

# Gut microbiome changes during pregnancy may influence immune system response

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During pregnancy, a woman's immune system changes dramatically but researchers don't yet understand all the underlying mechanisms. A new study shows how the gut microbiota may play a role.

In a paper published in *mSystems*, researchers in China report that during pregnancy, changes in levels of cytokines—[immune system proteins](#) important in inflammation—may be linked to specific alterations in the mother's gut microbiome and in plasma and fecal metabolites.

"To the best of our knowledge, these associations were first explored in our study," said first author Ting Huang, M.D., from the First Affiliated Hospital of Jinan University in Guangzhou, China.

Pregnancy brings a raft of changes, including fluctuations in hormones, changes to a woman's body structure, and variations in the immune system.

Previous studies have identified changes to the gut microbiome that can occur during pregnancy; they have also suggested that those changes may influence physiological processes in the mother through metabolites. Disturbances in the microbiota, for example, have been connected to the promotion of preeclampsia, a dangerous pregnancy complication characterized by [high blood pressure](#). However, it remains unclear how alterations in the gut microbiota during pregnancy affect maternal immunity.

To investigate these connections, the researchers at the First Affiliated Hospital of Jinan University compared the [gut microbiota](#), [metabolite](#) profiles and immune system status of 30 healthy [pregnant women](#) to 15 healthy women who weren't pregnant.

All the women in the study were between 18 and 34 years old. Fecal and [blood samples](#) were collected from the pregnant women during or after the 37th week of pregnancy, and samples from non-pregnant women were collected on the 14th day of the menstrual cycle.

Analyses of the fecal samples showed that Firmicutes was the most

dominant phylum of bacteria in both groups of women. However, Bacteroidota bacteria accounted for a smaller relative share of the microbial population in pregnant women than in non-pregnant women. Pregnant women also showed a higher relative abundance of both Actinobacteriota and Proteobacteria, compared to non-pregnant women.

The researchers similarly found a distinct difference in the [cytokine](#) levels in the two groups. Pregnant women had lower levels of cytokines that promote inflammation, and they showed higher levels of cytokines that act against inflammation. These results suggest that the immune system may be suppressed during pregnancy, the authors noted.

The researchers then identified hundreds of metabolites found in the plasma and fecal samples. They found that each group—pregnant and non-pregnant women—had its own distinct collection of metabolites, or metabolome. Notably, many of those metabolites were connected to bile acid secretion and metabolism, and bile acids have previously been tied to inflammation.

In further analyses, they found that some enriched metabolites in pregnant women were associated with lower levels of pro-inflammatory cytokines. Similarly, some of the depleted metabolites are associated with increases in pro-inflammatory cytokines.

Finally, the group identified a total of 46 connections among microbes, metabolites and cytokines. They found that some microbes enriched in pregnant women, for example, may inhibit an immune response by inhibiting pro-inflammatory metabolites. Overall, Huang said, the results support the idea that gut microbes interact with host metabolites to change cytokine levels in the blood.

However, Huang cautioned that the study doesn't establish causation. In addition, the [small sample size](#) may amplify errors arising from

[individual differences](#), and the design of the study doesn't account for confounding factors, like diet. More [clinical trials](#) are needed, the researchers said, to confirm and better elucidate the connections.

**More information:** Ting Huang et al, *mSystems* (2024)

Provided by American Society for Microbiology

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