0.0213, 0.0123, 0.01, 0.0096, | [[(-1, 1), (0, 1), (-1, 1), (-1, 0), (1, 0), (|
| 1 Female | 1572.307172 | 0.136395 | 0.054128 | 0.880509 | 145 | | [0.0519, 0.0418, 0.0266, 0.0134, 0.0104, 0.007 | |
| 2 Female | 1117.535253 | 0.147987 | 0.052652 | 0.918772 | 79 | 9382 | [0.0422, 0.0385, 0.0213, 0.0152, 6.0127, 0.008 | [[], [], [], [(1, 0), (1, 0), (1, 0), (1, 1), |
| 3 Female | 2481.296263 | 0.143295 | 0.073946 | 0.832857 | 169 | 9055 | [0.0523, 0.0424, 0.0256, 0.011, 0.0087, 0.0063 | [[(0, 1), (0, 1), (0, 1), (0, -1), (0, -1)], [|
| 4 Female | 2947.629293 | 0.133257 | 0.070236 | 0.806927 | 208 | 8994 | [0.062, 0.0507, 0.0259, 0.0125, 0.0065, 0.0045 | [[], [(0, 1), (0, 1), (0, 1), (0, -1), (0, -1) |

<mark>Female</mark>

Threshold =0.05 No similar rows Threshold = 0.5 No similar rows. Threshold= 1000 Row1, Row2, Row3, Row4 similar to Row



1.Gabor_filter A Gabor filter is a linear filter used in signal processing and image processing for tasks like texture analysis, edge detection, and feature extraction. It's particularly useful for analyzing textures within images.



