TOD TEN in gastroenterologia 14^ EDIZIONE

24-25 NOVEMBRE 20**23**

BERGAMO

HOTEL EXCELSIOR SAN MARCO Piazza della Repubblica, 6



MECCANISMO D'AZIONE DEI PROBIOTICI

Maurizio Koch MD







I have no disclosures to declare

Maurizio Koch M.D.

Kochmaurizio@gmail.com

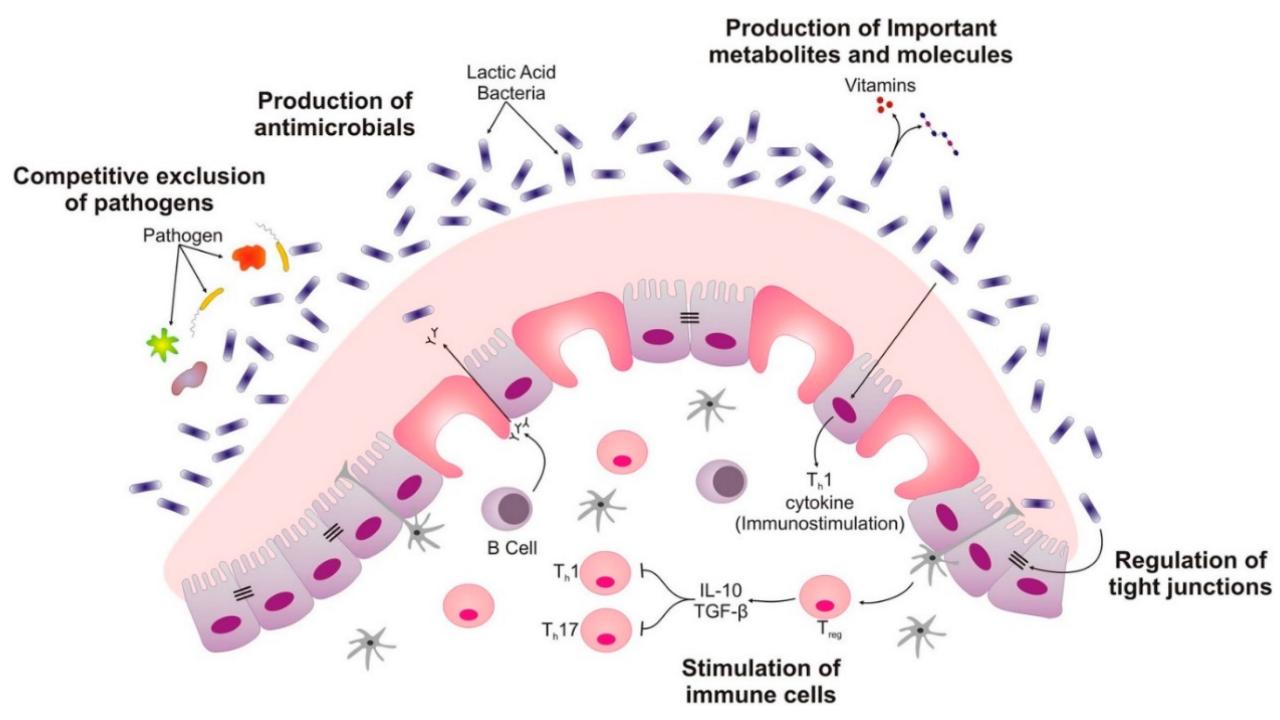
Senior Director
GI & LIver Unit
General Hospital S.Filippo Neri
Club for Evidence Based Gastroenterology & Hepatology (<u>www.EBGH.it</u>)
Rome Italy

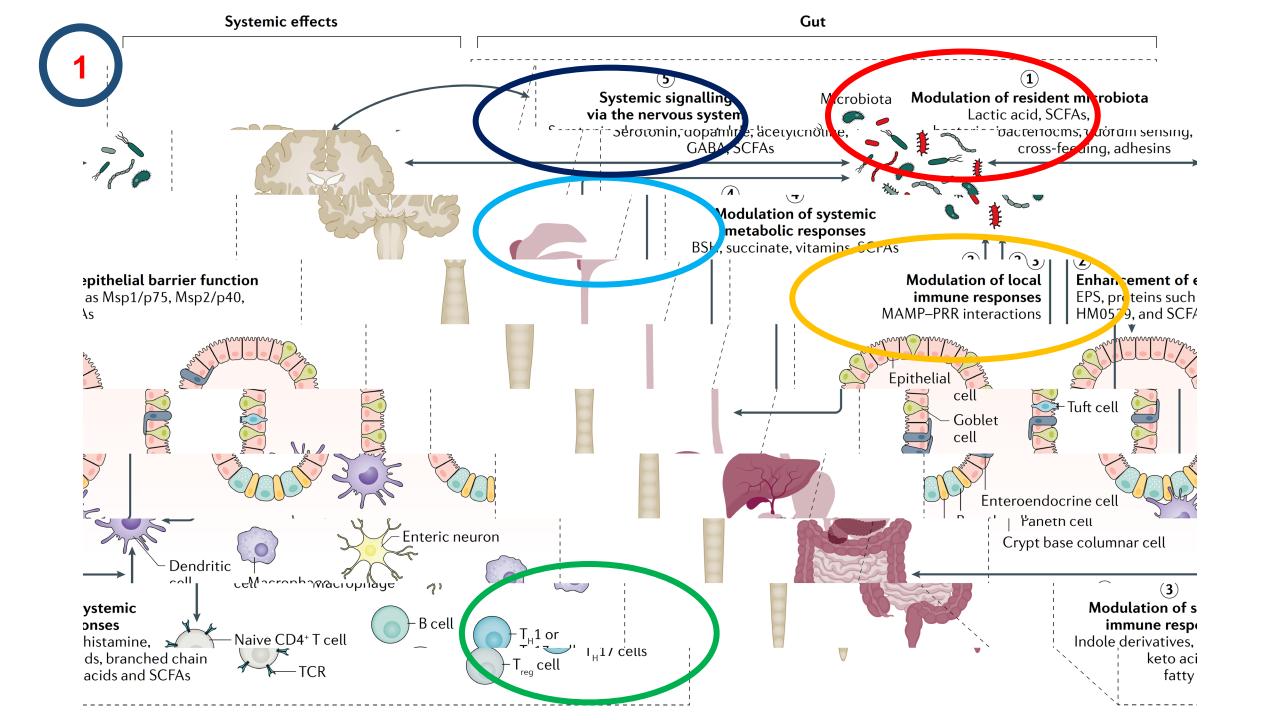






YOU SHOULD START TAKING PROBIOTICS NOW, BEFORE WE DISCOVER THAT THEY DON'T MAKE ANY DIFFERENCE

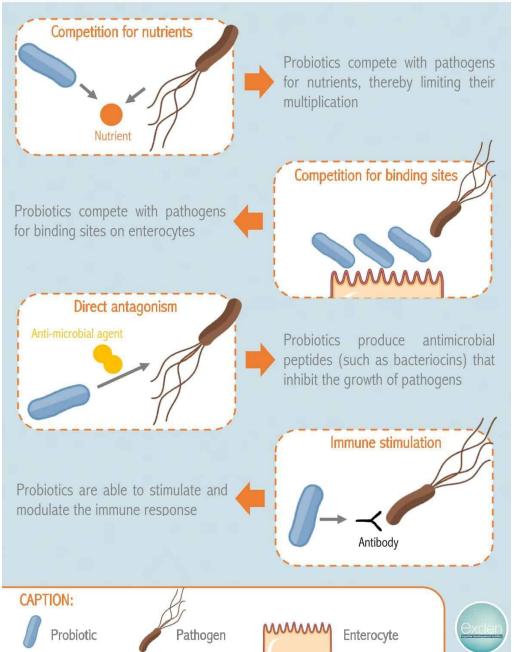




WHAT ARE THE MECHANISMS OF ACTION OF PROBIOTICS AGAINST PATHOGENS?



