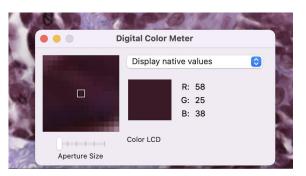
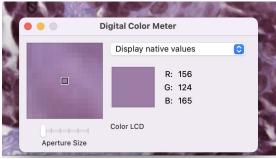
# Presentation 1: Critique of Spring 2025 MIL Model

STAT 390 | Project 1 | Fall 2025

# 1. Color Representation - Problems

- Challenges with Current Approaches Using RGB
  - **High Correlation between RGB Channels leads to inefficient feature learning**. e.g. If a color is darker, R, G, B channels decreases together





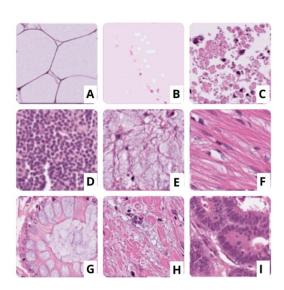
- Coupled Information: RGB combines luminance (brightness/intensity of light) and chrominance (color), obscuring the structural features critical for detecting melanocyte distribution
  - Human perception relies heavily on luminance to identify boundaries and shapes. Using a color space containing luminance info could potentially enhance model interpretability

# 1. Color Representation - Other Color Spaces

Color Space Name	Description	Channels
YCbCr	Used in digital video and image compression, it separates luma (brightness) from two chroma (color difference) components	Luma (Y), Blue-difference Chroma (CB), Red-difference Chroma (CR)
Lab (CIELAB)	Designed to be perceptually uniform, it defines color based on lightness and two color-opponent dimensions	Lightness (L), Green-Red Axis (a), Blue-Yellow Axis (b)
HSV	Represents colors in terms of their tint, purity, and brightness, often used in color pickers	Hue (H), Saturation (S), Value (V)
XYZ (CIEXYZ)	A device-independent color space based on how an average human observer perceives color, serving as a standard reference	Red-like (X), Green-like (Y), Blue-like (Z)

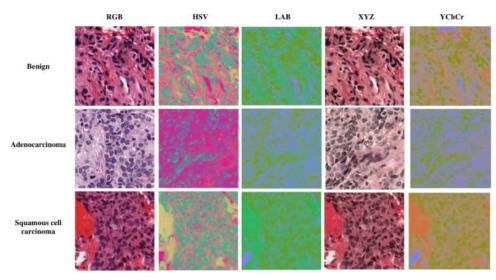
# 1. Color Representation - Literature Review

- Bishnoi & Goel (2023): impact of 5 color spaces on tissue slice lung cancer classification
  - Color spaces: RGB, HSV, LAB, XYZ, YCbCr
  - HSV consistently outperforms other color spaces in ResNet, DenseNet, and Inception V3, and their own model
- Velastegui & Pedersen (2023): 9 categories of colorectal cancer histological imaging
  - Color spaces: RGB, XYZ, HSV, YCbCr, CIELAB, LLL, AAA, BBB
  - HSV and CIELAB have the highest overall accuracy
  - HSV is the best for category D (lymphocytes) similar to our images



# 1. Color Representation - Action Plans

- Transform RGB into HSV and CIELAB and benchmark with MIL from last quarter
- Test whether channels that primarily contain brightness information (Y-channel in YCbCr,
   V-channel in HSV, and L-channel in CIELAB) perform well standalone
  - $\circ$  If so, we can consider using 3 channels across matched stains. Eg.  $(Y_{HE}, Y_{Mel}, Y_{Sox})$



### 1. Color Representation - Action Plans

Option A: Transform RGB into HSV, CIELAB



Option B: Transform RGB into HSV and CIELAB

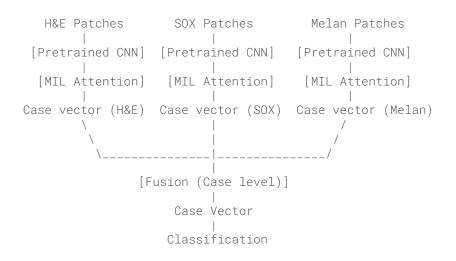
```
H&E (RGB) SOX (RGB)
                           Melan (RGB)
convert to convert to
                       convert to
brightness brightness
                       brightness
(L/Y/V) (L/Y/V)
                           (L/Y/V)
     \_____ stack as 3-ch ____/
         (L_HE, L_SOX, L_MEL)
           [Pretrained CNN]
            Patch Vectors
      [MIL Attention across patches]
             Case Vector
```