1 4 0 E # E | |











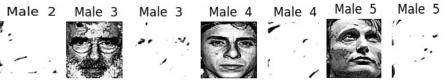




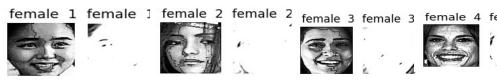
























| _ | | ale . | 4 |
|----|---|-------|---|
| 1. | | ·. · | |
| | - | , | |

| | Harali | ck_Contrast | Haralick_Homogeneity Ha | aralick_Energy | Haralick_Correlation Box_C | Counting_Values Area_V | alues | BP_Values | Chain_Code_Values | Male_Gabor_Pixel_Values_1 | Male_Gabor_Pixel_Values_2 | Male_Gabor_Pixel_Values_3 | Male_Gabor_Pixel_Values_4 | Male_Gabor_Pixel_Values_5 | = |
|---------|----------|-------------|-------------------------|----------------|----------------------------|------------------------|-------|---|---|--|--|--|--|--|------------------|
| | 0 | 4034.849495 | 0.196894 | 0.122726 | 0.768381 | 268 | | [0.0608, 0.0611, 0.0229, 0.0105, 0.0049, 0.004 | [[(1, 0), (1, 0), (-1, 0)], [(0, 1), (1, 1), (| [223, 255, 255, 255, 255, 255, 255, 255, | [223, 255, 255, 255, 255, 255, 255, 255, | [223, 255, 255, 255, 255, 255, 255, 255, | [223, 255, 255, 255, 255, 255, 255, 255, | [223, 255, 255, 255, 255, 255, 255, 255, | |
| | 1 | 2312.089192 | 0.181731 | 0.099928 | 0.850885 | 169 | 8827 | [0.0550 0.0437, 0.0305, 0.0146, 0.009, 0.0071 | [[], [], [], [(1, 0), (1, 0), (-1, 0)], [], [(| [70, 76, 108, 139, 166, 236, 255, 255, 255, 25 | [70, 76, 108, 139, 166, 236, 255, 255, 255, 25 | [70, 76, 108, 139, 166, 236, 255, 255, 255, 25 | [70, 76, 108, 139, 166, 236, 255, 255, 255, 25 | [70, 76, 108, 139, 166, 236, 255, 255, 255, 25 | |
| | 2 | 3983.521212 | 0.173836 | 0.108295 | 0.759076 | 268 | 8605 | [0.0652, 0.0518, 0.0264, 0.0132, 0.0088, 0.006 | [[], [], (0, 1)], [], [(1, 0)], [(0, 1), (0, | | [255, 255, 255, 255, 255, 255, 248, 138, 162, | [255, 255, 255, 255, 255, 255, 255, 248, 138, 162, | [255, 255, 255, 255, 255, 255, 248, 138, 162, | [255, 255, 255, 255, 255, 255, 248, 138, 162, | |
| | 3 | 2175.686061 | 0.166918 | 0.091632 | 0.859594 | 161 | | [0.0528, 0.0428, 0.0205, 0.0117, 0.0063, 0.005 | [[(0, 1), (0, 1)),(0, 1), (0, 1), (0, | | [58, 58, 90, 142, 176, 194, 167, 131, 110, 128 | [58, 58, 90, 142, 176, 194, 167, 131, 110, 128 | [58, 58, 90, 142, 176, 194, 167, 131, 110, 128 | [58, 58, 90, 142, 176, 194, 167, 131, 110, 128 | |
| | 4 | 1838.646970 | 0.163560 | 0.083060 | 0.869798 | 126 | 8957 | [0.0525, 0.0425, 0.0237, 0.0154, 0.0116, 0.009 | [[], [(-1, 1), (0, 1), (1, 1), (0, 1), (0, 1), (0, 1) | [255, 255, 255, 255, 255, 255, 255, 255, | [255, 255, 255, 255, 255, 255, 255, 255, | [255, 255, 255, 255, 255, 255, 255, 255, | [255, 255, 255, 255, 255, 255, 255, 255, | [255, 255, 255, 255, 255, 255, 255, 255, | |
| 0 | female_c | df | | | | | | | | | | | | | |
| <u></u> | | *** | Haralick Homogeneity Ha | aralick Energy | Haralick Correlation Box C | Counting Values Area V | alues | BP Values (| Chain Code Values F | emale Gabor Pixel Values 1 | Female Gabor Pixel Values 2 | Female Gabor Pixel Values 3 | Female Gabor Pixel Values | 4 Female Gabor Pixel Valu | ues 5 |
| | 0 | 1360.080505 | 0.153852 | 0.051972 | 0.897294 | 113 | 9291 | [0.0542, 0.0396, .0213, 0.0123,0.01, 0.0036, | [[(-1, 1), (0, 1), (-1, 1), (-1, 0), (1, 0), (| [25, 26, 29, 34, 37, 35, 37, 33, 35, 43, 55, 7 | [25, 26, 29, 34, 37, 35, 37, 33, 35, 43, 55, 7 | [25, 26, 29, 34, 37, 35, 37, 33, 35 43, 55, 7 | | 35, [25, 26, 29, 34, 37, 35, 37, 3 ' 43, 5 | 33, 35, 55, 7 |
| 8 | 1 | 1572.307172 | 0.136395 | 0.054128 | 0.880509 | 145 | 9276 | [0.0510, 0.0418] 0.0266, 0.0134, 0.0104, 0.007 | [[(1, 1), (1, 0), (1, 1), (1, 0), (1, 0), (1, | [255, 255, 255, 255, 255, 255, 255, | [255, 255, 255, 255, 255, 255, 255, 255, | [255, 255, 255, 255, 255, 255, 255, 255, | | | |
| | 2 | 1117.535253 | 0.147987 | 0.052652 | 0.918772 | 79 | 9382 | [0.0422, 0.0385, 0.0213, 0.0152, 0.0127, 0.008 | [], [], A [(1, 0), (1, 0), (1, 0), (1, 0), (1, 1), | [225, 222, 226, 225, 247, 242, 227, 206, 210, | [225, 222, 226, 225, 247, 242, 227, 206, 210, | [225, 222, 226, 225, 247, 242, 227 206, 210, | | | |
| | 3 | 2481.296263 | 0.143295 | 0.073946 | 0.832857 | 169 | 9055 | [0.0523, 0.0424, 0.0256, 0.011, 0.0087, 0.0063 | [[(0, 1), (0, 1), (0, 1), (0, -1), (0, -1), [| [255, 255, 255, 255, 255, 255, 255, | [255, 255, 255, 255, 255, 255, 255, 255, | [255, 255, 255, 255, 255, 255, 255, 255, | | | |
| | 4 | 2947.629293 | 0.133257 | 0.070236 | 0.806927 | 208 | 8994 | [0.062, 0.0507, 0.0259, 0.0125, 0.0065, 0.0045 | [[], [(0, 1), (0, 1), (0, 1), (0, -1), | [255, 255, 244, 188, 149, 127, 136, 157, 186, | [255, 255, 244, 188, 149, 127, 136, | [255, 255, 244, 188, 149, 127, 136 157, 186, | | | |