MODULATION OF RESIDENT BACTERIA



BENEFICIAL MODULATION OF MICROBIOTA

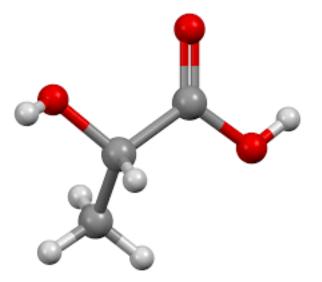
LACTIC ACID

BACTERIOCINS

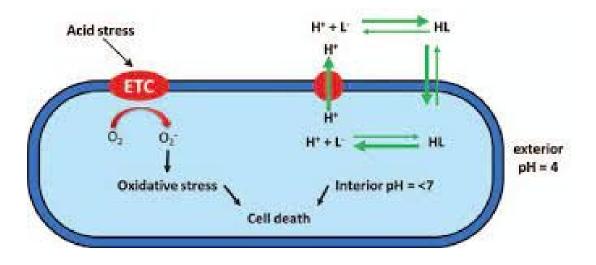
QUORUM SENSING AND QUORUM QUENCHING MOLECULES

SCFAS AND BUTYRATE

COMPETE WITH RESIDENT MICROORGANISMS FOR ADHESION SITES

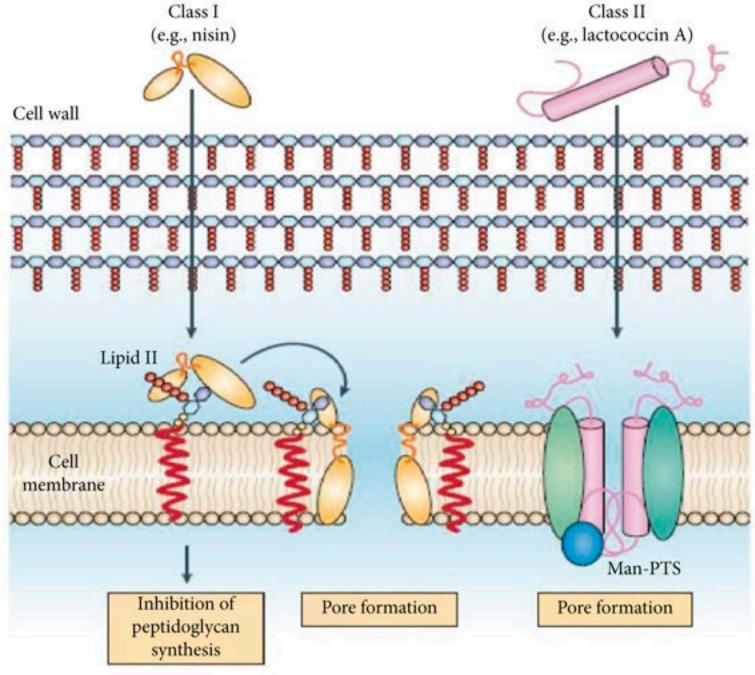


LACTIC ACID



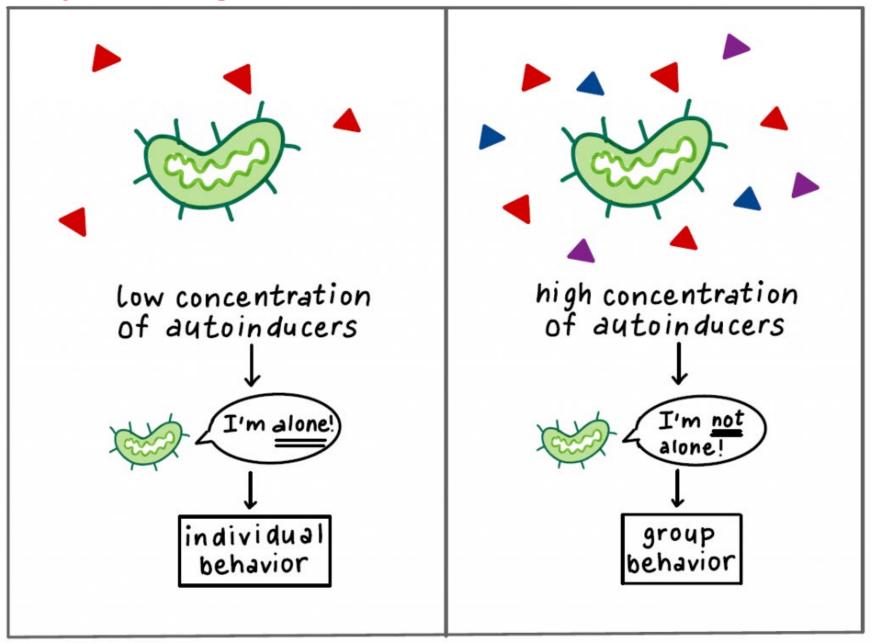
bacteriostatic effects by interfering with pathogen cell membrane functions, leading to membrane permeability, loss of cell contents, lysis, and death

