

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

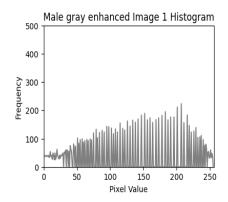
This project is about gender detection where 5 male and 5 female images are used for preprocessing and feature extraction. Finally, the correlation coefficient is used to measure the similarity between males and females respectively.

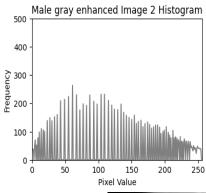
Images taken from Kaggle.com

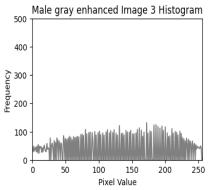


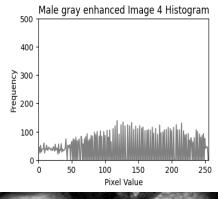
Enhanced Images with Histogram Equalization

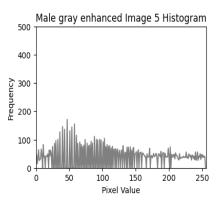


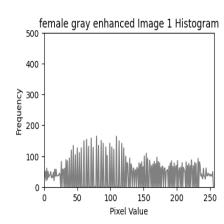


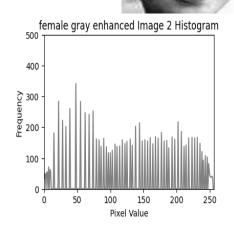


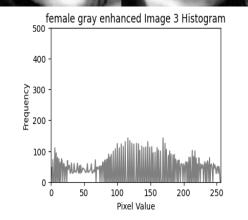


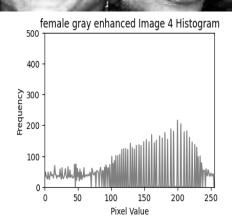


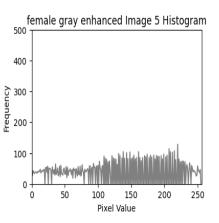






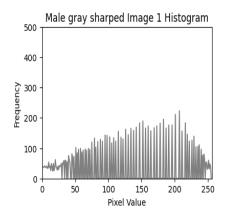




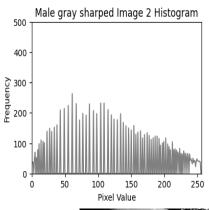


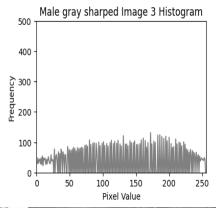
Sharpened Images Using

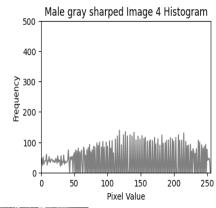
kernel = np.array([[-0.5,-.5,-.5], [-.5, 5,-.5], [-.5,-.5,-.5]])

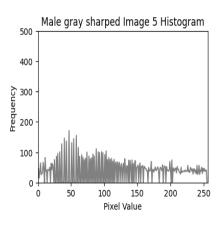


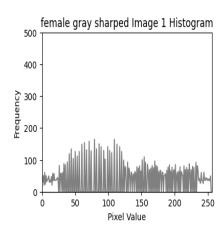


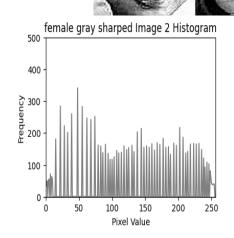


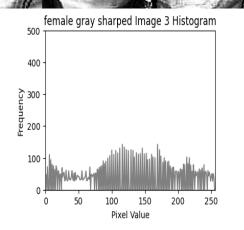


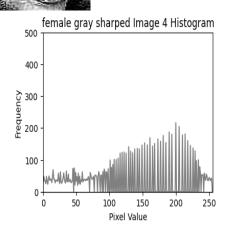


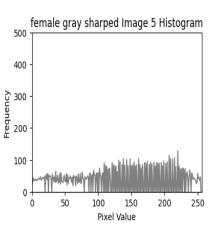












Images Preprocessing and feature generation

- 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

2

3

4



Female Image 1 Haralick values: contrast: 1360.080505050505 homogeneity: 0.15385161438812073 energy: 0.05197197921778521 correlation: 0.8972944219136472 Female Image 2 Haralick values: contrast: 1572.3071717171717 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

lmage 1



lmage 2



lmage 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.

