## A Novel Multiresolution-Statistical Texture Analysis Architecture: Radiomics-Aided Diagnosis of PDAC Based on Plain CT Images

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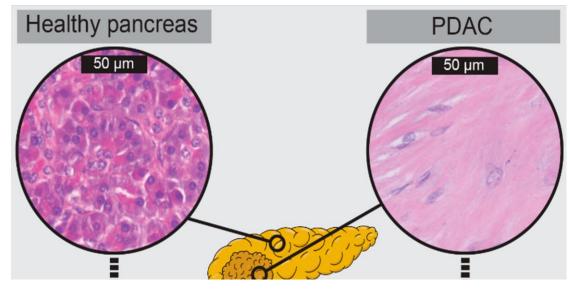
#### Outline

- Introduction to the problem
- Methodology
- Results
- Implementation details and results

## INTRODUCTION

#### **Problem Statement**

- PDAC Pancreatic ductal adenocarcinoma Type of malignant tumor
- CT imaging is a frequently used non-invasive examination method for PDAC
  - Plain CT
  - Contrast enhanced CT
- Limitations:
  - Subjective judgement
  - No access contrast enhancement
  - Allergic reactions or renal toxicity caused by contrast agent



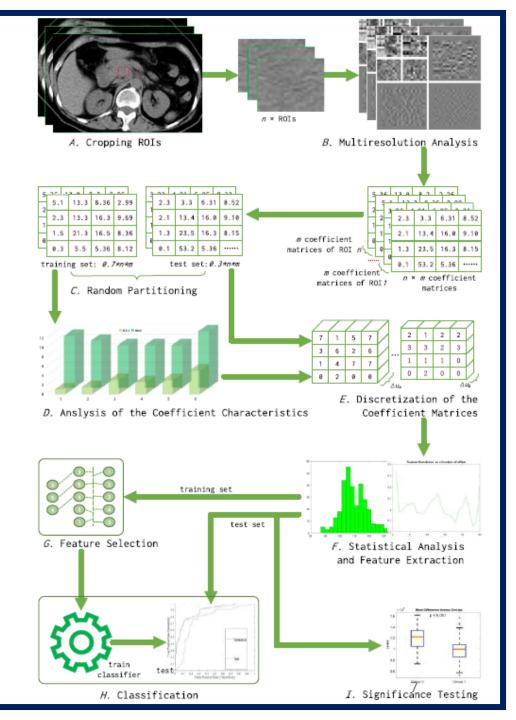
### Radiomics and Al

- Quantitative features of radiomics can provide interpretability.
- Radiomics + AI plays important role in early diagnosis of PDAC based on plain CT
- Radiomics Studies in previous works (pancreatic tumor related)
  - LoG (Laplacian of Gaussian), Histogram features
  - GLCM (Gray-level co-occurrence matrix), GLRLM (Gray-level run-length matrix), ACM (Angle co-occurrence matrix)
  - Wavelet transform features

## **METHODOLOGY**

#### Overview

- ✓ Cropping ROIs
- ✓ Multiresolution Analysis :
  - ✓ Wavelet transform
  - ✓ Wavelet packet transform
  - ✓ Contourlet transform
- ✓ Random Partitioning
- ✓ Analysis of Coefficient Characteristics
- ✓ Discretization of the coefficient matrices
- √ Statistical Analysis and Feature extraction
- √ Feature selection
- ✓ Classification and Significance testing



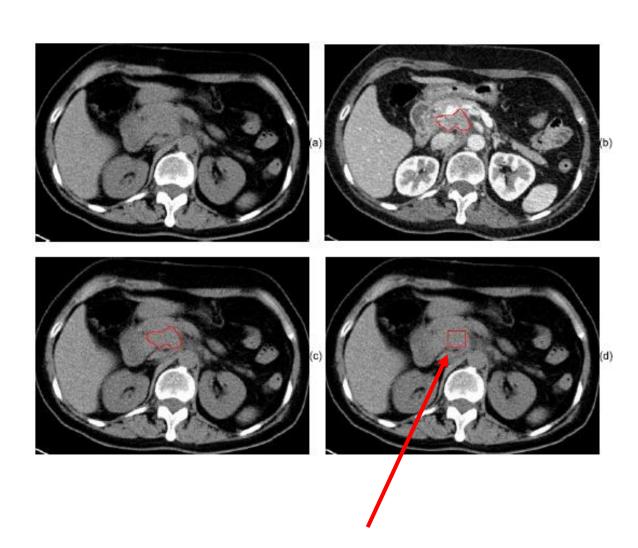
#### **Dataset**

- Subjects
  - 153 PDAC
    - 92 male; 61 female
    - Age: 30-84 (59.1y)
  - 159 HP
    - 104 male; 55 female
    - Age: 40 57 (45.9y)
- Random partition
  - Train set : Test set = 7:3

- Data acquisition setup
  - Plain CT slices (2 or 5mm thickness)
  - Window level: 40-50 HU
  - Window width: 300-350 HU
  - Enhanced CT slices may used as reference
  - Scanner: Siemens SOMATOM
     Definition AS+/Flash or Philips
     Brilliance 64 CT scanners.