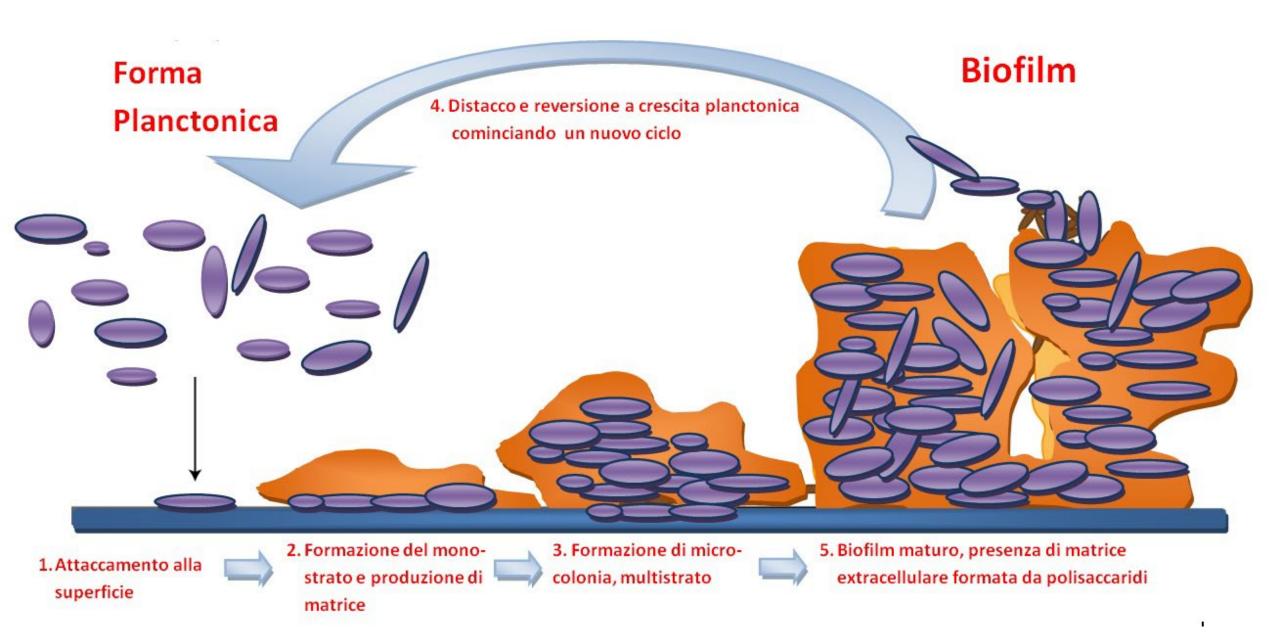
BACTERIOCINS



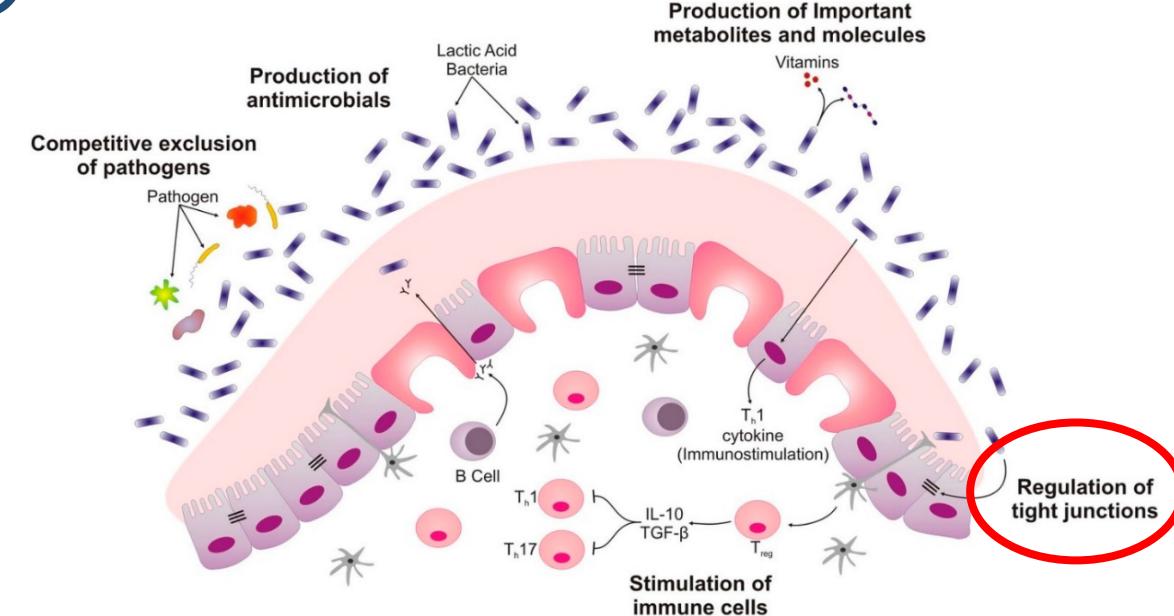
QUORUM SENSING/ quenching

Il quorum-sensing è una forma di comunicazione utilizzata dai batteri per scambiarsi informazioni l'uno con l'altro. Per comunicare, i batteri utilizzano particolari composti chimici (chiamati autoinduttori) che forniscono informazioni sulla popolazione batterica, fra cui informazioni sulle sue dimensioni.

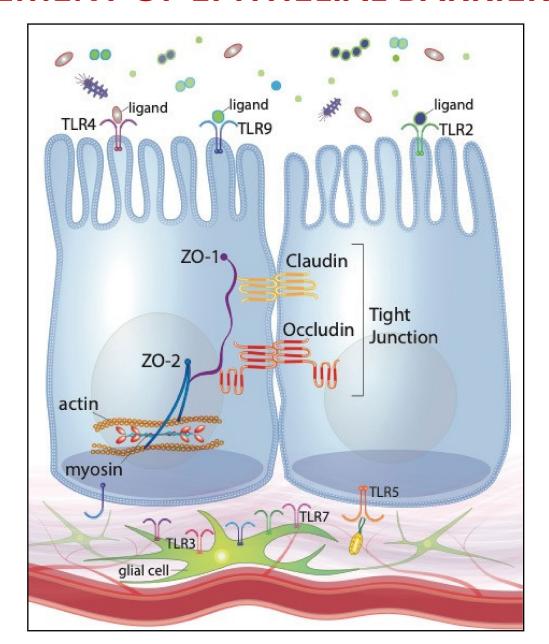




ENHANCEMENT OF EPITHELIAL BARRIER FUNCTION



ENHANCEMENT OF EPITHELIAL BARRIER FUNCTION



SECRETED PROTEINS

MSP1/P75

MSP1/P40

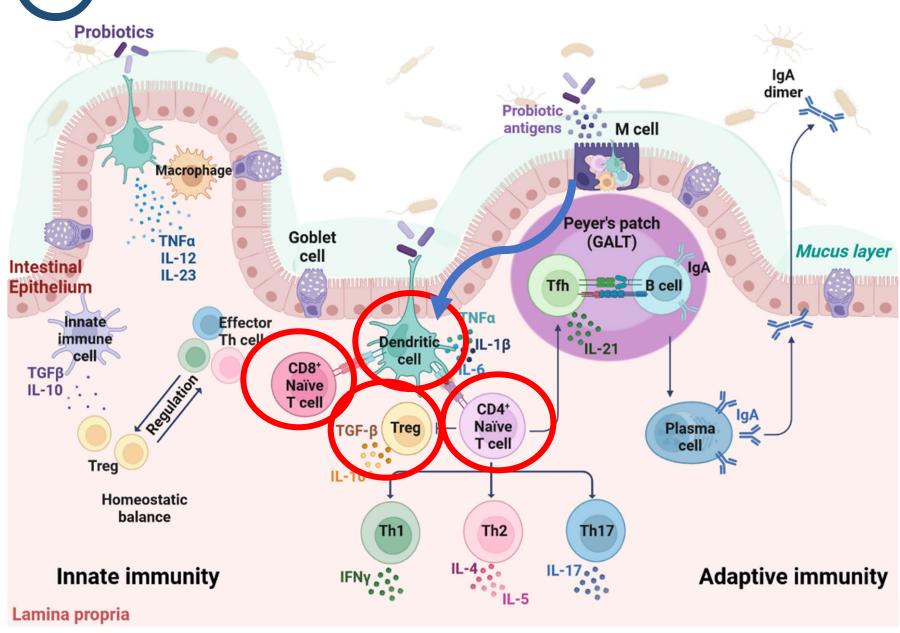
HM0539

EXOPOLYSACCHARIDES

SHORT CHAIN FATTY ACID!

4

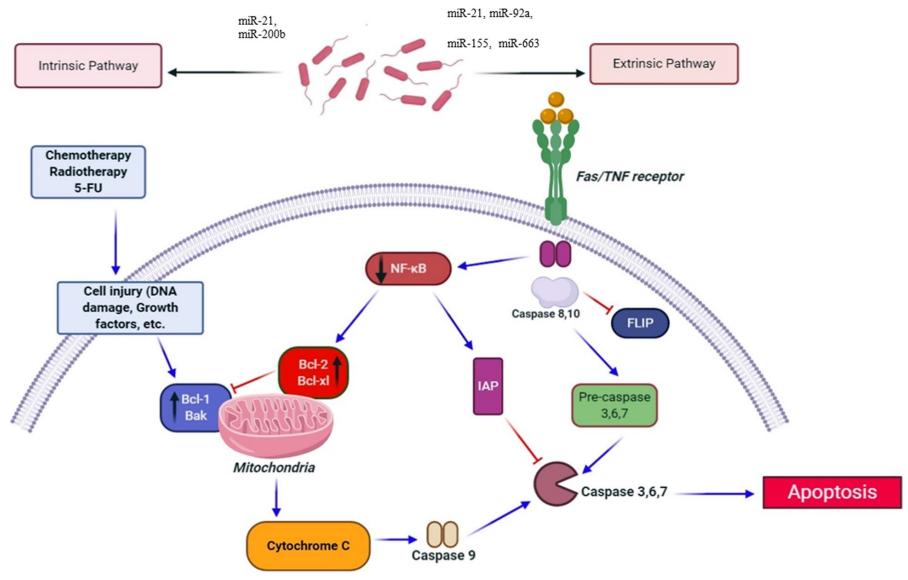
MODULATION OF IMMUNE RESPONSES



- A. ENHANCE SIGNALING IN HOST CELLS TO REDUCE INFLAMMATORY RESPONSE.
- B. SWITCH IN IMMUNE RESPONSE TO REDUCE ALLERGY.
- C. PROBIOTICS AS VEHICLES TO DELIVER ANTI-INFLAMMATORY MOLECULES TO THE INTESTINE.
- D. REDUCE THE PRODUCTION OF INFLAMMATORY SUBSTANCES.

5

EFFECTS OF THERAPEUTIC PROBIOTICS ON MODULATION OF MICRORNAS



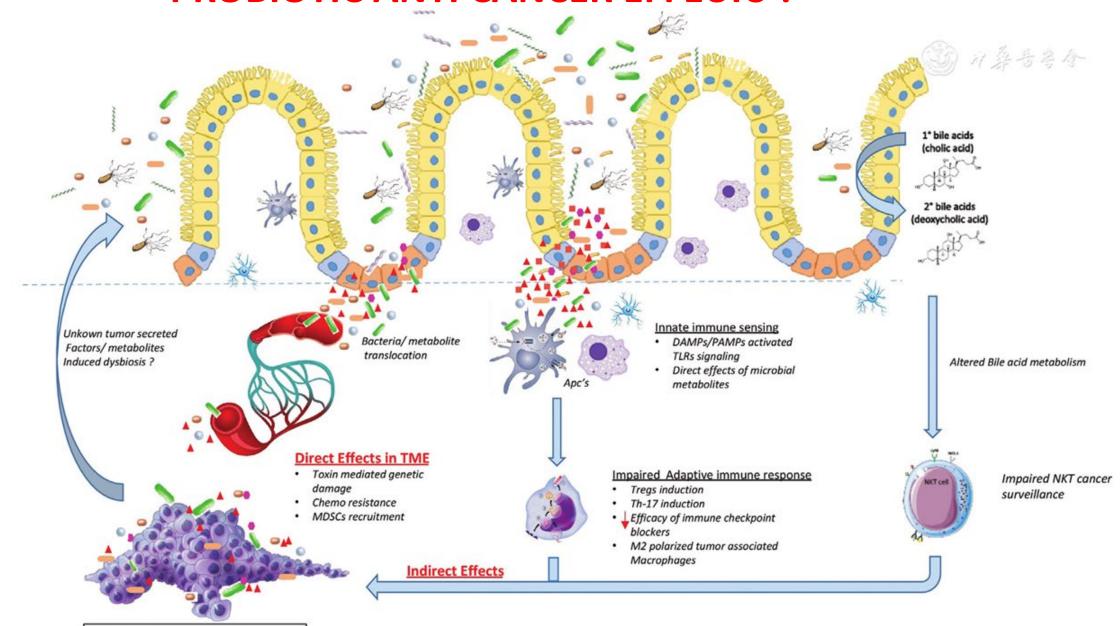
A schema of anti- apoptotic effects of probiotics. Various microRNAs i.e., miR-21, miR-200b and miR-21 can indirectly affect on apoptosis pathways

