

M. Carmen Collado

Institute of Agrochemistry and Food
Technology-National Research
Council (IATA-CSIC), Spain

Investigadora en el Departamento
de Biotecnología en el Instituto de
Agroquímica y Tecnología de los
Alimentos del Consejo Superior de
Investigaciones Científicas
(IATA-CSIC), España.

Breast milk microbiota and factors
influencing its composition.

La microbiota de la leche materna
y los factores que influyen en su
composición.



www.milkscienceconference.com

VALENCIA
24 · 25 OCTUBRE 2019



Instituto de Agroquímica
y Tecnología de Alimentos



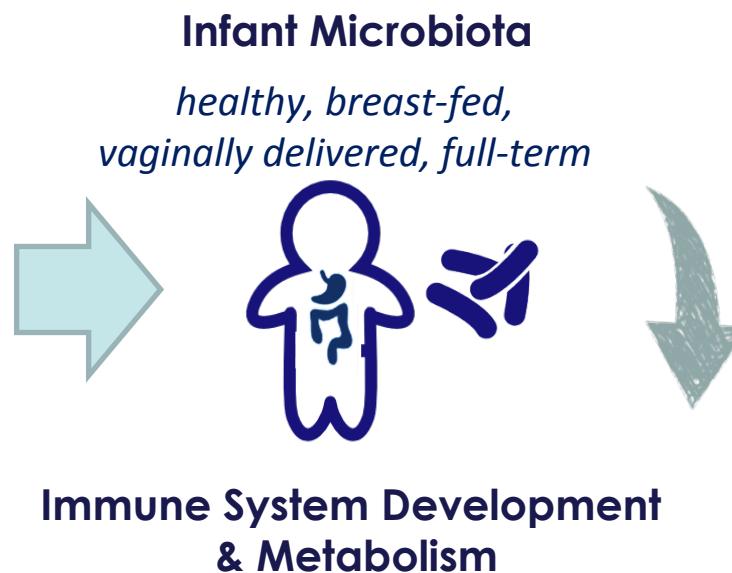
INTERNATIONAL
SCIENTIFIC CONFERENCE
ON RAW MILK

Breast milk microbiota and factors influencing its composition.

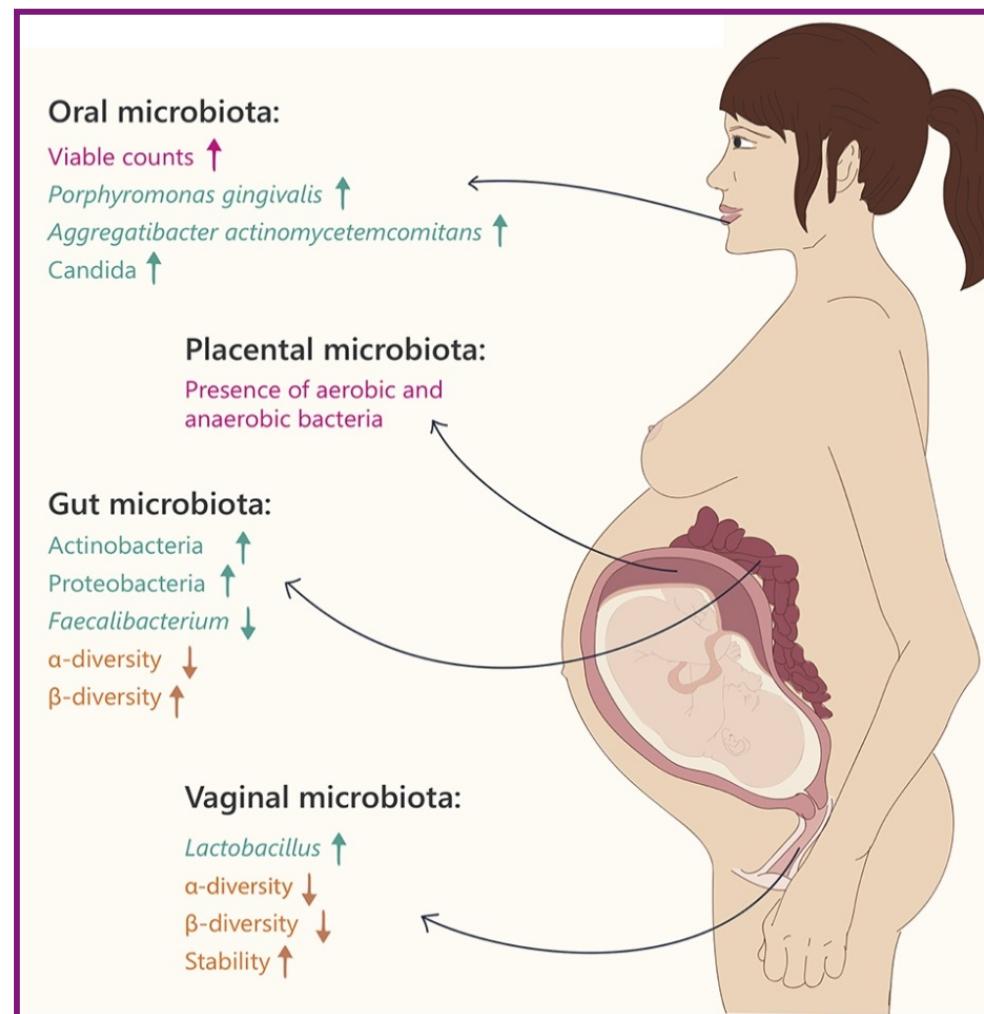
Maria Carmen Collado

6/11/19

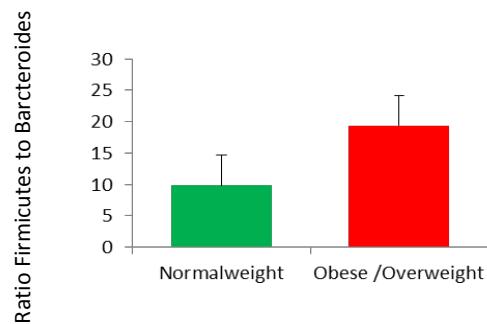
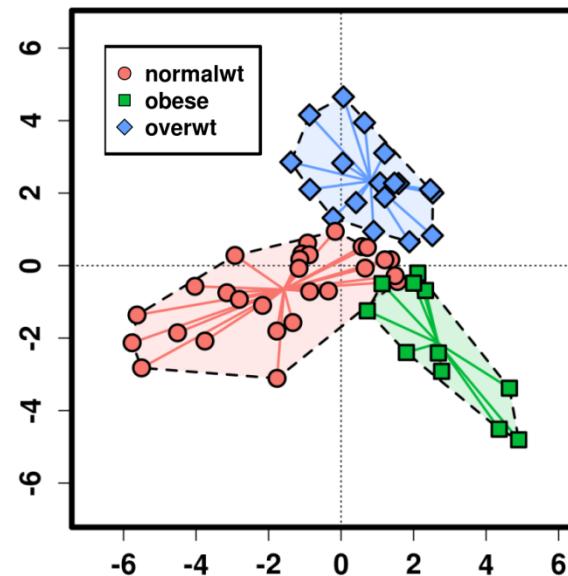
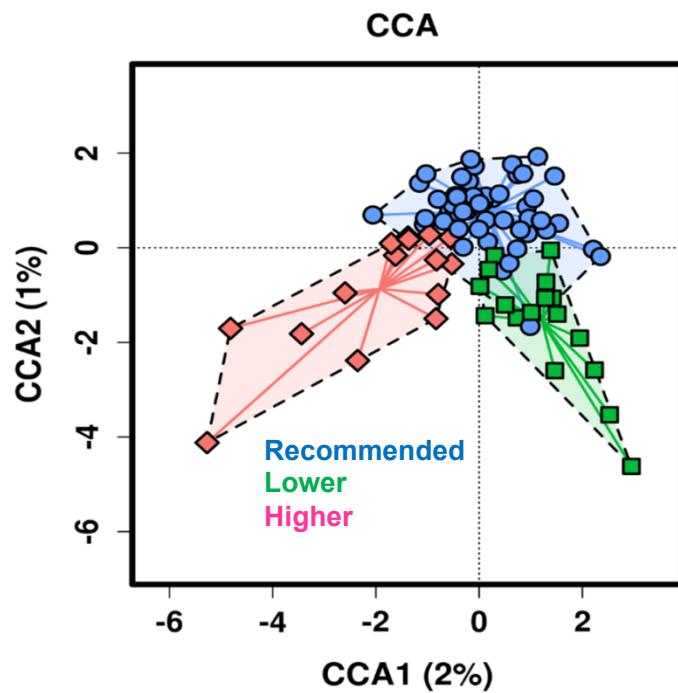
Maternal microbiota is being recognized as one of the essential factors determining maternal-child health outcomes, which would also be affected by perinatal factors



Shifts in the maternal microbiome have been implicated in metabolic adaptations to pregnancy



The maternal gut microbiota is modified by BMI and weight gain over pregnancy



Marta
Selma-Royo

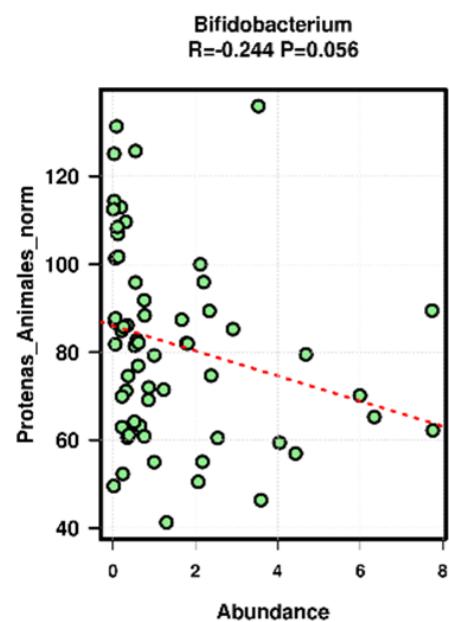
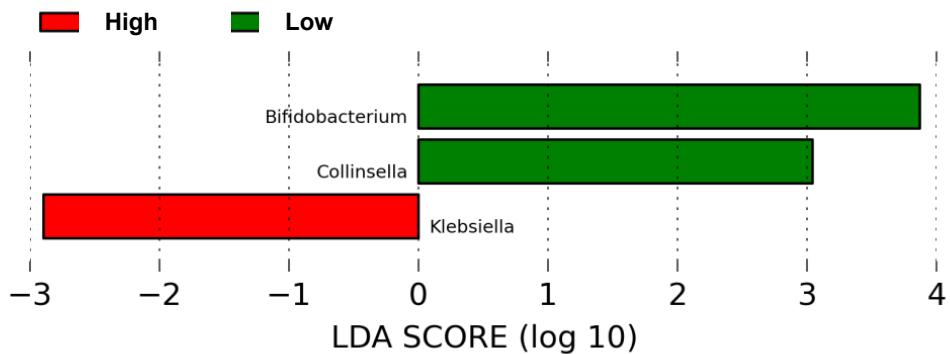
Associations with pre-gestational BMI