PCA

Transformed Images for Males:

Image 1 - Transformed Values: [-3.41003199e+03 -3.84031191e+03 -4.89872177e+03 -1.66046846e+03 2.44999530e-12]

Image 2 - Transformed Values: [-2.01860698e+03 6.12583677e+02 6.07860477e+02 6.28077781e+03 2.44999530e-12]

Image 3 - Transformed Values: [8.19002341e+03 -3.64009948e+02 -1.22320292e+03 5.27395010e+01 2.44999530e-12]

Image 4 - Transformed Values: [-1.00270469e+03 -3.38774879e+03 5.60569700e+03 -2.16166229e+03 2.44999530e-12]

Image 5 - Transformed Values: [-1.75867975e+03 6.97948696e+03 -9.16327783e+01 -2.51138657e+03 2.44999530e-12]



Transformed Images for Females:

Image 1 - Transformed Values: [7.63262172e+03 3.34708167e+03 1.05123081e+02 3.83509980e+02 2.11030588e-12]

Image 2 - Transformed Values: [1.29391428e+03 -4.45219382e+03 -1.51947379e+03 -4.03004876e+03 2.11030588e-12]

Image 3 - Transformed Values: [-5.64016238e+03 4.49466163e+03 -9.29661705e+02 -2.05250355e+03 2.11030588e-12]

Image 4 - Transformed Values: [-1.61902870e+03 -1.80431681e+03 -3.54873939e+03 4.41191379e+03 2.11030588e-12]

Image 5 - Transformed Values: [-1.66734492e+03 -1.58523267e+03 5.89275180e+03 1.28712855e+03 2.11030588e-12]



A combined data frame with all the features

2947.629293

0.133257

0.070236

Excluding the crossed parts they were affecting the similarity measures.

