

# WEB APPLICATION FOR AUTOMATIC NUCLEUS COUNTING IMMUNOFLUORESCENCE TISSUE BIOPSIES



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## ABSTRACT

Accurate nucleus counting is vital in pathology for disease diagnosis and treatment advancement. Pathologists typically employ Fluorescence Microscopy machines to image DAPI-stained tissue and rely on the Columbus platform for automated nucleus counting. However, the platform struggles to accurately count and locate nuclei in cases of overlapping, clustered, or indistinct shapes caused by overlapping fluorescent dyes. Consequently, pathologists are compelled to manually count nuclei, resulting in time-consuming processes prone to errors.

## OBJECTIVE

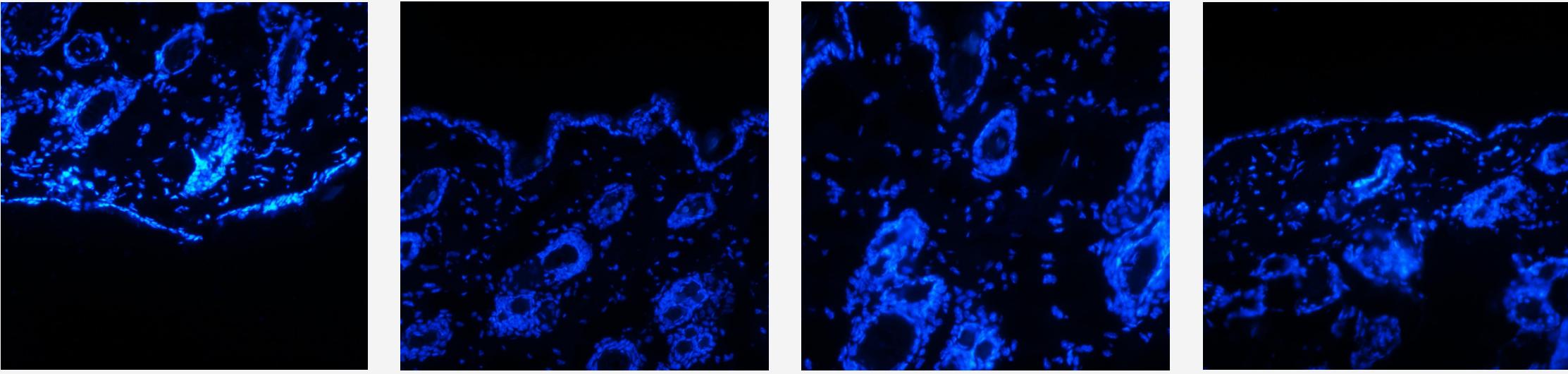
- 1

To develop a model for counting nuclei from histological images stained with a fluorescent dye.
- 2

To develop a web application that can perform nucleus counting from histological images stained with a fluorescent dye.

