

# Multiclass Classification and Semantic Segmentation of Colorectal Cancer Cells from Histopathology Images

Advanced Image Analysis | Machine Learning | Deep Learning  
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By

Mahdi Islam

Musarrat Tabassum

Sirada Kittipaisarnkul

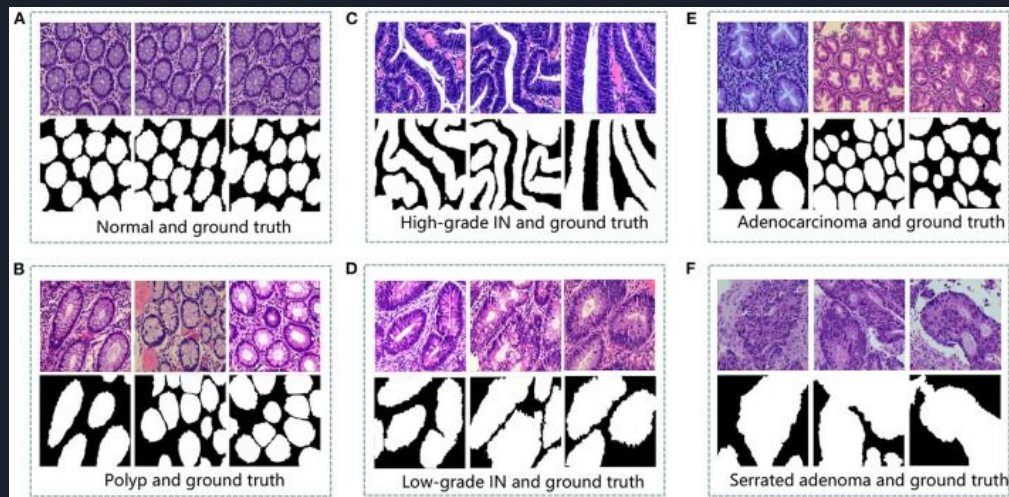
Sarita Mourya

Tuhinangshu Gangopadhyay



# Dataset

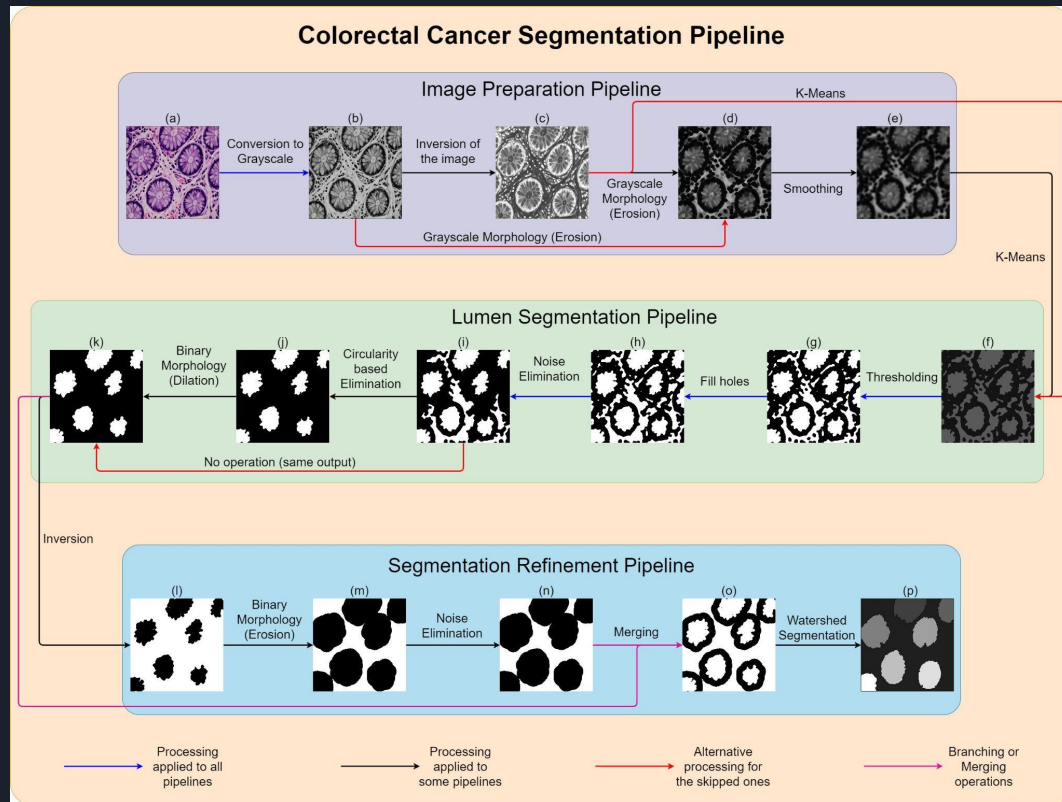
- Publicly available EBHI-Seg Dataset containing 2,228 colorectal histopathology images and their corresponding images.
- This dataset has the following classes:
  - Normal: 76 images
  - Polyp: 474 images
  - Low-grade IN: 637 images
  - High-grade IN: 186 images
  - Adenocarcinoma: 795 images
  - Serrated Adenoma: 58 images



