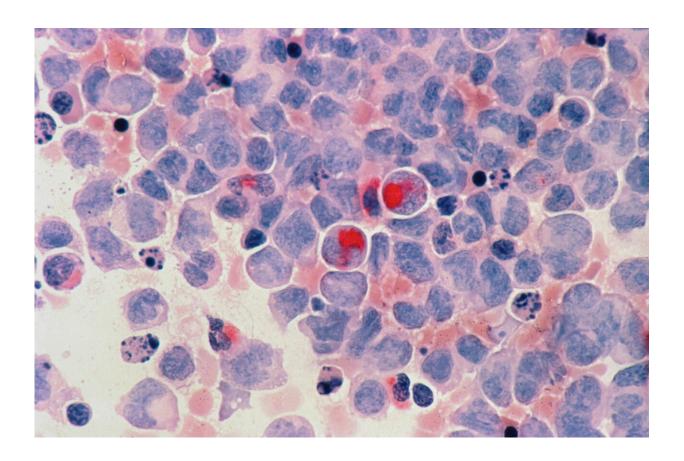


Researchers develop AI foundation models to advance pathology

March 19 2024



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Foundation models, advanced artificial intelligence systems trained on large-scale datasets, hold the potential to provide unprecedented advancements for the medical field. In computational pathology (CPath),



these models may excel in diagnostic accuracy, prognostic insights, and predicting therapeutic responses.

Researchers at Mass General Brigham have designed two of the largest CPath foundation models to date: UNI and CONCH. These foundation models were adapted to over 30 clinical and diagnostic needs, including disease detection, disease diagnosis, organ transplant assessment, and rare disease analysis.

The new models overcame limitations posed by <u>current models</u>, performing well not only for the clinical tasks the researchers tested but also showing promise for identifying new, rare and challenging diseases. Papers on <u>UNI</u> and <u>CONCH</u> have been published in *Nature Medicine*.

UNI is a foundation model for understanding <u>pathology</u> images, from recognizing <u>disease</u> in histology region-of-interests to gigapixel whole slide imaging. Trained using a database of over 100 million tissue patches and over 100,000 whole slide images, it stands out as having universal AI applications in anatomic pathology.

Notably, UNI employs transfer learning, applying previously acquired knowledge to new tasks with remarkable accuracy. Across 34 tasks, including cancer classification and organ transplant assessment, UNI outperformed established pathology models, highlighting its versatility and potential applications as a CPath tool.

CONCH is a foundation model for understanding both pathology images and language. Trained on a database of over 1.17 million histopathology image-text pairs, CONCH excels in tasks like identifying rare diseases, tumor segmentation, and understanding gigapixel images. Because CONCH is trained on text, pathologists can interact with the model to search for morphologies of interest.



In a comprehensive evaluation across 14 clinically relevant tasks, CONCH outperformed standard models and demonstrated its effectiveness and versatility.

The research team is making the code publicly available for other academic groups to use in addressing clinically relevant problems.

"Foundation models represent a new paradigm in medical artificial intelligence," said corresponding author Faisal Mahmood, Ph.D., of the Division of Computational Pathology in the Department of Pathology at Mass General Brigham.

"These models are AI systems that can be adapted to many downstream, clinically relevant tasks. We hope that the proof-of-concept presented in these studies will set the stage for such self-supervised models to be trained on larger and more diverse datasets."

More information: Lu MY et al. A visual-language foundation model for computational pathology. *Nature Medicine* DOI: 10.1038/s41591-024-02856-4

Chen, RJ et al. Towards a general-purpose foundation model for computational pathology. *Nature Medicine* DOI: 10.1038/s41591-024-02857-3

Provided by Mass General Brigham

Citation: Researchers develop AI foundation models to advance pathology (2024, March 19) retrieved 28 April 2024 from



https://medicalxpress.com/news/2024-03-ai-foundation-advance-pathology.html

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