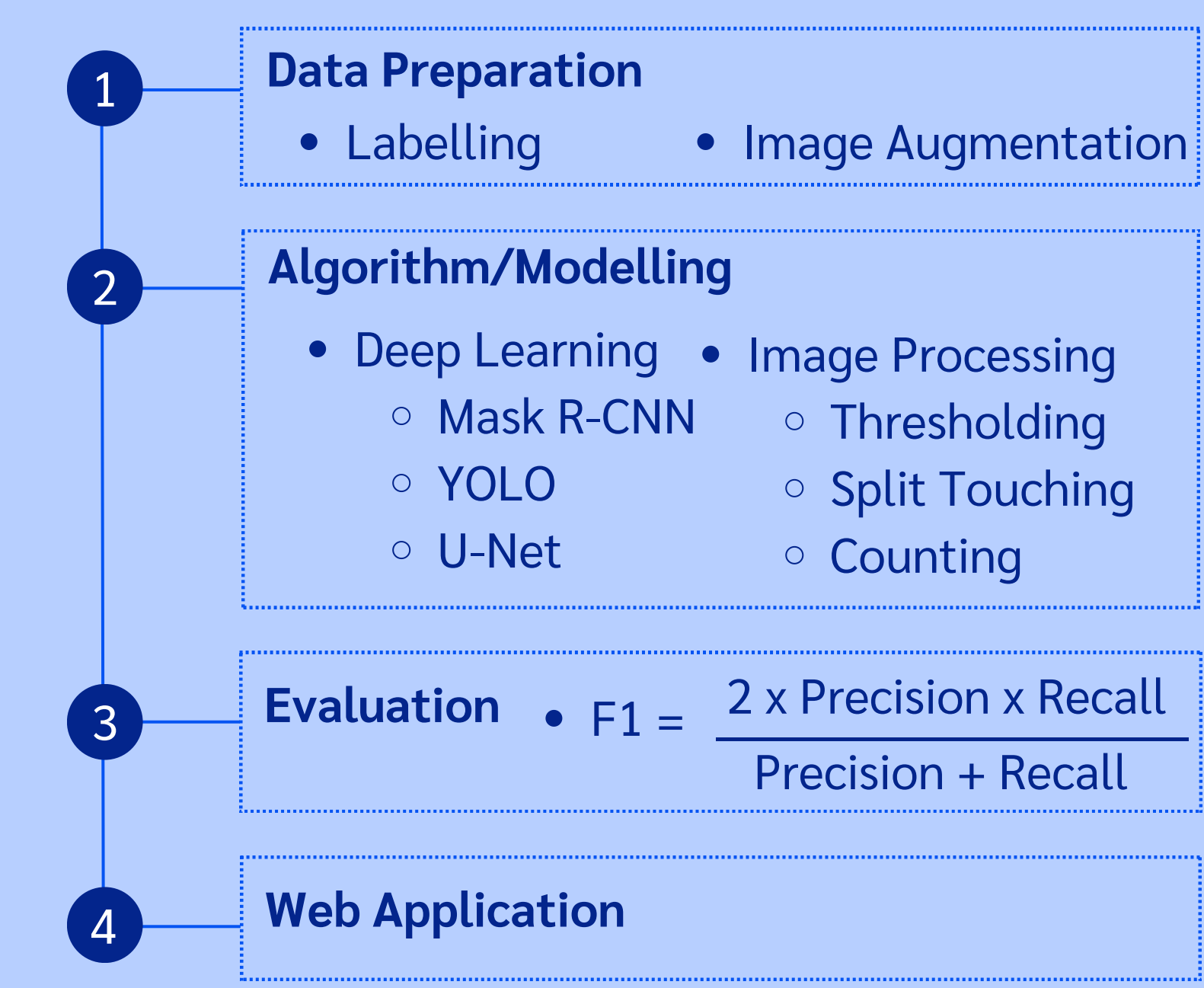
## DATASET

Immunofluorescence Tissue with 4',6-diamidino-2-phenylindole (DAPI), to precisely identify the locations of nuclei.