↓Diversity and richness ↑ BMI and weight gain

Selma-Royo et al. unpublished

The maternal gut microbiota is influenced by diet

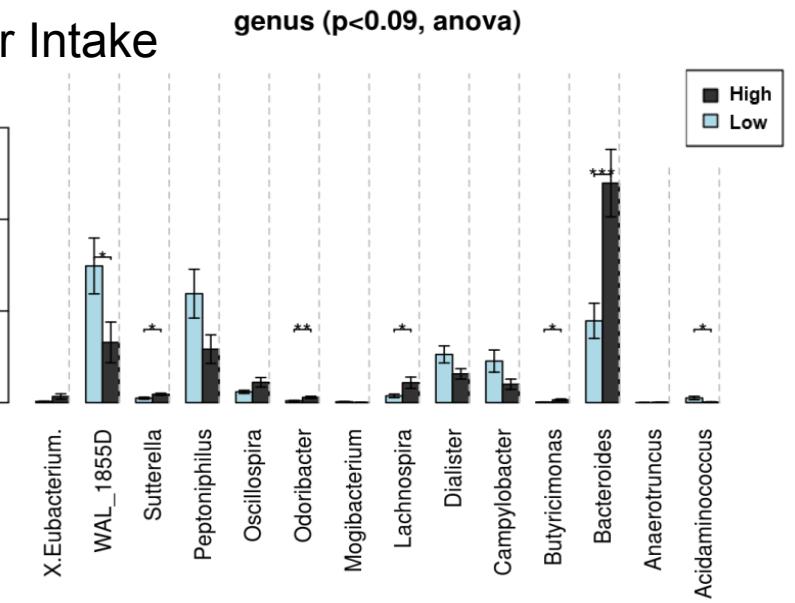
Total animal protein intake



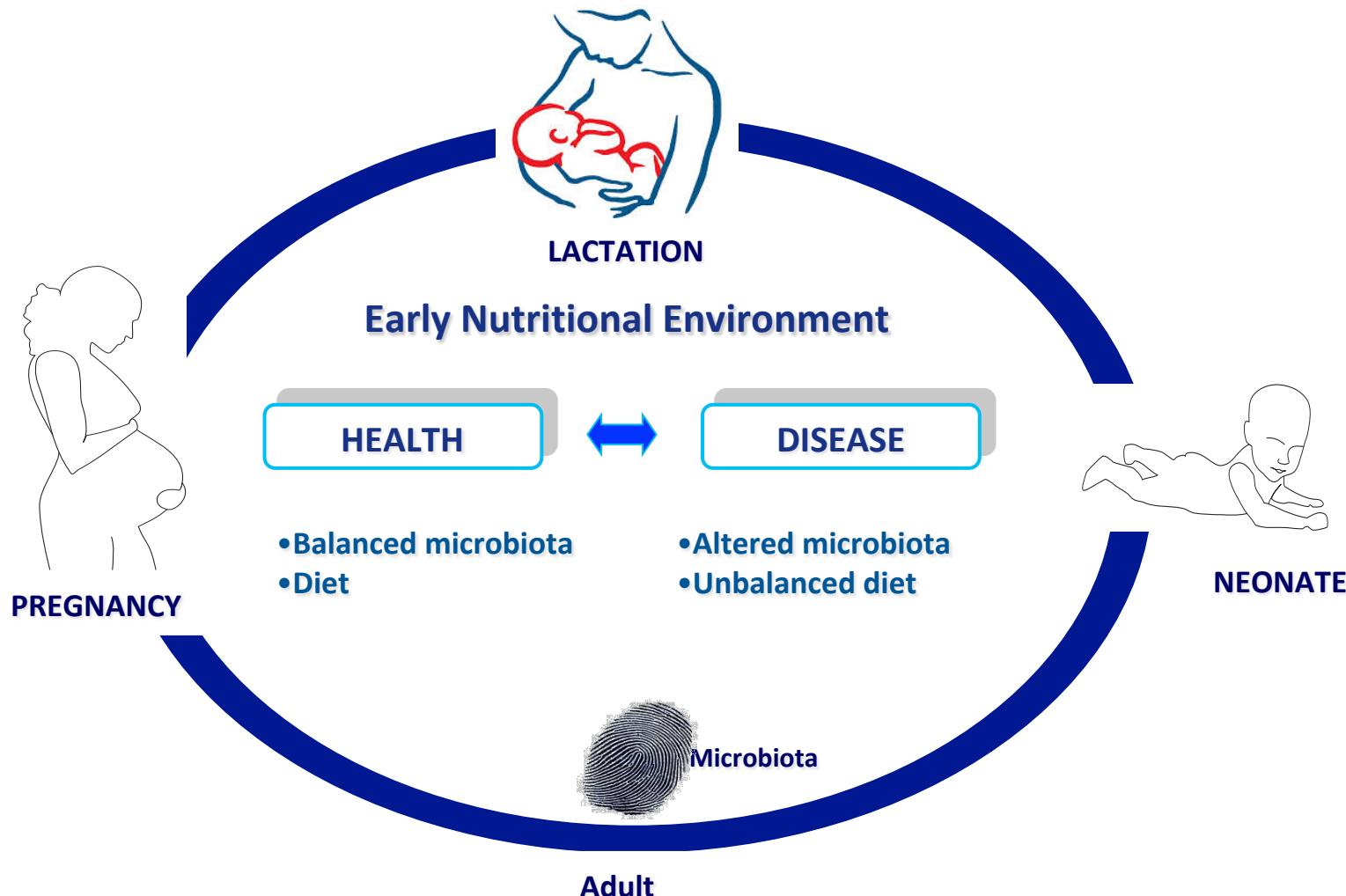
FFQ and
3d recall



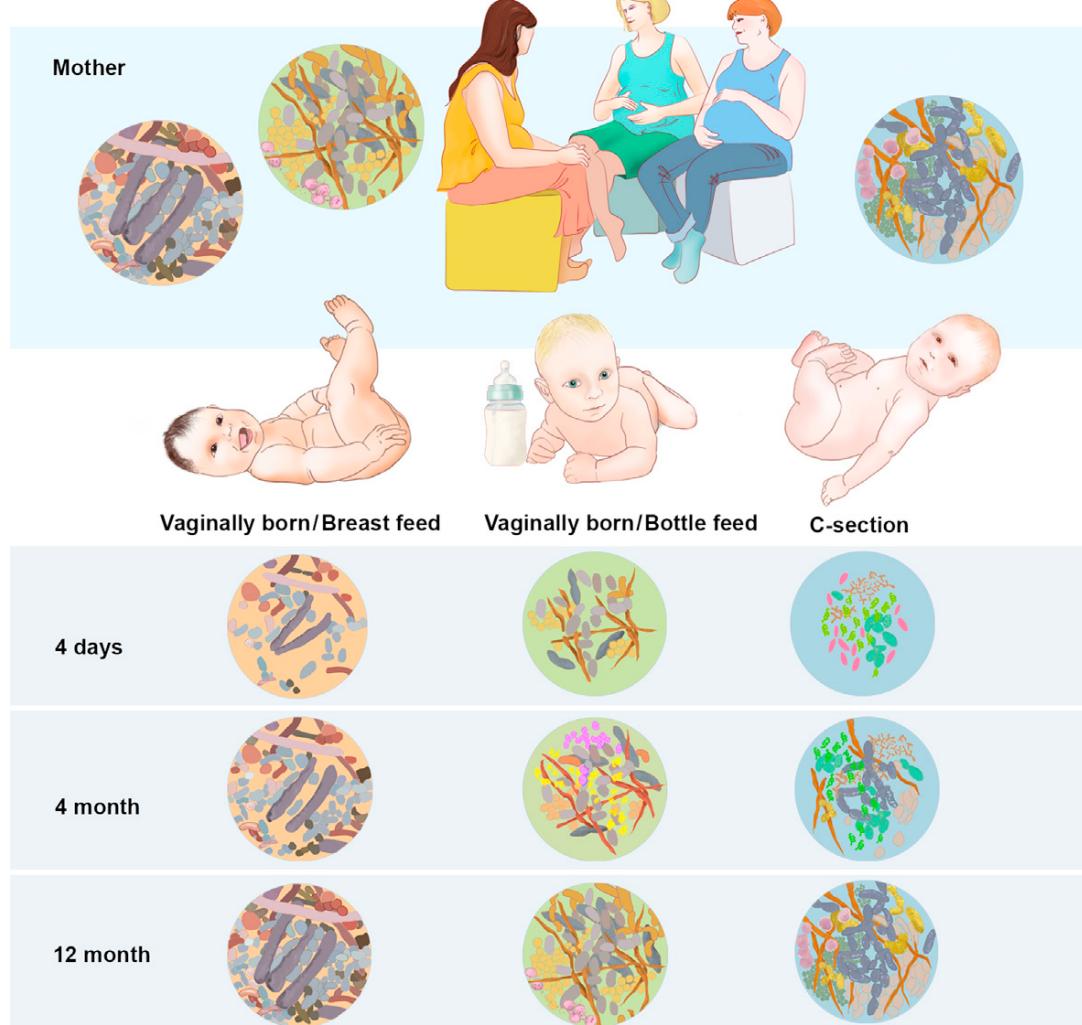
Fiber Intake



What extend influence of maternal microbial and nutritional environment to infant health?.



Mode of delivery and Breastfeeding practices: role in neonatal microbiome

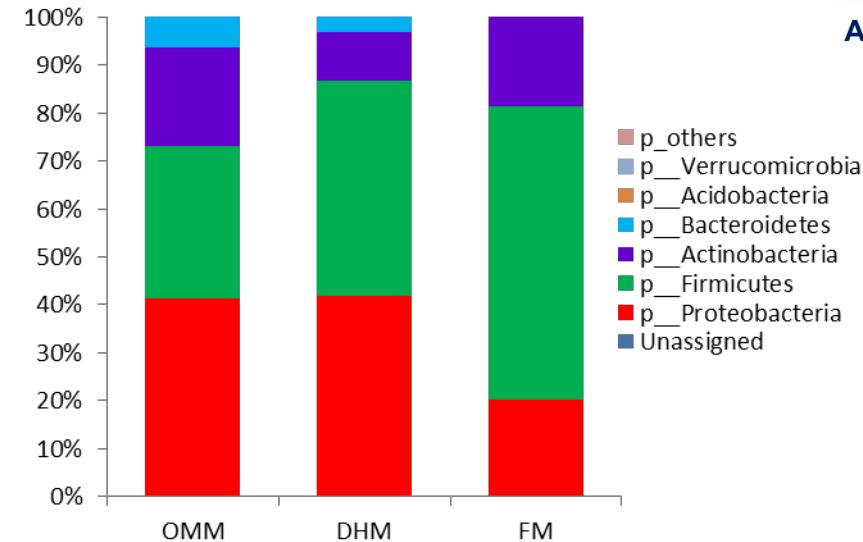
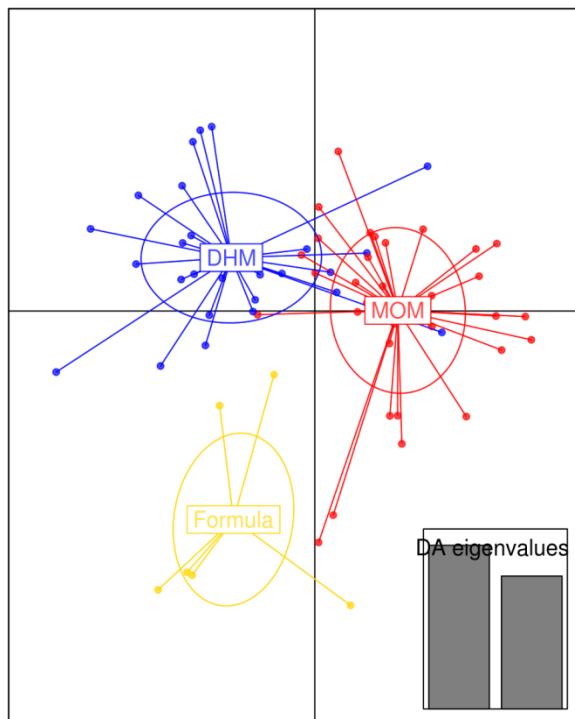


Diet shapes infant gut microbial community

Nutrition had a major impact on early microbiota composition and function, with cessation of breastfeeding, rather than introduction of solid food, being required for maturation into an adult-like microbiota.

PRETERM GUT MICROBIOME influenced by FEEDING TYPE

A prospective observational cohort study, paired, in NICU >32 weeks and birth weight ≤ 1.500 g



Anna Parra

Despite the higher variability, preterm microbiota composition and predictive functional profiles were significantly influenced by feeding type although no differences in microbial diversity and richness were found.

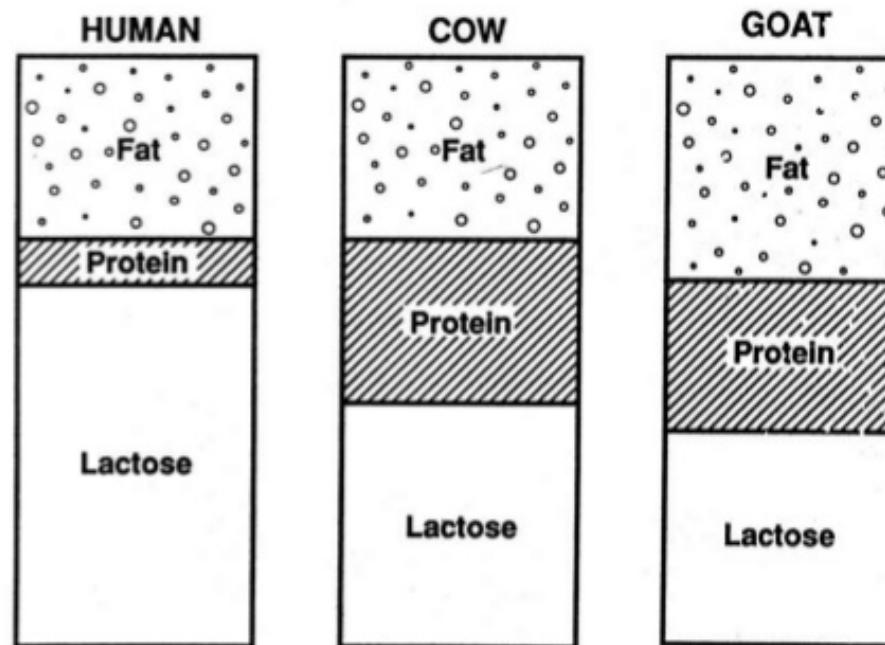
Breast milk composition: Beyond nutrition



BREASTMILK: *PERSONALIZED* INFANT NUTRITION

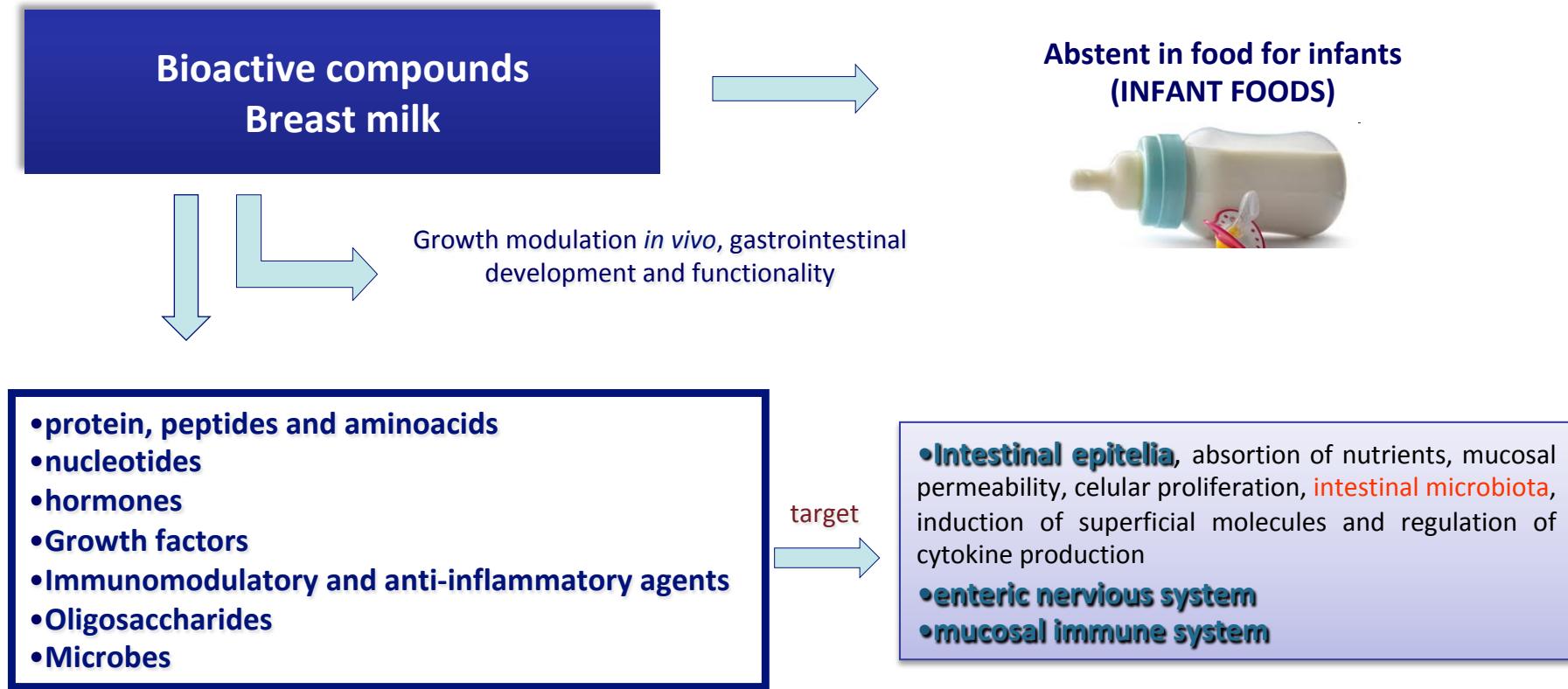
Nutrients in Human & Animal Milk

What are the differences between these milks?