EFFECTS OF THERAPEUTIC PROBIOTICS ON MODULATION OF MICRORNAS

MicroRNA	Probiotics	Probiotic concentrations	Expression	Target gene	Effects	Model	Sample (n)
miR-215-5p, miR-10b-5p, miR-21-5p, miR-26a-5p, miR-2-3p, miR-10a-5p, miR-148a-3p, miR-194, miR-92-3p, miR-30d, miR-181a-5p, miR-429-3p, let-7f-5p, miR-30a-5p, miR-133a-3p, miR-199-3p, miR-30c-5p, miR-200a-3p, miR-27b-3p	Lactobacillus plantarum Z01 (LPZ01)	1 × 10 ⁸ CFU/mL	Down-regulation of miR-215-5p, miR- 3525, miR-122-5p and up regulation of miR-193a-5p, miR- 375 and miR-215-5p	cAMP-dependent protein kinase activity, stress-activated MAPK cascade, MAPK and Wnt signaling pathways.	Decrease inflammation in S. typhimurium infection in neonatal broiler chicks	In vivo (Newly hatched chicks)	-
miR-135b, miR-155 miR-26b and miR- 18a	Lactobacillus acidophilus and Bifidobacterium bifidum (Bla/016P/M)	1 × 10 ⁹ CFU/g and 1 × 10 ⁹ CFU/g	Up regulation of miR- 135b, miR-155 and down regulation of miR-26b, miR-18a	APC, PTEN, KRAS, and PU.1		In vivo (Mice)	-
miR-423-5p	Enterococcus faecium NCIMB 10415	3.6 × 10 ⁶ CFU/g	Up regulation of miR- 423-5p	Immunoglobulin lambda light C region (IGLC) and immunoglobulin kappa constant (IGKC)	-	<i>In vivo</i> (Landrace pigs)	-
	Lactobacillus rhamnosus GG, Bifidobacterium animalis subsp. lactis Bb-12 and L. acidophilus La-5	5 × 10 ¹⁰ CFU, 5 × 10 ¹⁰ CFU and 5 × 10 ⁹ CFU				Human	54
miR-122a	Lactobacillus rhamnosus GG	1×10 ⁹ CFU	Down-regulation of miR-122a	?	Decrease ethanol- elevated miR122a levels and attenuate ethanol- induced liver injury	In vivo (Mice)	-

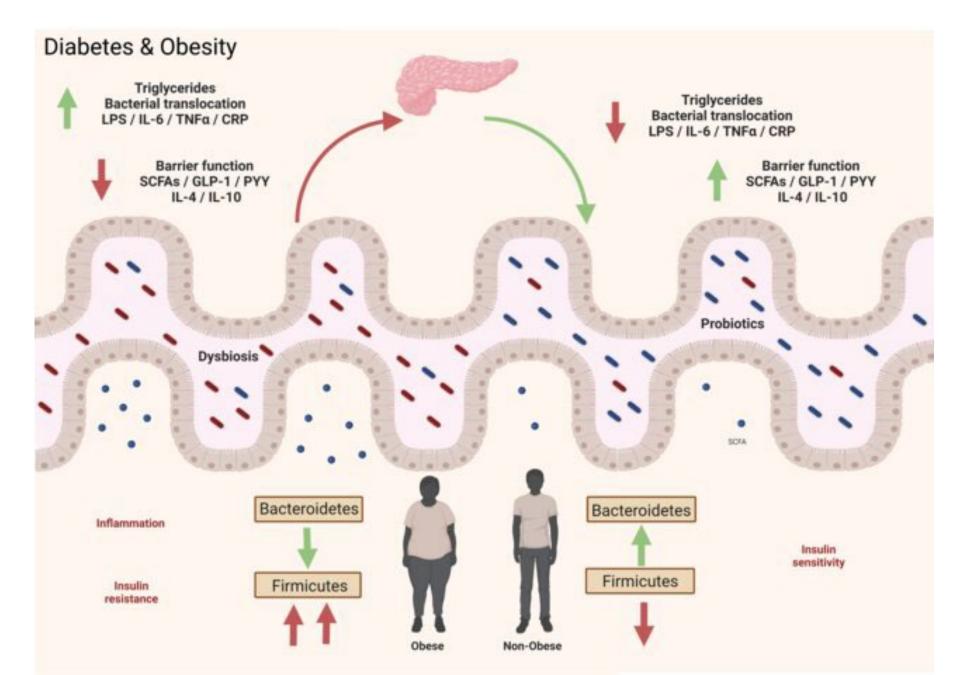
6

PROBIOTIC ANTI CANCER EFFECTS?



Tumor Progression

MODULATION OF SYSTEMIC METABOLIC RESPONSENS



MODULATION OF SYSTEMIC METABOLIC RESPONSENS

BILE SALT HYDROLASE



Glucose, lipid, energy metab (loss: *Clostridium Difficile*)

SUCCINATE



Intestinal gluconeogenesis

SCFA



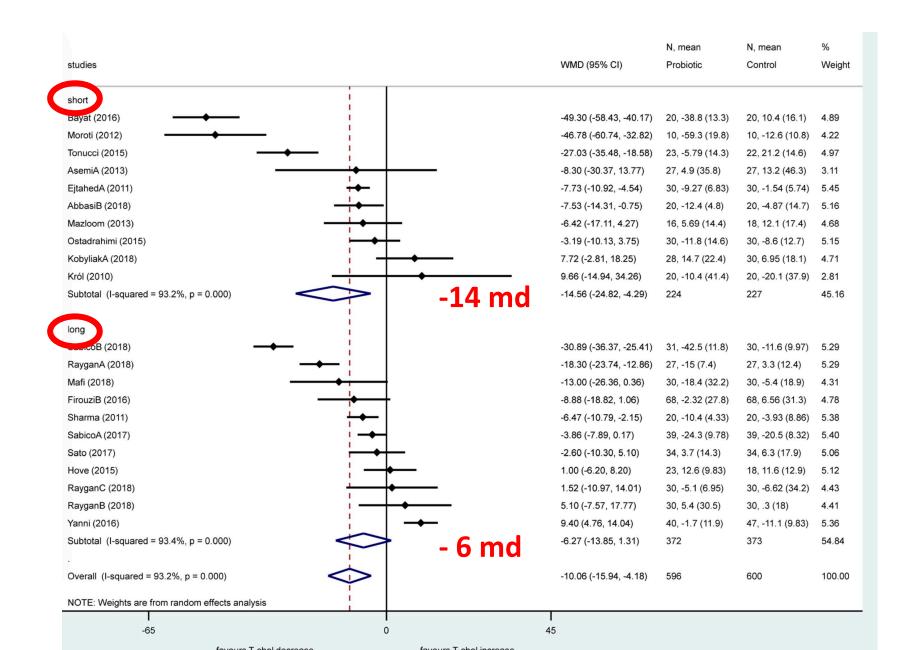
Insulin sensitivity, glucose tolerance; lipid metab.;
Oxidative stress

Probiotics have beneficial metabolic effects in patients with type 2 diabetes mellitus: a meta-analysis of randomized clinical trials

Tícia Kocsis¹, Bálint Molnár¹, Dávid Németh¹, Péter Hegyi¹,², Zsolt Szakács¹,³, Alexandra Bálint¹,⁴, András Garami¹, Alexandra Soós¹, Katalin Márta¹ & Margit Solymár¹⊠