# Image Processing based Segmentation

The general image Processing based Segmentation pipeline has the following steps involved:

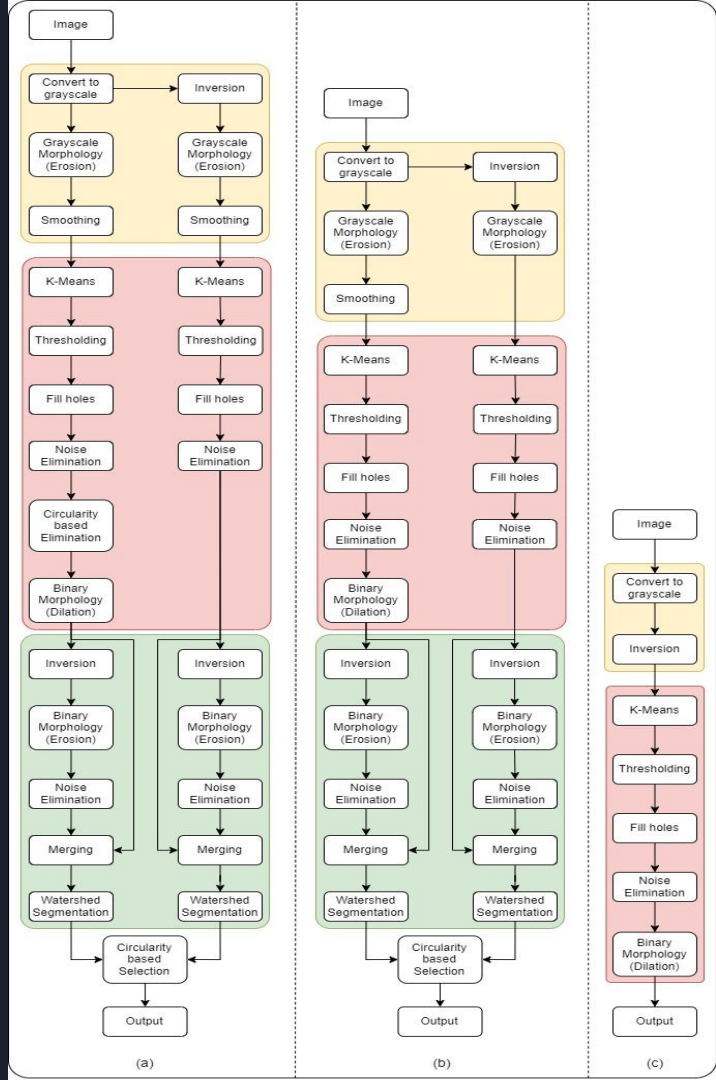
- Image Preparation Pipeline:
  - Conversion to Gray-scale
  - Inversion of Gray-scale image
  - Grayscale Morphology (Erosion)
  - Smoothing



# Image Processing based Segmentation (continued)

- Lumen Segmentation Pipeline:
  - K-Means
  - Thresholding
  - Filling Holes
  - Noise Elimination
  - Circularity based Elimination
  - Binary Morphology (Dilation)
- Segmentation Refinement Pipeline
  - Inversion
  - Binary Morphology (Erosion)
  - Noise Elimination
  - Merging
  - Watershed Segmentation

