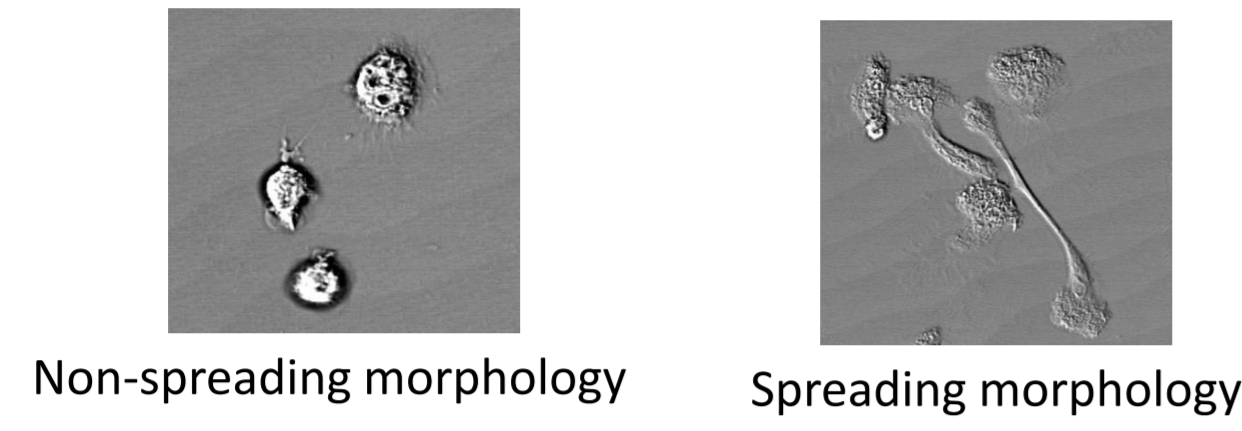


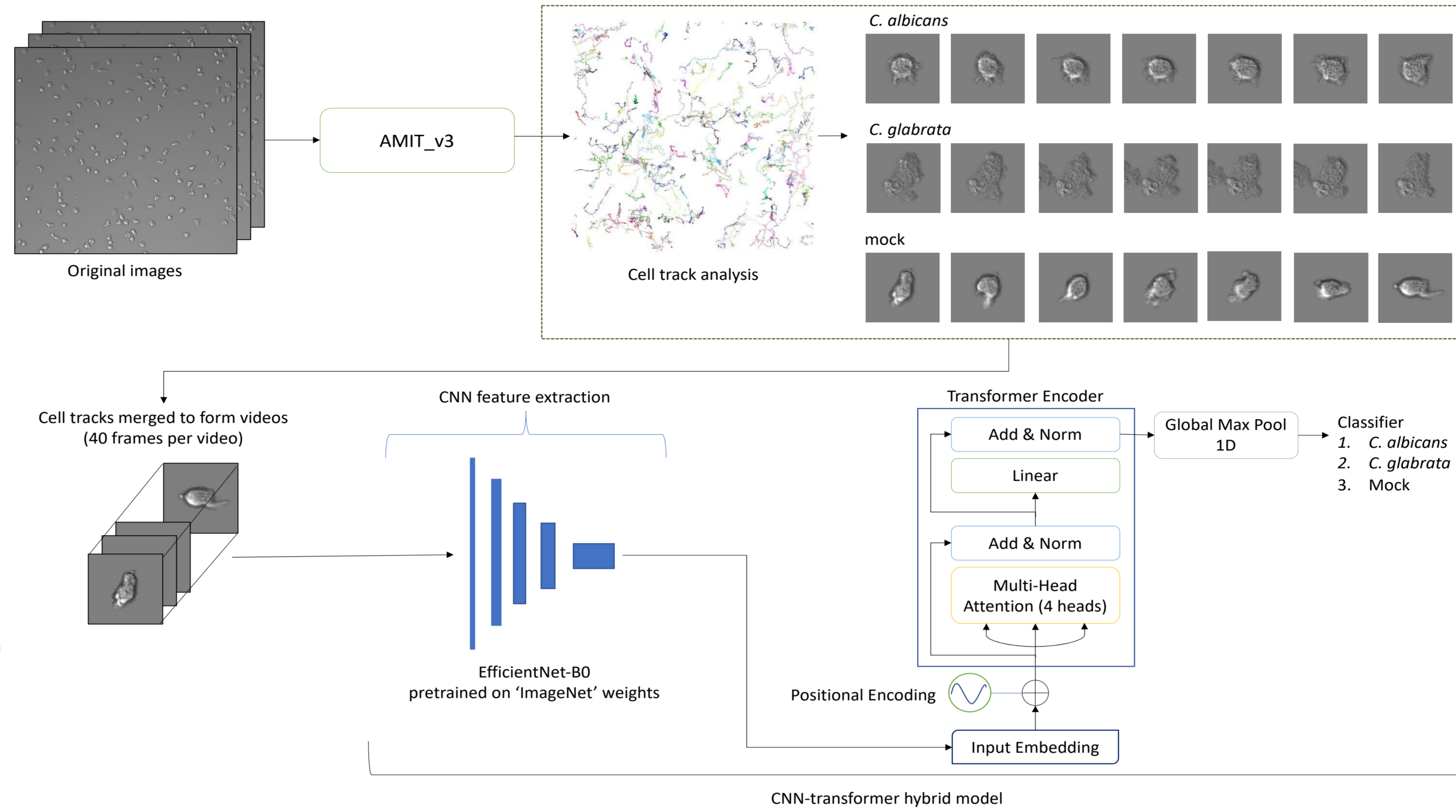
Single cell morphodynamics analysis

- Whole blood infection – 9 Donors [1]
- Time-lapse microscopy data
- 3 cases each (Mock, *C. albicans*, *C. glabrata*)
- Each video with 260 frames / images
- Video duration: 30 min, Time-step: 7 sec
- Neutrophils exhibit two morphological appearances



Single cell morphodynamics – classifier on cell tracks

- AMIT_v3 [2] (Algorithm for Migration and Interaction Tracking) for tracking and splitting cells
- Split cells merged across 40 frames to form videos
- EfficientNetB0 [3] (Convolutional Neural Network - CNN) for feature extraction
- Transformer network for classification
- LOOCV - Leave One Out Cross-Validation applied