Breastfeeding: What is in human milk?

BREASTMILK: *GOLD STANDARD OF INFANT NUTRITION*



Goldman, 2000; Baró et al., 2001

Human breast milk is a constant source of microbes which have an impact on infant's microbial colonization

- Culture dependent & independent-methods
- New technologies → increase the knowledge



Staphylococcus

Streptococcus

Bifidobacterium

Lactobacillus

Enterococcus

Lactococcus

Leuconostoc

Others:

Gram + : *Actinomyces* spp. / *Corynebacterium* spp./*Kocuria* spp./

Gram - : *Escherichia coli* / *Klebsiella* / *Ralstonia* spp



Structural

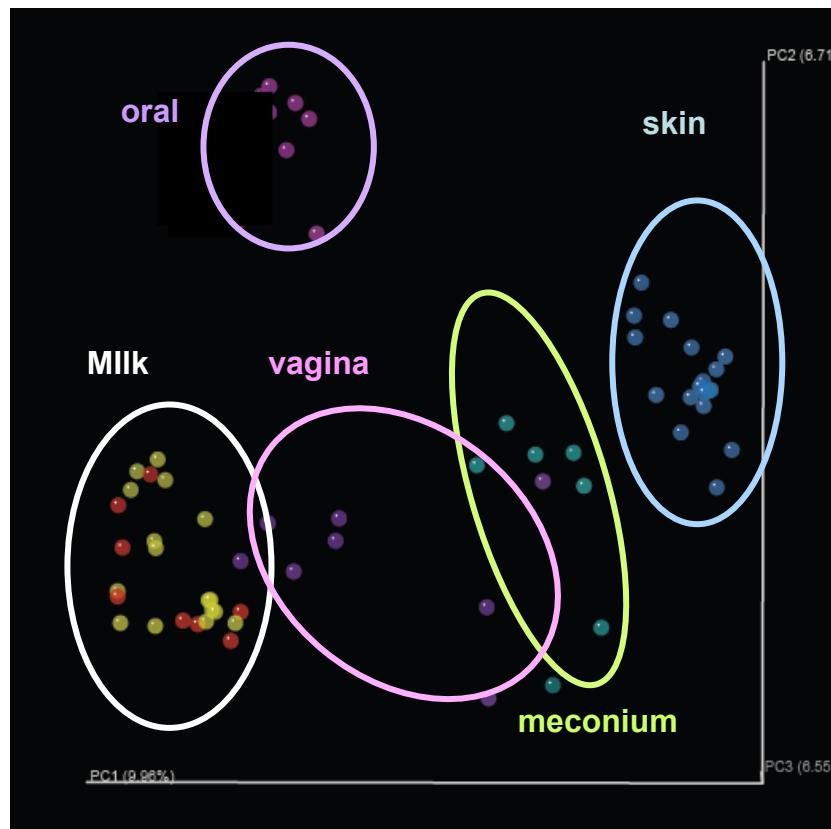
Metabolic

Protector

Breastfeeding

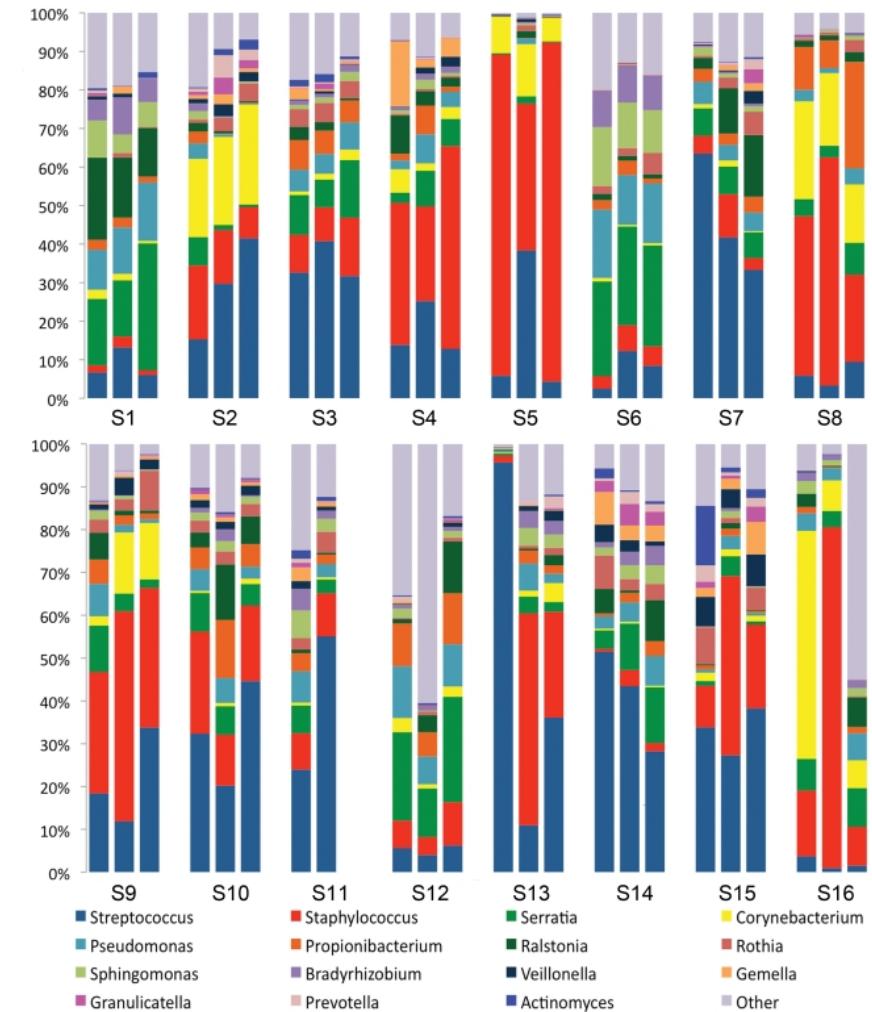
BREASTMILK: *source of microbes*

Great Variation from mother to mother



Cabrera-Rubio et al. 2011

Collado et al., 2008; 2010; 2012



Hunt et al., 2011



NIH-PA Author Man

H Public Access

Author Manuscript

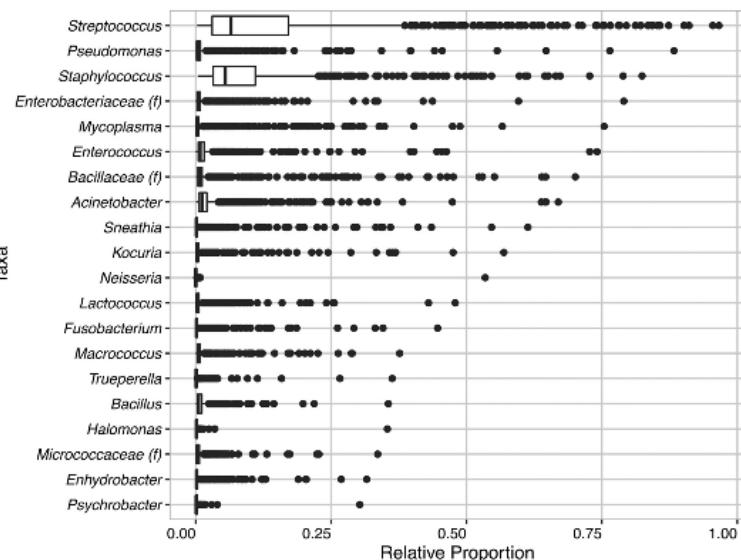
J Primatol. Author manuscript; available in PMC 2013 July 01.

Published in final edited form as:

J Primatol. 2011 February ; 40(1): 52–58. doi:10.1111/j.1600-0684.2010.00450.x.

Species diversity and relative abundance of lactic acid bacteria in the milk of rhesus monkeys (*Macaca mulatta*)

L. Jin¹, K. Hinde^{2,3}, and L. Tao¹



Journal of Dairy Research



Article Supplementary materials Metrics

Volume 83, Issue 3 August 2016, pp. 383-386

Isolation and characterisation of lactic acid bacteria from donkey milk

Maria de los Dolores Soto del Rio ^(a1), Christian Andriguetto ^(a2), Alessandra Dalmasso ^(a1), Angioletta Lombardi ^(a2) ... ⁺

DOI: <https://doi.org/10.1017/S0022029916000376> Published online: 07 September 2016

Food Microbiology



Volume 46, April 2015, Pages 121–131



Analysis of raw goat milk microbiota: Impact of stage of lactation and lysozyme on microbial diversity

Elizabeth A. McInnis^a, Karen M. Kalantra^b, David A. Mills^{b, c}, Elizabeth A. Maga^a,

^a Show more



RESEARCH ARTICLE



The Core and Seasonal Microbiota of Raw Bovine Milk in Tanker Trucks and the Impact of Transfer to a Milk Processing Facility

Mary E. Kable,^a Yanin Srisengfa,^a Miles Laird,^a Jose Zaragoza,^a Jeremy McLeod,^b Jessie Heidenreich,^b Maria L. Marco^a

Department of Food Science and Technology, University of California Davis, Davis, California, USA;^a Hilmar Cheese Company, Hilmar, California, USA^b



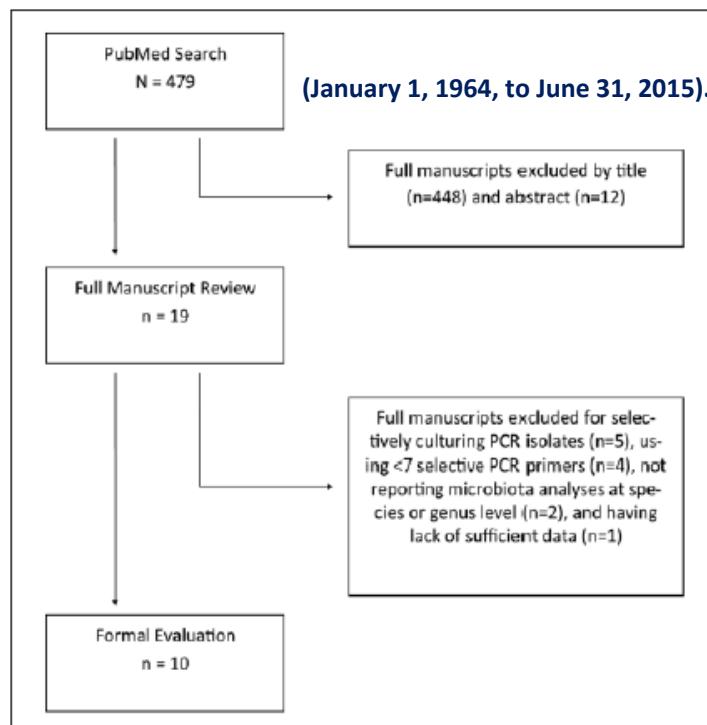
Review

Systematic Review of the Human Milk Microbiota

**John L. Fitzstevens¹; Kelsey C. Smith, MA^{2,3}; James I. Hagadorn, MD^{2,3,4};
Melissa J. Caimano, PhD^{5,6}; Adam P. Matson, MD^{3,4,7};
and Elizabeth A. Brownell, PhD^{2,3,4}**



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for Parenteral and Enteral Nutrition
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***Streptococcus* and *Staphylococcus* are the predominant genera in the human milk microbiota.**

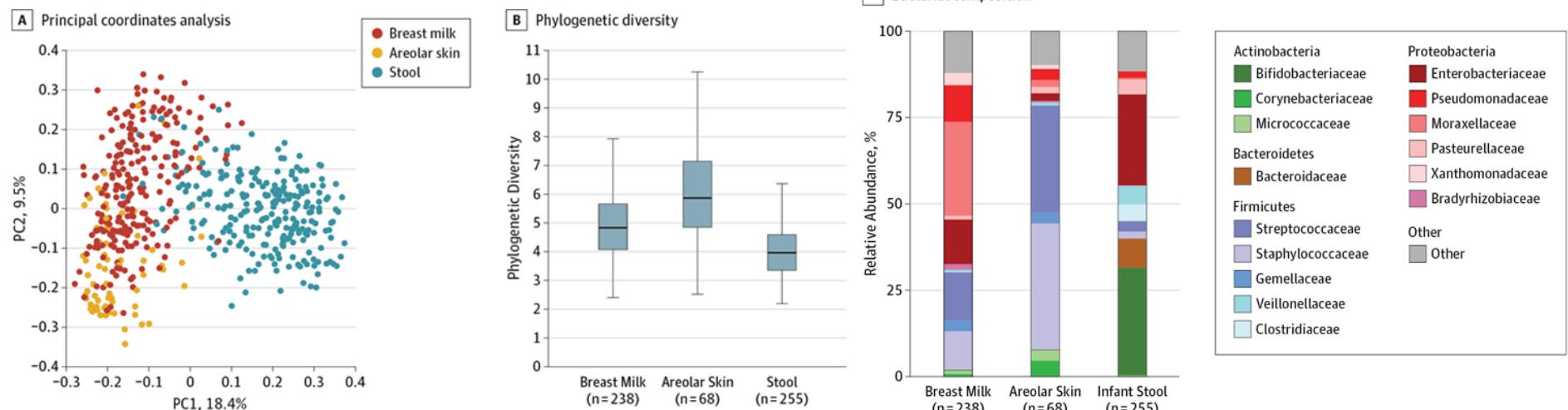
These 2 genera may be universally predominant in the human milk microbiota, independent of geographic location or milk collection technique and may have been underestimated in previous work using conventional PCR methods.

