

New AI method may boost Crohn's disease insight and improve treatment

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High magnification micrograph of Crohn's disease. Biopsy of esophagus. H&E stain. Credit: Nephron/Wikipedia

Scientists have developed a computer method that may help improve understanding and treatment of <u>Crohn's disease</u>, which causes inflammation of the digestive tract.



The Rutgers-led study, published in the journal *Genome Medicine*, used artificial intelligence to examine genetic signatures of Crohn's in 111 people. The method revealed previously undiscovered genes linked to the disease, and accurately predicted whether thousands of other people had the disease.

"Our method is not a clinical diagnosis tool, but it generates interesting observations that need to be followed up," said senior author Yana Bromberg, an associate professor in the Department of Biochemistry and Microbiology at Rutgers University-New Brunswick. "Further <u>experimental work</u> could reveal the molecular reasons behind some forms of Crohn's disease and, potentially, lead to better treatments of the disease."

Crohn's affects up to 780,000 people in the United States, the study notes. Chronic inflammation may occur in any part of the gastrointestinal tract, although symptoms may occur elsewhere. Crohn's can also cause joint pain and skin problems, and children with the disease may have growth problems, according to the U.S. National Library of Medicine.

The scientists evaluated genetic variants in the 111 people, including 64 with Crohn's disease, and used <u>artificial intelligence</u> techniques to pinpoint genes whose functions changed more in Crohn's patients than in healthy people, and vice versa. While the model's accuracy may improve by including more people, it could help reveal the origins of Crohn's and improve <u>early diagnosis</u> and accuracy, the study says.

"We believe that we can use the knowledge gained from this study to similarly model other genetically linked diseases," said Bromberg, who works in the School of Environmental Sciences and School of Arts and Sciences.



More information: Yanran Wang et al, Identifying Crohn's disease signal from variome analysis, *Genome Medicine* (2019). <u>DOI:</u> <u>10.1186/s13073-019-0670-6</u>

Provided by Rutgers University

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