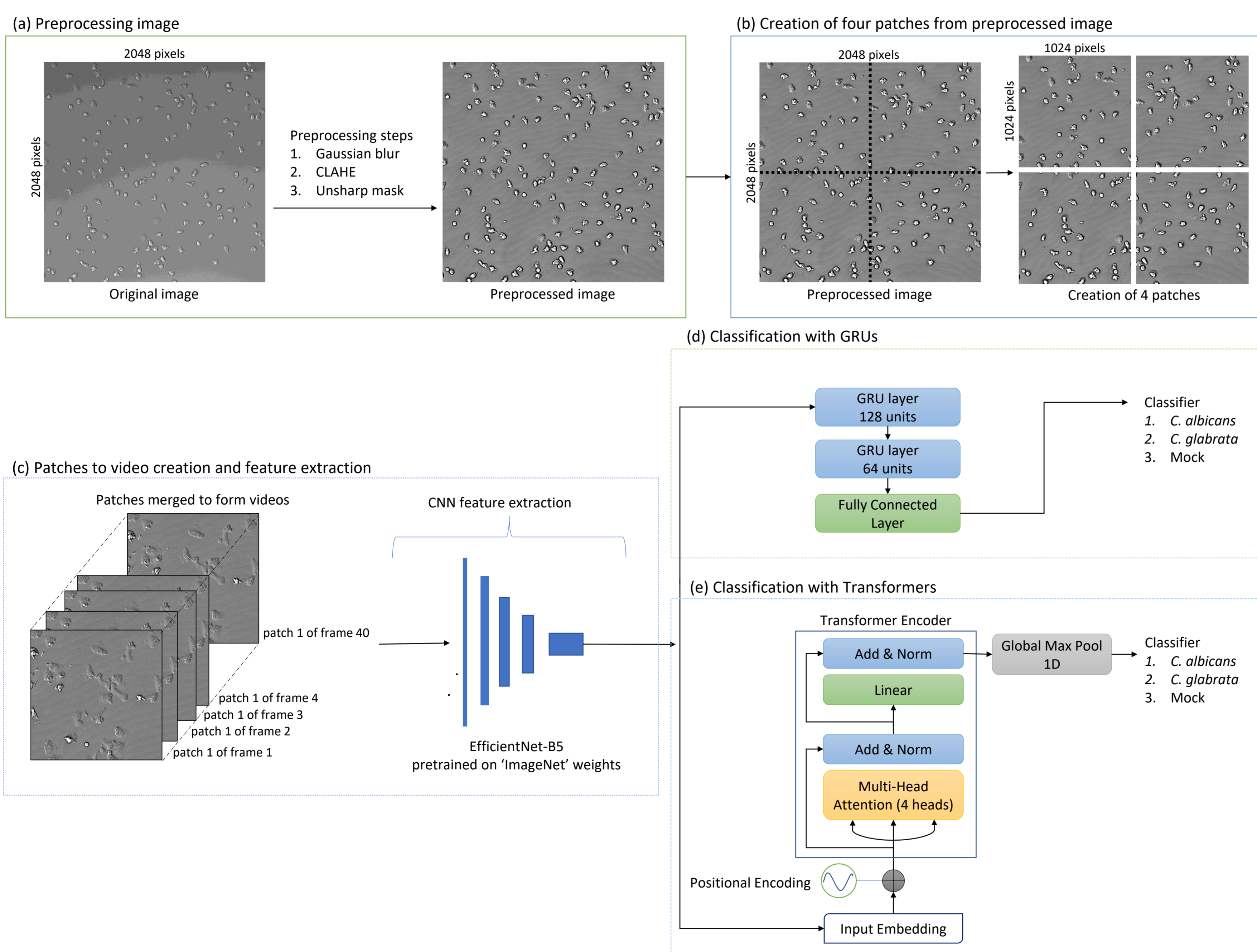
Results

- Mock classified correctly
- Network confuses between *C. albicans* and *C. glabrata*

Confusion matrix – transformers (LOOCV) on single cell videos

True Labels \ Predicted Labels	<i>C. albicans</i>	<i>C. glabrata</i>	Mock
<i>C. albicans</i>	5	2	2
<i>C. glabrata</i>	1	8	0
Mock	0	0	9

GRU (Gated Recurrent Unit) and Transformer based analysis of video data



Workflow

1. Preprocessing for perfect separation between background and neutrophils
 - Gaussian blur - reduce noise from the images
 - CLAHE (contrast-limited adaptive histogram equalization) - contrast enhancement
 - Unsharp Mask – sharpen image
2. Creating patches to increase dataset size
3. Patches merged across 40 frames to form videos
4. CNN used for feature extraction
 - CNN used - Pretrained EfficientNetB5 [3]
5. Classifier network
 - Extracted features through GRU [4] and Transformer [5] network

Results

- Pro: Transformer [5] classifies all videos correctly
- Con: Long inference time due to time-consuming video formation and feature extraction steps (up to 3 hours for each case)

