## Over 700 Bacteria Identified In Breast Milk

Spanish researchers have traced the bacterial microbiota map in breast milk, which is the **main source of nourishment** for newborns. The study has revealed a larger microbial diversity than originally thought: more **than 700 species.** 

The breast milk received from the mother is one of the factors determining how the bacterial flora will develop in the newborn baby. However, the composition and the biological role of these bacteria in infants remain unknown.

A group of Spanish scientists have now used a technique based on massive DNA sequencing to identify the set of bacteria contained within breast milk called **microbiome**. Thanks to their study, pre- and postnatal variables influencing the microbial richness of milk can now be determined.

**Colostrum** is the first secretion of the mammary glands after giving birth. In some of the samples taken of this liquid, more than 700 species of these microorganisms were found, which is more than originally expected by experts. The results have been published in the *American Journal of Clinical Nutrition*.

"This is one of the first studies to document such diversity using the pyrosequencing technique (a large scale DNA sequencing determination technique) on colostrum samples on the one hand, and breast milk on the other, the latter being collected after one and six months of breastfeeding," explain the coauthors, María Carmen Collado, researcher at the Institute of Agrochemistry and Food Technology (IATA-CSIC) and Alex Mira, researcher at the Higher Public Health Research Centre (CSISP-GVA).

The most common bacterial genera in the colostrum samples were <u>Weissella, Leuconostoc, Staphylococcus,</u> <u>Streptococcus</u> and <u>Lactococcus</u>. In the fluid developed between the first and sixth month of breastfeeding, bacteria typical of the oral cavity were observed, such as <u>Veillonella, Leptotrichia</u> and <u>Prevotella.</u>

"We are not yet able to determine if these bacteria colonise the mouth of the baby or whether oral bacteria of the breast-fed baby enter the breast milk and thus change its composition," outline the authors.

## The heavier the mother, the fewer the bacteria

The study also reveals that the milk of overweight mothers or those who put on more weight than recommended during pregnancy contains a lesser diversity of species.

The type of labour also affects the microbiome within the breast milk: that of mothers who underwent a planned caesarean is different and not as rich in microorganisms as that of mothers who had a **vaginal birth**. However, when the caesarean is unplanned (intrapartum), milk composition is very similar to that of mothers who have a vaginal birth.

These results suggest that the hormonal state of the mother at the time of labour also plays a role: "The lack of signals of physiological stress, as well as hormonal signals specific to labour, could influence the microbial composition and diversity of breast milk," state the authors.

## Help for the food industry

Given that the bacteria present in breast milk constitute one of initial instances of contact with microorganisms that colonise the infant's digestive system, the researchers are now working to determine if their role is metabolic (it helps the breast-fed baby to digest the milk) or **immune** (it helps to distinguish beneficial or foreign organisms).

For the authors, the results have opened up new doors for the design of child nutrition strategies that improve health. "If the breast milk bacteria discovered in this study were important for the development of the immune system, its addition to infant formula could decrease the risk of allergies, asthma and autoimmune diseases," conclude the authors.