

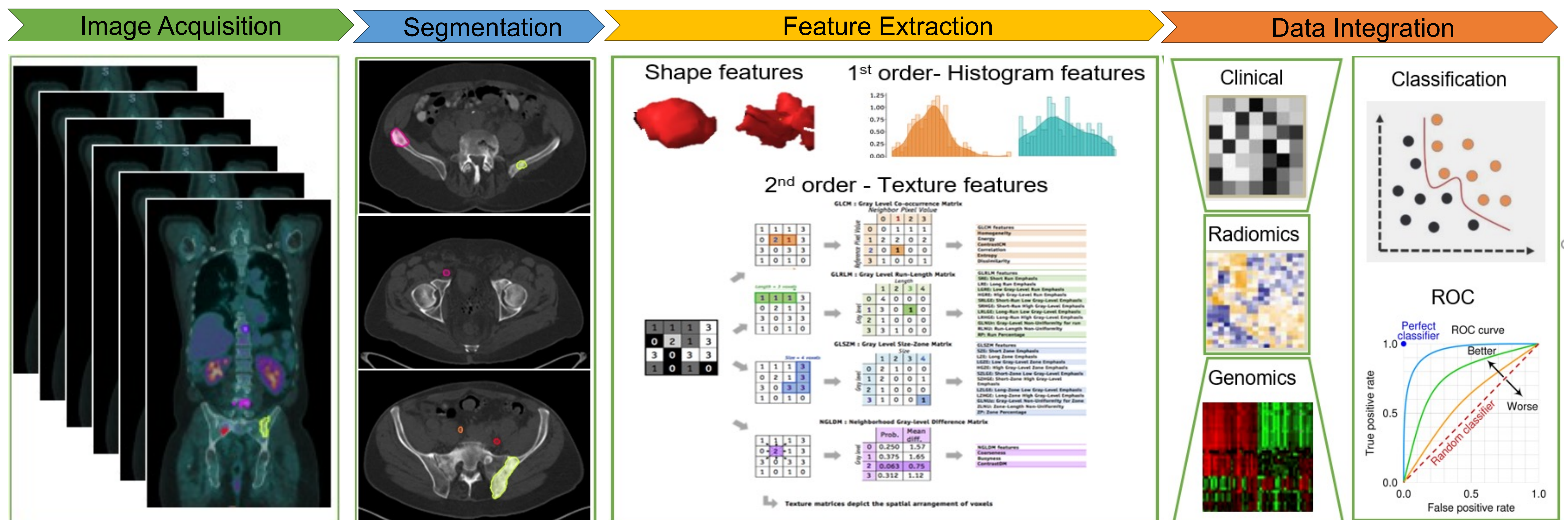
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# Introduction

- ❖ PSMA PET imaging is a sensitive modality for detection of early metastases in prostate cancer (PC)
- ❖ Optimal integration of PSMA scans in routine follow-up is unknown
- ❖ Hypothesis: Multi-omics (Radiomics + genomics + clinical details) could predict PSMA avidity in lesions seen on CT imaging