32 TRIALS vs PLACEBO

TOTAL CHOLESTEROL: SHORT vs LONG Tx > 8w



PROBIOTICS

COULD INFLUENCE

BLOOD TESTS,

STRONGLY CORRELATED

TO CLINICAL OUTCOMES

HEART



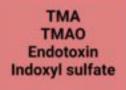


Hypertension

Atherosclerosis Coronary arteries disease

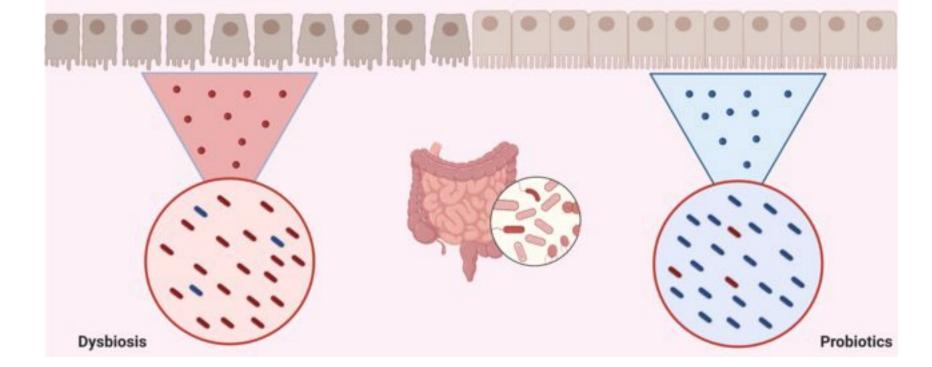


Homeostasis





SCFA Bile acids Anti-inflammatory metabolites





JOURNAL OF THE AMERICAN COLLEGE OF CARDIOLOGY



THE PRESENT AND FUTURE

2023

JACC REVIEW TOPIC OF THE WEEK

Gut Microbiome-Based Management of Patients With Heart Failure





JACC Review Topic of the Week

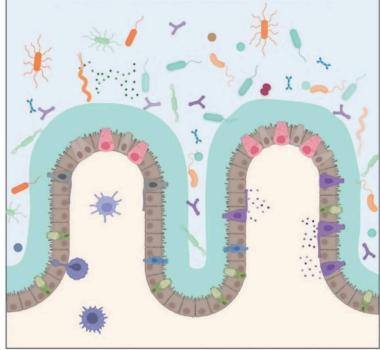
Petra Mamic, MD, a,b Michael Snyder, PhD,b W.H. Wilson Tang, MD^c

ABSTRACT

Despite therapeutic advances, chronic heart failure (HF) is still associated with significant risk of morbidity and mortality. The course of disease and responses to therapies vary widely among individuals with HF, highlighting the need for precision medicine approaches. Gut microbiome stands to be an important aspect of precision medicine in HF. Exploratory clinical studies have revealed shared patterns of gut microbiome dysregulation in this disease, with mechanistic animal studies providing evidence for active involvement of the gut microbiome in development and pathophysiology of HF. Deeper insights into gut microbiome-host interactions in patients with HF promise to deliver novel disease biomarkers, preventative and therapeutic targets, and improve disease risk stratification. This knowledge may enable a paradigm shift in how we care for patients with HF, and pave the path toward improved clinical outcomes through personalized HF care. (J Am Coll Cardiol 2023;81:1729–1739) © 2023 by the American College of Cardiology Foundation.

Healthy Gut

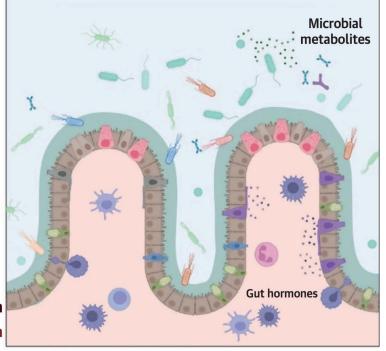
- Strong gut barrier function
- High microbiome α -diversity
- Potential pathogens suppressed
- Abundant anti-inflammatory microbes



Gut lumen
Intestinal
mucus layer
Gut epithelium
Lamina propria

Gut in Chronic Heart Failure

- Weakened gut barrier function
- Reduced microbiome α -diversity
- Pathogen overgrowth
- Depleted anti-inflammatory microbes



Key features of the heart failure-associated gut milieu and microbiome compared with healthy microbiome include impaired intestinal barrier function, reduced microbiome α -diversity, pathogen overgrowth, and loss of anti-inflammatory gut microbes.

HIGHLIGHTS

- Gut microbiome modulates HF pathophysiology, contributes to disease progression and therapeutic responses, and holds promise as a novel biomarker.
- Interactions among the gut microbiome, diet, and medications offer potentially innovative modalities for management of patients with HF.
- Interdisciplinary collaboration will facilitate translation of precision gut microbiomics to the clinical evaluation and management of patients with HF.

Microbiome-derived metabolites comprise about 10% of all circulating metabolites among humans.

- •The review highlighted particular metabolites that affect HF:
 - **Lipopolysaccharides** promote production of inflammatory cytokines and induce insulin resistance and atherothrombosis, leading to cardiac dysfunction
 - In contrast, **short-chain fatty acids** reduce systemic inflammation and augment the intestinal barrier. These molecules are also associated with benefits in cardiac remodeling; however, the concentration of short-chain fatty acids is reduced in cases of HF
 - TMAO is a metabolite related to the consumption of dietary L-carnitine and (phosphatidyl) choline, which are commonly found in animal products. TMAO has atherogenic and thrombotic effects, as well as negative effects on cardiac remodeling in mouse models.
 - Phenacetylglutamine reduces cardiomyocyte contractility and induces natriuretic gene expression, and adults with HF have higher mean levels of this metabolite as well

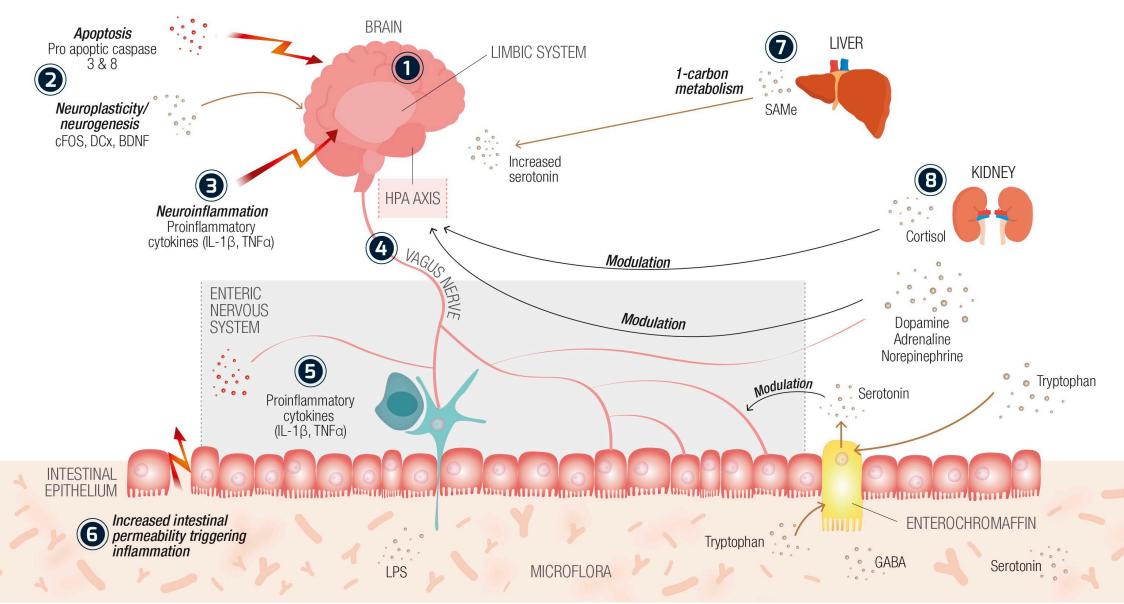
TABLE 2 Design of Future HF Microbiome Studies				
Study Feature	Rationale			
General study design	Combining population-based and N-of-1 approach (in which each person serves as his or her own control) enables both group- and individual-level analyses, which is necessary to gain insights into highly			

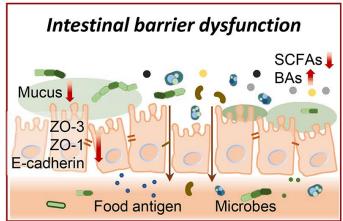
Combining population-based and N-of-1 approach (in which each per serves as his or her own control) enables both group- and individ level analyses, which is necessary to gain insights into highly personalized systems such as microbiome.