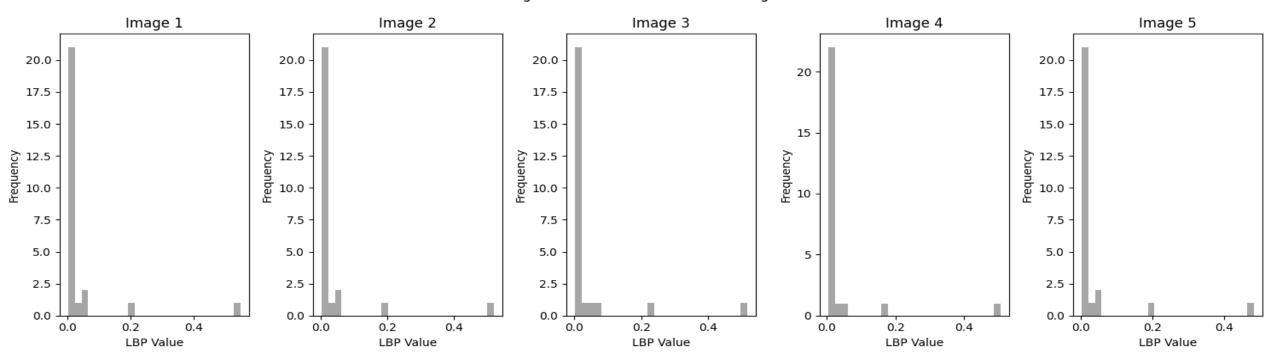
5.Extract lbp(local binary pattern)

- 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

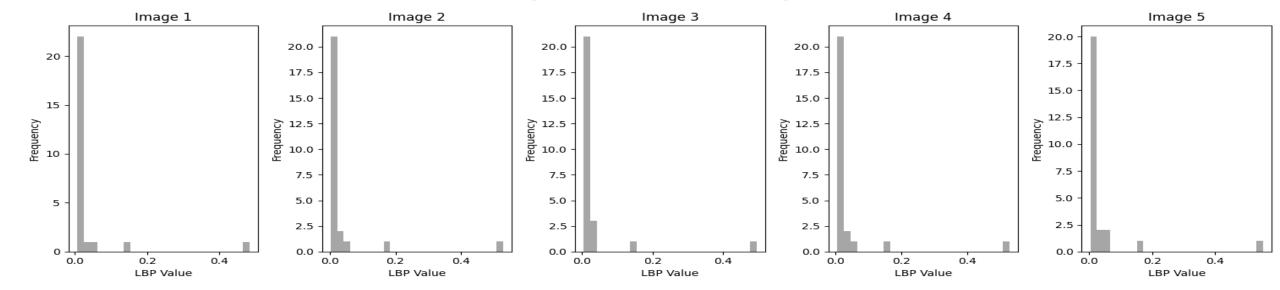


Histograms of LBP Values for Male Images





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 0.0085 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

	Go	ender Haral	ick_Contrast Haralick_Ho	omogeneity Harali	.ck_Energy Haralick_Co	rrelation Box_Counting_Va	lues Area_Va	lues	LBP_Values	Chain_Code_Values
	0 F	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 F	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 F	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 F	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 0 	4 F	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 	Gend	ler Haralich	k_Contrast Haralick_Hom	ogeneity Haralic	k_Energy Haralick_Cor	relation Box_Counting_Va	lues Area_Va	Lues	LBP_Values	Chain_Code_Value
 Haralick_Correlation 0 	0 M	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), (.
. Pay Counting Values 0										
Box_Counting_Values 0	1 M	Male 2	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				0.181731 0.173836	0.099928 0.108295	0.850885 0.759076		8827 [0.0556, 0.0437, 0.0305, 0.0146, 8605 [0.0652, 0.0518, 0.0264, 0.0132,		[[], [], [], [(1, 0), (1, 0), (-1, 0)], [], [([[], [], [(0, 1)], [], [(1, 0)], [(0, 1), (0,
	2 Ma	fale 3	3983.521212				268	•	0.0088, 0.006	[[], [], [(0, 1)], [], [(1, 0)], [(0, 1), (0,

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

0	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	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	Similarity	Dissimilarity
0.91	row1, row2, row3, row4	None
0.95	row1, row2, row3, row4	None
0.99	row1, row2, row3	row4
1	None	row1, row2, row3, row4



Applied chi-square

	Gender	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, 0.0888, 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	${\tt Haralick_Homogeneity}$	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, 0.0396, 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, 8.0127, 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

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.













