

MENA Microbiome

From Biomarkers Discovery
to Microbiota-based Therapeutics

Date: 28-29 September 2024

Venue: [The Plaza Doha](#)



1:00 – 3:30 pm	<p>Session V: Microbiome in Women and neonatal Health</p> <p>Learning objectives:</p> <ul style="list-style-type: none"> ■ List the major vaginal microbes in healthy women ■ Explain the major microbiome changes during pregnancy ■ Describe the role of the microbiome in prematurity <p>Session Chairs:</p> <ul style="list-style-type: none"> ■ Prof Johnny Awwad, Chair of Women's Services, Division Chief, Reproductive Medicine, Sidra Medicine ■ Dr. Lolwa Al Ansari, Senior Consultant and Head of OB/GYN, Al Wakra Hospital, Hamad Medical Corporation 	
2:20 – 2:45 pm	<p>Dr. Jaime Garcia Mena</p> <p>Department of Genetics and Molecular Biology</p> <p>Center for Research and Advanced Studies of the National Polytechnic Institute, Mexico</p>	<p>Maternal immunoglobulins differentially bind the bacterial community in human colostrum and stool of breastfed neonates.</p>



MENA Microbiome From Biomarkers Discovery to Microbiota-based Therapeutics

28-29 September 2024



CERTIFICATE OF APPRECIATION

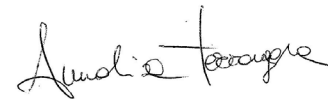
Presented to

Dr. Jaime Garcia Mena

*In recognition of their valuable contribution as a Speaker
at the **MENA Microbiome conference: From Biomarker Discovery to Microbiota-based Therapeutics** ,
held on 28-29 September in Doha, Qatar.*



Dr. Souhaila Al Khodor
Event Co -Chair



Dr. Annalisa Terranegra
Event Co -Chair

<https://www.sidra.org/events/research/mena-microbiome-conference-2024/speakers-2/>

MENA Microbiome

From Biomarkers Discovery to Microbiota-based Therapeutics

Date: 28-29 September 2024

Venue: [The Plaza Doha](#)



Add to Calendar



 **Overview / Agenda**

Agenda

Search



All Sessions

Day 1: September 28, 2024

Day 2: September 29, 2024



8:00 am – 12:00 pm	<div>Across Organs</div> <div>Learning objectives:</div> <div><div>Describe the microbiome composition of a healthy gut.</div><div>Summarize the different members of the gut microbiome.</div><div>Evaluate the application of systems biology approaches to study the microbiome.</div></div> <div>Session Chairs:</div> <div><div>Prof. Hilal Lashuel, Advisor to the Chairperson & Executive Director of RDI, Qatar Foundation</div><div>Dr. Khalid Fakhro, Chief Research Officer, Sidra Medicine</div></div>	
8:00 – 8:45 am	<div>Prof. Peer Bork</div> <div>Director of EMBL</div> <div>Heidelberg, Germany</div>	<div>Keynote Lecture II:</div> <div>Gut microbiome analysis for human health and wellbeing</div>
8:45 – 9:10 am	<div>Prof. Emmanuelle Maguin</div> <div>Director, French National Institute for Agriculture, Food, and Environment (INRAE), France</div>	<div>The food and microbiome interplay for health</div>
9:10 – 9:35 am	<div>Dr. Amelia McGuinness</div> <div>Associate Research</div>	<div>The Microbiome-Gut-Brain Axis: The Bacteriome and Beyond</div>

9:35 – 10:00 am	<p>Attending Pediatric Urologist, Division of Urology</p> <p>Children’s Hospital of Philadelphia, PA, USA</p>	<p>The Gut-Kidney Axis of Kidney Stone Disease: A Pathway to Novel Diagnostics and Therapeutics</p>
10:00 – 10:30 am	<p>Coffee break</p> <p>Poster viewing and judging</p>	
	<p>Session Chairs:</p> <p>Dr. Mamoun Elawad, Division Chief of Pediatric Gastroenterology and Nutrition, Sidra Medicine</p> <p>Dr. Amine Zorgani, Consultant, Biocodex Microbiota Institute, Belgium</p>	
10:30 – 10:55 am	<p>Prof. Nahla Mansour</p> <p>Professor of Microbial Molecular Biology</p> <p>National Research Center, Egypt</p>	<p>Gut Microbiome Analysis in Type 2 Diabetes Egyptians: Towards Personalized Medicine</p>
10:55 – 11:00 am	<p>Miss Shaikha Al Abduljabbar</p> <p>Research Specialist, Precision Nutrition, Sidra Medicine, Qatar</p>	<p>Manipulation of the microbiome through diet can improve clinical outcomes: A story from an IBD-Autistic patient at Sidra Medicine</p>

		UAE University, UAE	Approaches
11:25 – 11:50 pm	Prof Mohammad Issa El Mouzan Professor of Pediatrics, College of Medicine, King Saud University, Saudi Arabia		Gut virome and Mycobiome in Saudi pediatric population
11:50 – 12:00 pm	Ms. Daliya Abubakar Research specialist Microbiome and Biomarker Discovery Sidra Medicine, Qatar		Short oral talk: Profiling of the Blood Virome in Children with Idiopathic Nephrotic Syndrome
12:00 – 1:00 pm	Lunch Break Poster viewing and judging		
1:00 – 3:35 pm	Session V: Microbiome in Women and neonatal Health Learning objectives: List the major vaginal microbes in healthy women Explain the major microbiome changes during pregnancy Describe the role of the microbiome in prematurity Session Chairs: Prof Johnny Awwad, Chair of Women's Services, Division Chief, Reproductive Medicine, Sidra Medicine		

