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Abstract #: 1159

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BACKGROUND

- Determining pathological T (pT) category in invasive urothelial carcinoma from transurethral resection of bladder (TURB) specimens** is challenging due to tissue fragmentation and artifacts, and its study for deep-learning based approach has been less studied than radical cystectomy samples.
- Deep learning (DL) based applications for pathology diagnosis typically require burdensome manual annotations, and **multiple instance learning (MIL)** has been widely used in pathology, with many variations improving its performance.
- We tried to use MIL-based approach to not only discern carcinoma, but categorize pT category by implementing MIL-based techniques, and implemented modifying some of the proposed techniques in the literature.

Related Works

- Attention-based Deep Multiple Instance Learning (Ilse et al., 2018):**
 - Neural network-based attention mechanism for MIL that provides insight into each instance's contribution to bag classification.
 - Outperformed other methods on histopathology datasets while maintaining interpretability with learnable attention mechanism.
- Clinical-grade Computational Pathology (Campanella et al., 2019):**
 - A weakly supervised DL system with fixed top-K instance selection validated on 44,732 WSIs from 15,187 patients.
 - Achieved AUCs above 0.98 for prostate cancer, basal cell carcinoma, and breast cancer metastases detection.
- Joint Breast Neoplasm Detection and Subtyping (Casson, Liu et al., 2023):**
 - A multi-resolution network using top-K MIL with adaptive K selection via Otsu's thresholding method on over 125,000 WSIs.
 - Utilized transformer architecture to integrate features across magnification levels for comprehensive breast neoplasm analysis.

METHODS

Patient data selection and experimental setup

- Analyzed 5,673 matched H&E-stained megapatches from bladder cancer TURB whole slide image (WSI) specimens from 55 patients (mean images per patient: 56.73; min: 7; max: 175), scanned with PANNORAMIC™ Digital Slide Scanner (3DHISTECH)
- Used megapatches (1024×1024 pixels) instead of whole slide images for computational efficiency
- Dataset distribution: T1 (2978 patches, 52.5%), T2 (2695 patches, 47.5%)
- Experiments conducted on NVIDIA A100 80GB GPU with CUDA 11.3

Model architecture

1. Bi-Resolution Top K-MIL Approach (Figures 1&2)

- Processed megapatches at dual magnifications (400x and 50x) separately
- Used SEResNet18 for feature extraction with adaptive Top-K patch selection
- Employed Otsu thresholding to automatically determine optimal K value
- Integrated features with transformer architecture to combine information across magnifications

2. Weighted Attention MIL (Figures 1&2)

- CNN backbone at 200x magnification with fully convolutional network layers
- FCN with attention layer with automatic feature weighting