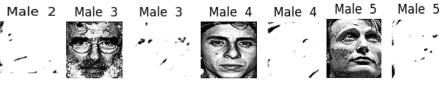


























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1 T V G E # 1 1 1 1

	Harali	ck_Contrast	Haralick_Homogeneity H	Haralick_Energy	Haralick_Correlation Box_Count	ting_Values Area	_Values	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	田
	0	4034.849495	0.196894	0.122726	0.768381	268		[0.0608, 0.0611, 0.0229, 0.0105, 0.0049, 0.004			[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	[0.055] 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, 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			[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_d	if													
→		~	Haralick Homogeneity F	Jaralick Energy	Haralick_Correlation Box_Count	ting Values Area	Values	▲RP 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	ies 5
				,				[0.0542,							
,	0	1360.080505	0.153852	0.051972	0.897294	113	9291	0.0396, 0.0213, 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			33, 35, 55, 7
	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, 255,			
	2	1117.535253	0.147987	0.052652	0.918772	79	9382	[0.0422, 0.0385, 0.0213, 0.0152, 0.0127, 0.008	[[], [], h, [(1, 0), (1, 0), (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), (0, -1)	[255, 255, 244, 188, 149, 127, 136, 157, 186,	[255, 255, 244, 188, 149, 127, 136, 157, 186,	[255, 255, 244, 188, 149, 127, 136, 157, 186,			



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

0.052652

0.073946

0.070236

0.147987

0.143295

0.133257

0.918772

0.832857

0.806927

79

169

208

1117.535253

2481.296263

2947.629293

Excluding the crossed parts they were affecting the similarity measures.