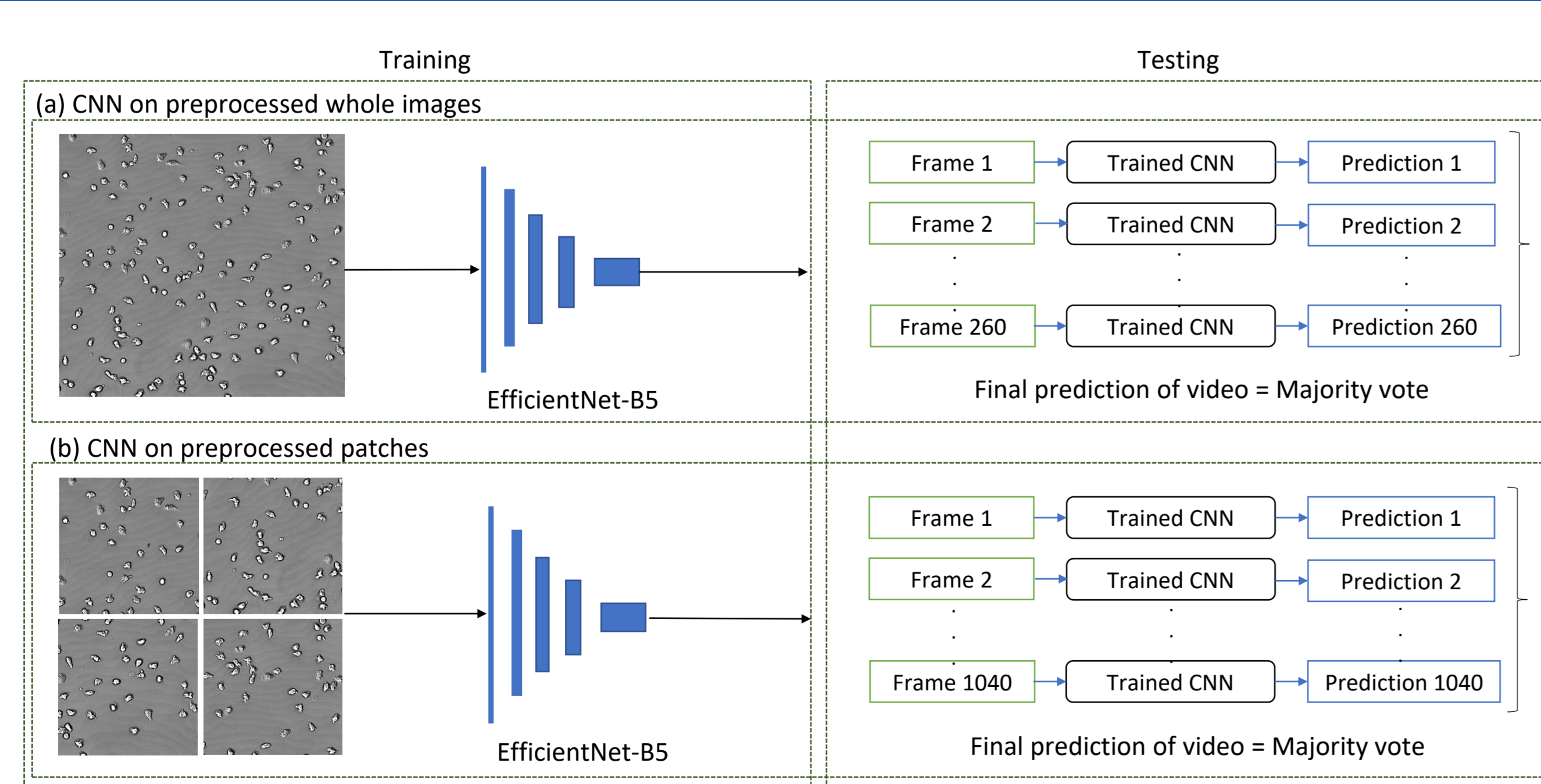
Confusion matrix – (d) GRUs (LOOCV) on preprocessed patch videos

True Labels \ Predicted Labels	<i>C. albicans</i>	<i>C. glabrata</i>	Mock
<i>C. albicans</i>	8	1	0
<i>C. glabrata</i>	0	9	0
Mock	0	0	9

Confusion matrix – (e) Transformers (LOOCV) on preprocessed patch videos

True Labels \ Predicted Labels	<i>C. albicans</i>	<i>C. glabrata</i>	Mock
<i>C. albicans</i>	9	0	0
<i>C. glabrata</i>	0	9	0
Mock	0	0	9