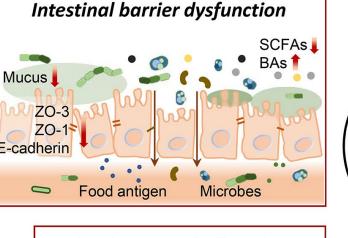
	through multiomic approaches. Simultaneous clinical assessments—including HF functional status, cardiac imaging, biomarkers, and data on outcomes—are equally important to accurately relate microbiome activity to HF physiology.
Confounders	Given their impact on the microbiome community, relevant microbiome confounders—including comorbidities, medications, dietary, and lifestyle habits—need to be catalogued carefully to interpret study findings correctly.
Feasibility	To scale HF-microbiome research efforts, existing HF biobanks, consortia, and clinical trials could be leveraged.
HF = heart failure.	

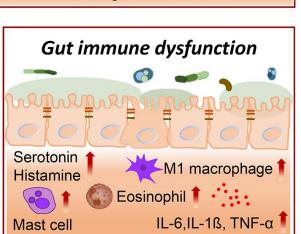
SYSTEMATIC SIGNALLING VIA THE NERVOUS SYSTEM

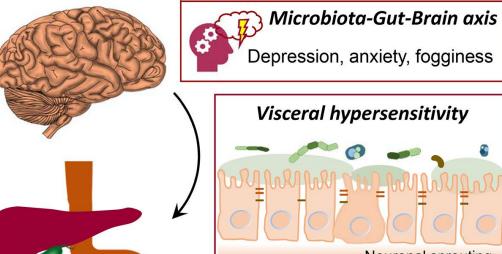




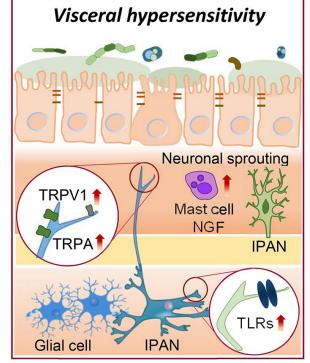


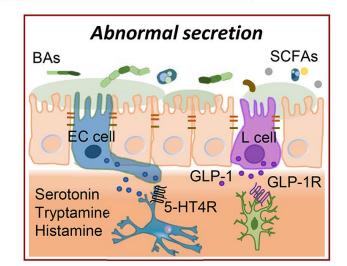


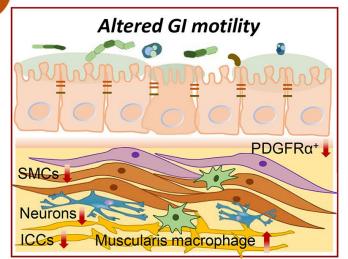




--







"Most diseases are characterized by a pathobiome"

This term is unfortunately overly simplistic and inherently flawed. Microorganisms and their metabolites are neither 'good' nor 'bad', they merely exist. Their impacts on us as hosts are heavily dependent on context. Microorganisms or metabolites that are deleterious in one context may cause no harm in another. As examples, Clostridioides difficile can be carried asymptomatically throughout life, and only cause problems in older age when the host is immunocompromised and treated with antibiotics. Similarly, a strain of E. coli may be relatively harmless in the colon, but cause a urinary tract infection if it invades the urethra.

It is true, however, that numerous human conditions have been shown to correlate with alterations in microbiota composition. This is sometimes referred to as 'dysbiosis', which is also a vague term with limited clinical applicability.

to disease progression in some conditions, including inflammatory bowel diseases: however, such alterations are rarely consistent and the microbiota is hugely variable between individuals, both in health and disease. This makes it extremely difficult to identify gut microbiota configurations with the required specificity and reproducibility for clinical practice



REPLICABILITY

Developing a new conceptual framework and applying it to the human microbiome will require much more collaboration between investigators working across disparate fields, including evolution, ecology, microbiology, biomedicine and computational biology. It will also demand significant changes in how data and other resources are distributed between scientists, and in how currently disparate areas of microbiome research inter-relate.

Data standards. Microbiome researchers have not yet broadly embraced quality-control practices for their data in a way that would make results more reproducible, and that would facilitate the analysis and interpretation of data across multiple studies.

Studies based on characterizing genetic material, proteins or metabolites using high-throughput analyses will remain the norm for the foreseeable future. To produce useful results, however, researchers must adopt better data-sharing practices.

MEDICINA DEI PROBIOTICI

HOME / AREE TEMATICHE / MEDICINA DEI PROBIOTICI

ATTENZIONE
RICERCATORI: OCCHIO
ALLA REPLICABILITA'!
COME RENDERLA PIU'
SISTEMATICA E
MIGLIORARE LE
EVIDENZE! DA"NATURE"
(FREE)

GASTROENTEROLOGIA, LINEE GUIDA EBM.

MEDICINA DEI PROBIOTICI,

METODOLOGIA, NEWSLETTER

Di Redazione • 27 Settembre 2023

Replication games: how to make reproducibility research more systematic Abel Brodeur, Anna Dreber, Fernando Hoces de la Guardia & Edward Miguel...

Approfondisci... >

PROBIOTICI E COLON IRRITABILE. ULTIMISSIMA META-ANALISI DA "GASTROENTEROLOGY" (FREE). PECCATO CHE NON SI DISTINGUA TRA FORME STIPSI E DIARREA!

GASTROENTEROLOGIA, IBS,
MALATTIA DIVERTICOLARE,
MEDICI DI FAMIGLIA,
MEDICINA DEI PROBIOTICI,

MEDICINA INTERNA, NEWSLETTER

Di Redazione • 5 Agosto 2023

Efficacy of Probiotics in Irritable Bowel Syndrome: Systematic Review and Meta-analysis Vivek C. Goodoory et al.

BACKGROUND & AIMS: Some ...

TRAPIANTO FECALE: LA
PRIMA CONFERENZA
INTERNAZIONALE PER
TENTARE LA
STANDARDIZZAZIONE
DELLA TECNICA NELLE
IBD! FINALMENTE!
DA"BMJ" (FREE)

GASTROENTEROLOGIA, IBD, MEDICINA DEI PROBIOTICI,

MEDICINA INTERNA, NEWSLETTER, Senza categoria

Di Redazione • 7 Luglio 2023

The first international Rome consensus conference on gut microbiota and faecal microbiota transplantation in inflammatory bowel disease Loris Riccardo...

Approfondisci... ▶

NEWSLETTER

Email

Iscriviti

I NOSTRI CORSI

Torgiano a Roma 2019

CONTRIBUTI VIDEO - BROCHURE

Notizie da EBGH.it - Gastroenterologia in evidenza

Dante, Paradiso, XXII, 67-69

Se siete interessati ad un accesso unlimited segnalate la vs. email alla Segreteria.