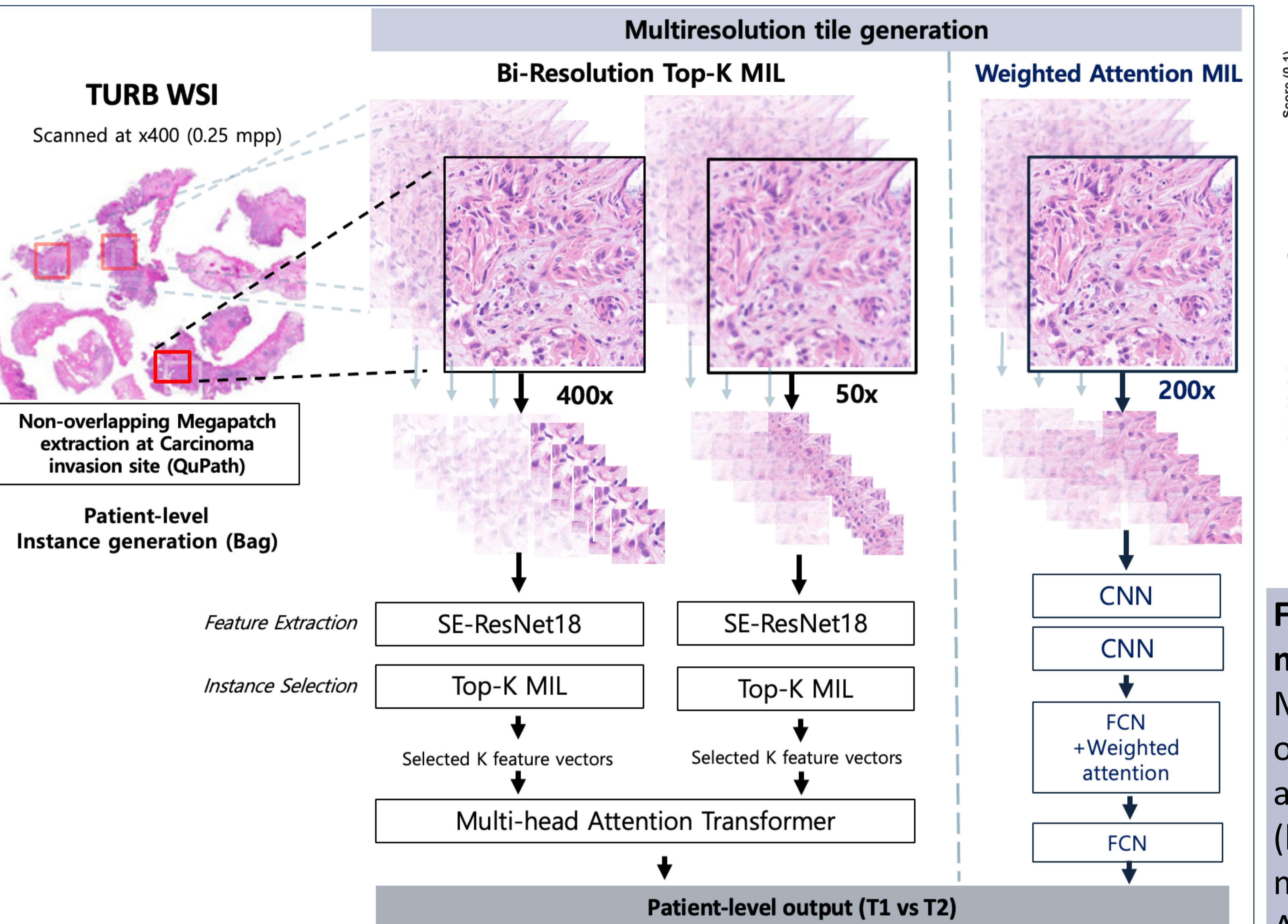


Figure 1. Schematic View of the Comparative MIL approaches for pathological tumor staging using multiresolution Megapatches. WSI, Whole Slide Image; TURB, Transurethral Resection of Bladder; mpp, microns per pixel; MIL, Multiple Instance Learning; CNN, Convolutional Neural Network; FCN, Fully Connected Network; SE-ResNet18, Squeeze-and-Excitation ResNet (18 layers); QuPath, Open-source software for digital pathology analysis.