1:00 -1:25 pm	Director of Institute of Reproductive and Developmental Biology, Professor of Obstetrics and Gynaecology Imperial College London, UK	Microbiome aspects of fertility, miscarriage, and preterm birth
1:25 – 1:50 pm	Dr. Maha Al Asmakh Department Head of Biomedical Sciences Associate Professor, Qatar University, Qatar	Salivary Microbiome in pregnancy
1:50 – 2:00 pm	Mrs Maysa Niaz Department of Pathobiology, University of Guelph, Canada	Short oral talk: Know thy neighbours: Latent topics for studying vaginal microbial communities
2:00 – 2:25 pm	Prof. Erika Isolauri Professor of Pediatrics Head of the Department of Clinical Medicine, Faculty of Medicine University of Turku, Finland	Resilience to risk exposures in the preterm child through the microbiome

	Children's Hospital of Philadelphia, PA, USA	Microbiome Analysis to Understand Infectious Diseases
2:50 – 3:15 pm	Dr. Ibrahim Hassan Senior attending Physician, Microbiology and Virology, Sidra Medicine, Qatar	Antimicrobial stewardship and microbiome in the pediatric population: Where are we?
3:15 – 3:25 pm	Miss Noora Al Mohannadi Research specialist Microbiome and Biomarker Discovery Sidra Medicine, Qatar	Short oral talk: Oral microbiome and immunity in women with gestational diabetes mellitus: A pilot study
3:25- 3:35 pm	Dr. Jaime Garcia Mena Department of Genetics and Molecular Biology Center for Research and Advanced Studies of the National Polytechnic Institute, Mexico	Short oral talk: Maternal immunoglobulins differentially bind the bacterial community in human colostrum and stool of breastfed neonates.
3:35 – 4:00 pm	Coffee Break Poster viewing and judging	



4:00 – 5:30 pm	Screening: The Invisible Extinction	School, NJ, USAMaria Gloria Dominguez Bello Professor, Department of Biochemistry and Microbiology, Rutgers School of Environmental and Biological Sciences, NJ, USA
5:30 – 5:45 pm	Closing remarks for Day 2 & Posters Award Announcement	MENA Microbiome conference Chairs

Sidra Medicine

Al Gharrafa Street,
Ar-Rayyan, Doha, Qatar
+974 4003 3333



Maternal immunoglobulins differentially bind the bacterial community in human colostrum and stool of breastfed neonates.

Jaime García-Mena^{*,#a}, Karina Corona-Cervantes^a, Erick Sánchez-Salguero^b, Paola Berenice Zárate-Segura^c, Aparna Krishnakumar^a, Alberto Piña-Escobedo^a, Martín Noé Rangel-Calvillo^d, Tito Ramírez-Lozada^e, Gustavo Acosta-Altamirano^e, Noemí del Socorro Lázaro-Pérez^a, Mónica Sierra-Martínez^e, Leopoldo Santos Argumedo^b.

a) Departamento de Genética y Biología Molecular, Centro de Investigación y de Estudios Avanzados, México City, Mexico.

b) Departamento de Biomedicina Molecular, Centro de Investigación y de Estudios Avanzados, México City, Mexico.

c) Escuela Superior de Medicina, Instituto Politécnico Nacional, México City, Mexico.

d) Hospital General “Dr. José María Rodríguez”, Ecatepec de Morelos, Estado-de-México, Mexico.

e) Hospital Regional de Alta Especialidad de Ixtapaluca, Estado-de-México, Mexico.

*Presenting author

Corresponding author

In newborns, the successional establishment of the primordial microbiota strains in the gut is an interesting topic of investigation, where the IgA1, IgA2, IgM, and IgG immunoglobulins provided by the mother during breastfeeding play a primordial role. The objective of this work was to explore the functional role of the colostrum's maternal immunoglobulins, which bind differently, a diverse bacterial community in the intestine of breastfed neonates. We sequenced V3-16S rRNA gene libraries prepared with DNA extracted from single IgA1, IgA2, IgM, and IgG fluorescence-activated cell sorting fractions from meconium or colostrum. Our results show that in colostrum, the bacteria are already differentially bound by these immunoglobulins. We determined that IgA2 and IgM bind alpha and beta Proteobacteria at early breastfeeding stages, which might stimulate the immune system in the gut of neonates. In addition, it was found that IgG mostly binds facultative anaerobes of the Firmicutes phylum, which are reported as part of the human milk microbiota and pioneer elements of the neonatal gut. In the case of the neonatal stool, the immunoglobulins supplied by the mother, bind a wide diversity of bacteria. For example, IgA2 and IgM bind more bacteria of the phylum Bacteroidetes in comparison to what IgG binds. Bacteroidetes and some Firmicutes have been reported as late colonizers in the successional population of the neonatal gut since they can produce short-chain fatty acids like propionate and butyrate. Our results support the current view that joint microbial and immunoglobulin transference is fundamental for the normal development of the neonate's immune system and the establishment of a functional gut microbiota. Work financed by Fondo SEP-Cinvestav, No. 174, Consejo Nacional de Ciencia y Tecnología CONACYT-163235, INFR-2011-01, and CONACyT FORDECYT-PRONACES/6669/2020_Programa Presupuestario F003-Ciencia de Frontera 2019.