#### 2. Multi-Stain - Overview

- Current baseline: Prior MIL used a single stain (H&E)
  - Risks missing complementary information available in different stains
  - Robustness issue when H&E is noisy (less serious when used case-level)
- Our upgrade: MIL + Cross-stain
  - Build three parallel stain branches (H&E / SOX / Melan)
  - Two ways to implement this
    - Option A: Each branch has its own MIL attention
      - Outputs per-stain case-level embeddings
    - Option B: Each branch does feature extraction only
      - Then, patch embeddings "fused" together; single MIL attention for final case vector

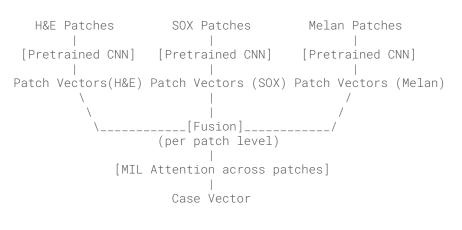
#### 2. Multi-Stain - Architecture

# Option A: Per-stain Case Embeddings



(Three parallel MIL pipelines)

# Option B: Per-stain Patch-Level Features



(Cross-stain feature fusion; requires matching patches)

# 2. Multi-Stain - Comparison of Two Options

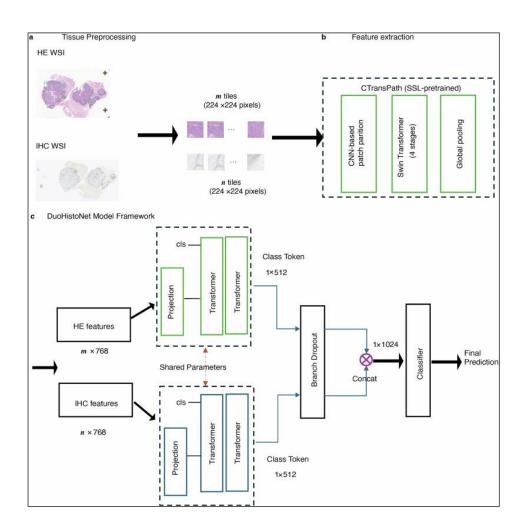
Method	Option A: Per-stain Case Embeddings	Option B: Per-stain Patch-Level Features
Implementation	MIL attention in each branch	MIL attention after per-stain feature extraction
Advantages	<ul> <li>Robust to patch mismatches</li> <li>Robust when a stain is missing</li> <li>Interpretable per-stain attention maps</li> </ul>	- Captures cross-stain interaction for a specific cell area - Efficient: only one MIL attention module
Disadvantages	- Cross-stain interactions captured only after case summarization - Heavy (3 * MIL), potential redundancy	<ul> <li>Not robust to patch mismatches</li> <li>Less per-stain interpretability (no per-stain attention map)</li> </ul>

#### 2. Multi-Stain - Literature Review

- Foersch et al., Nature Medicine (2023) <u>Link</u>
  - Built a multi-stain deep learning model using H&E + multiple IHC stains for colorectal cancer.
  - Does not implement MIL attention; uses gated attention instead
    - Otherwise similar to *Option B*: patch-level feature extraction and fusion
  - Multi-stain architecture improves...
    - Accuracy: better accuracy than using one single stain
    - Interpretability: attention weights aligned with known biology (e.g., high weights for certain cell features).

# 2. Multi-Stain -Literature Review

- Cheng et al., Communications Medicine (2025) (DuoHistoNet) <u>Link</u>
  - Dual-branch, per-stain transformer
    - patch-level feature extraction and attention-based MIL aggregation
    - Each branch outputs a case-level embedding
    - Resembles Option A
  - Outputs from the two branches are concatenated and fed into a case-level classifier
  - Result: Significantly outperformed single-stain models

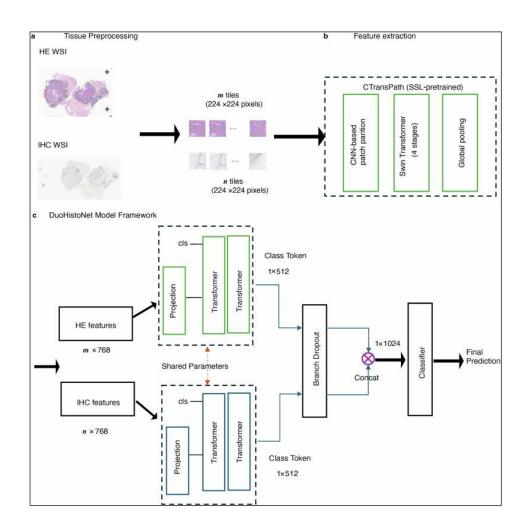


# 2. Multi-Stain - Literature Review

Cheng et al., Communications Medicine (2025) (DuoHistoNet) *Link* 

#### Other highlights of this model

- Transformer aggregator (MHSA):
   authors of this article claim that this is
   superior to other MIL approaches
- Each tile is passed through a CNN +
   Swin Transformer hybrid base model first
  - 768-dim feature embedding
- Then, per-stain MIL