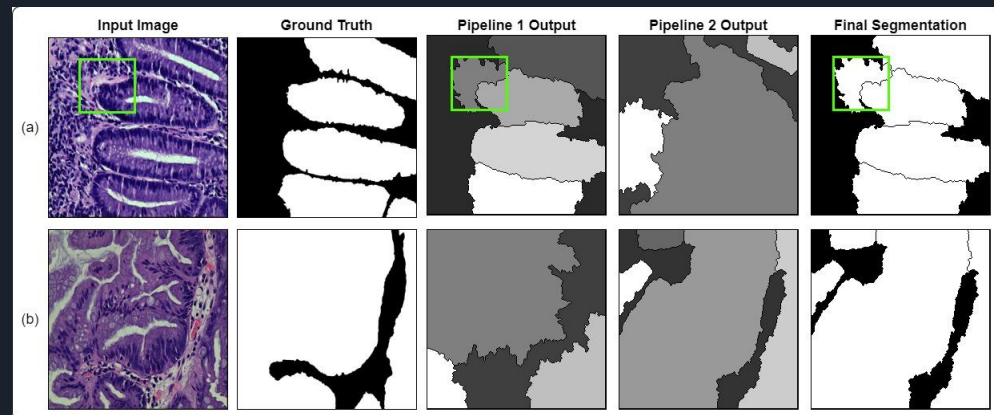
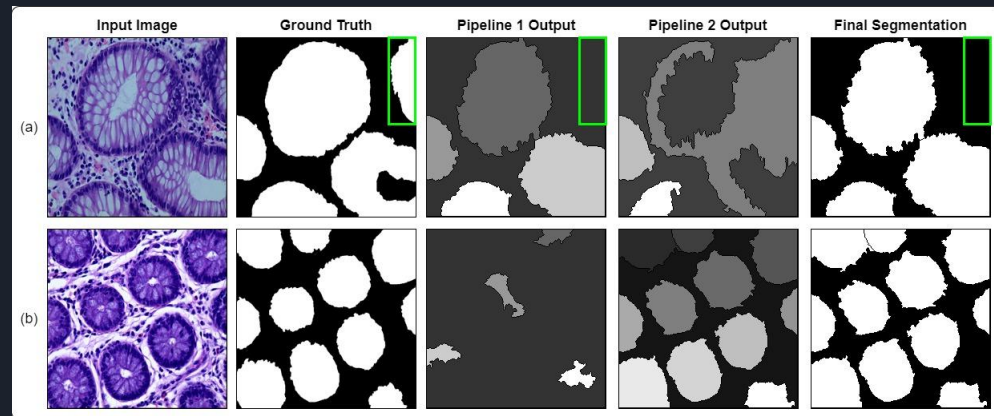
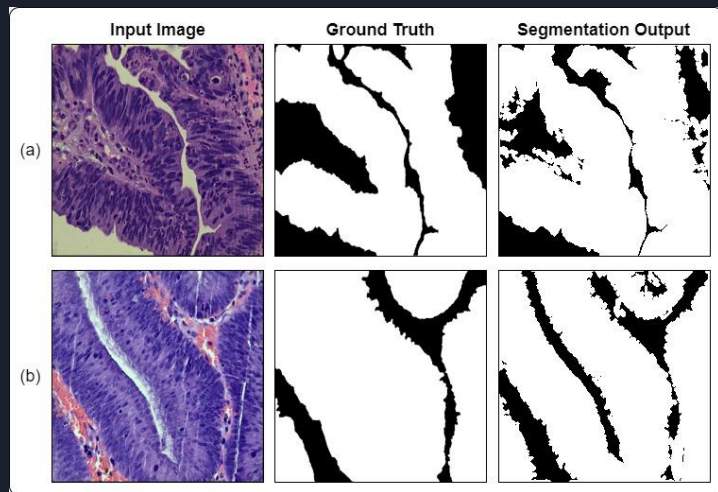
This general pipeline was further updated class wise to capture the specific patterns observed in the image.