CNN based analysis on image data

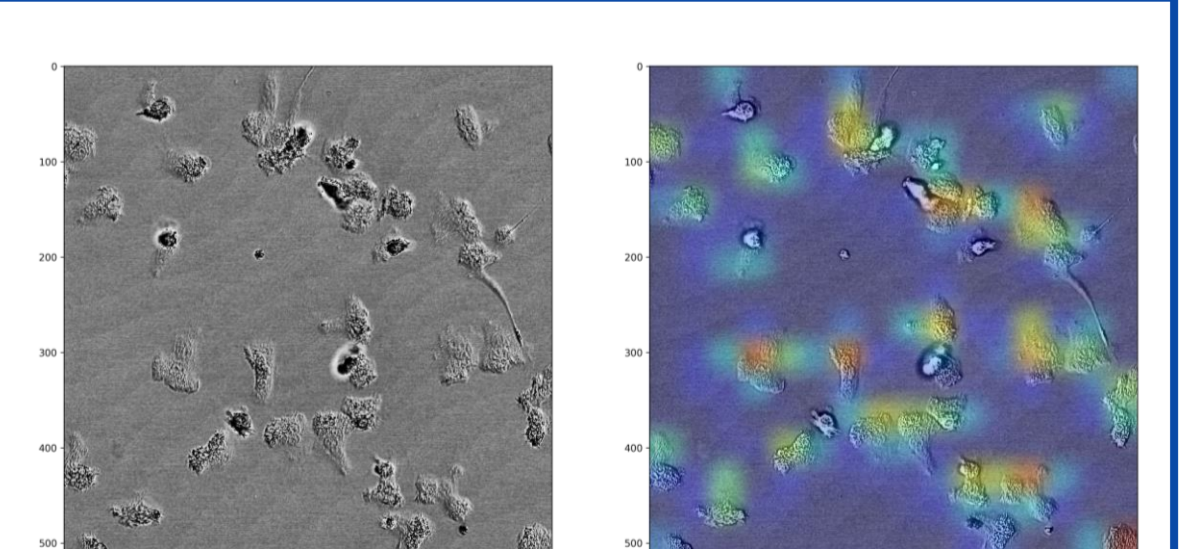


Confusion Matrix (a) CNN (LOOCV) on preprocessed whole images

True Labels \ Predicted Labels	<i>C. albicans</i>	<i>C. glabrata</i>	Mock
<i>C. albicans</i>	8	1	0
<i>C. glabrata</i>	0	9	0
Mock	0	0	9

Confusion Matrix (b) CNN (LOOCV) on preprocessed patches

True Labels \ Predicted Labels	<i>C. albicans</i>	<i>C. glabrata</i>	Mock
<i>C. albicans</i>	9	0	0
<i>C. glabrata</i>	0	9	0
Mock	0	0	9



Grad-CAM on *C. glabrata* shows CNN focusing on the spreading cells to make predictions

Results

- CNN on patches classifies all videos correctly
- Very fast inference time (15 minutes)
- Grad-CAM [6] for explainability
- Needs fewer images for inference
- Ensemble learning for best results

Workflow

1. Preprocessed images and patches trained using CNN – EfficientNetB5 [3]
2. Majority vote across 260 frames gives final prediction

References

- [1] Belyaev, I., & Praetorius, JP. *et al.* 2022. *Comput struct biotechnol j.*,20, (2297-2308)
- [2] Belyaev, I. *et al.* 2021. *Cytom Part A. cyto a.*24466
- [3] Tan, M., & Le, Q. 2019. *ArXiv, abs/1905.11946.*
- [4] Chung, J., *et al.* 2014, *ArXiv:1412.3555*
- [5] Vaswani, A., *et al.* 2017, *ArXiv:1706.03762*
- [6] Selvaraju, RR., *et al.* 2017, *ArXiv:1610.02391*