May 8, 2017

JAMA Pediatrics | Original Investigation

Association Between Breast Milk Bacterial Communities and Establishment and Development of the Infant Gut Microbiome

Pia S. Pannaraj, MD, MPH; Fan Li, PhD; Chiara Cerini, MD; Jeffrey M. Bender, MD; Shangxin Yang, PhD; Adrienne Rollie, MS; Helty Adisetyo, PhD; Sara Zabih, MS; Pamela J. Lincez, PhD; Kyle Bittinger, PhD; Aubrey Bailey, MS; Frederic D. Bushman, PhD; John W. Slezasman, MD; Grace M. Aldrovandi, MD



Breastfed infants received 27.7% of their gut bacteria from breast milk and 10.4% from areolar skin during the first month of life



Contents lists available at SciVerse ScienceDirect

International Journal of Food Microbiology

journal homepage: www.elsevier.com/locate/ijfoodmicro



Short communication

Fungal diversity in cow, goat and ewe milk

Emilie Delavenne, Jerome Mounier, Katia Asmani, Jean-Luc Jany, Georges Barbier, Gwenaelle Le Blay *

Université Européenne de Bretagne, France

Université de Brest, EA3882 Laboratoire Universitaire de Biodiversité et Ecologie Microbienne, IFR148 ScInBioS, ESMISAB, Technopôle de Brest Iroise, 29280 Plouzané, France



Fungal species in dairy animals' milk: *Candida*, *Cryptococcus*, *Debaryomyces*, *Malassezia*, *Pichia*, *Rhodotorula*, *Aspergillus*, etc.

SCIENTIFIC REPORTS



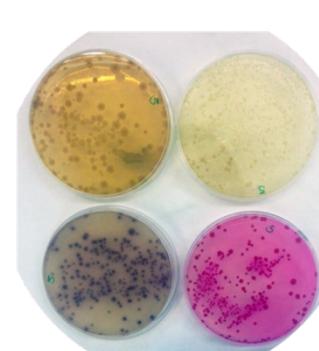
OPEN

Multiple Approaches Detect the Presence of Fungi in Human Breastmilk Samples from Healthy Mothers

Received: 10 May 2017

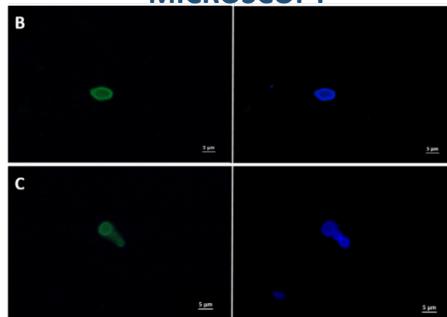
Accepted: 21 September 2017

Published online: 12 October 2017

Alba Boix-Amorós^{1,2}, Cecilia Martínez-Costa³, Amparo Querol¹, María Carmen Collado¹ & Alex Mira²

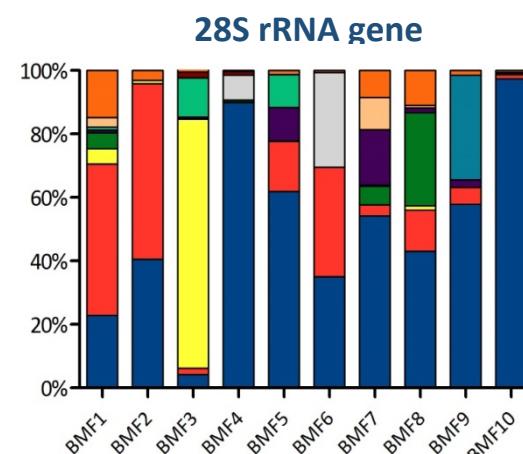
Alba Boix-Amorós

FISH HYBRIDIZATION AND MICROSCOPY



FISH probe especific for 18S rRNA gene

Calcofluor White Stain



- *Malassezia*
- *Saccharomyces*
- *Candida*
- *Alternaria*
- *Rhodotorula*
- *Christiansenia*
- *Cystobasidium*
- *Debaryomyces*
- *Cladosporium*
- *Lodderomyces*
- *Others*

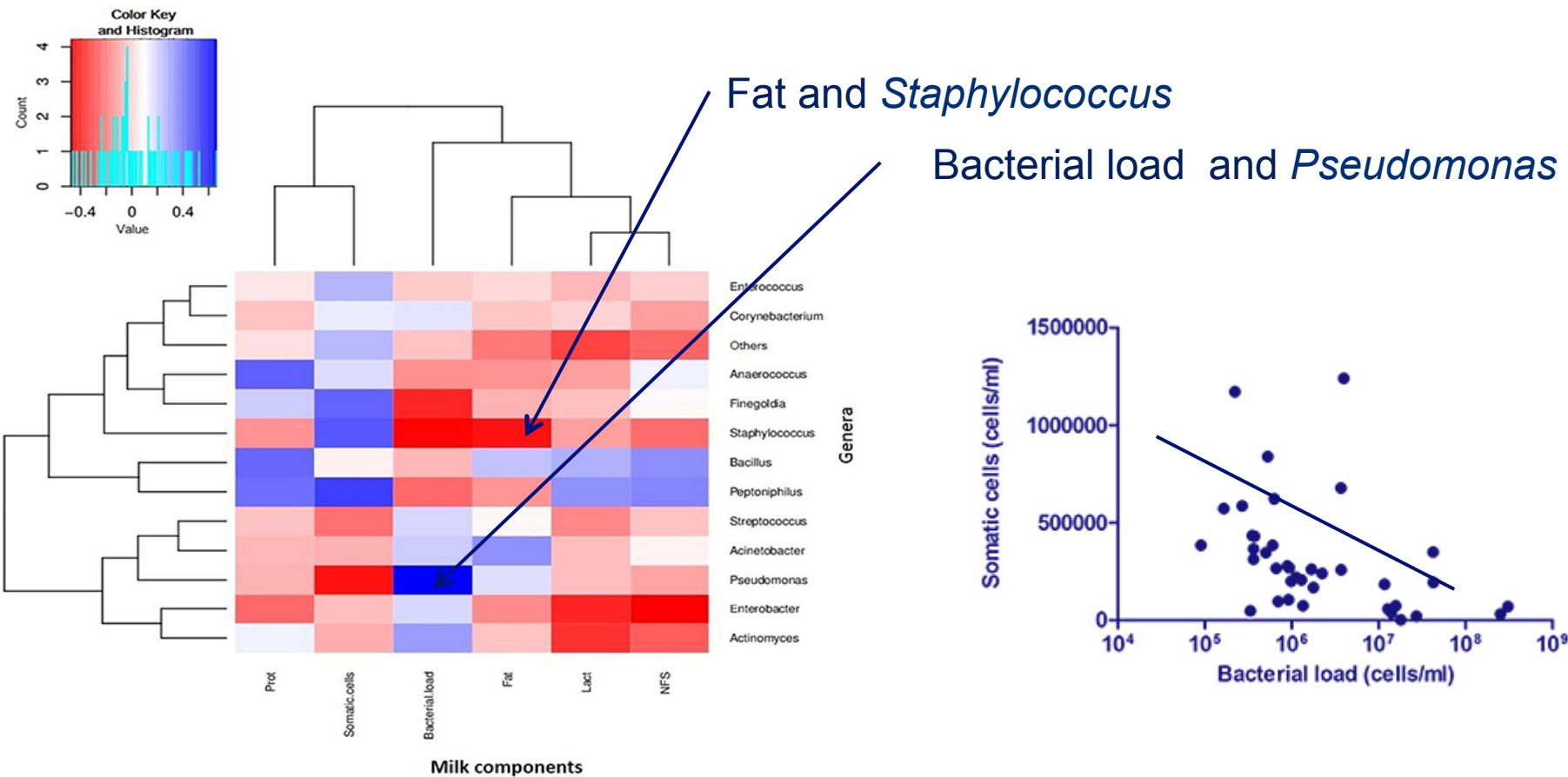
Species	%	Prevalence
<i>Malassezia globosa</i>	34,63	6/10
<i>Candida</i> sp. HA1671	19,54	3/10
<i>Saccharomyces cerevisiae</i>	14,56	7/10
<i>Malassezia restricta</i>	9,07	10/10
<i>Alternaria arborescens</i>	4,82	4/10
uncultured <i>Candida</i>	3,48	2/10
<i>Cladosporium bruhnei</i>	3,28	1/10
<i>Alternaria</i> sp. NT-2015a	2,67	5/10
<i>Candida sake</i>	1,93	2/10
<i>Alternaria tenuissima</i>	1,57	5/10
<i>Debaryomyces hansenii</i>	1,24	3/10
<i>Cystobasidium</i> sp. CBS7295	1,20	1/10
uncultured <i>Debaryomyces</i>	0,63	3/10

Boix-Amorós et al., 2017

Factors influencing Milk Microbiome

Milk microbiome is associated to macronutrients profile in breast milk ?

Relationships between bacterial composition and nutritional or cellular content of HM



Boix-Amoros et al., 2016

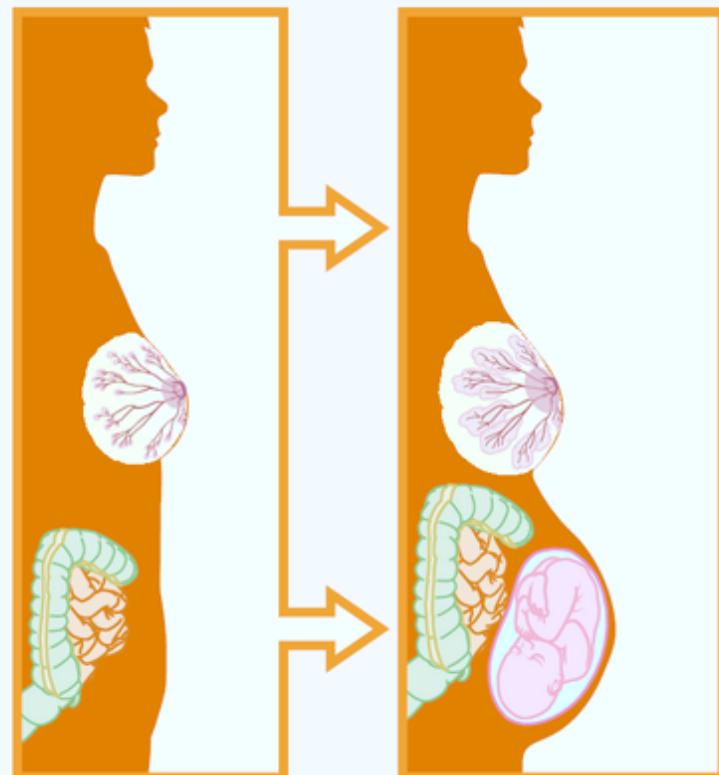
**Breast milk microbiota:
where does it come from?**

Origin and Source of breast milk microbes

Physiological changes during pregnancy

A

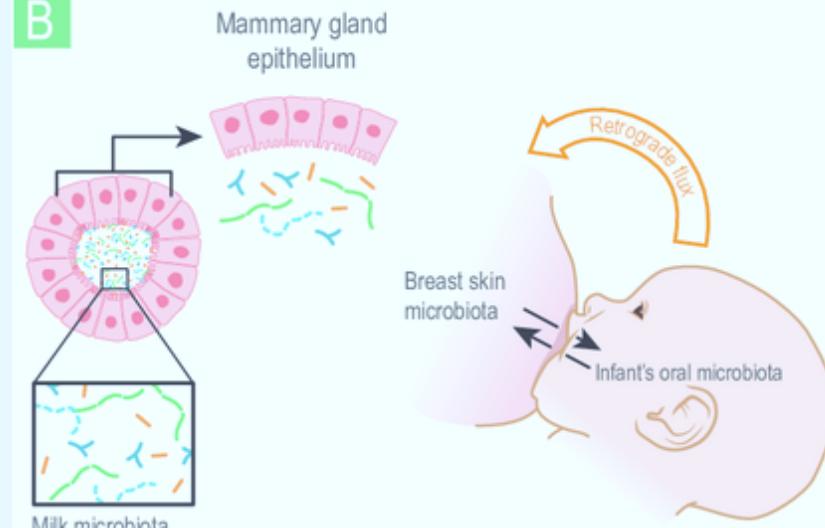
Differentiation of the mammary gland
Massive migration of immune cells



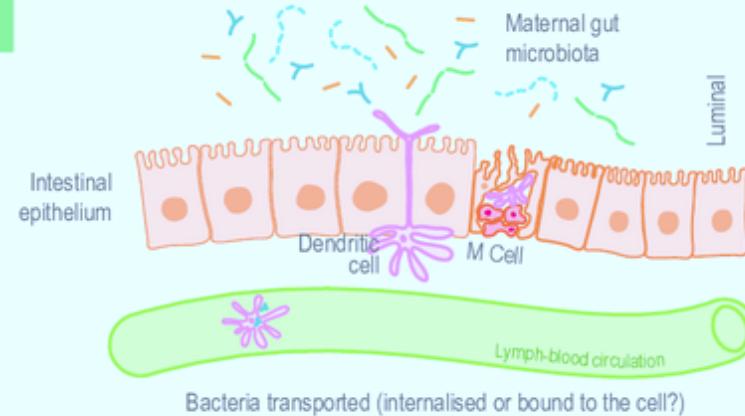
Increased permeability?
Hormonal changes?