### **Cropping ROIs**

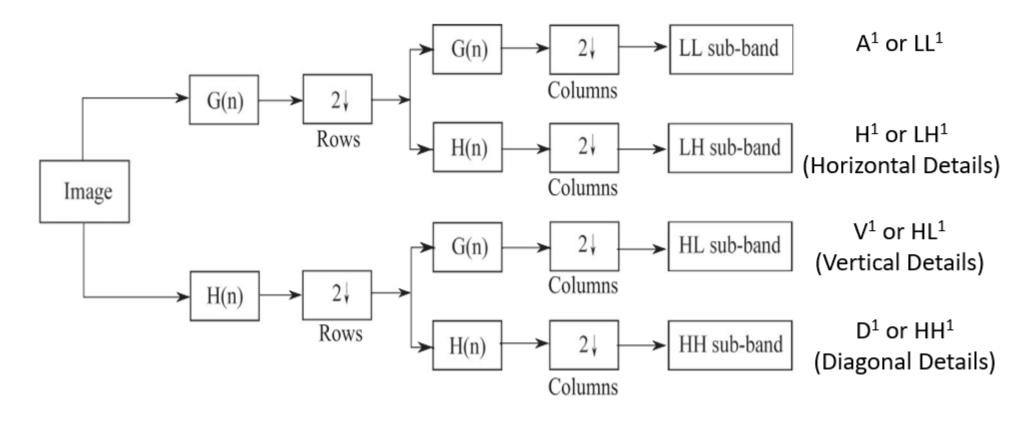
- Cropping strategy
  - PDAC Slice with largest cross-sectional area of tumor
  - HP Slice with largest crosssectional area of HP.
- ROI acquisition
  - By 3 senior radiologists (>6y experience)



### Multiresolution Analysis

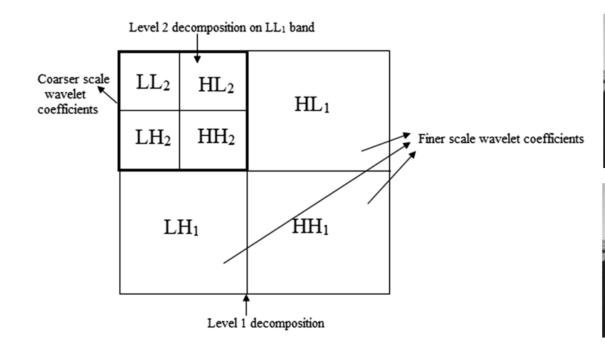
Wavelet transform, Wavelet packet transform and Contourlet transforms

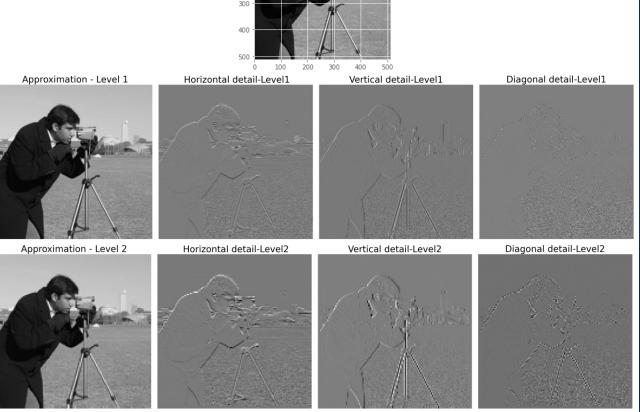
- Decompose an ROI into multiple sub-band components.



## Multiresolution Analysis

- Wavelet Name: bior1.3
- 2 level decomposition





original image

## Analysis of Coefficient Characteristics (Based on trainset)

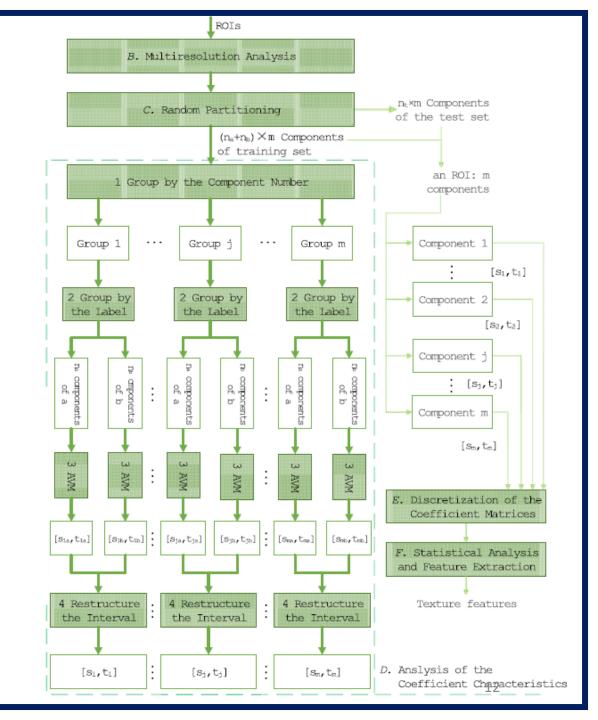
- Group by component number
- Group by label
- Find averages of min and max
- Structure the interval