	aralick_lontrast Haralick_	nomogeneity Harau	LCK_Energy HaraLlCK_LOF	relation Box_Lounti	ng_values Area_va	Lues	mate_wapor_rixet_vatues_i	male_wapor_rixe	ı_vaıues_z	mate_wapor_rixet_vat	.ues_s mate_wapo	r_rixet_vatues_4	male_wabor_rixel_	values_5 r	male_rua_i	male_rLA_Z	male_rta_s	male_rta_4	male_rLA_5
0	4034.849495	0.196894	0.122726	0.768381	268	8606	[223, 255, 255, 255, 255, 255, 255, 255,	[223, 255, 255, 255, 2	55, 255, 255, 255, 255,	[223, 255, 255, 255, 255, 25 255,	5, 255, [223, 255, 25 255,	5, 255, 255, 255, 255, 255, 255,		, 255, 255, 255, 255,	3410.031991	-3840.311905	-4898.721772	-1660.468456	2.449995e- 12
1	2312.089192	0.181731	0.099928	0.850885	169	8827	70, 76, 108, 139, 166, 236, 255, 255, 25	[70, 76, 108, 139, 166, 2	36, 255, 255, 255, 25	[70, 76, 108, 139, 166, 236, 25 25	5, 255, [70, 76, 108, 13 5, 25	9, 166, 236, 255, 255, 255, 25	[70, 76, 108, 139, 166, 236	, 255, 255, 255, 25	2018.606979	612.583677	607.860477	6280.777813	2.449995e- 12
2	3983.521212	0.173836	0.108295	0.759076	268	8605	[255, 255, 255, 255, 255, 255, 248, 138, 162,		55, 255, 248, 138, 162,	[255, 255, 255, 255, 255, 25 138,	5, 248, [255, 255, 25 162,	5, 255, 255, 255, 248, 138, 162,	[255, 255, 255, 255, 255	i, 255, 248, 38, 162,	8190.023413	-364.009948	-1223.202923	52.739501	2.449995e- 12
3	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, 1	94, 167, 131, 110, 128	[58, 58, 90, 142, 176, 194, 16 110	7, 131, [58, 58, 90, 14 , 128	2, 176, 194, 167, 131, 110, 128		, 167, 131, 110, 128	1002.704694	-3387.748787	5605.696996	-2161.662287	2.449995e- 12
4	1838.646970	0.163560	0.083060	0.869798	126	8957	[255, 255, 255, 255, 255, 255, 255, 255,		55, 255, 255, 255, 255,	[255, 255, 255, 255, 255, 25 255,	5, 255, [255, 255, 25 255,	5, 255, 255, 255, 255, 255, 255,	[255, 255, 255, 255, 255, 255, 255]	i, 255, 255, 255, 255,	1758.679749	6979.486963	-91.632778	-2511.386572	2.449995e- 12
	mbined_female_with_pcal= co mbined_female_with_pcal	mbined_female_with	_pca.drop(["LBP_Values",	, "Chain_Code_Values	"], axis=1)													↑ ↓ ⊝	ធ¢ឦ៖
lara	ick_Contrast Haralick_Homo	geneity Haralick_E	nergy Haralick_Correlat	tion Box_Counting_V	alues Area_Values	Femal	e_Gabor_Pixel_Values_1 Fema	le_Gabor_Pixel_Value	es_2 Female	_Gabor_Pixel_Values_3	Female_Gabor_Pixel_	Values_4 Female_	Gabor_Pixel_Values_5	Female_PCA	_1 Female_	PCA_2 Femal	e_PCA_3 Fer	nale_PCA_4 F	emale_PCA_5
	1360.080505	0.153852 0.0	051972 0.89	7294	113 9291	[25,	26, 29, 34, 37, 95, 37, 33, 35, [2 43, 55, 7	5, 26, 29, 34, 37, 35, 37, 33 43, 55	, 35, [25, 2 , 7	6, 29, 34, 37, 35, 37, 33, 35, 43, 55, 7	[25, 26, 29, 34, 37, 35,	37, 33, 35, [25, 26 43, 55, 7	i, 29, 34, 37, 35, 37, 33, 35, 43, 55, 7	7632.6217	719 3347.0	81673 10	5.123081	383.509980	2.110306e-12
	1572.307172	0.136395 0.0	054128 0.880	0509	145 9276	[255,	255, 255, 255, 255, 255, 255, [25	5, 255, 255, 255, 255, 255,	255, [255, 2	55, 255, 255, 255, 255, 255,	[255, 255, 255, 255, 255	, 255, 255, [255, 25	5, 255, 255, 255, 255, 255,	1293.9142	279 -4452.1	93821 -151	9.473788 -4	1030.048762	2.110306e-12

255, 255, .

157, 186, ...

 $[225, 222, 226, 225, 247, 242, 227, \\ [225, 222, 226, 225, 247, 242, 227, \\$

255, 255, ...

157, 186, ...

255, 255, ...

255, 255, .

157, 186, ...

[255, 255, 255, 255, 255, 255, 255,

 $[255, 255, 244, 188, 149, 127, 136, \\ [255, 255, 244, 188, 149, \\ [255, 255, 244, 188, 149, \\ [255, 255, 244, 188, 149, \\ [255, 255, 244, 188, 149, \\ [255, 255, 244, 188, 149, \\ [255, 255, 244, 188, 149, \\ [255, 255, 244, 188,]]$

255, 255, .

206, 210, ...

157, 186, ...

206, 210, ...

255, 255, ...

-929.661705

-3548,739389

5892.751801

-1804.316812

[225, 222, 226, 225, 247, 242, 227, [225, 222, 226, 225, 247, 242, 227, [225, 222, 226, 225, 247, 242, 227,

