

Remodeling of the maternal gut microbiome during pregnancy is shaped by parity

Alexander S. F. Berry

Meghann K. Pierdon

Ana M. Misic

Megan C. Sullivan

Kevin O'Brien

Samuel J. Murray

Ying Chen

Lydia A. Ramharack

Robert N. Baldassano

Thomas D. Parsons

Daniel Beiting

Video Byte

Keywords: pig, pregnancy, parity, gut microbiome, neonate microbiome, early-life microbiota, 16S rRNA sequencing, shotgun metagenomics, Microbiome, microbiota, maternal gut, microbiome remodeling, infant microbiome, swine, *Treponema bryantii*, *Lactobacillus amylovorus*, *Lactobacillus reuteri*, *Akkermansia muciniphila*, *Prevotella stercorea*, *Campylobacter coli*

Posted Date: October 13th, 2021

DOI: <https://doi.org/10.21203/rs.3.rs-967794/v1>

License:  This work is licensed under a Creative Commons Attribution 4.0 International License.
[Read Full License](#)

Abstract

In mammals, the maternal microbiome influences both mother and infant health and can be used to predict the likelihood of preterm birth. The mother's number of previous pregnancies, or parity, can also affect preterm birth risk, but whether parity influences the maternal or infant microbiome is unclear. A recent study used marker gene sequencing and metagenomics analyses of stool samples to investigate the microbiomes of pregnant pigs of different parities and their offspring. Microbiome "maturity index" and Dirichlet multinomial mixtures (DMM) model analyses indicated that the sow microbiome changed in predictable ways throughout pregnancy and that the changes occurred more rapidly in sows with higher numbers of previous pregnancies. Parity (high vs. zero) was linked to the levels of specific types of bacteria in the maternal gut at the end of pregnancy. Sow parity also affected the abundances of certain metabolism-related bacteria in the piglet gut 10 days after birth. Although samples were not collected in early pregnancy due to difficulties in reliable pregnancy detection, the results show that pregnancy history affects the maternal and infant gut microbiomes in pigs and suggest that the influence of parity should be considered in studies on the human microbiome during pregnancy and infancy.