

Automated Identification of Necrotic Regions in Digital Images of Multiplex Immunofluorescence Stained Tissue Using Deep Learning



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Background

- Necrosis:** A spectrum of morphological changes that follow cell death in living tissue, largely resulting from the progressive degradative action of enzymes on the lethally injured cell.

Types of necrosis:

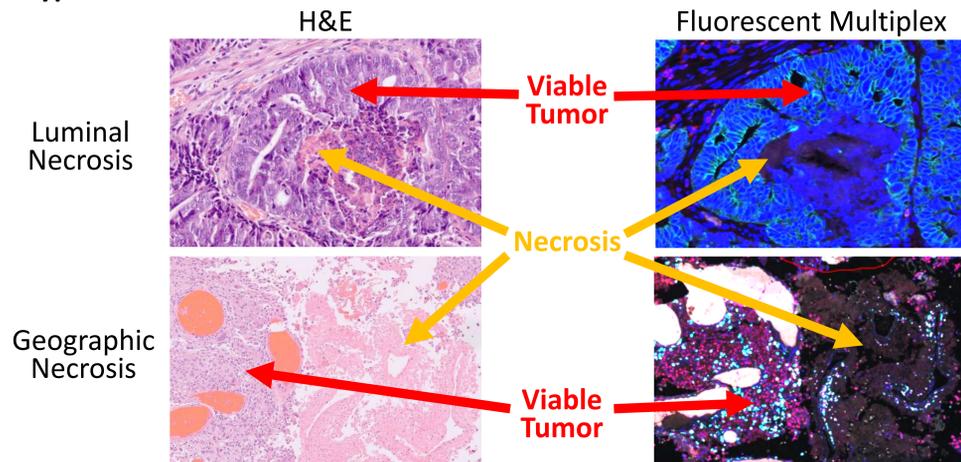


Figure 1: Luminal and geographic necrosis examples in H&E and Fluorescent multiplex stained images.

Necrosis Identification and Challenges

- Excluding necrosis allows us to focus on the Tumor Microenvironment (TME) and get better insight into responses to immunotherapy approaches.
- Manual identification and annotation is the most common practice.

Challenges in manual detection:

- Variation with respect to size, shape, number, region of occurrence.
- Time consuming.
- Reliance on H&E stained image.

Result of manual annotation:

- Expensive process.
- Annotations can vary across pathologists.
- Susceptible to errors and inaccuracies.

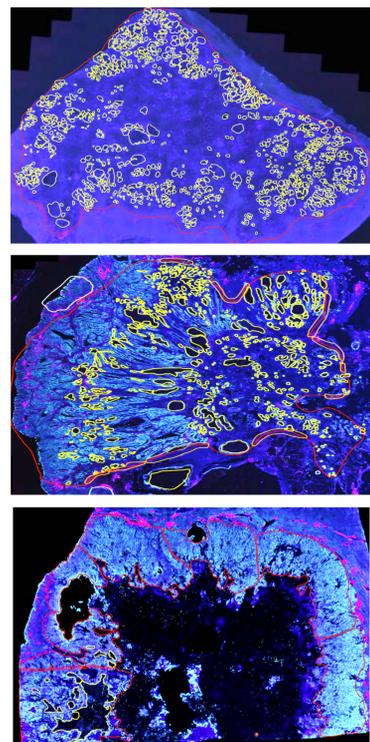


Figure 2: Variation in necrotic regions (yellow) within TME (red).

Methods

Automated Workflow

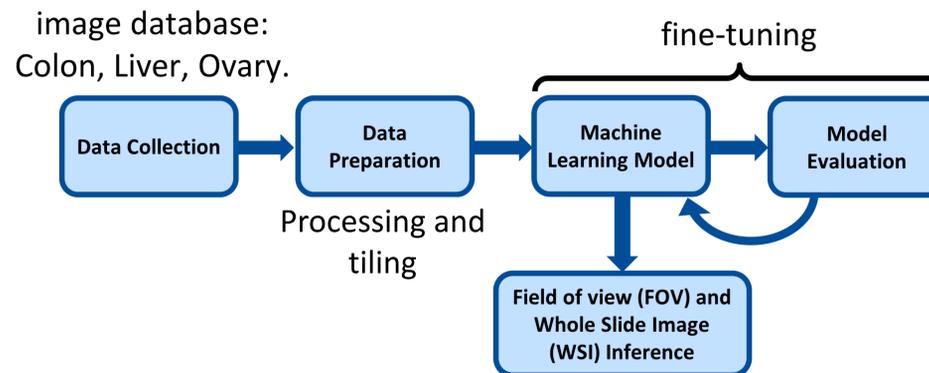


Figure 3: Illustration of workflow for an automated framework.

Gamma Correction

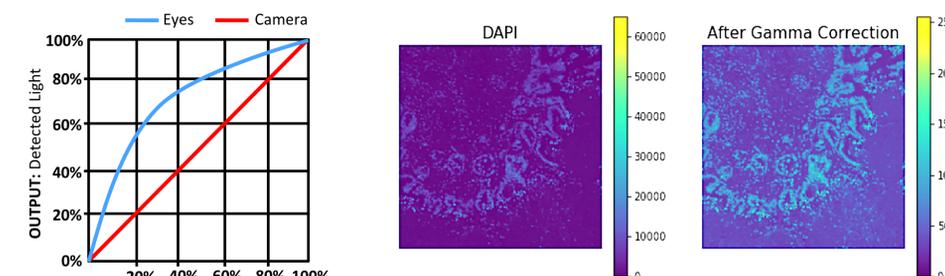


Figure 4: Gamma correction with $\gamma < 1$.

Figure 5: Amplification of lower intensity pixels using gamma correction with $\gamma = 0.5$.