Potential routes

B



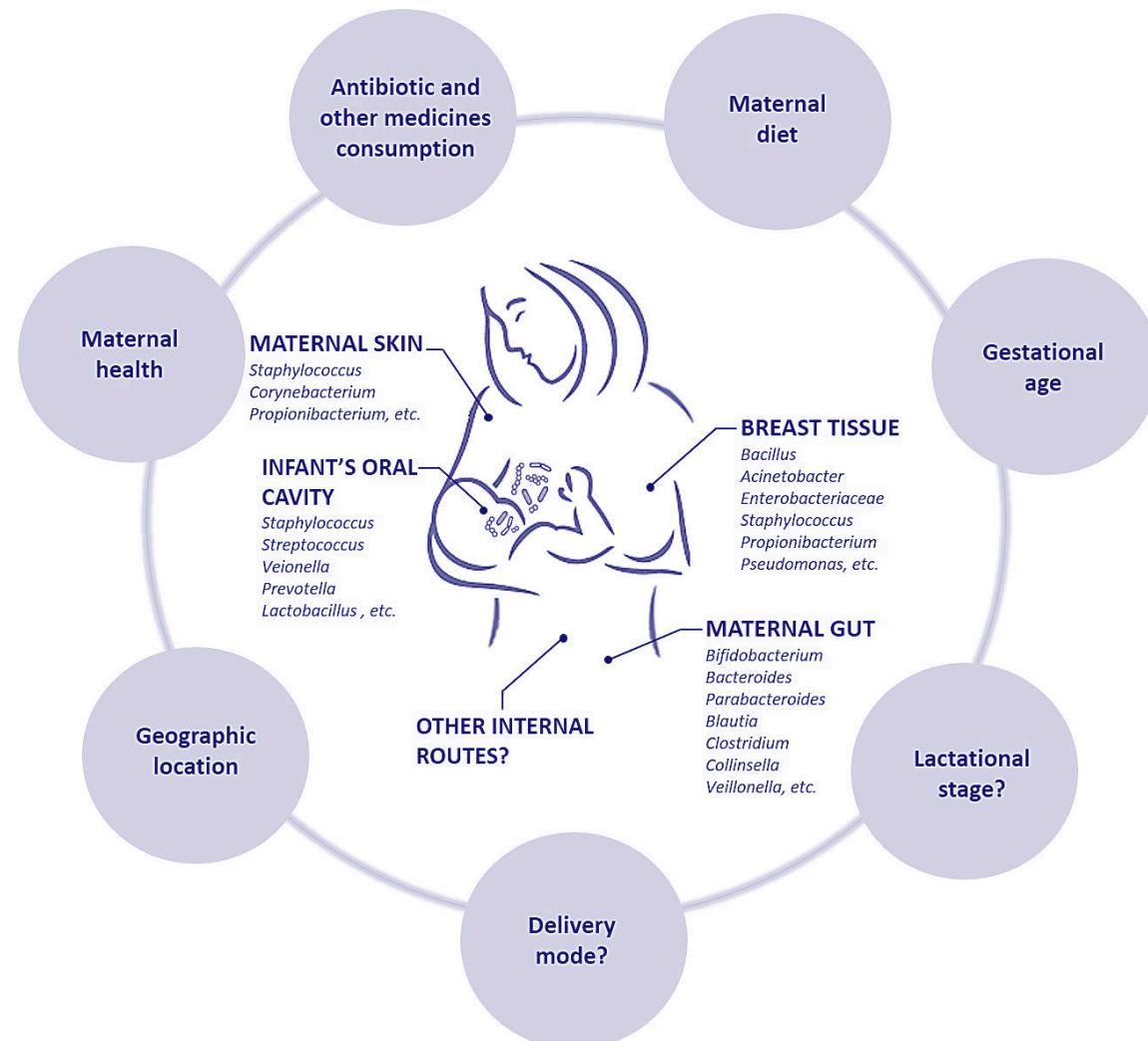
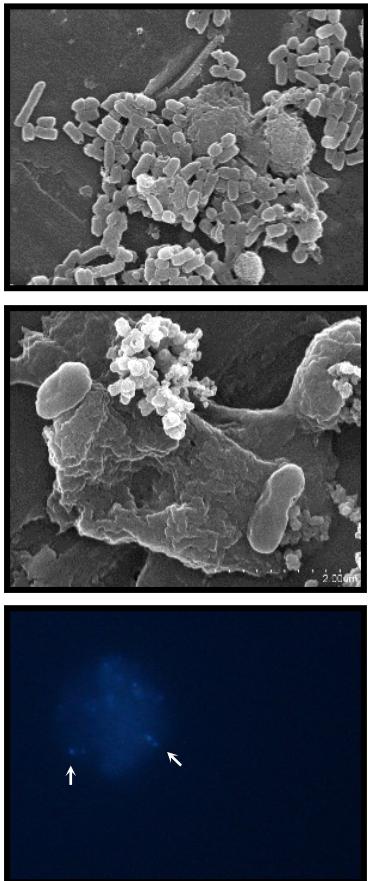
C



Jehuhrick et al. 2013

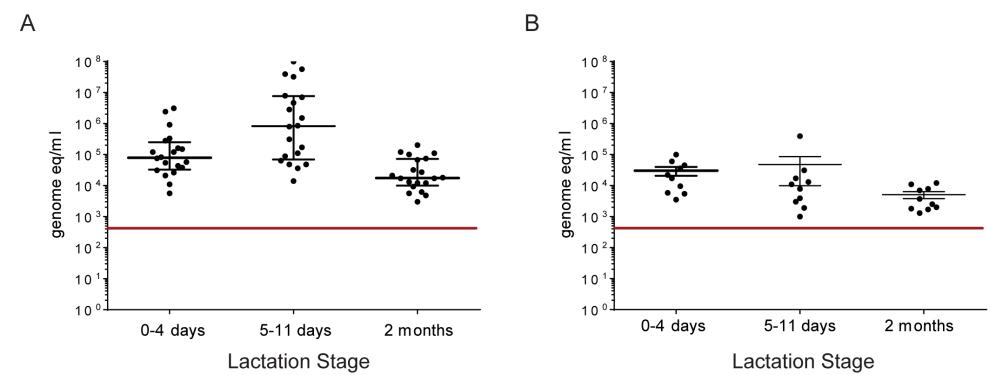
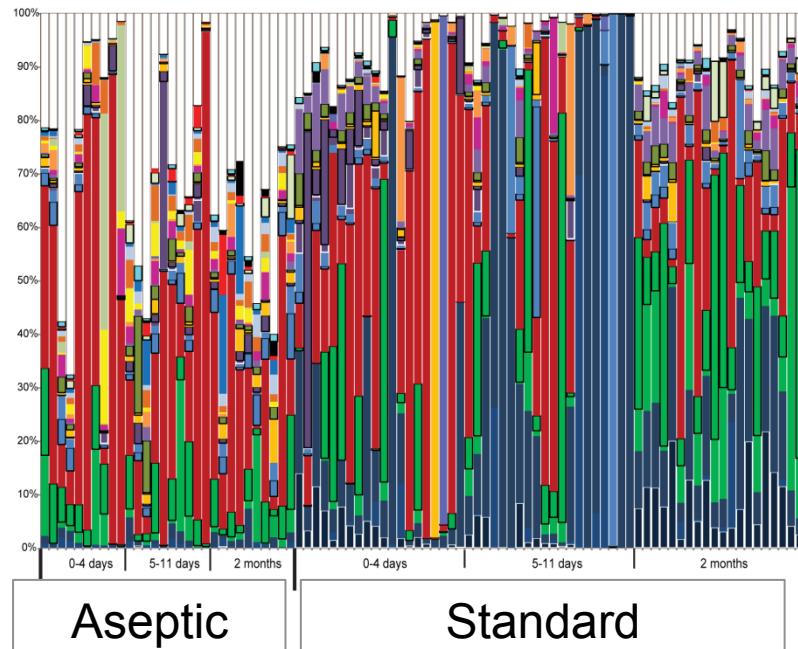
Factors influencing Breast milk composition

FACTORS AFFECTING HUMAN BREAST MILK COMPOSITION



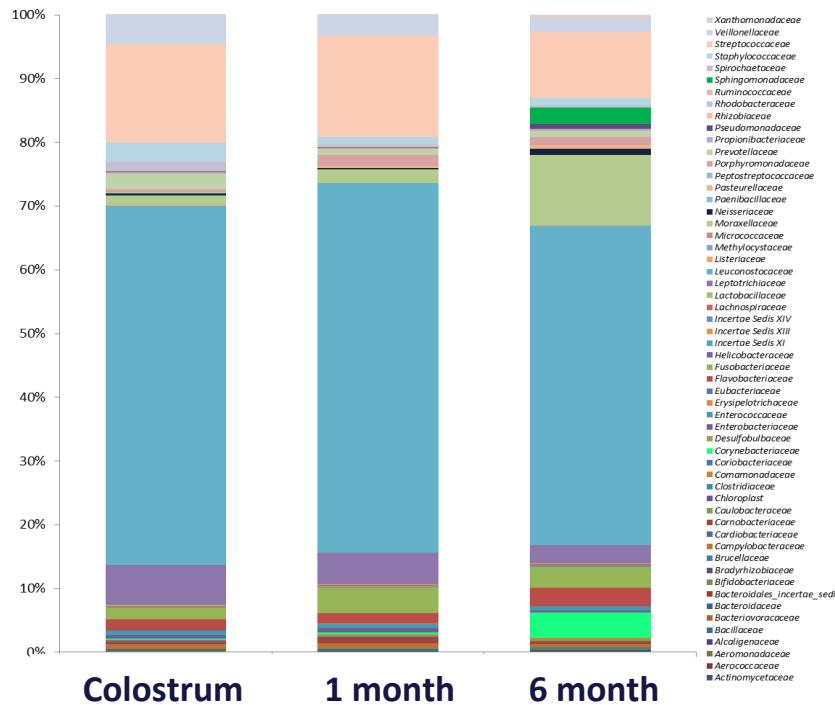
Milk microbiome is influenced by sampling protocol?

Staphylococcus and *Streptococcus* spp. were equally abundant in both sample-type,
↑*Acinetobacter* sp. in standard protocol, but not in strict aseptic protocol



Time

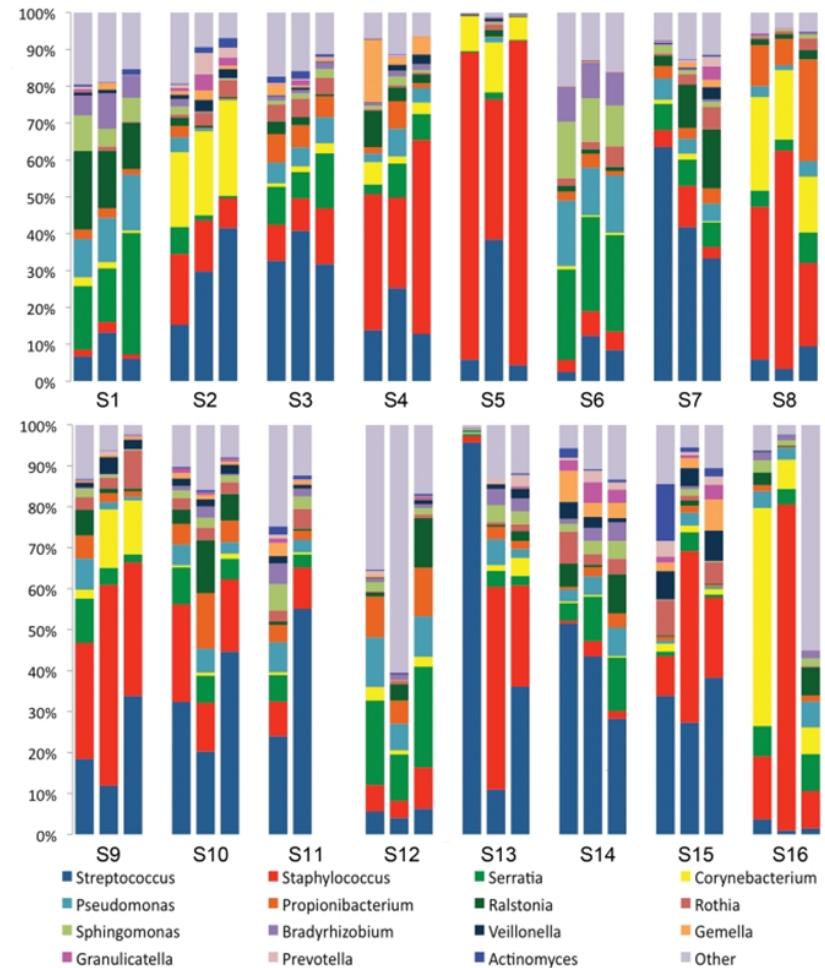
Milk microbiome is influenced by lactation stage



Colostrum: *Weisella* y *Leuconostoc* (*Lactobacillales*)

Staphylococcus, *Streptococcus* and *Lactococcus*.

Mature milk: Lactic Acid bacteria but increase of oral bacteria as *Veillonella*, *Leptotrichia*, *Prevotella*.