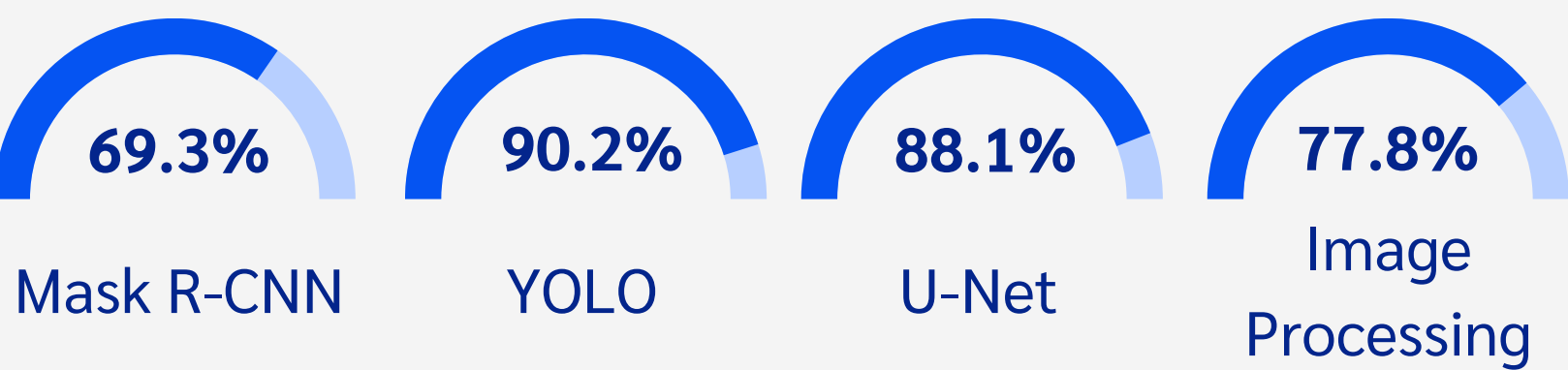


Target Users is  
" Pathologist "