Riceverete ID e PASSWORD



top ten in gastroenterologia

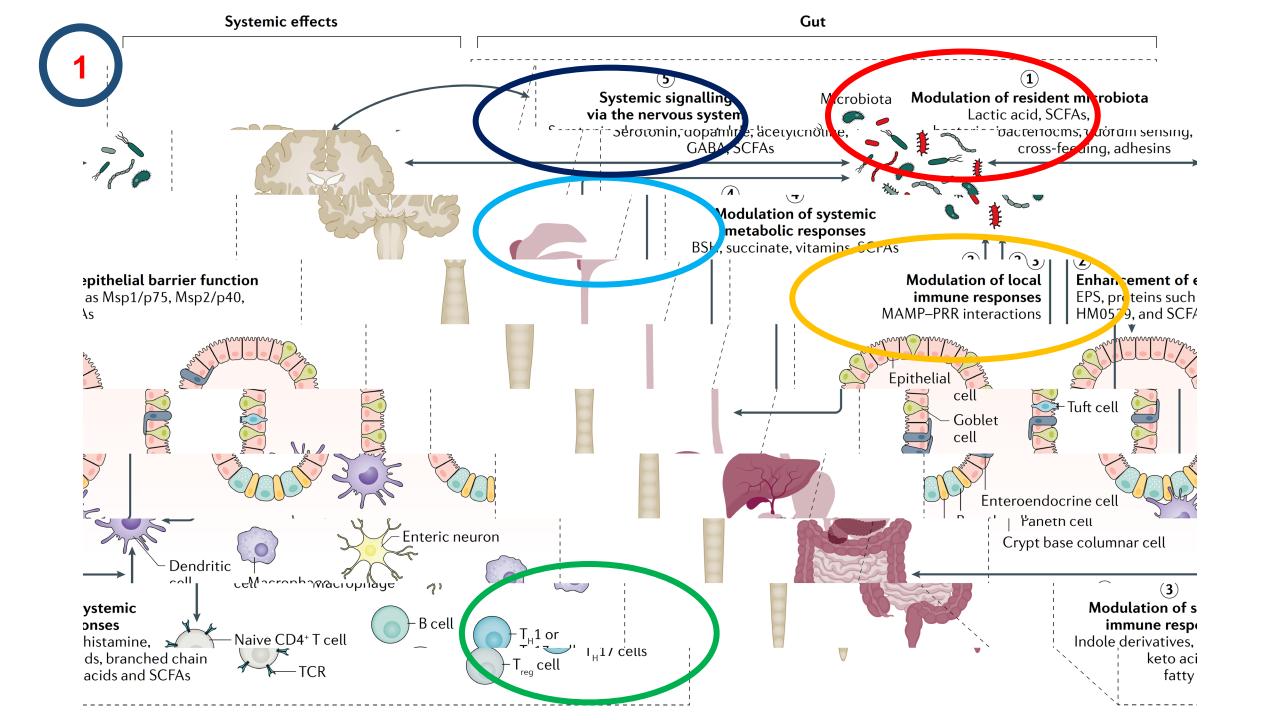
14[^] EDIZIONE

24-25 NOVEMBRE 20**23**

BERGAMO

HOTEL EXCELSIOR SAN MARCO Piazza della Repubblica, 6

TOP TEN Slides



BENEFICIAL MODULATION OF MICROBIOTA

LACTIC ACID

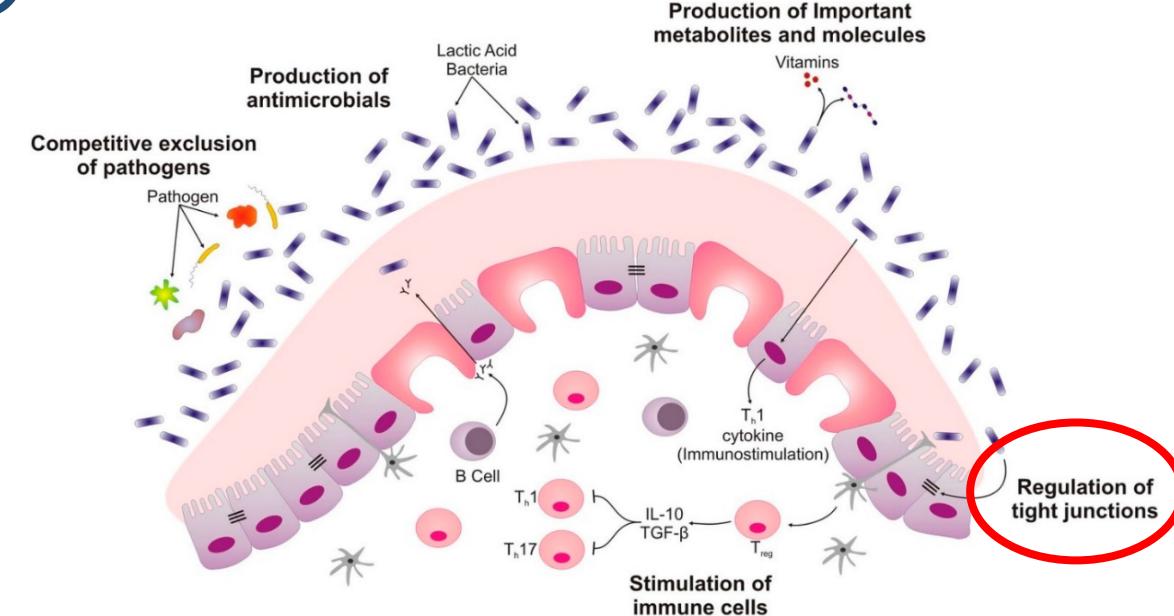
BACTERIOCINS

QUORUM SENSING AND QUORUM QUENCHING MOLECULES

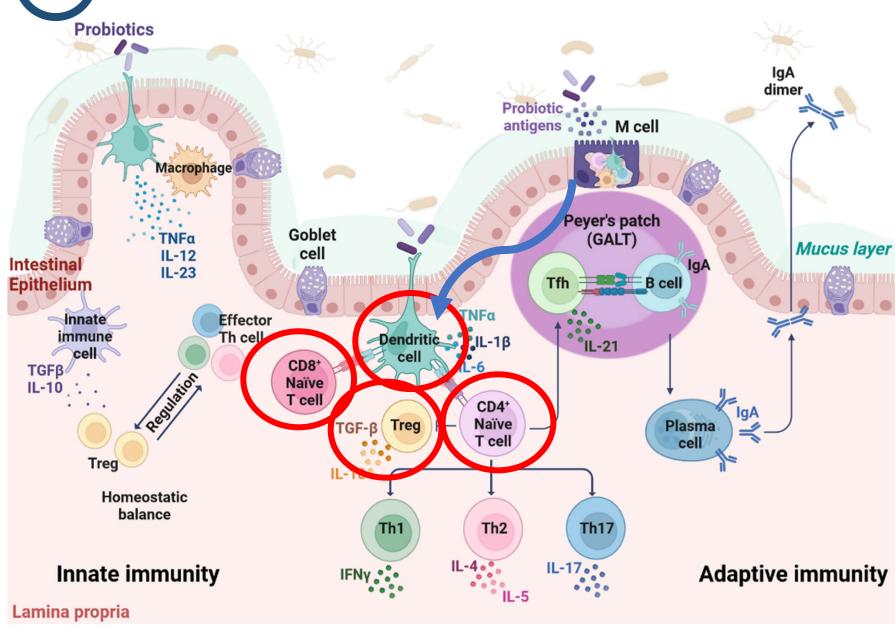
SCFAS AND BUTYRATE

COMPETE WITH RESIDENT MICROORGANISMS FOR ADHESION SITES

ENHANCEMENT OF EPITHELIAL BARRIER FUNCTION

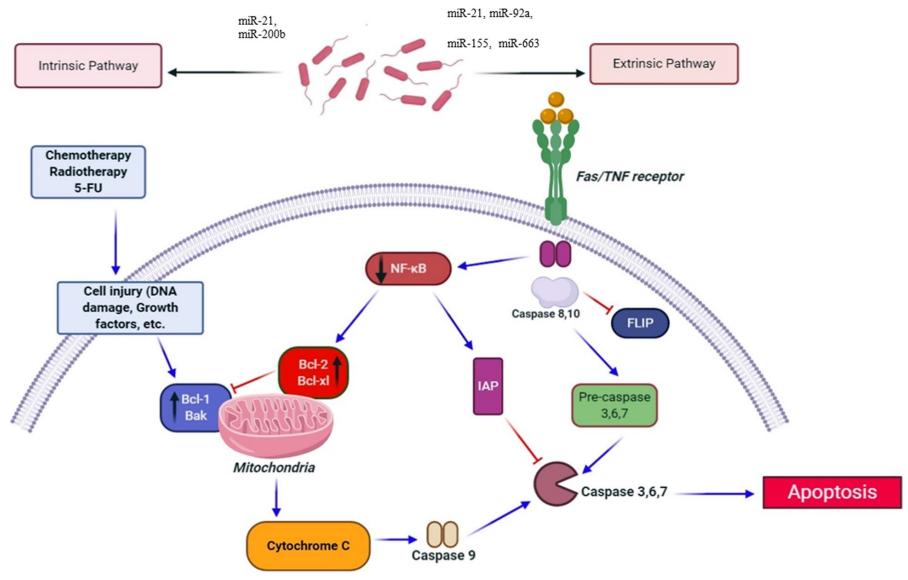


4 MODULATION OF IMMUNE RESPONSES



5

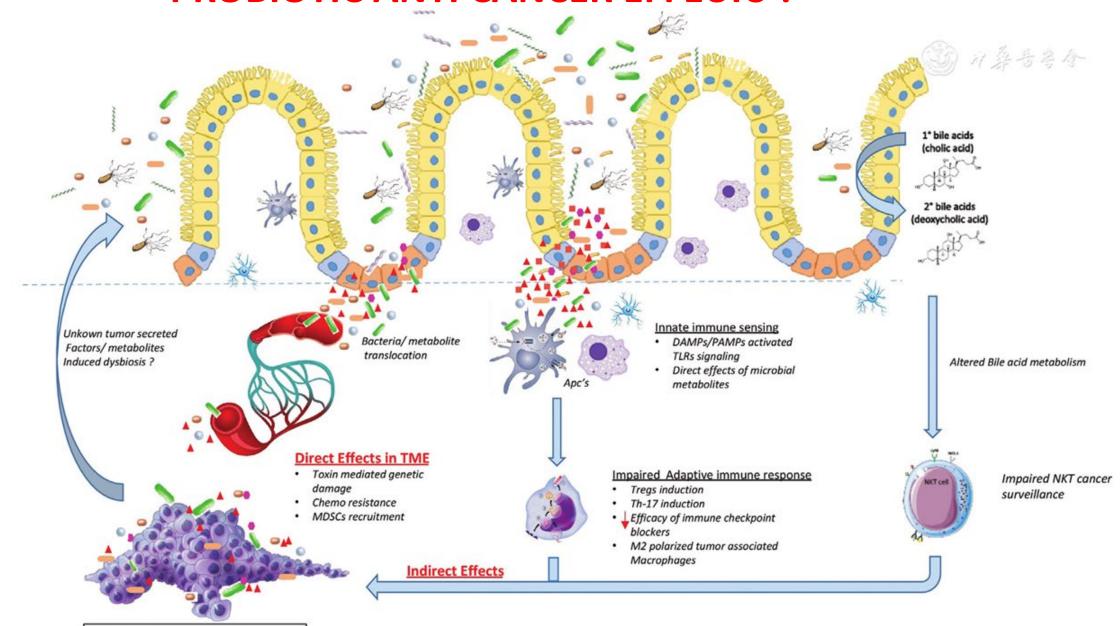
EFFECTS OF THERAPEUTIC PROBIOTICS ON MODULATION OF MICRORNAS



A schema of anti- apoptotic effects of probiotics. Various microRNAs i.e., miR-21, miR-200b and miR-21 can indirectly affect on apoptosis pathways

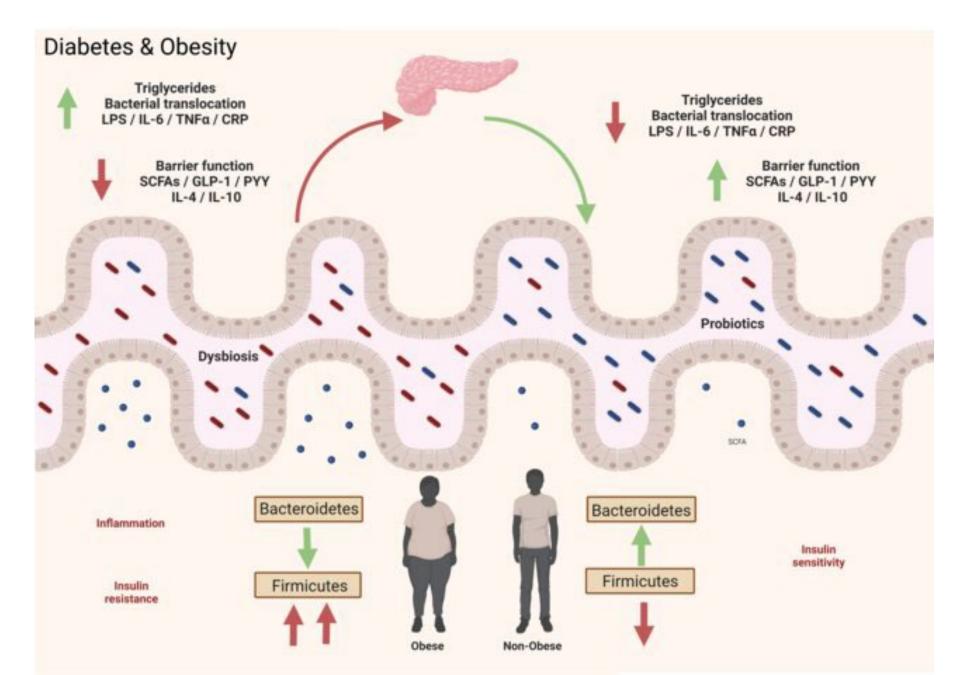
6

PROBIOTIC ANTI CANCER EFFECTS?



Tumor Progression

MODULATION OF SYSTEMIC METABOLIC RESPONSENS





JOURNAL OF THE AMERICAN COLLEGE OF CARDIOLOGY



THE PRESENT AND FUTURE

2023

JACC REVIEW TOPIC OF THE WEEK

Gut Microbiome-Based Management of Patients With Heart Failure





JACC Review Topic of the Week

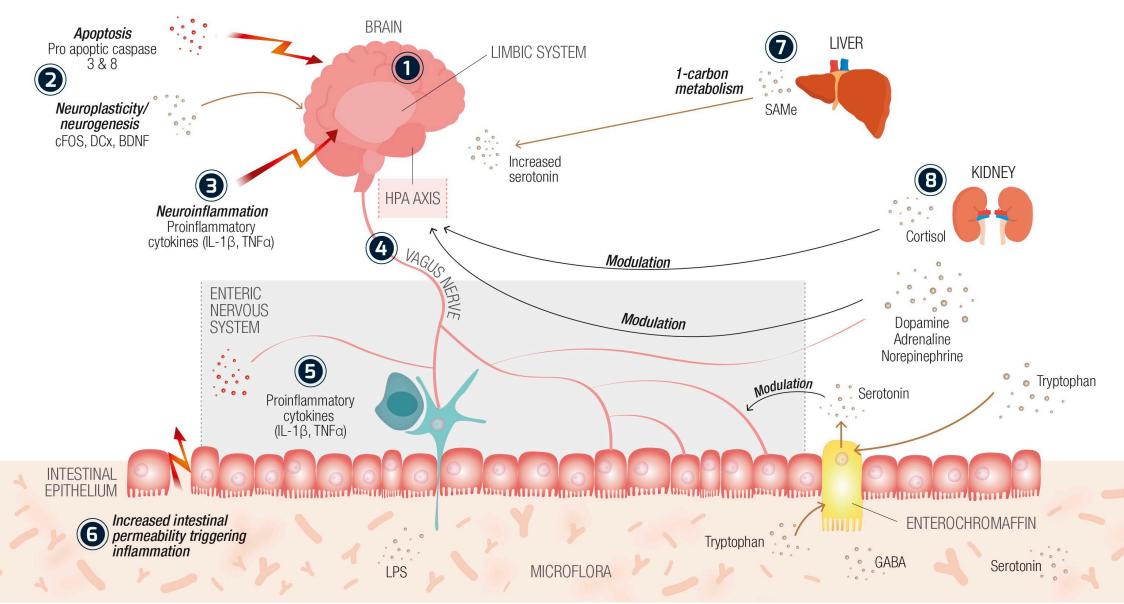
Petra Mamic, MD, a,b Michael Snyder, PhD,b W.H. Wilson Tang, MD^c

ABSTRACT