Taking row 0 as the main row to compare with others

<mark>Male</mark>

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 H	aralick_Homogeneity H	aralick_Energy Har	alick_Correlation Box_C	Counting_Values A	rea_Values	LBP_Values	s Chain_Code_Val
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, 0.0805, 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, 0.9088, 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
	Threshold 0.91 0.95 0.99		Similarity row1, row2, row3, row4 row1,row2,row3,row4 row2 None			Dissimilarity None None row1, row3, row4 row1,row2,row3,row4			
				None		row1,rov	vz,row3	,10W4	
	r Harali			k_Energy Haralick_(Correlation Box_Counting	_Values Area_Valu	ies	LBP_Values	Chain_Code_Values
Fema	r Harali e	1360.080505	0.153852	k_Energy Haralick_0	0.897294	_Values Area_Valu	ues 291 [0.0542,	LBP_Values 8.0396, 0.0213, 0.0123, 0.01, 0.0096, [[(-1, 1), ((0, 1), (-1, 1), (-1, 0), (1, 0), (
ema ema	r Harali de	1360.080505 1572.307172	0.153852 0.136395	k_Energy Haralick_0 0.051972 0.054128	0.897294 0.880509	113 9.	291 [0.0542, 276 [0.0519,	LBP_Values 8.0396, 0.0213, 0.0123, 0.01, 0.0096, [[(-1, 1), (0.00418, 0.0266, 0.0134, 0.0104, 0.007 [[(1, 1), (1.00418, 0.0266, 0.0134, 0.0104, 0.007 [[(1, 1), (1.00418, 0.0266, 0.0134, 0.0104, 0.007 [[(1, 1), (1.00418, 0.0266, 0.0134, 0.0104, 0.007 [[(1, 1), (1.00418, 0.0266, 0.0134, 0.0104, 0.007 [[(1, 1), (1.00418, 0.00418, 0.00418, 0.00418, 0.00418]]]	(0, 1), (-1, 1), (-1, 0), (1, 0), (1, 0), (1, 1), (1, 0), (1, 0), (1,
Fema Fema Fema	r Harali de de	1360.080505 1572.307172 1117.535253	0.153852 0.136395 0.147987	k_Energy Haralick_0 0.051972 0.054128 0.052652	0.897294 0.880509 0.918772	113 9. 145 9. 79 9.	291 [0.0542, 276 [0.0519, 382 [0.0422,	LBP_Values 8.0396, 0.0213, 0.0123, 0.01, 0.0096, [[(-1, 1), (0.00418, 0.0266, 0.0134, 0.0104, 0.007 [[(1, 1), (1.00385, 0.0213, 0.0152, 0.0127, 0.008 [[], [], [],	(0, 1), (-1, 1), (-1, 0), (1, 0), (1, 0), (1, 1), (1, 0), (1, 0), (1, , [(1, 0), (1, 0), (1, 0), (1, 1),
Fema Fema Fema Fema	r Harali de de de	1360.080505 1572.307172 1117.535253 2481.296263	0.153852 0.136395 0.147987 0.143295	k_Energy Haralick_0 0.051972 0.054128 0.052652 0.073946	0.897294 0.880509 0.918772 0.832857	113 9. 145 9. 169 9.	291 [0.0542, 276 [0.0519, 382 [0.0422, 055 [0.0523,	LBP_Values 8.0396, 0.0213, 0.0123, 0.01, 0.0096, [[(-1, 1), (0.00418, 0.0266, 0.0134, 0.0104, 0.007 [[(1, 1), (1.00385, 0.0213, 0.0152, 0.0127, 0.008 [[], [], [], [], (0.0424, 0.0256, 0.011, 0.0087, 0.0063 [[(0, 1), (0.0087, 0.0063 [[(0, 1), (0.0087, 0.0063])]]	(0, 1), (-1, 1), (-1, 0), (1, 0), (1, 0), (1, 1), (1, 0), (1, 0), (1, , [(1, 0), (1, 0), (1, 0), (1, 1), 0, 1), (0, 1), (0, -1), (0, -1)], [
Fema Fema Fema	r Harali de de de	1360.080505 1572.307172 1117.535253	0.153852 0.136395 0.147987	k_Energy Haralick_0 0.051972 0.054128 0.052652	0.897294 0.880509 0.918772	113 9. 145 9. 169 9.	291 [0.0542, 276 [0.0519, 382 [0.0422, 055 [0.0523,	LBP_Values 8.0396, 0.0213, 0.0123, 0.01, 0.0096, [[(-1, 1), (0.00418, 0.0266, 0.0134, 0.0104, 0.007 [[(1, 1), (1.00385, 0.0213, 0.0152, 0.0127, 0.008 [[], [], [],	(0, 1), (-1, 1), (-1, 0), (1, 0), (1, 0), (1, 1), (1, 0), (1, 0), (1, , [(1, 0), (1, 0), (1, 0), (1, 1), 0, 1), (0, 1), (0, -1), (0, -1)], [