

AI- ASSISTED ANNOATIONS FOR HISTOLOGY PRACTICALS

Sharmila Saran Rajendran, Helen Christian, Mary McMenamin, Damion Young, Rumyana Smilevska, Jon Mason

Context & Importance

- Studying cells and tissues core part of Year 1 pre-clinical medicine & biomedical science
- Understanding microscopic tissue structures is key to learning cellular structure/function



Current Tools: CSlide

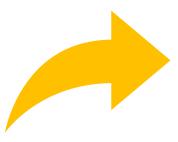
- A digital platform developed by Medical Sciences Division's Learning Technologies team
- Provides remote access to scanned histological slides
- Enables students to pan, zoom, & manually annotate regions of interest
- Supports distance learning, revision & independent study



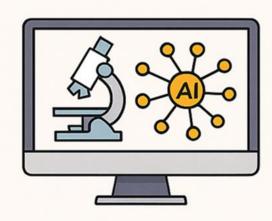
Challenges identified

- Difficult to locate labelled areas on large, high-resolution slides
- Manual annotation is timeconsuming and heavily reliant on expert input
- Limits scalability and flexibility in creating annotated resources





Objectives



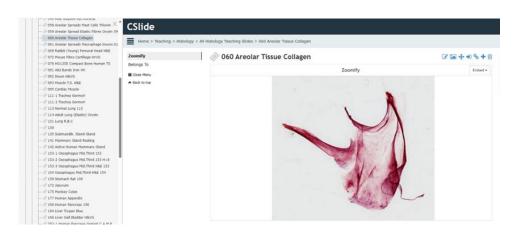
- Enhance histology teaching & learning for Year 1 medical and biomedical science students
- Leverage artificial intelligence to support the exploration & annotation of digital histological slides

Automate key processes:

- · Cell identification
- Segmentation
- Clustering
- Labelling
- Develop sustainable, scalable tool adaptable to different tissue types & educational contexts.

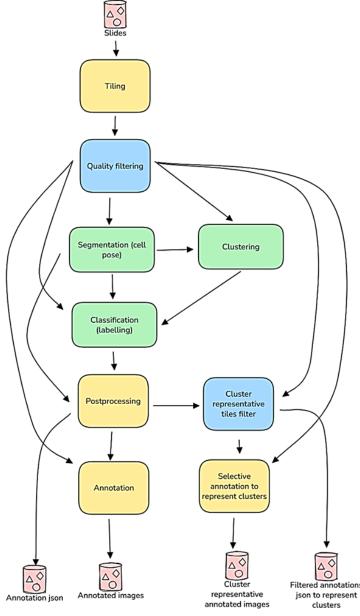
Data sources & Methodologies

Data set source Existing digital histology
slide collection
accessible through
CSlide platform, is key
focus for annotation
work.

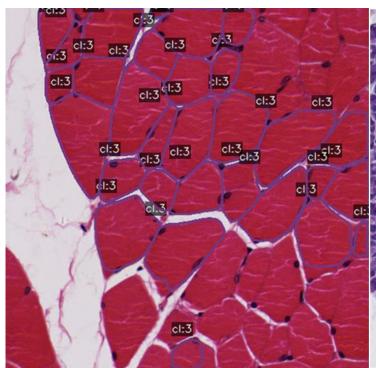


Manually annotated digital slides – A limited set of these already available served as a benchmark for qualitatively evaluating performance of AI model.



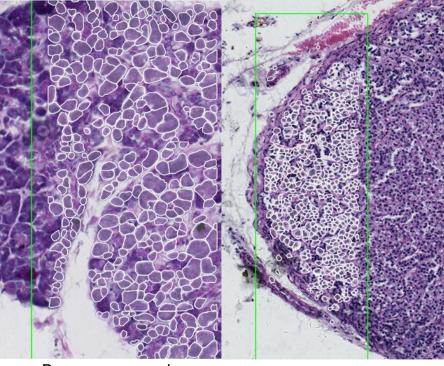


Examples of models use for segmentation



Skeletal muscle - excellent

Al segmentation model successfully delineated individual muscle fibres and each segmented fibre is assigned a cluster ID [cl:3], indicating that the model has grouped these muscle fibres into a single morphological or cluster.



Pancreas - good

fine. smooth Shows segmentation outlines around nearly all visible cluster of cells. Acinar cells, which are densely packed, sometimes irregular shape, clearly distinguished, demonstrating strong model generalization.

Adrenal gland - good

Segmentation model effectively identified cell boundaries in regions where cells are larger, more isolated, & well-spaced—typical of glomerulosa & fasciculata. These segmented cells are uniform in size & distribution, consistent with adrenal cortical cells, which tend to have clear borders & rounded nuclei

Thyroid gland - fair

A thyroid gland image showed partial success, primarily detecting individual cuboidal follicular epithelial nuclei around colloid-filled follicles. In regions with flattened or low cuboidal cells some were missed where contrast is minimal.

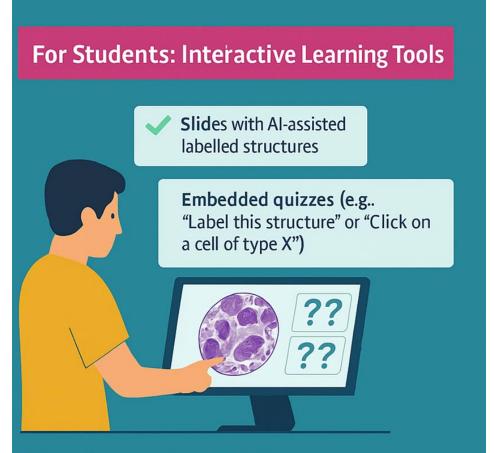
Pituitary gland - poor

Polygon outlines
generated around areas
that appear empty or
sparsely structured; no
obvious cellular
structures detected by the
model in a region known
to be rich in nerve fibers
and supporting (glial) cells
–posterior pituitary gland

How Trained AI Model Will be Used in CSlide V2



- Enhancing detection and grouling of similar cells/structures
- Testing model selection for various tissue types
- Exploring integration of the Al pipeline into CSlidide V2 to support large-scale slide processing
- Managing compute and cost challenges



Future enhancements may include:

- Arrows and interactive lists to guide user through slide content
- Self-assessment toois such as "Click on a cell of type X" or "Identify this highlighted cell," using Al-generated polygons and expert labelling to support active learning