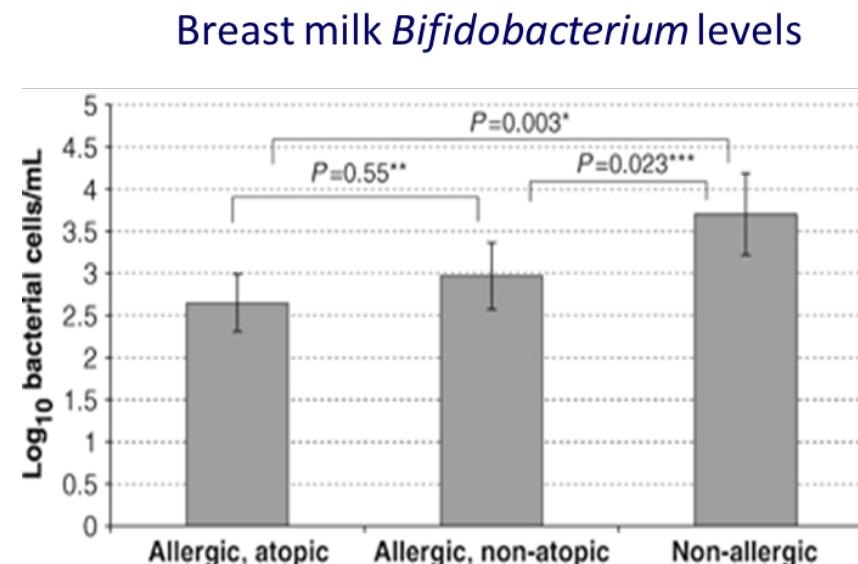
Hunt et al., 2011

Cabrera-Rubio et al., 2012 AJCN

Maternal Allergy influence milk and infant microbiota composition

- ↓ *Bifidobacterium* group numbers in milk samples of **allergic mothers** than non-allergic mothers

- 61 mother-infant pairs
- Breast milk samples 1 month
- All infants colonized with bifidobacteria, exception;
 - 1 born by CS
 - 1 having received antibiotics



Gronlund et al., 2007

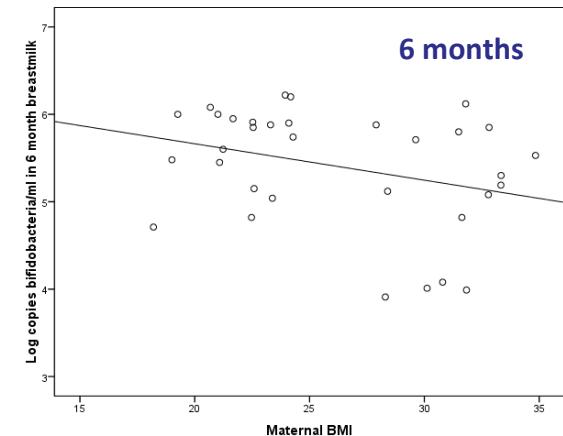
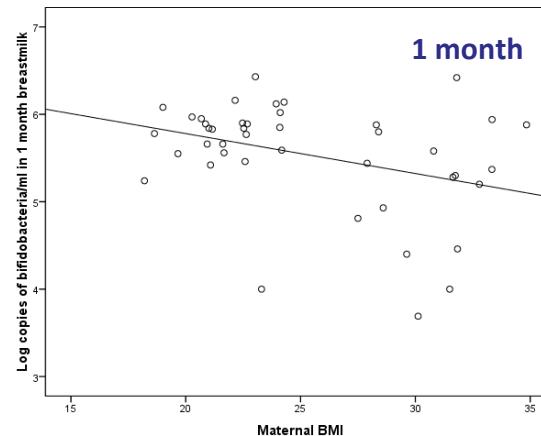
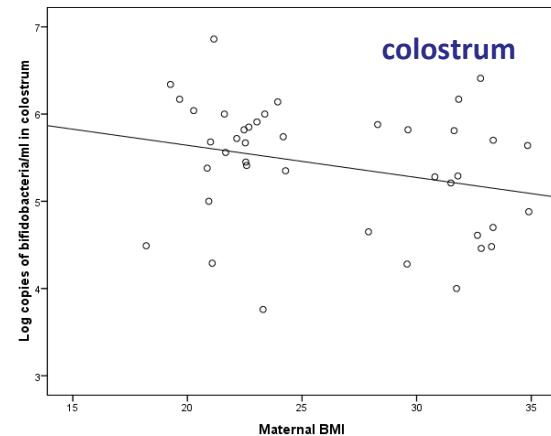
- Maternal allergy and gestational exposures can alter the concentration of Type-1/Type-2/T-regulatory markers in breast milk

Bursch et al., 2013



Maternal obesity influence milk and infant microbiota composition

- ↓ *Bifidobacterium* group & ↑ *Staphylococcus* group numbers in milk samples of obese mothers than normal-weight mothers



- TGF- β 2 and sCD14 levels in the breast milk of overweight mothers tended to be lower than the levels in that of normal-weight mothers

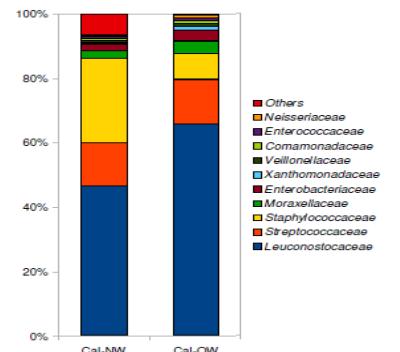
Collado et al., 2012

- Infants born to obese mothers were exposed to 2-fold ↑ HM insulin and leptin ($P < 0.01$) and showed a significant reduction in the early pioneering bacteria Gammaproteobacteria

Lemas et al., 2016 AJCN

FACTORS AFFECTING HUMAN BREAST MILK COMPOSITION

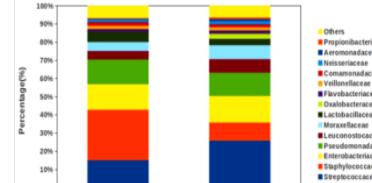
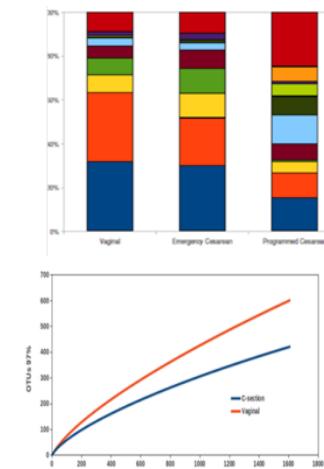
BMI



MODE of DELIVERY



Cabrera-Rubio et al., 2012



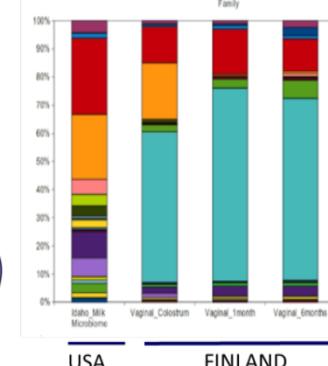
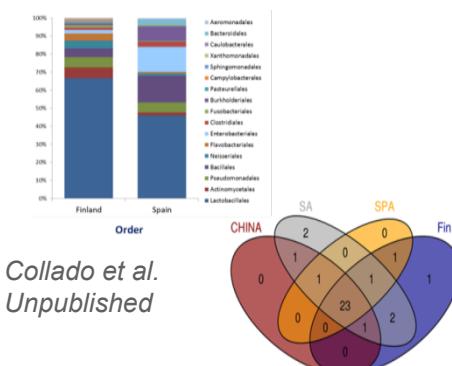
Cabrera-Rubio et al., 2015

Collado et al., 2010 & 2012
Garcia-Mantrana et al. 2017
Gomez-Gallego et al. 2017

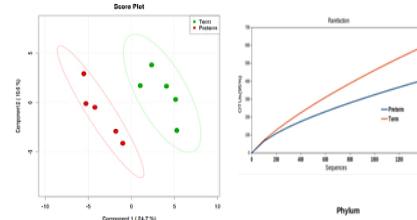


MEDICATION CHEMOTHERAPY

GEOGRAPHICAL LOCATION

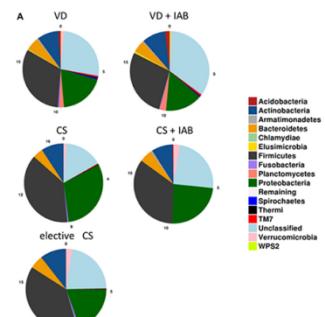


GESTATIONAL AGE



Cabrera-Rubio et al., 2015
Khodayar-Pardo et al. 2012
Alcantara-Baena et al. submitted

ANTIBIOTICS

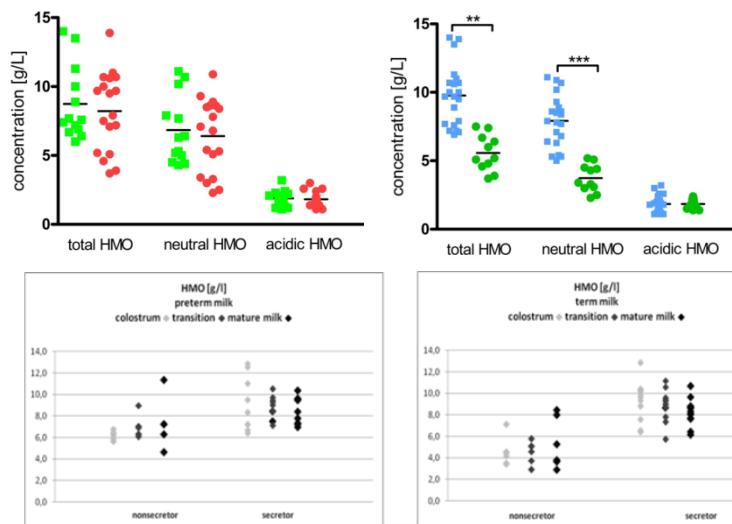


Hermansson et al. 2019
Frontiers Nutrition

Breast Milk Bioactive compounds

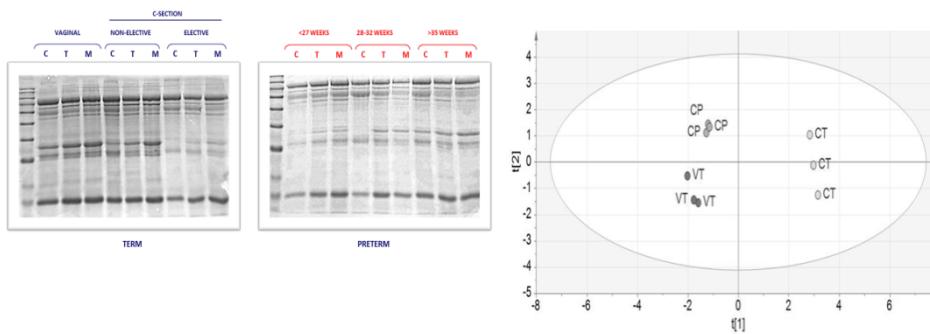
Human Milk Oligosaccharides

Hypothesis: HMO specifically interact with milk microbes



Protein & Peptides

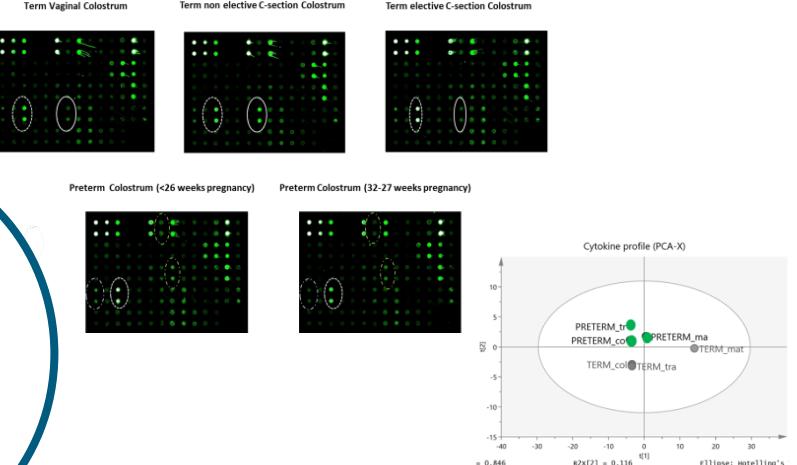
Multi-level approach : protein-peptide strategies followed by state-of-the-art Mass Spectrometry-based methodologies



Growth Factors, Cytokine & other

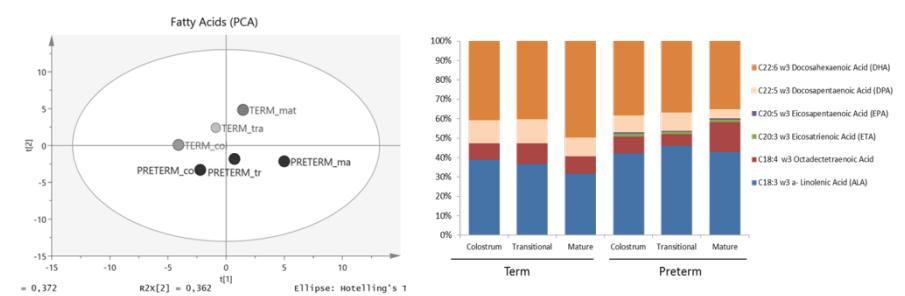
Complex interactions of cytokines and microbiota in breast milk

Collado et al., 2012



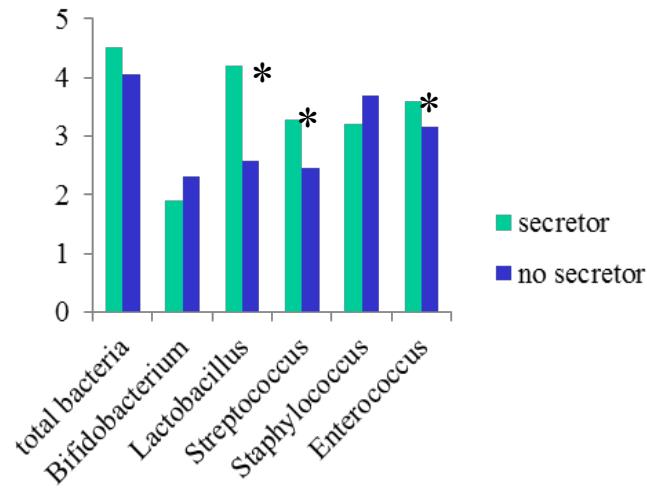
Fatty ACids

Omega fatty acids measured gas chromatography (GC)
Fatty acids have been shown to be modified by time and diet

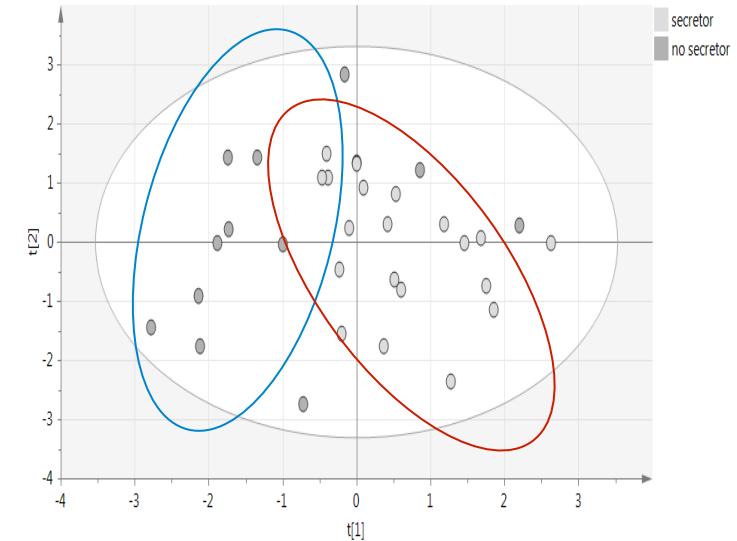
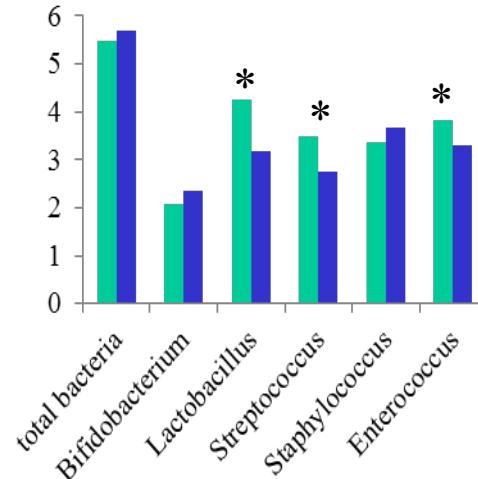


Milk microbiome is influenced by maternal genotype?

