

Tissue remodeling as a primary mechanism for diverticula formation

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Colonic diverticulosis is a prevalent condition among older adults, marked by the presence of thin-walled pockets in the colon wall that can



become inflamed and infected; they can then hemorrhage or rupture. A <u>new study</u>, published in the journal *Gut* now suggests that tissue remodeling is a primary mechanism for diverticula formation.

The study, led by corresponding author Anne Peery, MD, associate professor of medicine in the Division of Gastroenterology and Hepatology at the UNC School of Medicine, aimed at identifying the genetic and cellular determinants underlying colonic diverticulosis and the relationship with other gastrointestinal disorders.

Researchers conducted DNA and RNA sequencing on colonic tissue from 404 patients. Results showed 38 genes with differential expression and 17 with varied transcript usage linked to diverticulosis. Additionally, diverticulosis severity was positively correlated with <u>genetic</u> <u>predisposition</u> to diverticulitis.

Peery and colleagues linked the formation of diverticula to stromal and <u>epithelial cells</u> in the colon, particularly <u>endothelial cells</u>, myofibroblasts, fibroblasts, goblet, tuft, enterocytes, neurons, and glia. The researchers also analyzed genes to find fiveâ€"CCN3, CRISPLD2, ENTPD7, PHGR1 and TNFSF13â€"that showed potential causal effects on diverticulosis. Interestingly, the researchers confirmed that the gene ENTPD7 was over-expressed in diverticulosis cases.

In addition, the study showed that individuals with an increased genetic proclivity to diverticulitis show larger numbers of diverticula upon colonoscopy.

More information: Jungkyun Seo et al, Genetic and transcriptomic



landscape of colonic diverticulosis, *Gut* (2024). DOI: <u>10.1136/gutjnl-2023-331267</u>

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