$$x_{ji} = \min \left( C_{ji} = \begin{bmatrix} c_{11} & \cdots & c_{1l} \\ c_{21} & \ddots & \vdots \\ \cdots & \cdots & c_{kl} \end{bmatrix} \right) \qquad s_{ja} = \frac{\sum_{i=1}^{n_a} x_{ji}}{n_a}, \quad s_{jb} = \frac{\sum_{i=1}^{n_b} x_{ji}}{n_b}$$

$$y_{ji} = \max \left( C_{ji} = \begin{bmatrix} c_{11} & \cdots & c_{1l} \\ c_{21} & \ddots & \vdots \\ \cdots & \cdots & c_{kl} \end{bmatrix} \right) \qquad t_{ja} = \frac{\sum_{i=1}^{n_a} y_{ji}}{n_a}, \quad t_{jb} = \frac{\sum_{i=1}^{n_b} y_{ji}}{n_b}$$

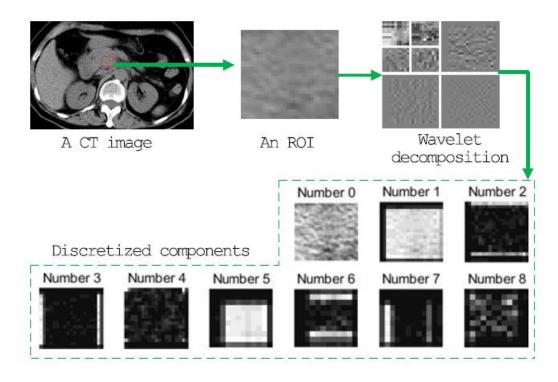
$$s_j = \min \left( s_{ja}, s_{jb} \right)$$

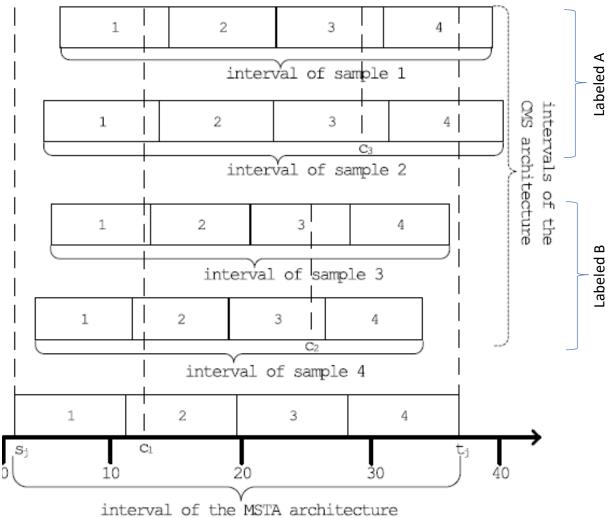
$$t_j = \max \left( t_{ja}, t_{jb} \right)$$



# Discretization of the coefficient matrices

• Number of bins = 8/16/24





## Statistical Analysis and Feature extraction

Coefficient statistics (3)	Histograms (7)	Gray-level cooccurrence matrix (5)	Gray level run-length matrix (11)
Mean	Mean	Contrast	Short-run emphasis
Variance	Standard deviation	Correlation	Long-run emphasis
Average Energy	Smoothness	Energy	Gray-level non-uniformity
	Cubic moment	Homogeneity	Run-length non-uniformity
	Uniformity	Entropy	Run percentage
	Entropy		Low gray-level run emphasis
	Fourth moment		High gray-level run emphasis
			Short run low gray-level emphasis
			Short run high gray-level emphasis
			Long run low gray-level emphasis
			Long run high gray-level emphasis

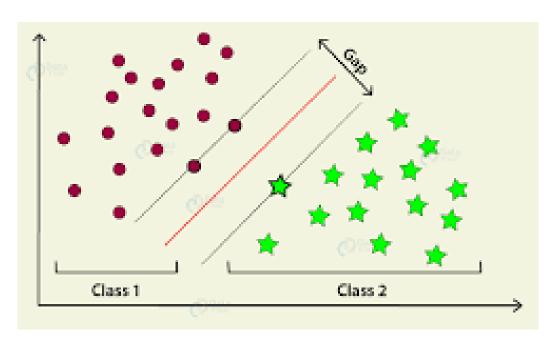
#### Feature selection

- ILFS (Infinite Latent Feature Selection) algorithm proposed by G.Roffo et al.
- Ranking of all the features based on trainset and selected the first k features.
  - K < 20 (empirically limited)