## METHODOLOGY



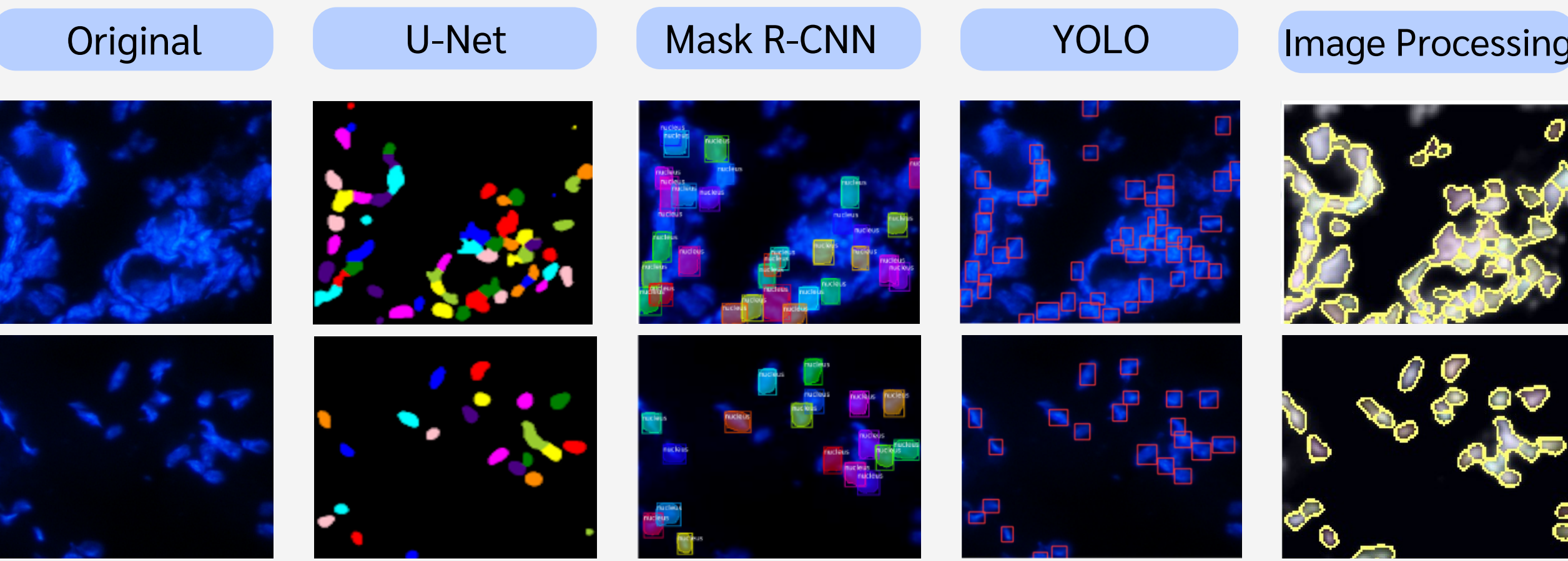
## EVALUATION with F1-score



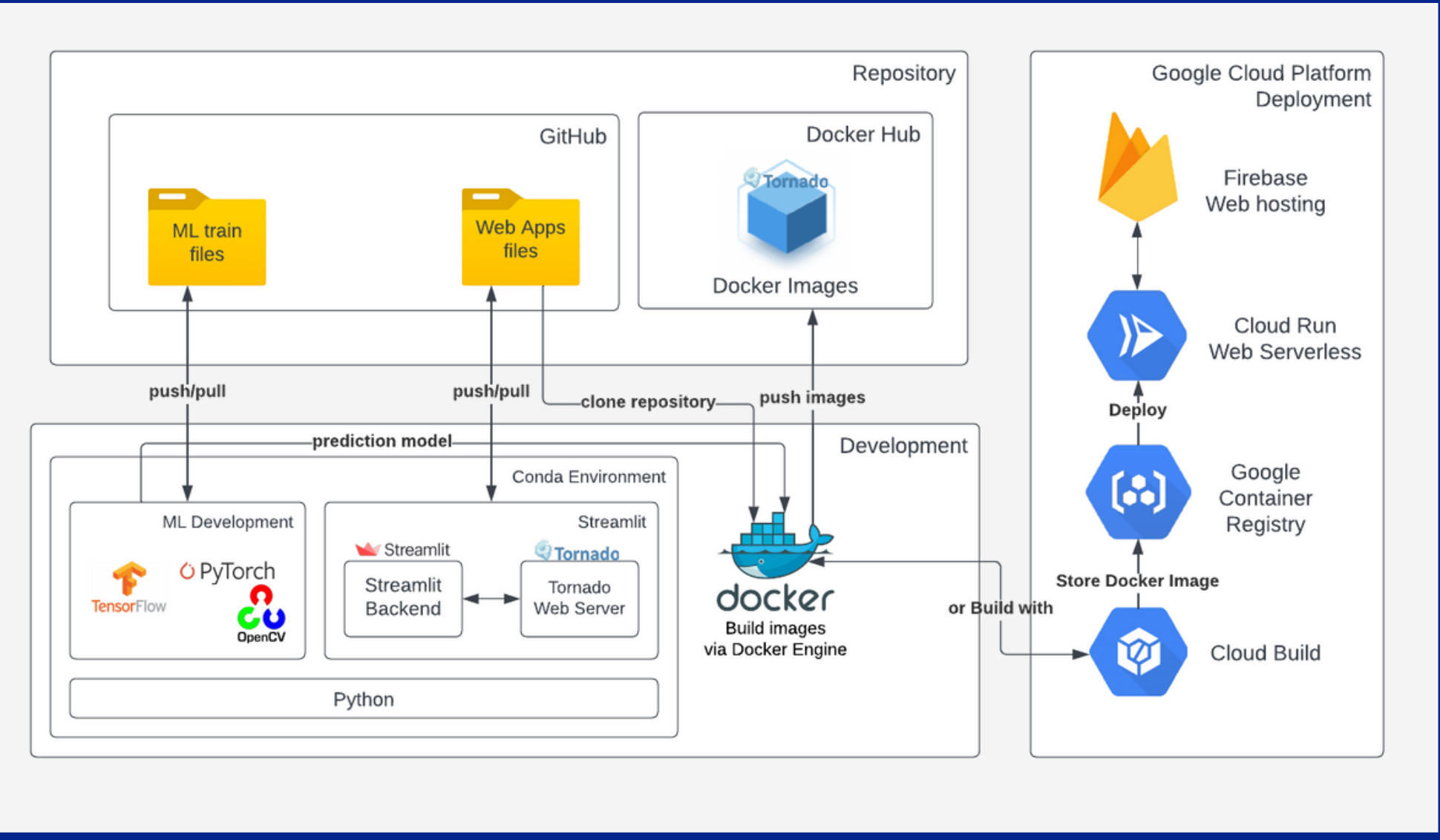
## RESULT

Based on the results obtained from comparing all the models, it was found that the YOLO model can accurately identify the positions of nuclei the best, with adjustments made to six parameters: Dropout, Leaky, Hardswish, momentum, scale image, and object loss.