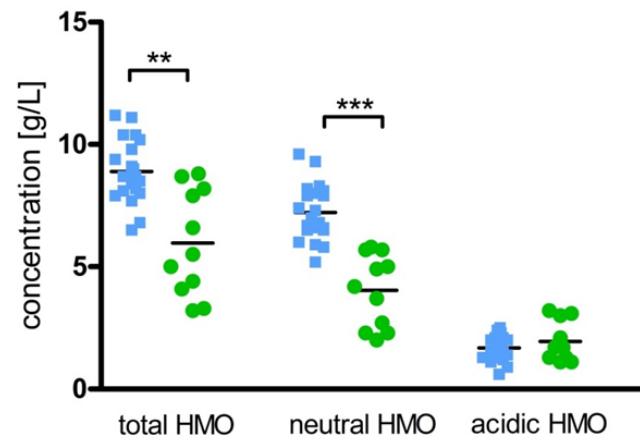
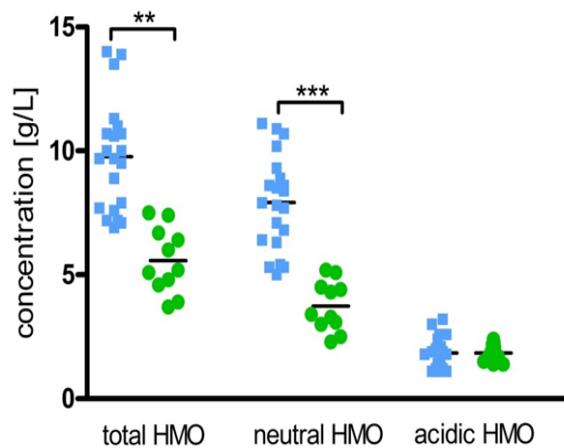
Colostrum



Mature milk

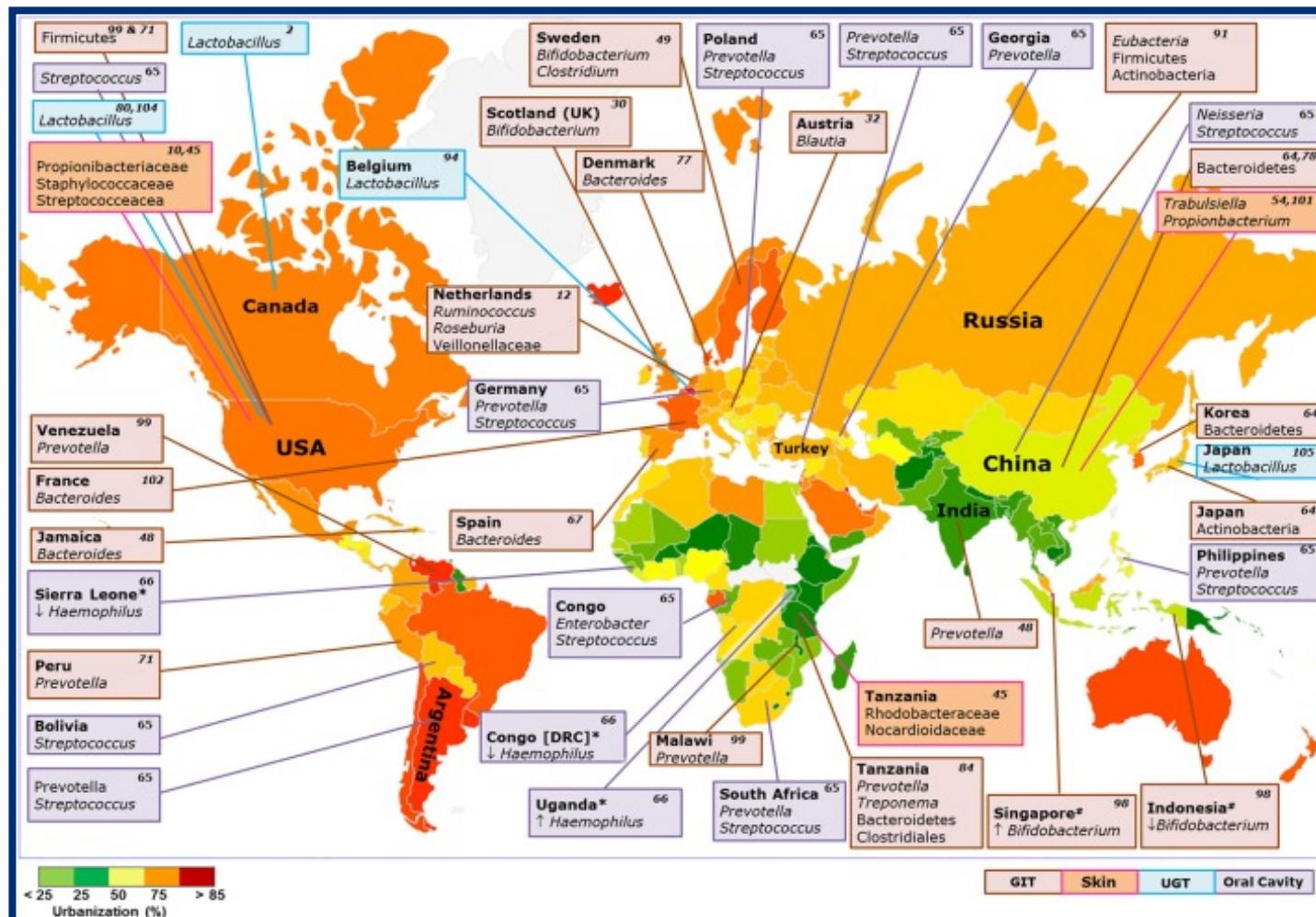


Cabrera-Rubio et al. 2019 JPGN



Kunz et al. 2016 JPGN

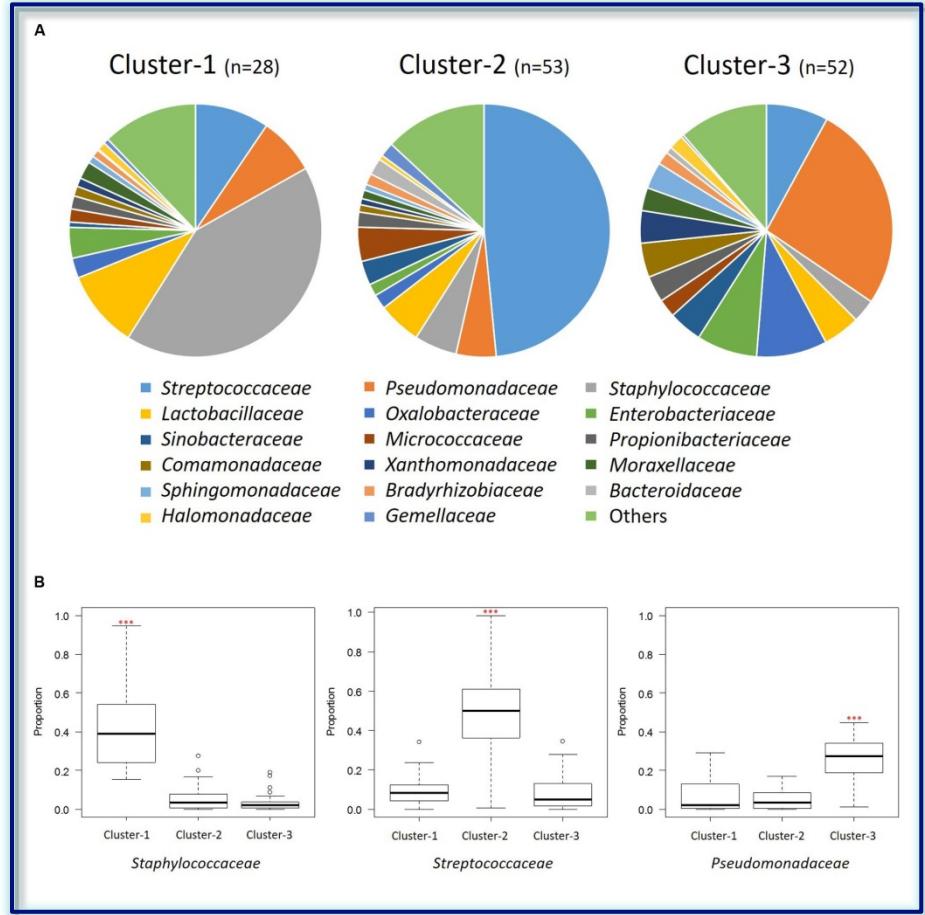
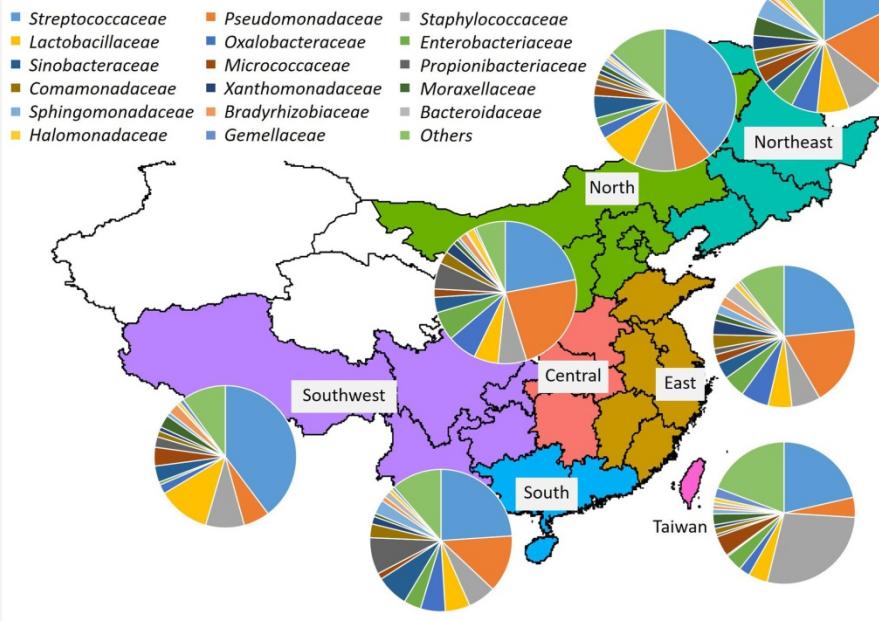
Geography, Ethnicity or Subsistence-Specific Variations in Human Microbiome Composition and Diversity