## Classification and Significance testing

- Experimented with multiple ML methods and selected SVM (with linear kernel).
  - Outstanding performance with small sample size
  - Widely used in radiomics studies
  - Interpretability
  - Generalization
- 10-fold cross validation
- Mann-Whitney U tests
  - Non-parametric test



## **RESULTS**

### Classification - Results

#### • On test set

**TABLE IV** 

TEST RESULTS OF THE CLASSIFICATIONS: ACC (%), ACCURACY; SEN (%), SENSITIVITY; SPE (%), SPECIFICITY. THE BOLD AND ITALIC TEXTS HIGHLIGHT THE HIGHEST PERFORMANCES OBTAINED FOR AN INDICATOR

No.	Method	ACC	SEN	SPE	AUC	No.	Method	ACC	SEN	SPE	AUC
1	GLS	74.47	73.91	75	0.726	6	ACM	62.77	56.52	68.75	0.671
2	GLH	70.21	63.04	77.08	0.772	7	LOG	70.21	67.39	72.92	0.693
3	GLCM	61.7	47.83	75	0.729	8	WT	71.28	69.57	72.92	0.715
4	GLRLM	72.34	69.57	75	0.767	9	WPT	72.34	71.74	72.92	0.701
5	GL-M	73.4	65.22	81.25	0.744	10	CT	69.15	67.39	70.83	0.727
11	CMS-W-CS	71.28	69.57	72.92	0.747	16	MSTA-W-CS	74.47	71.74	77.08	0.743
12	CMS-W-H	73.4	71.74	75	0.749	17	MSTA-W-H	73.4	69.57	77.08	0.746
13	CMS-W-COM	63.83	54.35	72.92	0.726	18	MSTA-W-COM	71.28	65.22	77.08	0.753
14	CMS-W-RLM	72.34	69.57	75	0.743	19	MSTA-W-RLM	77.66	78.26	77.08	0.778
15	CMS-W-M	73.4	71.74	75	0.752	20	MSTA-W-M	77.66	78.26	77.08	0.792
21	CMS-WP-CS	73.4	71.74	75	0.738	26	MSTA-WP-CS	72.34	73.91	70.83	0.73
22	CMS-WP-H	73.4	76.09	70.83	0.744	27	MSTA-WP-H	74.47	76.09	72.92	0.747
23	CMS-WP-COM	51.06	0	100	0.5	28	MSTA-WP-COM	68 09	56 52	79.17	0.757
24	CMS-WP-RLM	72.34	69.57	75	0.753	29	MSTA-WP- RLM	77.66	78.26	77.08	0.787
25	CMS-WP-M	75.53	73.91	77.08	0.791	30	MSTA-WP-M	/5.55	80.43	/0.83	0.765
31	CMS-C-CS	61.7	47.83	75	0.647	36	MSTA-C-CS	71.28	71.74	70.83	0.768
32	CMS-C-H	71.28	73.91	68.75	0.781	37	MSTA-C-H	71.28	67.39	75	0.769
33	CMS-C-COM	63.83	54.35	72.92	0.703	38	MSTA-C-COM	65 96	52 17	79.17	0.742
34	CMS-C-RLM	72.34	69.57	75	0.752	39	MSTA-C-RLM	77.66	78.26	77.08	0.777
35	CMS-C-M	69.15	65.22	72.92	0.771	40	MSTA-C-M	75.53	71.74	79.17	0.781