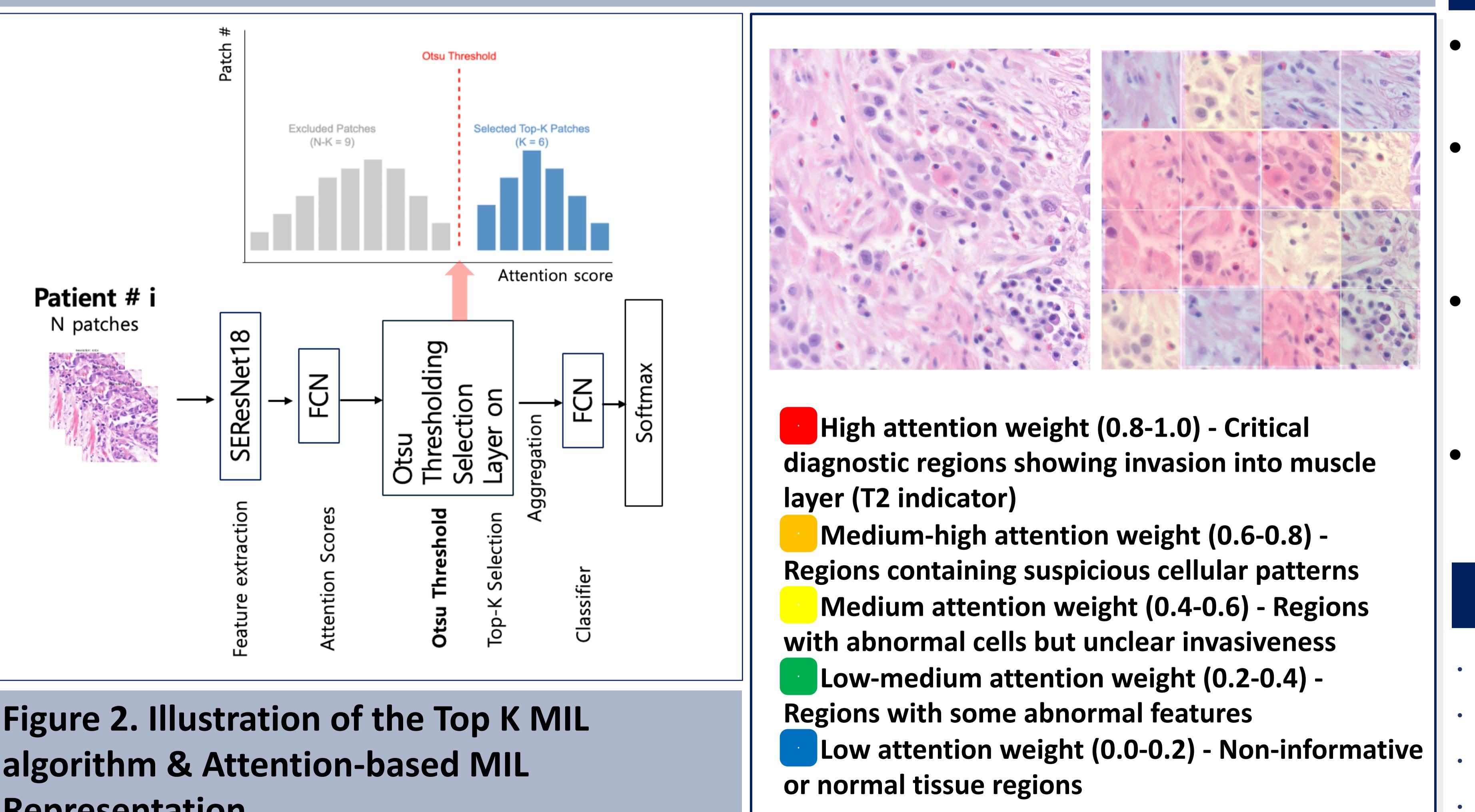


Figure 2. Illustration of the Top K MIL algorithm & Attention-based MIL Representation