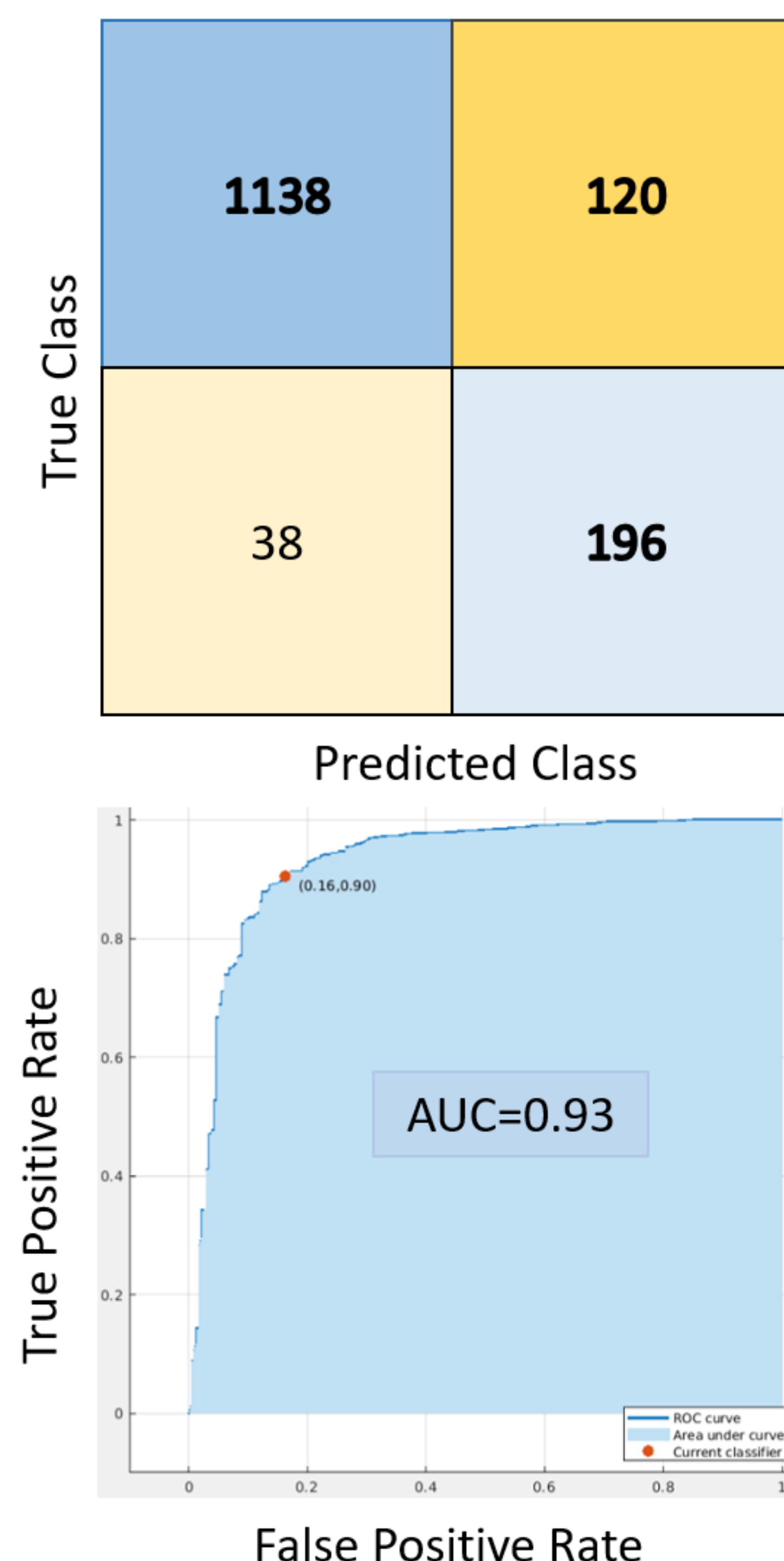
## Material and Methods

- ❖ Oligometastatic castrate-sensitive PC patients with PSMA PET, genomic mutational analysis and clinical details (n=84)
- ❖ High-risk mutations was defined as pathogenic mutations in ATM, BRCA1/2, Rb1 or TP53)
- ❖ Both PSMA avid and non-avid lesions in lymph node and bone were contoured on CT images.
- ❖ Radiomic features were extracted for all lesions using PyRadiomics.
- ❖ ML algorithms [support vector machine (SVM) and decision trees] were used to predict PSMA avidity in CT anatomic correlates
- ❖ Five-fold cross validation



## Results

Patient and Lesion Characteristics	
Total Number of patients	84
Total Lesions	1492
• PSMA + (avid)	• 234
• PSMA - (non-avid)	• 1258
LN Lesions	1016
• PSMA + (avid)	• 164 (16.14%)
• PSMA - (non-avid)	• 852
Bone Lesions	332
• PSMA + (avid)	• 47 (14.2%)
• PSMA - (non-avid)	• 285
Lesion per pt (Median / IQR)	2(1 - 3)
Low volume disease	96.4%
High Risk Mutations	34.5%
• TP53	• 20.2%
• PTEN	• 14.3%
• BRCA 2	• 8.3%
• ATM	• 4.8%
• BRCA 1	• 2.4%
• CHEK2	• 2.4%
• ATRX	• 2.4%
• RB1	• 1.3%



Best Predictive Model	
Top 5 Features	<ul style="list-style-type: none"> <li>• Site of lesion</li> <li>• Volume of lesion</li> <li>• Total number of lesions seen on CT</li> <li>• Gray level non-uniformity</li> <li>• High Risk mutation</li> </ul>
Other Features	<ul style="list-style-type: none"> <li>• Large area Gray level emphasis</li> <li>• Coarseness</li> <li>• Non-uniformity</li> <li>• CHEK2 mutation</li> <li>• FANCD2 mutation</li> <li>• Pre-metastatic PSA.</li> </ul>
Sensitivity	<ul style="list-style-type: none"> <li>• <b>83.7%</b></li> <li>• 196 of 234 lesions</li> </ul>
Specificity	<ul style="list-style-type: none"> <li>• <b>90.4%</b></li> <li>• 1138 of 1258 lesions</li> </ul>
AUC	<ul style="list-style-type: none"> <li>• <b>0.88 – 0.93</b></li> </ul>

## Conclusions

- ❖ Multi-omics including radiomics, genomics and clinical factors can predict PSMA avidity in early metastatic lesions in LN and bones on CT imaging in oligometastatic castrate-sensitive Prostate Cancer
- ❖ This approach can be applied for better personalization and decision-making regarding PSMA surveillance imaging.
- ❖ This multi-omics model needs to be further validated in a larger independent cohort.