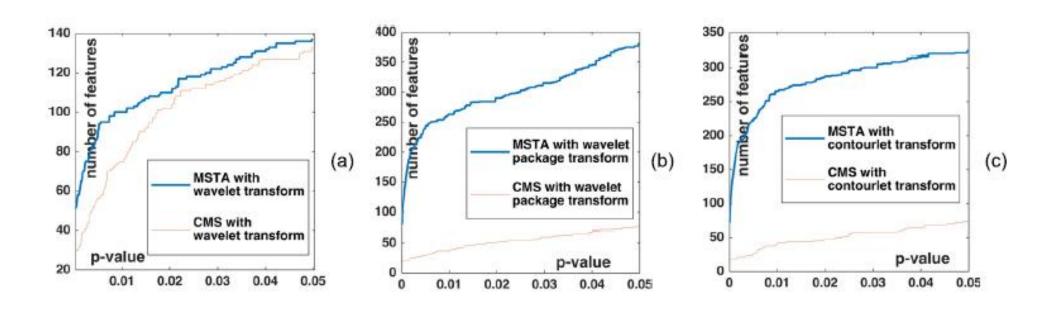
### Classification - Results

TEXTURE ANALYSIS	1.1	1.2	1.3	1.4	1.5	1.6	1.7	1.8	1.9	1.10	1.11	1.12	1.13	1.14	1.15	1.16
TEXTURE ANALYSIS	1.17	1.18	1.19	1.20	1.21	1.22	1.23	1.24	1.25							
MOTA W. CC	64.7	64.7	70.2	72.9	68.3	72.0	73.4	71.6	72.9	69.7	64.2	61.0	73.4	72.0	67.0	66.1
MSTA-W-CS	71.6	70.2	64.7	64.7	73.9	70.2	72.0	65.1	67.9							
MCTA WII	70.6	70.6	72.5	70.2	67.4	70.6	75.7	72.5	75.2	70.6	65.1	55.0	73.4	74.3	62.4	70.2
MSTA-W-H	75.2	70.2	67.9	69.7	72.9	72.5	74.3	72.0	71.6							
MSTA W COM	74.8	74.8	73.4	70.6	Failed	73.4	69.3	69.3	73.4	78.0	72.9	70.6	75.2	72.9	69.3	70.6
MSTA-W-COM	70.2	72.9	71.1	72.0	76.1	73.9	71.1	67.4	75.2							
MCTA W DI M	69.3	70.2	75.7	75.7	70.2	71.6	75.7	73.4	75.7	72.0	63.3	72.5	75.2	75.2	70.6	72.0
MSTA-W-RLM	71.6	74.3	72.9	73.9	73.9	74.3	78.4	74.3	72.5							
MCTA W M	75.2	75.2	71.1	75.7	Failed	76.6	77.5	76.6	79.8	76.1	68.3	74.3	76.6	76.1	67.4	75.2
MSTA-W-M	75.2	74.8	75.7	76.1	74.3	78.9	75.7	67.4	74.3							

<sup>1.1</sup> fine tree; 1.2 medium tree; 1.3 coarse tree; 1.4 linear discriminant; 1.5 quadratic discriminant; 1.6 logistic regression; 1.7 Gaussian naïve Bayes; 1.8 kernel naïve Bayes; 1.9 linear SVM; 1.10 quadratic SVM; 1.11 cubic SVM; 1.12 fine Gaussian SVM; 1.13 medium Gaussian SVM; 1.14 coarse Gaussian SVM; 1.15 fine KNN; 1.16 medium KNN; 1.17 coarse KNN; 1.18 cosine KNN; 1.19 cubic KNN: 1.20 weighted KNN; 1.21 ensemble boosted trees; 1.22 ensemble bagged trees; 1.23 ensemble subspace discriminant; 1.24 ensemble subspace KNN; 1.25 ensemble RUSBoosted tree.

### Significance Tests - Results

- Statistical tests were performed based on test set.
- Number of features with statistically significant difference (MSTA vs CMS)



### Significance Tests - Results

- x: median for PDAC; y: median for HP.
- The alternative hypothesis of a left-tailed test states that x < y</li>
- The alternative hypothesis of a right-tailed test states that x > y.

LEFT-TAILED AND RIGHT-TAILED TESTS: DIFFERENCE IN THE TEXTURE
FEATURE VALUES FOR PDACS AND HPS; A, H, V, AND D ARE
APPROXIMATE, HORIZONTAL, VERTICAL, AND DIAGONAL
DIRECTIONS, RESPECTIVELY

No.	Texture analysis method	Component	Feature name	$\begin{array}{c} Two\text{sided} \\ p \leq \end{array}$	Left-tailed p≤	Right-tailed p≤
f1	MSTA-W-CS	D in 2-level	variance	6.66×10 <sup>-6</sup>	3.33×10 <sup>-6</sup>	
f2	MSTA-W-CS	D in 2-level	average energy	7.16×10 <sup>-6</sup>	3.58×10 <sup>-6</sup>	
f3	MSTA-W-H	A in 1-level	entropy	0.25×10 <sup>-6</sup>	1.27×10 <sup>-7</sup>	
f4	MSTA-W-H	A in 1-level	uniformity	1.78×10 <sup>-6</sup>		8.92×10 <sup>-7</sup>
f5	MSTA-W-COM	A in 1-level	d=1, homogeneity	1.09×10 <sup>-6</sup>	5.45×10 <sup>7</sup>	
f6	MSTA-W-COM	A in 1-level	d=1, energy	1.78×10 <sup>-6</sup>		8.99×10 <sup>-7</sup>
f7	MSTA-W-RLM	D in 2-level	long-run high level emphasis	2.32×10 <sup>-6</sup>	1.16×10 <sup>-6</sup>	
f8	MSTA-W-RLM	D in 2-level	short-run emphasis	1.39×10 <sup>-6</sup>		6.95×10 <sup>-6</sup>

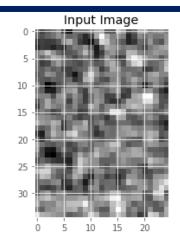


## IMPLEMENTATION

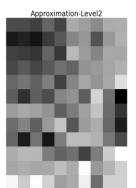
https://github.com/Mithunjha/PDAC Prediction