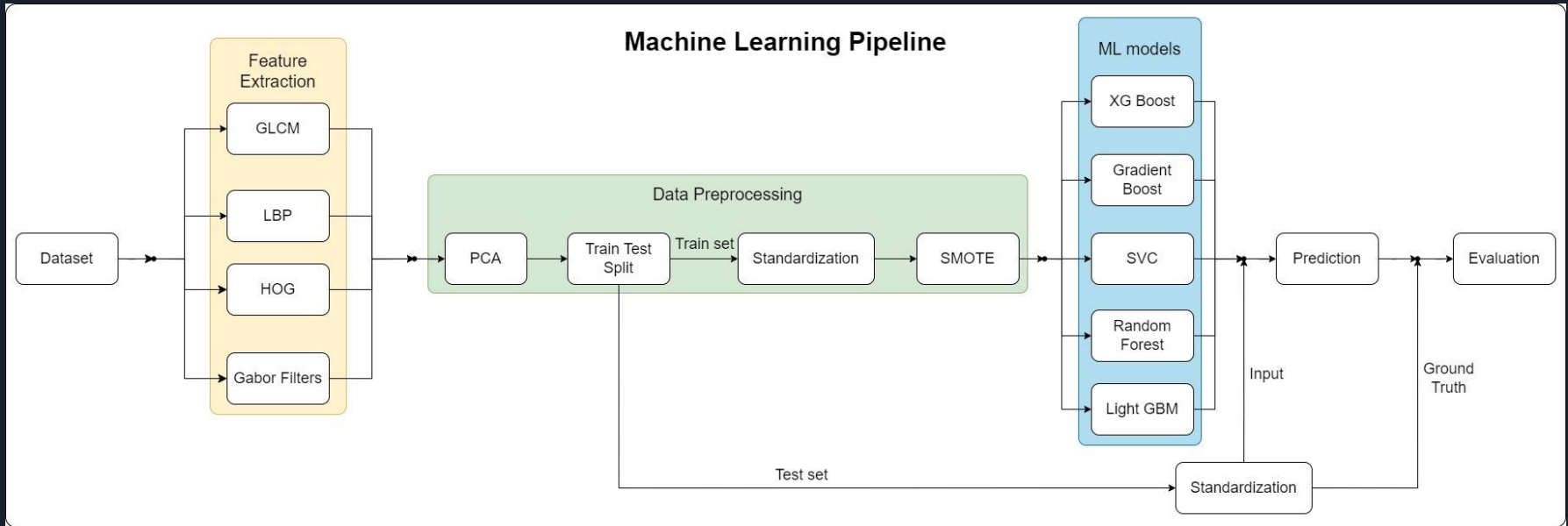
# Segmentation Result

	Precision (P)	Recall (R)	Jaccard Similarity (J)	Dice Score (D)	Accuracy (A)
Normal	0.90	0.78	0.73	0.83	0.81
Polyp	0.89	0.78	0.72	0.83	0.82
Low-grade IN	0.86	0.71	0.63	0.76	0.72
High-grade IN	0.85	0.93	0.80	0.88	0.83
Adenocarcinoma	0.90	0.82	0.74	0.84	0.79
Serrated Adenoma	0.78	0.85	0.67	0.80	0.73
<b>Combined Performance over all classes</b>	<b>0.86</b>	<b>0.81</b>	<b>0.71</b>	<b>0.82</b>	<b>0.78</b>

# Segmentation Result



# Machine Learning for Classification



# Classification Result

Features	Number of Features	Models	Precision (P)	Recall (R)	Accuracy (A)
LBP1 + LBP2 + LBP3	36	XGB	0.73	0.74	0.74
		GB	0.72	0.73	0.73
		SVC	0.76	0.77	0.77
		RF	0.70	0.71	0.71
		LGBM	0.75	0.75	0.75
LBP1 + LBP2 + LBP3 + HOG1 + PCA (n=400)	436	XGB	0.67	0.71	0.71
		GB	0.69	0.72	0.72
		SVC	0.56	0.64	0.64
		RF	0.63	0.64	0.64
		LGBM	0.70	0.71	0.71
LBP1 + LBP2 + LBP3 + HOG2 + PCA (n=400)	436	XGB	0.70	0.73	0.73
		GB	0.69	0.72	0.72
		SVC	0.60	0.64	0.64
		RF	0.61	0.64	0.64
		LGBM	0.71	0.73	0.73
<b>LBP1 + LBP2 + LBP3 + GLCM + Gabor filters</b>	<b>252</b>	<b>XGB</b>	<b>0.83</b>	<b>0.83</b>	<b>0.83</b>
		<b>GB</b>	<b>0.79</b>	<b>0.80</b>	<b>0.80</b>
		<b>SVC</b>	<b>0.82</b>	<b>0.81</b>	<b>0.81</b>
		<b>RF</b>	<b>0.79</b>	<b>0.79</b>	<b>0.79</b>
		<b>LGBM</b>	<b>0.81</b>	<b>0.82</b>	<b>0.82</b>
LBP1 + LBP2 + LBP3 + GLCM + Gabor filters + SMOTE	252	XGB	0.82	0.81	0.81
		GB	0.82	0.82	0.82
		SVC	0.82	0.82	0.82
		RF	0.78	0.78	0.78
		LGBM	0.81	0.81	0.81