| maralick_Lontrast mar | alick_Homogeneity Ha | aralick_Energy Hara | strck_rolleration Rox_ro | ounting_values Area_ | values | mate_wapor_rixet_values_i | male_wapor_rixel_value | s_z mate_wapor_rixet_vatues | s_3 male_Gapor_rixel_value | s_4 male_wapor_rixel_va | .ues_> male_r | CA_1 Male_PCA_ | Z male_rta_3 | male_rtA_4 | male_rta_o |
|-------------------------|----------------------|---------------------|--------------------------|----------------------|----------|--|--|---|---|---|---------------------------|-------------------|----------------|--------------|------------------|
| 4034.849495 | 0.196894 | 0.122726 | 0.768381 | 268 | 8606 | [223, 255, 255, 255, 255, 255, 255, 255, | [223, 255, 255, 255, 255, 255, 255, 255, | | | | 5, 255, 255,3410.03 | 31991 -3840.31190 | 5 -4898.721772 | -1660.468456 | 2.449995e- 12 |
| 2312.089192 | 0.181731 | 0.099928 | 0.850885 | 169 | 8827 | [70, 76, 108, 139, 166, 236, 255, 255, 255, 255, 255, 25 | [70, 76, 108, 139, 166, 236, 255, 2 255, 2 | 55, [70, 76, 108, 139, 166, 236, 255, 2 5 255, 2 | 255, [70, 76, 108, 139, 166, 236, 255, 2 25 255, 2 | 255, [70, 76, 108, 139, 166, 236, 25 25 25 | 55, 255, 55, 252018.60 | 06979 612.58367 | 7 607.860477 | 6280.777813 | 2.449995e- 12 |
| 3983.521212 | 0.173836 | 0.108295 | 0.759076 | 268 | 8605 | [255, 255, 255, 255, 255, 255, 248, 138, 162, | [255, 255, 255, 255, 255, 255, 2 138, 162 | | | | 55, 248, 162, 8190.02 | 23413 -364.00994 | 8 -1223.202923 | 52.739501 | 2.449995e- 12 |
| 2175.686061 | 0.166918 | 0.091632 | 0.859594 | 161 | 8760 | [58, 58, 90, 142, 176, 194, 167, 131, 110, 128 | [58, 58, 90, 142, 176, 194, 167, 1 110, 12 | | 131, [56, 58, 90, 142, 176, 194, 167, 1 28 110, 12 | | 67, 131, 0, 1281002.70 | 04694 -3387.74878 | 7 5605.696996 | -2161.662287 | 2.449995e- 12 |
| 1838.646970 | 0.163560 | 0.083060 | 0.869798 | 126 | 8957 | [255, 255, 255, 255, 255, 255, 255, 255, | [255, 255, 255, 255, 255, 255, 2 255, 255 | | | 255, [255, 255, 255, 255, 255, 255, 255, | 55, 255, 255,1758.67 | 79749 6979.48696 | 3 -91.632778 | -2511.386572 | 2.449995e- 12 |
| combined female with po | al | | _Values", "Chain_Code_Va | | _ 16 | | | V 17 14 V 17 1 1 | 0.00.0000 | | | | | | 251 |
| lick_Contrast Haralic | k_Homogeneity Harali | .ck_Energy Haralick | _Correlation Box_Countin | ng_Values Area_Value | es Fema | le_Gabor_Pixel_Values_1 Femal | Le_Gabor_Pixel_Values_2 Fe | male_Gabor_Pixel_Values_3 Fem | nale_Gabor_Pixel_Values_4 Fema | le_Gabor_Pixel_Values_5 Fe | male_PCA_1 Fe | emale_PCA_2 Fem | ale_PCA_3 Fer | male_PCA_4 F | male_PCA_5 |
| 1360.080505 | 0.153852 | 0.051972 | 0.897294 | 113 92 | 91 [25 | 5, 26, 29, 34, 37, 95, 37, 33, 35, [25, 43, 55, 7 | , 26, 29, 34, 37, 35, 37, 33, 35, 43, 55, 7 | [25, 26, 29, 34, 37, 35, 37, 33, 35, [2 43, 55, 7 | 25, 26, 29, 34, 37, 35, 37, 33, 35, [2: 43, 55, 7 | 5, 26, 29, 34, 37, 35, 37, 33, 35, 43, 55, 7 | 7632.621719 | 3347.081673 | 105.123081 | 383.509980 | 2.110306e-12 |
| 1572.307172 | 0.136395 | 0.054128 | 0.880509 | 145 92 | 76 [255 | 5, 255, 255, 255, 255, 255, 255, 255, 2 | , 255, 255, 255, 255, 255, 255, 255, 25 | 255, 255, 255, 255, 255, 255, 255, [26 | 55, 255, 255, 255, 255, 255, 255, 255, | 5, 255, 255, 255, 255, 255, 255, 255, | 1293.914279 | -4452.193821 -1 | 519.473788 - | 4030.048762 | 2.110306e-12 |
| 1117.535253 | 0.147987 | 0.052652 | 0.918772 | 79 93 | 182 [225 | 5, 222, 226, 225, 247, 242, 227, [225, 206, 210, | , 222, 226, 225, 247, 242, 227, 206, 210, | 225, 222, 286 , 225, 247, 242, 227, [22 206, 210, | 25, 222, 226, 225, 247, 242, 227, [22: 206, 210, | 5, 222, 226, 225, 247, 242, 227, 206, 210, | -5640.162380 | 4494.661634 | 929.661705 - | 2052.503553 | 2.110306e-12 |
| | | | | | | | | | | | | | | | |
| 2481.296263 | 0.143295 | 0.073946 | 0.832857 | 169 90 | 55 [255 | 5, 255, 255, 255, 255, 255, 255, 255, 2 | , 255, 255, 255, 255, 255, 255, 255, [3 | 255, 255, 255, 255, 255, 255, 255, 255, | 55, 255, 255, 255, 255, 255, 255, 255, | 5, 255, 255, 255, 255, 255, 255, 255, | -1619.028699 | -1804.316812 -3 | 548.739389 | 4411.913786 | 2.110306e |

157, 186, ...

Taking row 0 as the main row to compare with others

Male

Threshold Similarity Dissimilarity
0.30 row1, row2, row3, row4 None
0.45 row1, row2, row3, row4 None

Female

Threshold Similarity Dissimilarity
0.30 row1, row2, row3, row4 None
0.45 row1 row2, row3, row4

Tried removing and adding other features but got no better results than above slides



Conclusion:

Having many features doesn't mean that the model will train or work better, we need to check for the best features that are highly correlated and important in the dataset. Also, we saw correlation coefficient worked better for the first 3 features than Ch--Square Haralick, Box counting and Area but Ch—Square worked very weakly.