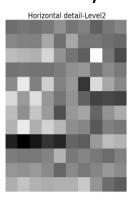


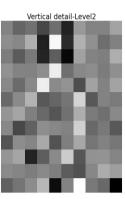
#### 1. Extraction of ROI

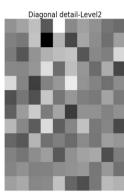


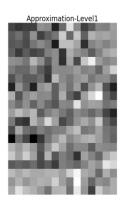
#### 2. Multi resolution analysis : Wavelet Transform

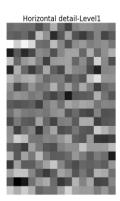


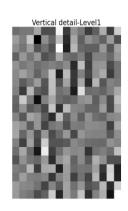


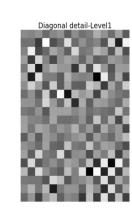












#### 3. Random partitioning

Train set: Test set = 7:3

**Train data** 

Class 1 (PDAC) = 107

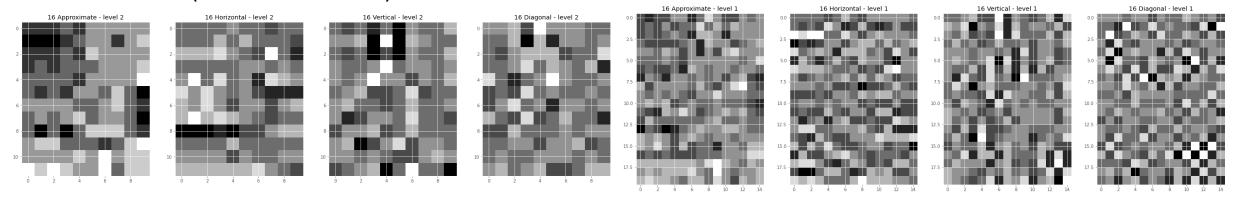
Class 2 (HP) = 111

Test data

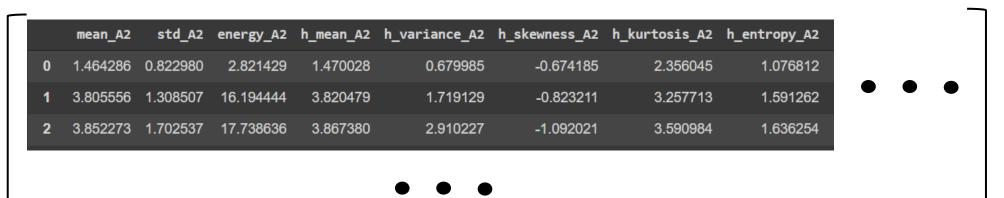
Class 1 (PDAC) = 46

Class 2 (HP) = 48

#### 4. Discretization (Number of bins = 8)



#### 5. Feature Extraction



312 x 208

3 coefficient statistics features, 5 histogram features, 4 gray level co-occurrence matrices features and 11 run length matrices features

#### 6. Feature selection – based on trainset (218)

03.133088e+143.133088e+142.049790e+142.049790e+1414.367321e+144.367321e+144.990584e+144.990584e+1422.121932e+152.612274e+152.612274e+1538.220777e+148.220777e+141.021465e+151.021465e+15		ShortLowGrayLevelEmphasis_A2	LowGrayLevelRunEmphasis_A2	LowGrayLevelRunEmphasis_A1	ShortLowGrayLevelEmphasis_A1
2 2.121932e+15 2.121932e+15 2.612274e+15 2.612274e+15	0	3.133088e+14	3.133088e+14	2.049790e+14	2.049790e+14
2 2.1213326+13 2.1213326+13 2.0122146+13 2.0122146+13	1	4.367321e+14	4.367321e+14	4.990584e+14	4.990584e+14
<b>3</b> 8.220777e+14 8.220777e+14 1.021465e+15 1.021465e+15	2	2.121932e+15	2.121932e+15	2.612274e+15	2.612274e+15
	3	8.220777e+14	8.220777e+14	1.021465e+15	1.021465e+15
<b>4</b> 1.055106e-01 1.189265e-01 1.181645e-01 9.655603e-02	4	1.055106e-01	1.189265e-01	1.181645e-01	9.655603e-02

218 x # features

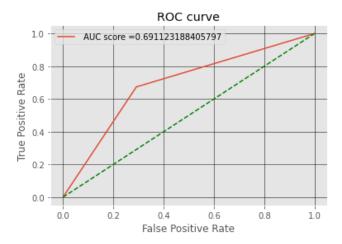
#### 7. Classification and significance testing

Experimented with: Linear SVM, Decision tree, KNN, Poly SVM.

Significance test: Mann U Whitney test Results are reported in following slides.