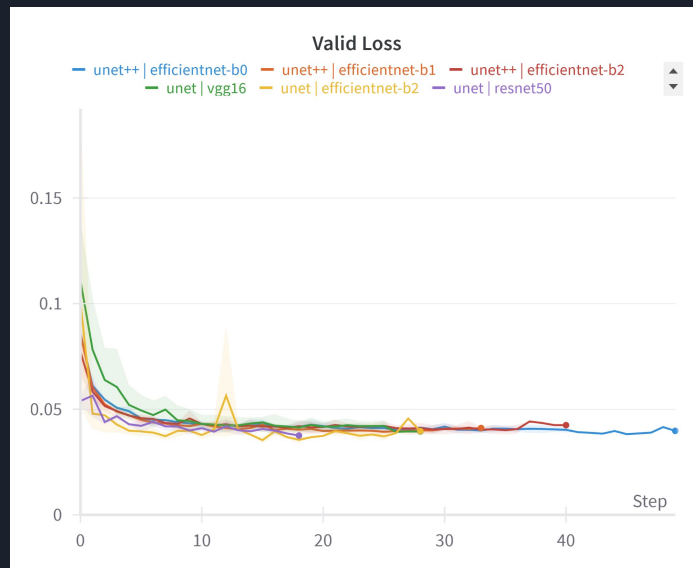
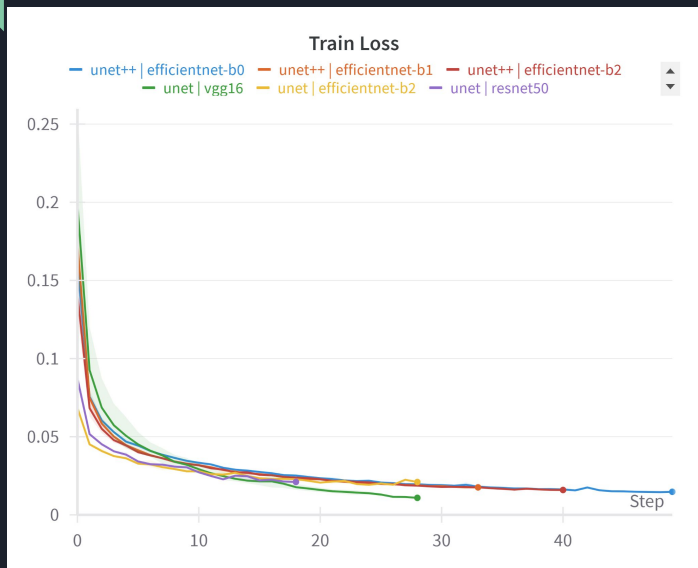




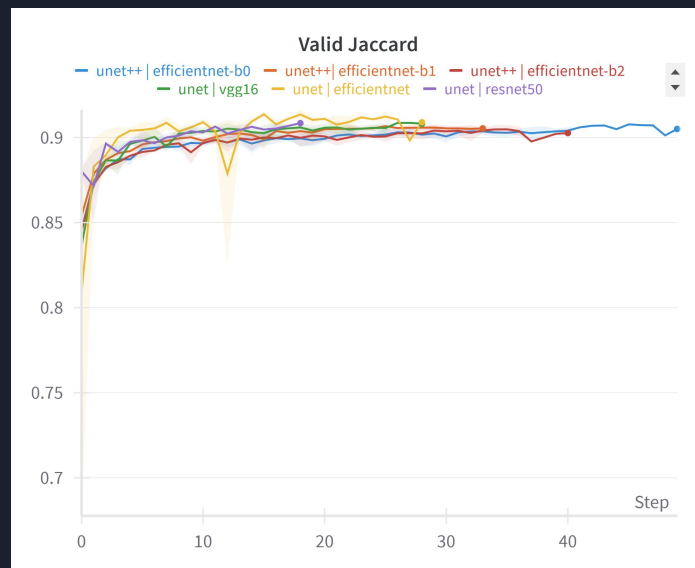
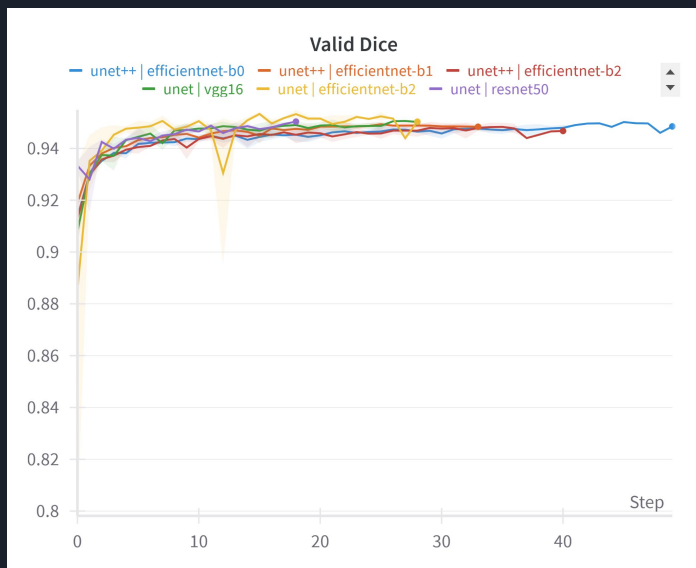
# Deep Learning for segmentation: Pipeline