Best Worked Measure

| , | Gender Ha | aralick_Contrast Har | alick_Homogeneity Ha | ralick_Energy Har | alick_Correlation Box_C | ounting_Values Are | a_Values | LBP_Va1 | lues Chain_Code_ |
|----------------|-------------------------------|---|--|----------------------------------|---------------------------------------|---|--|--|--|
| 0 | Male | 4034.849495 | 0.196894 | 0.122726 | 0.768381 | 268 | 8606 | [0.0608, 0.0611, 0.0229, 0.0105, 0.0049, 0.0 | 004 [[(1, 0), (1, 0), (-1, 0)], [(0, 1), (|
| 1 | Male | 2312.089192 | 0.181731 | 0.099928 | 0.850885 | 169 | 8827 | [0.0556, 0.0437, 0.0305, 0.0146, 0.009, 0.00 | 071 [[], [], [], [(1, 0), (1, 0), (-1, 0 |
| 2 | Male | 3983.521212 | 0.173836 | 0.108295 | 0.759076 | 268 | 8605 | [0.0652, 0.0518, 0.0264, 0.0132, 0.9088, 0.0 | 006 [[], [], [(0, 1)], [], [(1, 0)], [(0, |
| 3 | Male | 2175.686061 | 0.166918 | 0.091632 | 0.859594 | 161 | 8760 | [0.0528, 0.0428, 0.0205, 0.0117, 0.0063, 0.0 | 005 [[(0, 1), (0, 1), (0, 1), (0, 1), (0, |
| 4 | Male | 1838.646970 | 0.163560 | 0.083060 | 0.869798 | 126 | 8957 | [0.0525, 0.0425, 0.0237, 0.0154, 0.0116, 0.0 | 009 [[], [(-1, 1), (0, 1), (1, 1), (0, 1), |
| | Thres 0.91 0.95 0.99 | hold | | • | ity w2, row3, row4 v2,row3,row4 | Dissimilari None None row1, row row1,row2 | , 3, row | | |
| Gende | r Harali | -l. C++ U11 | | | | | | | |
| | | ck_contrast Haralick | _Homogeneity Haralick | _Energy Haralick_(| Correlation Box_Counting | _Values Area_Values | 1 | LBP_Values | Chain_Code_Values |
| Femal | | 1360.080505 | | _Energy Haralick_0 0.051972 | Correlation Box_Counting 0.897294 | _Values Area_Values 113 9291 | | LBP_Values 0396, 0.0213, 0.0123, 0.01, 0.0096, [[(-1, | |
| Femal | le | | 0.153852 | | | 113 9291 | [0.0542, 6 | 0396, 0.0213, 0.0123, 0.01, 0.0096, [[(-1, | |
| Femal Femal | le le | 1360.080505 | 0.153852 0.136395 | 0.051972 | 0.897294 | 113 9291 145 9276 | [0.0542, 6 | 0396, 0.0213, 0.0123, 0.01, 0.0096, [[(-1, 0418, 0.0266, 0.0134, 0.0104, 0.007 [[(1, 1 | 1), (0, 1), (-1, 1), (-1, 0), (1, 0), (|
| Femal | le le | 1360.080505 1572.307172 | 0.153852 0.136395 0.147987 | 0.051972 0.054128 | 0.897294 0.880509 | 113 9291 145 9276 | [0.0542, 6] [0.0519, 0] [0.0422, 0] | | 1), (0, 1), (-1, 1), (-1, 0), (1, 0), (1), (1, 0), (1, 1), (1, 0), (1, 0), (1, |
| Femal | le le le | 1360.080505 1572.307172 1117.535253 | 0.153852 0.136395 0.147987 0.143295 | 0.051972 0.054128 0.052652 | 0.897294 0.880509 0.918772 | 113 9291 145 9276 79 9382 169 9055 | [0.0542, 6] [0.0519, 0] [0.0422, 0] [0.0523, 0] | | 1), (0, 1), (-1, 1), (-1, 0), (1, 0), (1), (1, 0), (1, 1), (1, 0), (1, 0), (1, [], [], [(1, 0), (1, 0), (1, 0), (1, 1), 1), (0, 1), (0, 1), (0, -1), (0, -1)], [|