### Results

No.	Method	ACC	SEN	SPE	AUC
16	MSTA-W-CS	74.47	71.74	77.08	0.743
17	MSTA-W-H	73.4	69.57	77.08	0.746
18	MSTA-W-COM	71.28	65.22	77.08	0.753
19	MSTA-W-RLM	77.66	78.26	77.08	0.778
20	MSTA-W-M	77.66	78.26	77.08	0.792



**ROC curve of SVM - Linear** 

- Multi resolution analysis method : wavelet transform
- Selected number of features = 20; Number of bins = 8
- Model = SVM

Features	Accuracy	Sensitivity	Specificity	AUC	Precision	F1
CS	72.34	73.91	70.83	0.7237	70.83	0.7234
Н	64.89	60.87	68.75	0.6481	65.12	0.6292
СОМ	68.09	67.39	68.75	0.6807	67.39	0.6739
RLM	70.21	65.22	75.0	0.7011	71.43	0.6818
ALL	69.15	67.39	70.83	0.6911	68.88	0.6813

#### Results

- Selected number of features = 20
- Feature extraction = ALL

Train Accuracy: 0.7752293577981652 Test Accuracy: 0.6914893617021277 Precision: 0.6888888888888888 Sensitivity: 0.6739130434782609 Specificity: 0.70833333333333334 f1: 0.6813186813186812

Area Under Curve (AUC): 0.691123188405797

lassificati	lon Repo	rt:					
	prec	ision	recall	f1-9	core	suppo	rt
,		0.60	0.74	_	. 70		
6	,	0.69	0.71	٤	70	4	ಶ
1		0.69	0.67	6	68	4	6
accuracy	/			6	69	9,	4
macro ava	5	0.69	0.69	6	<b>69</b>	94	4
eighted av	5	0.69	0.69	6	<b>69</b>	9	4

Train Accuracy: 0.8302752293577982 Test Accuracy: 0.7021276595744681 Precision: 0.7142857142857143 Sensitivity: 0.6521739130434783 Specificity: 0.75

f1: 0.6818181818181819

Area Under Curve (AUC): 0.7010869565217391

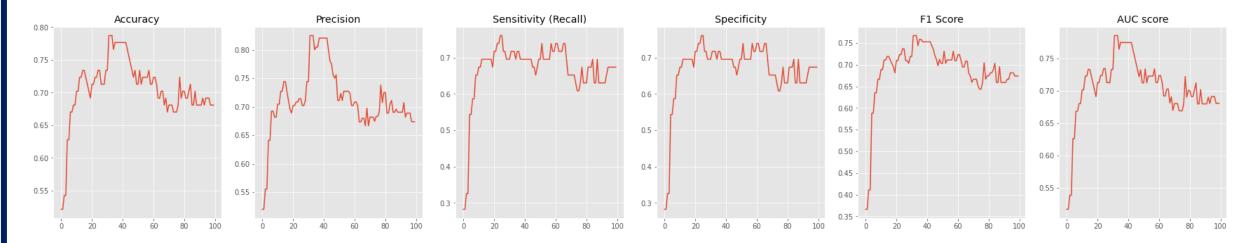
Classification Report : recall f1-score support precision 0.75 0.69 0.72 0.71 0.65 0.68 0.70 accuracy macro avg 0.70 0.70 0.70 94 weighted avg 0.70 0.70 0.70

**Results of Linear SVM** 

**Results of Poly SVM** 

Models	Accuracy	Sensitivity	Specificity	AUC	Precision	F1
Linear SVM	69.15	67.39	70.83	0.6911	68.88	0.6813
Decision Tree	62.77	63.04	62.5	0.6277	61.70	0.6237
KNN	58.51	56.52	60.42	0.5847	57.77	0.5714
Poly SVM	70.21	65.22	75	0.7011	71.43	0.6818

#### • Variance of performance metrics along with number of features



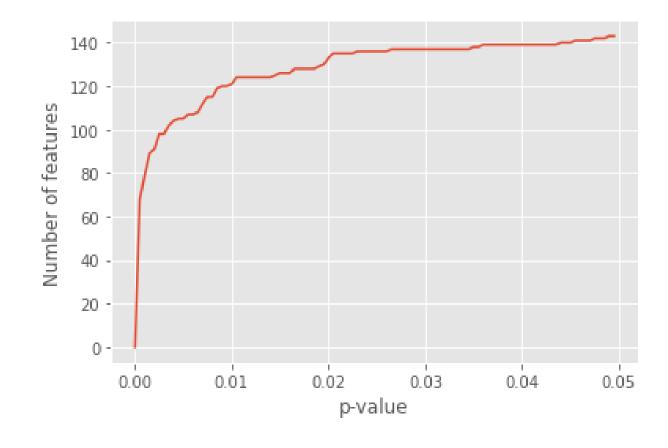
#### • Selected number of features = 32; Feature extraction = ALL