- Dataset Preparation Pipeline:
  - Image Normalization
  - Mask Thresholding
  - Image Augmentation (not used during inference)
- Segmentations Architectures
  - UNet: Backbones-
    - VGG16
    - Resnet50
    - EfficientNet-B2
  - UNet++: Backbones-
    - EfficientNet-B0
    - EfficientNet-B1
    - EfficientNet-B2
- Experiments:
  - Detailed Performance of DL models
  - Class wise performance for each model
  - Sample Image Visualization for checking output masks

# Deep Learning for segmentation: Training Phase



# Deep Learning for segmentation: Training Phase cont.



# Deep Learning for segmentation: Inference Phase

## Detailed Performance of DL models

Models	Dice Score	Jaccard Similarity	Precision (P)	Recall (R)	Accuracy (A)
Baseline Unet Backbone: VGG16	0.9337	0.8921	0.9305	0.9397	0.9402
UNet Backbone: ResNet50	0.9353	0.8952	0.9308	0.9429	0.9421
<b>UNet</b> <b>Backbone: EfficientNet-B2</b>	<b>0.9372</b>	<b>0.8986</b>	<b>0.9323</b>	<b>0.9448</b>	<b>0.9448</b>
UNet++ Attention: scSE Backbone: EfficientNet-B0	0.9328	0.8909	0.9315	0.9377	0.9387
UNet++ Attention: scSE Backbone: EfficientNet-B1	0.9337	0.8926	0.9305	0.9408	0.9403
UNet++ Attention: scSE Backbone: EfficientNet-B2	0.9289	0.8846	0.9211	0.9417	0.9330

# Deep Learning for segmentation: Inference Phase

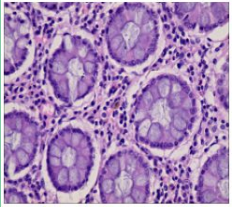
## Class wise performance of best model

	Dice Score	Jaccard Similarity	Precision (P)	Recall (R)	Accuracy (A)
<b>Normal</b>	0.9337	0.8921	0.9305	0.9397	0.9402
<b>Polyp</b>	0.9353	0.8952	0.9308	0.9429	0.9421
<b>Low-grade IN</b>	0.9372	0.8986	0.9323	0.9448	0.9448
<b>High-grade IN</b>	0.9328	0.8909	0.9315	0.9377	0.9387
<b>Adenocarcinoma</b>	0.9337	0.8926	0.9305	0.9408	0.9403
<b>Serrated Adenoma</b>	0.9289	0.8846	0.9211	0.9417	0.9330

# Deep Learning for segmentation: Inference Phase

## Sample Segmentation Outputs

Image  
Label: Normal



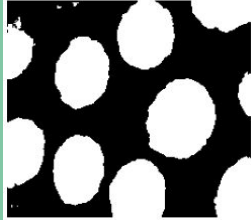
Ground Truth Mask



Unet|Vgg16|Baseline



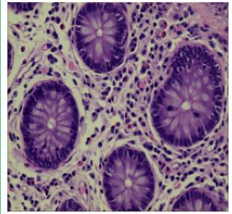
Unet|ResNet50



Unet|EfficientNet-B2



Image  
Label: Polyp



Ground Truth Mask



Unet|Vgg16|Baseline



Unet|ResNet50



Unet|EfficientNet-B2





THANK YOU