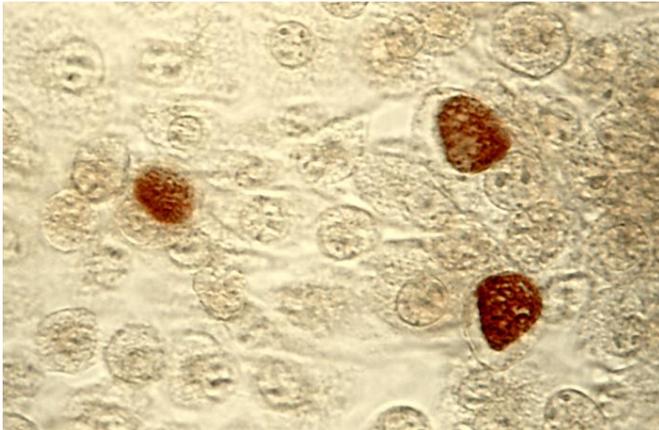


Researchers identify how vaginal microbiome can elicit resistance to chlamydia

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Chlamydia trachomatis inclusion bodies (brown) in a McCoy cell culture. Credit: public domain

The vaginal microbiome is believed to protect women against *Chlamydia trachomatis*, the etiological agent of the most prevalent sexually transmitted infections (STIs) in developed countries. New research by the University of Maryland School of Medicine (UMSOM) shows how the microbiome can either protect or make a woman more susceptible to these serious infections. The research is important amid a rising number of cases of chlamydia worldwide. In the U.S. alone, 1.7 million cases of chlamydia were reported in 2017, a 22% increase since 2013, according to data from the Centers for Diseases Control and Prevention (CDC).

"Chlamydia is a major growing health issue in the U.S., and more work is needed to understand why some women are apparently naturally protected while other are not," commented Jacques Ravel, Ph.D., Professor of Microbiology and Immunology, Associate Director and Senior Scientist at the Institute for Genome Sciences (IGS) at UMSOM.

Dr. Ravel is also a Principal Investigator for this research.

"Our novel research aims to decipher the mechanistic and functional underpinnings of communication between the host and the cervicovaginal [microbiome](#) to better understand resistance and susceptibility to this infection."

An Important Mechanism in Vaginal Microbiome

While *Lactobacillus*-dominated microbiota in a woman's vagina has long been suspected to provide a protective barrier against STIs like [chlamydia](#), investigators at IGS and the University of Maryland School of Dentistry (UMSOD) are reporting for the first time a mechanism enabling specific types of cervicovaginal microbiome to predispose cells in the vagina and cervix to resist chlamydial infection.

"We will now be able to leverage these microbiomes to identify women at risk of infections, but more importantly to develop improved strategies to restore an optimal protection when it is lacking. Unlike our genes, the [vaginal microbiome](#) can be modulated to increase protection against chlamydia, but also against other sexually transmitted infections, including HIV," said Dr. Ravel of the research, which was published today in *mBio*, "Cervicovaginal Microbiota-Host Interaction Modulates *Chlamydia trachomatis* Infection."

The investigators have shown previously that five major types of vaginal microbiome exist, four of which are dominated by a different species of *Lactobacillus*, while the fifth has very low numbers of *Lactobacillus* bacteria and is associated with an increased risk of adverse outcomes including STIs, such as HIV, and even premature births.

The current research showed that *Lactobacillus iners*, a bacterium actually commonly found in the vagina did not optimally protect [human cells](#) against chlamydial infection, while products of *Lactobacillus crispatus*, another *Lactobacillus* species frequently found in the vagina, did.

new stepping stone on which future translational research to exploit the microbiome in the fight against chlamydial [infection](#) and disease, can be based."

More information: Vonetta L. Edwards et al. The Cervicovaginal Microbiota-Host Interaction Modulates Chlamydia trachomatis Infection, *mBio* (2019). [DOI: 10.1128/mBio.01548-19](https://doi.org/10.1128/mBio.01548-19)

Previously published research has hinted at *L. iners* being a risk factor for STI; however, the mechanism by which these bacteria were specifically suboptimal at protecting women against STI has remained elusive. Like other *Lactobacillus*, *L. iners* produces lactic acid, but only the L isoform. The researchers found that D-lactic acid, not L-[lactic acid](#), down-regulates cell cycling through epigenetic modifications thus blocking *C. trachomatis* entry into the cell, one of the pathogen key infectious process, among other processes.

Provided by University of Maryland School of Medicine

Thus, a rather unexpected result of this study is that the vaginal microbiome does not affect the pathogen per se, but drives susceptibility or resistance to infection, by modifying the cells that line up the cervicovaginal epithelium. The researchers further demonstrated that exposure to optimal vaginal microbiota provided long term protection, which has major implication on how a woman is protected. These mechanisms are now being exploited to develop strategies to optimize protection against *C. trachomatis* infections but also other STIs.

Patrik Bavoil, Ph.D., Professor & Chair, Department of Microbial Pathogenesis, University of Maryland School of Dentistry, a well-known expert in *C. trachomatis* biology and pathogenesis, is a Co-Principal Investigator with Dr. Ravel on the NIH funding that supported this study. The investigators also collaborated with Larry Forney, Ph.D. at the University of Idaho.

"Chlamydia is reputed to be a most difficult microorganism to study. By hiding inside cells, the pathogen routinely avoids antimicrobial host defenses. By causing mostly asymptomatic infection, it often escapes detection by both the infected host and the physician alike," said Dr. Bavoil. "What we have done in this study through several years of hard work by dedicated researchers is to provide, for the first time, a huge,

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