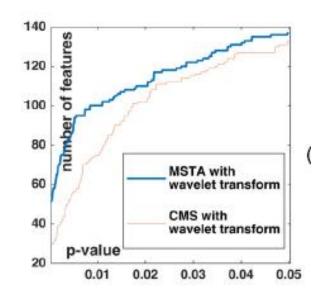
Models	Accuracy	Sensitivity	Specificity	AUC	Precision	F1
Linear SVM	78.72	71.74	71.74	0.7858	82.5	0.7674

No.	Method	ACC	SEN	SPE	AUC
16	MSTA-W-CS	74.47	71.74	77.08	0.743
17	MSTA-W-H	73.4	69.57	77.08	0.746
18	MSTA-W-COM	71.28	65.22	77.08	0.753
19	MSTA-W-RLM	77.66	78.26	77.08	0.778
20	MSTA-W-M	77.66	78.26	77.08	0.792

## Statistical Testing

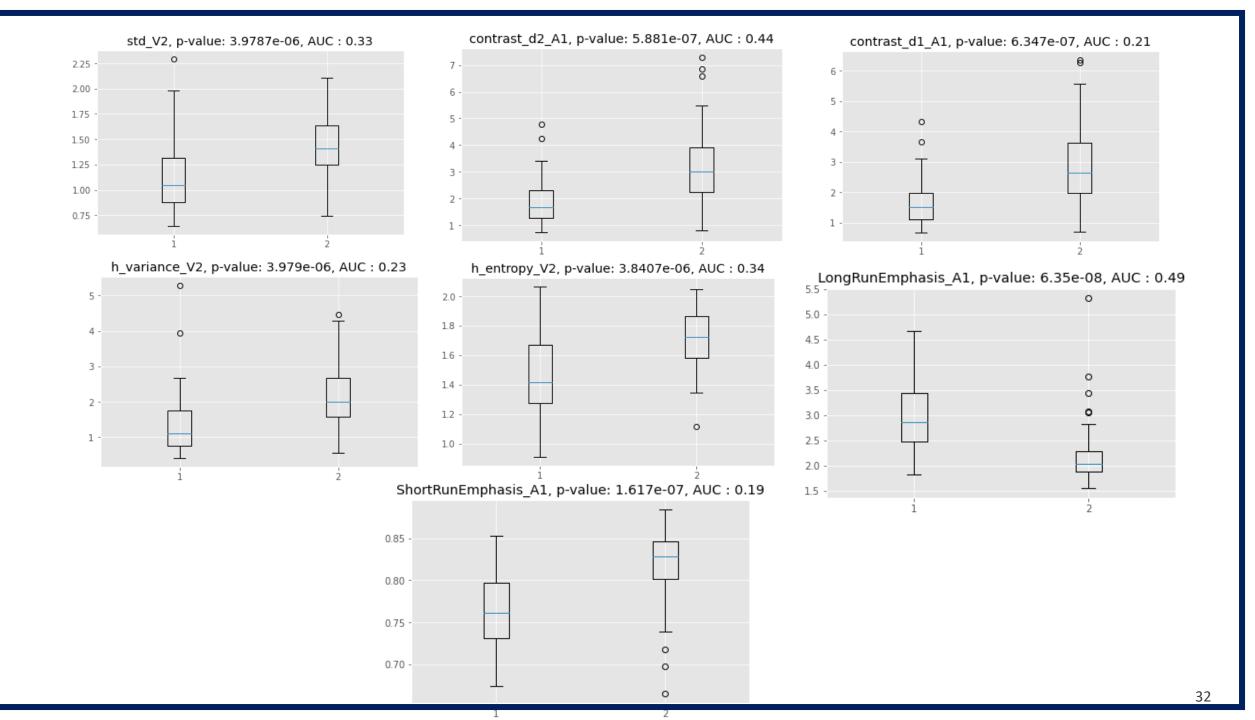
• Number of significant features (for different p-value)





## LEFT-TAILED AND RIGHT-TAILED TESTS: DIFFERENCE IN THE TEXTURE FEATURE VALUES FOR PDACs AND HPs;

Feature name	Statistical	Component	Two sided	Less PDAC <hp< th=""><th>Greater PDAC&gt;HP</th></hp<>	Greater PDAC>HP
Std_V2	CS - STD	Vertical – level 2	7.95e-06	3.97e-06	-
Mean_V1	CS – Mean	Vertical – level 1	2.92e-05	-	-
H_entropy_V2	H - Entropy	Vertical – level 2	7.68e-06	3.84e-06	-
H_variance_V2	H - variance	Vertical – level 2	7.96e-06	3.98e-06	-
Contrast_d2_A1	Contrast – Distance 2	Approximate – level 1	1.18e-06	5.88e-07	-
Contrast_d1_A1	Contrast – Distance 1	Approximate – level 1	1.27e-06	6.35e-07	-
LongRunEmphasis_A1	RLM – Long Run Emphasis	Approximate – level 1	1.27e-07	-	6.35e-08
ShortRunEmphasis_A1	RLM – Short Run Emphasis	Approximate – level 1	3.23e-07	1.62e-07	-



## THANK YOU

