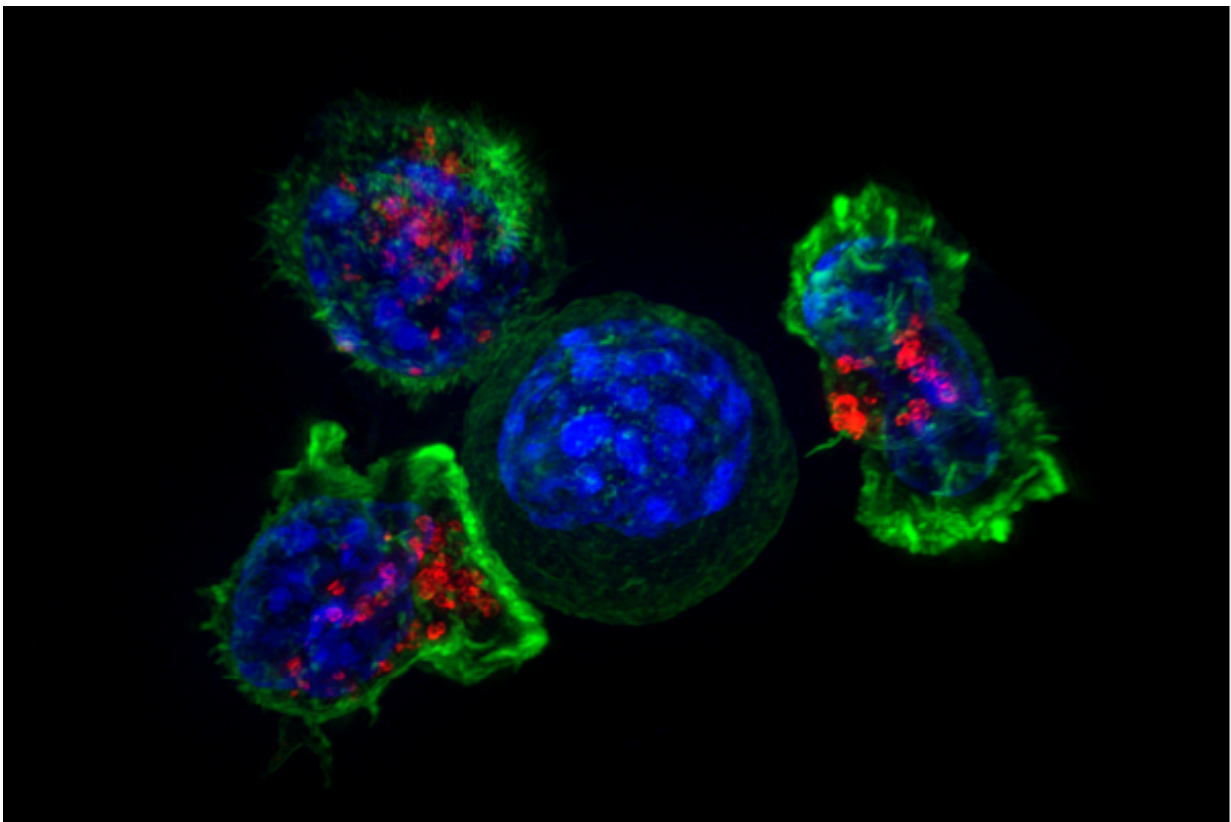


# Researchers identify breast microbiome/bacterial differences between healthy and cancerous tissue

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Killer T cells surround a cancer cell. Credit: NIH

A team of Mayo Clinic researchers has identified evidence of bacteria in sterilely-obtained breast tissue and found differences between women

with and without breast cancer. The findings are published in the Aug. 3 issue of *Scientific Reports*.

"Our research found that breast tissue samples obtained in the operating room under sterile conditions contain bacterial DNA, even when there is no sign of infection. Furthermore, we identified significant differences in the breast tissue microbiome of women with cancer versus women without cancer," says Tina Hieken, M.D., a breast surgical oncologist at Mayo Clinic. "Our work confirmed the presence of a distinct [breast tissue](#) microbiome and that it is different than the microbiome of the overlying breast skin."

According to the National Cancer Institute, a microbiome is a collection of microorganisms and viruses that live in a specific environment in the human body.

Dr. Hieken says, globally, [breast cancer](#) accounts for nearly one quarter of all cancers and is the leading cause of cancer death among women. She says that while there are established risk factors for breast cancer, at least 70 percent of breast cancer cases occur in women of average risk, and current prediction models are poor at identifying risk for individual women.

"Differences in the microbiome have been implicated in cancer development at a variety of body sites including stomach, colon, liver, lung and skin," says co-investigators Amy Degenim, M.D. a breast surgical oncologist at Mayo Clinic. "There is mounting evidence that changes in the breast microbiome may be implicated in cancer development and the aggressiveness of cancer and that eliminating dangerous microorganisms or restoring normal microbiota may reverse this process," adds Nick Chia, Ph.D., a microbiome researcher at Mayo Clinic.

Dr. Hieken says it remains unclear whether small shifts in microbial communities, the presence of a virulent pathogenic strain or the absence of a beneficial one might be responsible for promoting the development of cancer in the breast microbiome. However, she says the findings of this study will spur further research to identify potential causes of [breast cancer](#) development and new microbial-based prevention therapies.

Provided by Mayo Clinic

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