U-Net

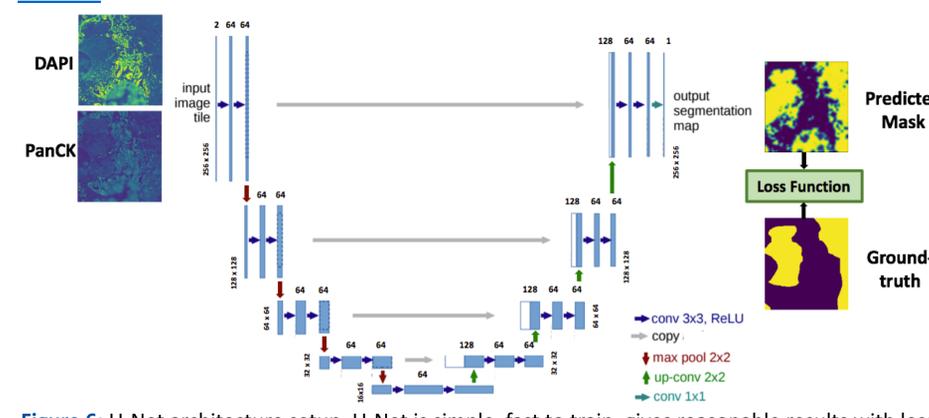


Figure 6: U-Net architecture setup. U-Net is simple, fast to train, gives reasonable results with less amount of training data. U-Net and its variants have achieved state-of-the-art results for medical image segmentation.

Results

- Loss functions:** Binary cross-entropy, Dice similarity coefficient.
- Optimizers:** Momentum, Adam.
- Data augmentation:** Rotation, Translation, Zoom, Shear, Horizontal flip.

Hyperparameters	Dice Similarity
Loss: Binary cross-entropy, Optimizer: Momentum, Learning rate: 1e-3	0.82
Loss: Dice Similarity Coefficient, Optimizer: Adam, Learning rate: 1e-4	0.82

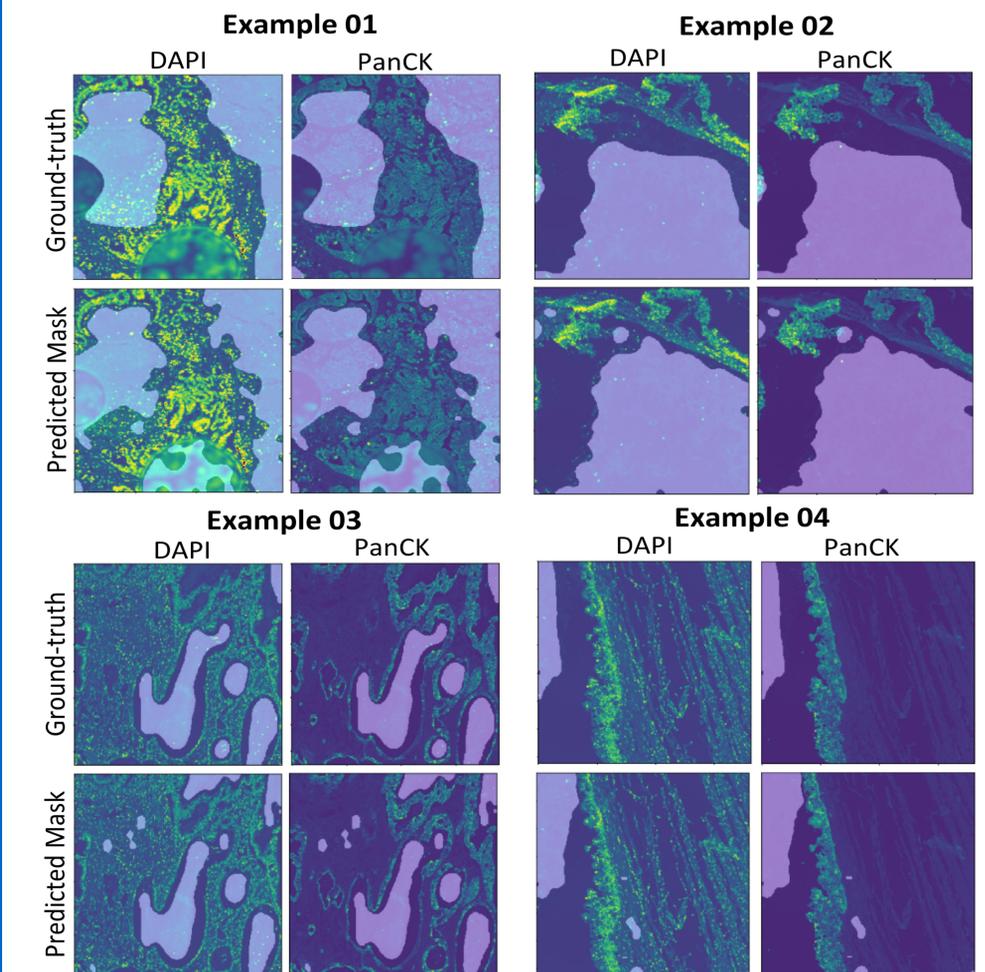


Figure 7: Comparison of ground-truth and predicted masks for different FOVs.

References

- Ronneberger et al., "U-Net: Convolutional networks for biomedical image segmentation", MICCAI, 2015.
- Moen et al., "Deep learning for cellular image analysis", Nature Methods, 2019.
- R. M. Levenson, "Spectral Imaging Perspective on Cytomics", ISAC, 2006.
- Lahiani et al., "Generalising multistain immunohistochemistry tissue segmentation using end-to-end colour deconvolution deep neural networks", IET Image Processing, 2019.