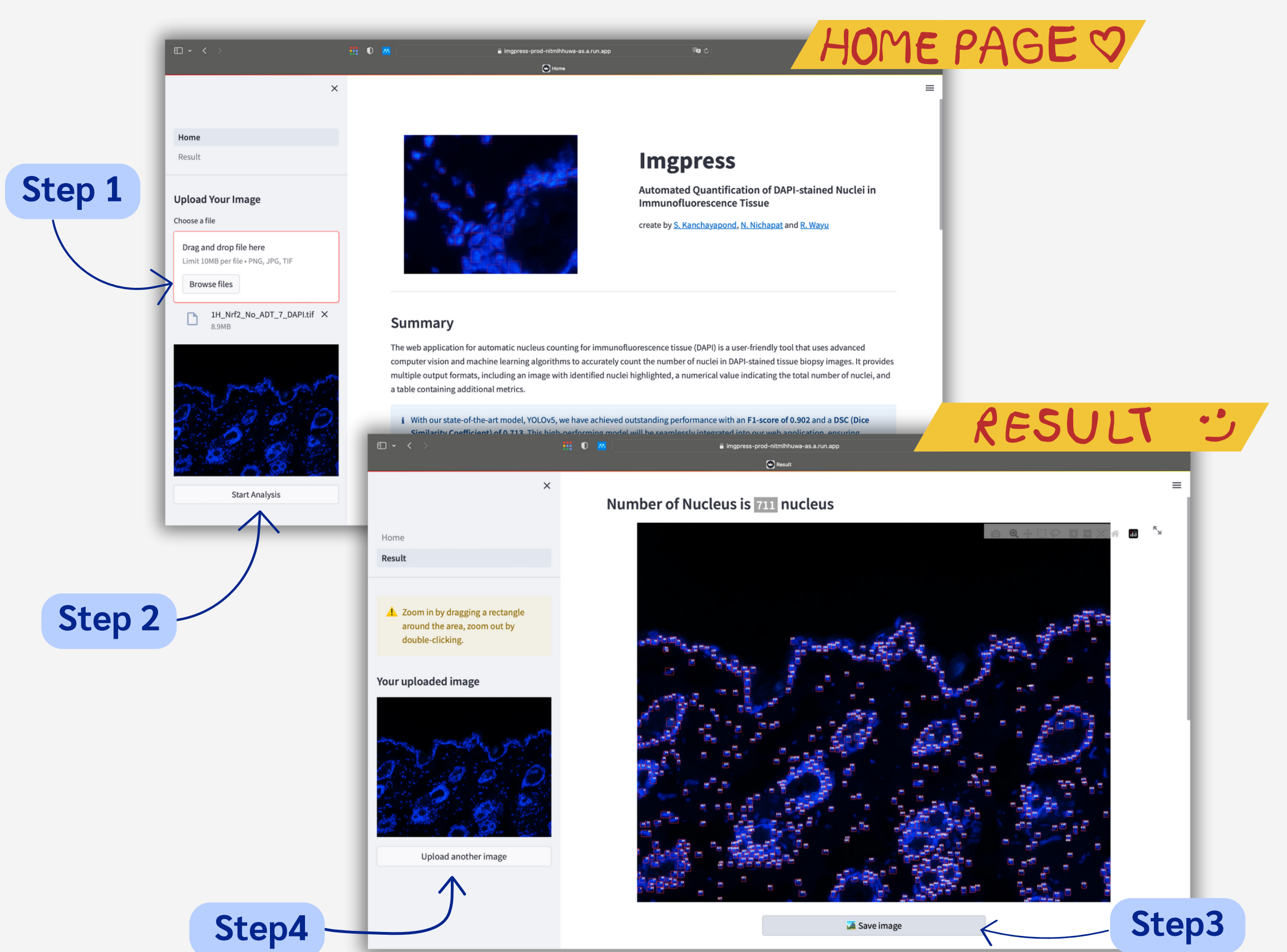
Algorithm/Model	Precision	Recall	F1 Score	ENr	DSC
Mask R-CNN	0.857	0.581	0.693	0.659	0.509
YOLO	<b>0.896</b>	0.908	<b>0.902</b>	0.823	0.713
U-Net	0.840	0.926	0.881	0.908	<b>0.740</b>
Image Processing	0.657	<b>0.953</b>	0.778	<b>1.083</b>	0.670



## SYSTEM ARCHITECTURE



## WEB APPLICATION



## USER SATISFACTION



Best Performance  
**YOLO**  
with F1-score  
**90.2%**