#### 2. Multi-Stain - Discussion

- Should we preserve stain-specific MIL attention (option A), or should we apply MIL attention on cross-stain feature embeddings?
- Do we expect synergy mainly at the patch level (feature level) or the case level?

# 3. Image Size – Current Approaches

#### Adaptive Pooling (without resize)

- Use ResNet built-in adaptive
   pooling layer with batch size = 4
  - Only worked well for Sox10 stain (recall of 0.88)
- Increasing batch size from 4 to 8 decreases validation performance volatility and improves performance

#### Resizing

- Images are resized to 224x224 and can distort information
- Use built-in ResNet adaptive pooling layer
  - Consistent results for all 3 stains, but
     Sox10 still the strongest
- Resizing + adaptive pooling + increased batch size shows most promising results

Source: Veer's Spring 2025 Findings

# 3. Image Size – Proposed Solutions

#### Areas to explore

- Continue testing additional combinations of image resizing methods
  - o adaptive pooling, resizing, cropping, random horizontal flip

- Test higher batch sizes fed into the model given computational restraints
  - Ex. batch size = 16 or 32 for KimiaNet and ResNet

- Minimize padding (currently number buckets = batch size)
  - Higher number of buckets and lower batch size had best results (David and Veer)
  - Ideas: Number of buckets follow distribution of image sizes, find optimal buckets given computational restraints

# 4. Training, Testing, & Validation Sets

#### **Problems**

- Training set might not have all categories / patterns of high-grade patches
- We don't have a categorization of high-grade patterns to refer to in stratifying the train set

#### **Proposed Solutions**

- Calculate variation in accuracy in under-performing cases when different cases are in the training set
- Develop categorization to ensure all high-grade patterns are present in training set

# 4. Training, Testing, & Validation Sets

#### **Proposed method:**

**Step 1:** Identify which cases underperform (are misclassified often)

→ Run experiment using ~8-fold CV and 2-3 different seeds to identify which cases' classification success vary by training set

**Step 2:** Extract patch embeddings from model and cluster high-grade patches to find distinct pattern types

→ Use k-means or HDBSCAN for clustering

Step 3: Identify which clusters patches in underperforming cases fall into

Step 4: Oversample or add similar cases in training set accordingly

#### References

Madusanka, N., Jayalath, P., Fernando, D., Yasakethu, L., & Lee, B.-I. (2023). Impact of H&E Stain Normalization on Deep Learning Models in Cancer Image Classification: Performance, Complexity, and Trade-Offs. Cancers, 15(16), 4144. <a href="https://doi.org/10.3390/cancers15164144">https://doi.org/10.3390/cancers15164144</a>

Bishnoi, Vidhi, and Nidhi, Goel. (2023). A Color-Based Deep-Learning Approach for Tissue Slide Lung Cancer Classification. Biomedical Signal Processing and Control. <a href="https://www.sciencedirect.com/science/article/abs/pii/S1746809423005840">www.sciencedirect.com/science/article/abs/pii/S1746809423005840</a>

# **Questions for Krish on Tuesday**

- Did the MIL group use the resized images? If so, why? (does their model mandate that?) Update: because they used densenet. Did DeIT or Kimeanet seem as promising? <u>Yes, resized to 224\*224</u>
- Total presentation time <u>"Time doesn't matter"</u>
- Can we get a categorization from doctors? Or get them to vibe check our unsupervised clustering results
  - <u>He will ask (i don't think we can get the categories from the doctors)</u>
  - He suggested that we test whether changing the training set, for example by using a different combination of cases, would affect the model's performance, particularly the accuracy on a specific case.
    - For example, if model performance on case #4 (in the test set) is poor, see if changing the combination of the training set will boost/worsen the performance on case #4
  - <u>Unsupervised clustering might be a good approach, but what it does is finding the difference between cases, rather than finding what features/patterns the model struggles with</u>