

Prostate Cancer Diagnosis and quantification using Al-enabled Software (SW)

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Background

Gleason scoring system is widely used in prostate cancer grading. Cancer volume is also known to predict patient's outcome and is used for decisionmaking process [1] Accurate quantification of each Gleason Pattern (GP) is important and is demanded in pathology practice. [2,3] There are some challenges:

- > Evaluation of prostate biopsy slides is timeconsuming
- \succ High inter-observer variability [4,5]
- > Discordance of diagnosing a minor component of GP5 in prostate biopsy is reported at 48.7% between general pathologists an expert GU pathologists [6]

Our goal: Develop a universal and standardized platform for Gleason grading and GP quantification trained by GU pathologists to achieve accurate and reproducible diagnosis

Materials

1000 H&E prostate biopsy slides from the University of Wisconsin-Madison pathology archive were scanned with Aperio CS2 (Leica) at 40x.

- Slides were split into training set (800) and test set (200).
- Training slides were annotated by GU pathologist
- Balanced dataset of varied morphologies, including GP3, GP4, GP5 cancer, high-grade prostatic intraepithelial neoplasia (HGPIN), perineural invasion (PNI), vessels and lymphocytes

Disclosures

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Methods

> Deep Convolutional Neural Networks

- ➢Hybrid Architecture optimized for Grading
- Combination of Classification and Segmentation Networks
- \succ Multiple nets:
 - ➢Gland segmentation, Epithelial detection, Glandbased and nuclei-based grading.
- \succ Multi-scale model: multiple patch sizes at 5x to 40x resolution to capture nuclear detail as well as glandular context



Fine-Tuned Model

Sensitive to very small amount of high-grade cancer Intelligent data selection for training >ensures balanced learning across various pattern types within and across imbalanced labels >Annotation Assistant: Pathologist only needs to review

<5% of the data for annotations

			Group 1	Group 2	Group 3	Group 4	Grou
			<= 6	3 + 4	4 + 3	8	9 - 1
	Group 1	<= 6	29	2	0	0	0
	Group 2	3+4	2	64	1	0	0
	Group 3	4 + 3	0	0	21	0	0
	Group 4	8	0	0	2	20	1
	Group 5	9 - 10	0	0	1	0	57



Summary

- > Deep learning enabled cancer-grading software offers objectivity, greater efficiency and precision in prostate cancer scoring and quantification
- > Potential to help pathologists to minimize inter-observer variability and to increase efficiency and accuracy in their practice

References

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- 4. Allsbrook, W.C., Jr., et al., Interobserver reproducibility of Gleason grading of prostatic carcinoma: urologic pathologists. Hum Pathol, 2001. 32(1): p. 74-80.
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Software Predicted Grade

Results





SW assisted diagnosis

Time	<1 min		
Accuracy	95+%		
Reproducibility	High		
Quantification	Precise		