RESULTS

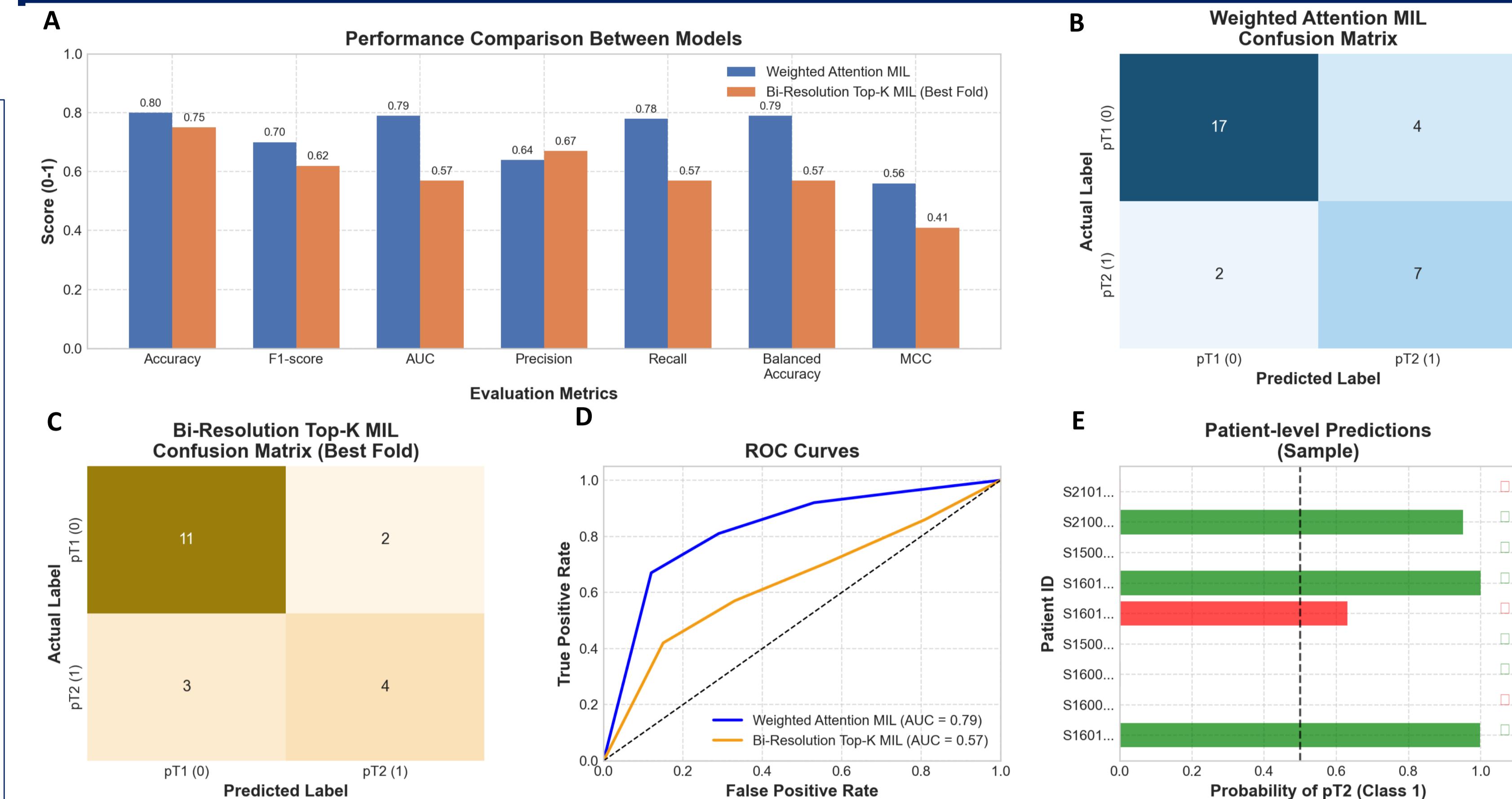


Figure 3. Performance evaluation of bladder cancer staging classification models. (A) Comparative performance metrics between Weighted Attention MIL and Bi-Resolution Top-K MIL (best fold) approaches. (B) Confusion matrix of Weighted Attention MIL showing the distribution of predictions across actual pT1 and pT2 classes. (C) Confusion matrix of Bi-Resolution Top-K MIL (best fold) demonstrating prediction distribution. (D) ROC curves for both models showing discriminative capability with AUC of 0.79 for Weighted Attention MIL and 0.57 for Bi-Resolution Top-K MIL. (E) Patient-level prediction probabilities for selected cases, with green bars indicating correct predictions and red bars indicating misclassifications (dashed line represents the classification threshold at 0.50).

DISCUSSION AND CONCLUSION

- Our approach demonstrates effectiveness beyond simple cancer vs. normal classification, successfully distinguishing between T1 and T2 staging.
- Tile-level processing offers computational efficiency while maintaining discriminative power, avoiding the extensive GPU resources and background processing required for full WSI analysis.
- Rescaled images don't provide the same advantages as true multi-scanning approaches due to missing contextual information: Low-resolution scaled images may potentially confuse rather than enhance the model.
- Findings suggest selective attention to discriminative regions may be more valuable, particularly when working with limited training datasets.

CONTACTS

- Data and code availability:** Please e-mail us. We are happy to provide details!
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