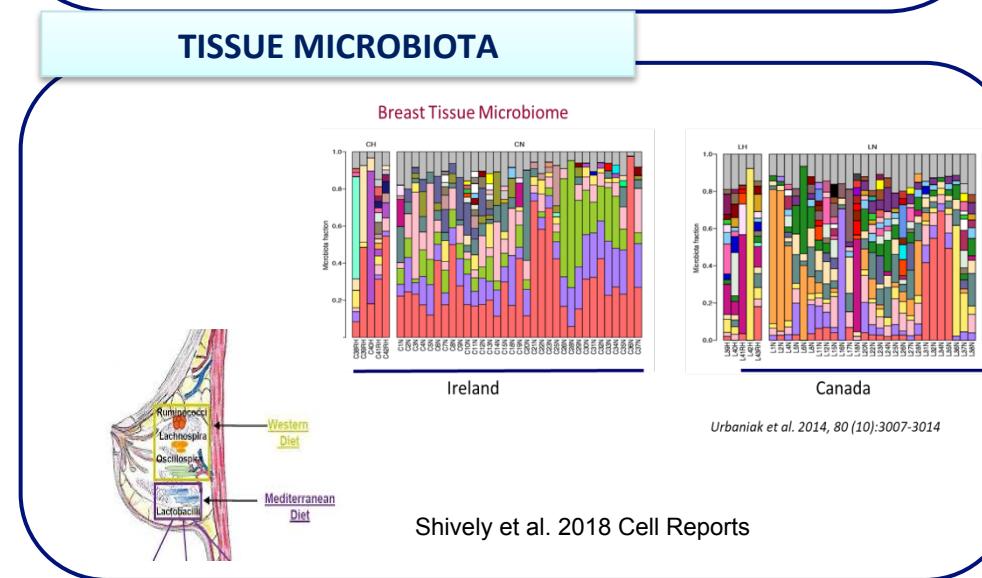
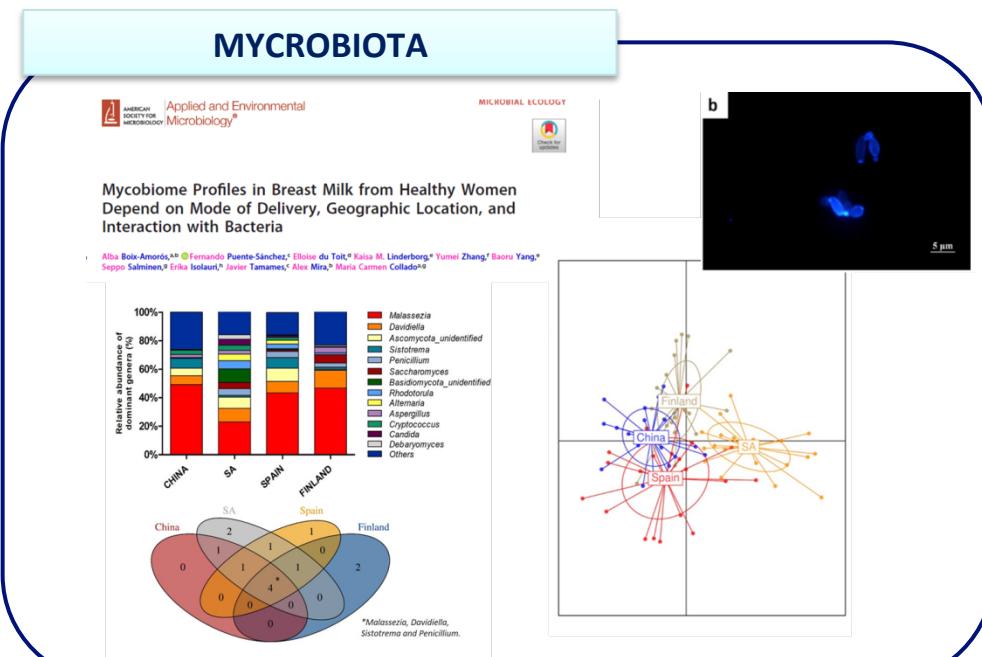
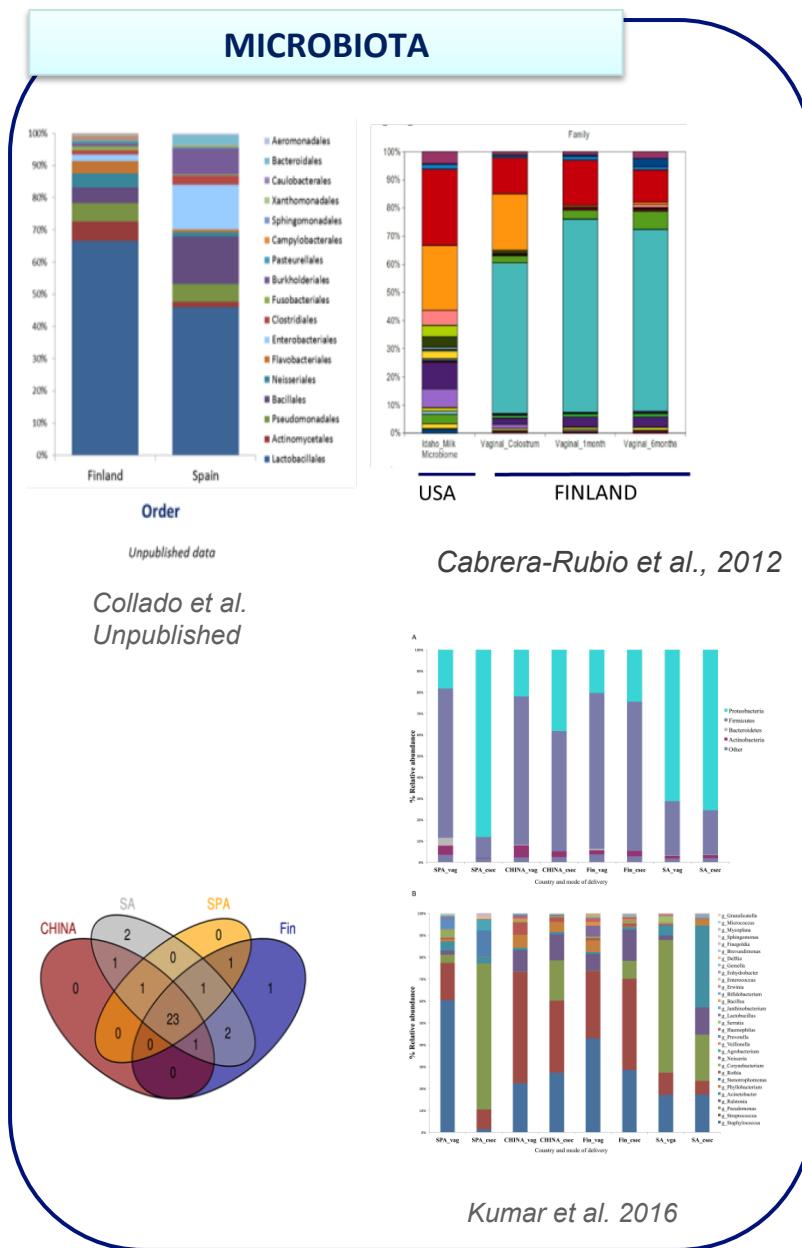
Factors influencing Milk Microbiome

Milk microbiome is influenced by geographical location or diet?

133 healthy mothers in Taiwan and in six regions of mainland China (Central, East, North, Northeast, South, and Southwest China)



Breast milk



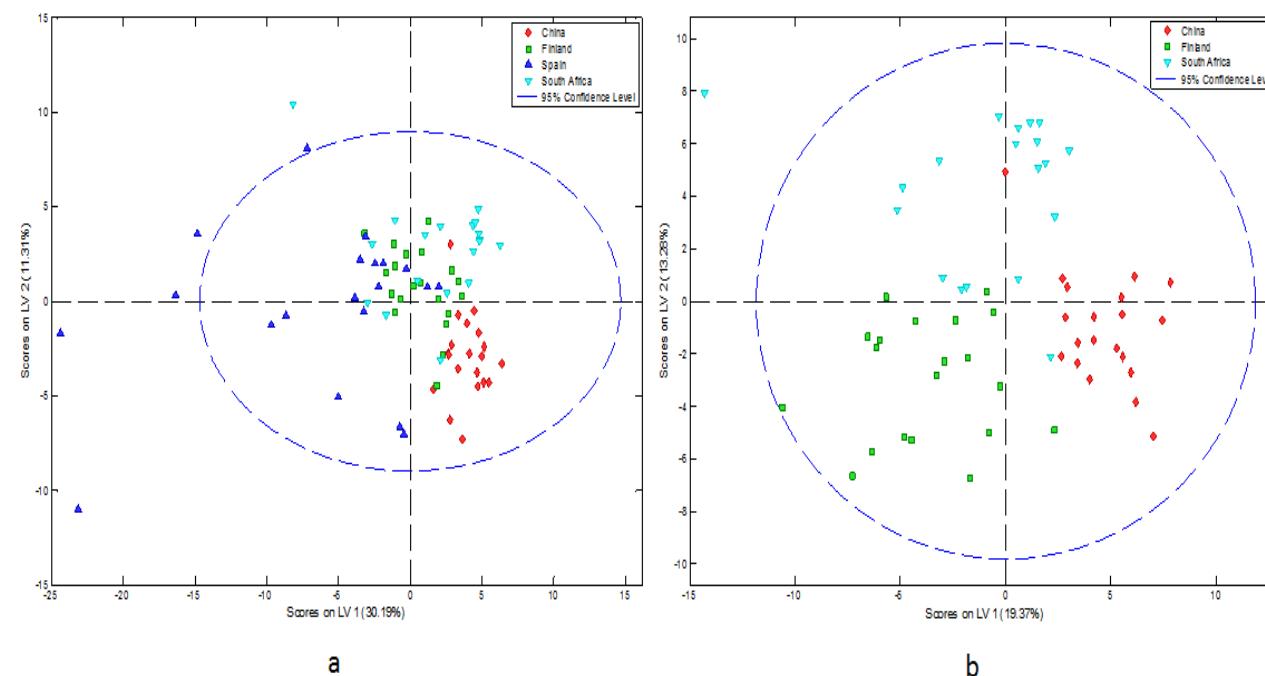
HUMAN BREAST MILK METABOLOMIC PROFILE DIFFERS ACROSS COUNTRIES



Article

Human Breast Milk NMR Metabolomic Profile across Specific Geographical Locations and Its Association with the Milk Microbiota

Carlos Gómez-Gallego ^{1,*}, Jose Manuel Morales ^{2,3,4}, Daniel Monleón ^{2,4}, Eloise du Toit ⁵,
Himanshu Kumar ¹, Kaisa M. Linderborg ⁶, Yumei Zhang ⁷, Baoru Yang ⁶, Erika Isolauri ⁸,
Seppo Salminen ¹ and María Carmen Collado ^{1,9,*}



Maternal diet-geographical location

What's normal? Oligosaccharide concentrations and profiles in milk produced by healthy women vary geographically^{1,2}

Michelle K McGuire,^{3,4} Courtney L Meehan,⁵ Mark A McGuire,⁶ Janet E Williams,^{6,7} James Foster,⁸ Daniel W Sellen,¹⁰ Elizabeth W Kamau-Mbuthia,¹¹ Egidioh W Kamundia,¹¹ Samwel Mbugua,¹¹ Sophie E Moore,^{12,13,21} Andrew M Prentice,¹⁴ Linda J Kvist,¹⁵ Gloria E Otoo,¹⁶ Sarah L Brooker,^{6,7} William J Price,⁹ Bahman Shafii,⁹ Caitlyn Placek,^{5,22} Kimberly A Lackey,³ Bianca Robertson,^{17,18} Susana Manzano,¹⁹ Lorena Ruiz,¹⁹ Juan M Rodríguez,¹⁹ Rossina G Pareja,²⁰ and Lars Bode^{17,18*}*

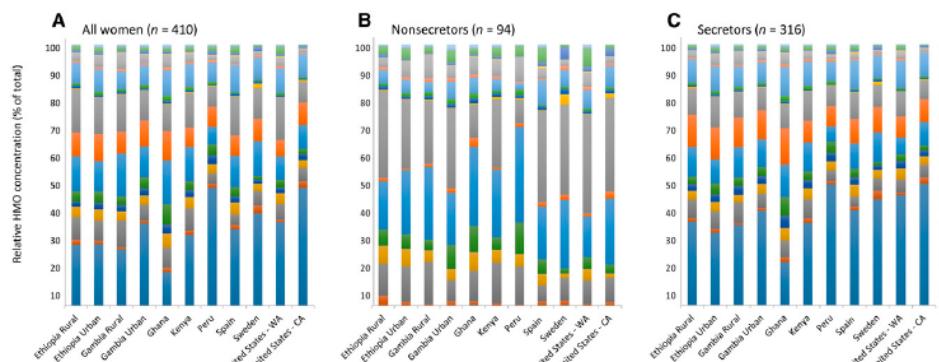


FIGURE 2 Mean \pm SEM relative abundance of HMO concentrations of all women combined (A), nonsecretors (B), and secretors (C) in each cohort. Note that there was only one nonsecretor subject each in Peru and United States - CA, CA, California; DFLac, difucosylactose; DFLNH, difucosylacto-*N*-hexose; DFLNT, difucosylacto-*N*-tetrose; DSLNH, disialylacto-*N*-hexose; DSLNT, disialylacto-*N*-tetrose; FDSLNH, fucosidialacto-*N*-hexose; FLNH, fucosylacto-*N*-hexose; HMO, human milk oligosaccharide; LNFP, lacto-*N*-fucopentaose; LNH, lacto-*N*-hexose; LNTn, lacto-*N*-neotetraose; LNT, lacto-*N*-tetrose; LSTb, sialyl-lacto-*N*-tetraose b; LSTc, sialyl-lacto-*N*-tetraose c; WA, Washington; 2'FL, 2'-fucosyllactose; 3FL, 3-fucosyllactose; 3'SL, 3'-sialyllactose; 6'SL, 6'-sialyllactose.

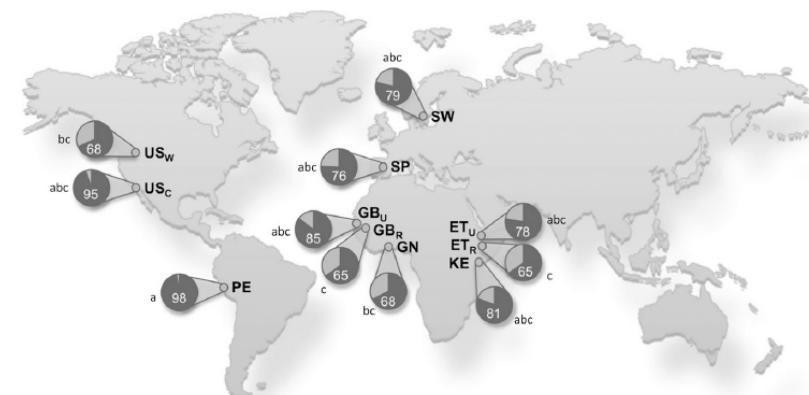


FIGURE 3 Percentages of women in each cohort categorized as securtors. Cohorts that do not share a common lowercase letter differ ($P < 0.05$) in terms of their percentages of women who were securtors with the use of a chi-square test with Benjamini and Hochberg false-discovery rate corrections. ET_R, rural Ethiopia; ET_U, urban Ethiopia; GB_R, rural Gambia; GB_U, urban Gambian; GN, Ghana; KE, Kenya; PE, Peru; SP, Spain; SW, Sweden; US_R, United States (California); US_W, United States—Washington.



Article

Colostrum and Mature Human Milk of Women from London, Moscow, and Verona: Determinants of Immune Composition

Daniel Munblit ^{1,2,3,*}, Marina Treneva ^{2,4}, Diego G. Peroni ^{2,5}, Silvia Colicino ⁶, LiYan Chow ¹, Shobana Dissanayake ⁷, Priya Abrol ¹, Shreya Sheth ¹, Alexander Pampura ^{2,4}, Attilio L. Boner ⁸, Donna T. Geddes ^{2,9}, Robert J. Boyle ^{1,2,†} and John O. Warner ^{1,2,†}



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What's Normal? Immune Profiling of Human Milk from Healthy Women Living in Different Geographical and Socioeconomic Settings

Lorena Ruiz ^{1*†}, Irene Espinosa-Martos ^{1,2*†}, Cristina García-Carral ¹, Susana Manzano ¹, Michelle K. McGuire ^{3,4}, Courtney L. Meehan ⁵, Mark A. McGuire ⁶, Janet E. Williams ⁶, James Foster ⁷, Daniel W. Sellen ⁸, Elizabeth W. Kamau-Mbuthia ⁹, Egidio W. Kamundia ⁹, Samwel Mbugua ⁹, Sophie E. Moore ^{10,11}, Linda J. Kvist ¹², Gloria E. Otoo ¹³, Kimberly A. Lackey ³, Katherine Flores ⁵, Rossina G. Pareja ¹⁴, Lars Bode ¹⁵ and Juan M. Rodríguez ^{1*}

To increase our understanding of the role of maternal bacteria in infant microbiota as it will help to develop new dietary strategies based on microbial modulation, aimed at the beneficial microbiological, immunological and metabolic programming of child health.

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