Despite therapeutic advances, chronic heart failure (HF) is still associated with significant risk of morbidity and mortality. The course of disease and responses to therapies vary widely among individuals with HF, highlighting the need for precision medicine approaches. Gut microbiome stands to be an important aspect of precision medicine in HF. Exploratory clinical studies have revealed shared patterns of gut microbiome dysregulation in this disease, with mechanistic animal studies providing evidence for active involvement of the gut microbiome in development and pathophysiology of HF. Deeper insights into gut microbiome-host interactions in patients with HF promise to deliver novel disease biomarkers, preventative and therapeutic targets, and improve disease risk stratification. This knowledge may enable a paradigm shift in how we care for patients with HF, and pave the path toward improved clinical outcomes through personalized HF care. (J Am Coll Cardiol 2023;81:1729–1739) © 2023 by the American College of Cardiology Foundation.

SYSTEMATIC SIGNALLING VIA THE NERVOUS SYSTEM







REPLICABILITY

Developing a new conceptual framework and applying it to the human microbiome will require much more collaboration between investigators working across disparate fields, including evolution, ecology, microbiology, biomedicine and computational biology. It will also demand significant changes in how data and other resources are distributed between scientists, and in how currently disparate areas of microbiome research inter-relate.

Data standards. Microbiome researchers have not yet broadly embraced quality-control practices for their data in a way that would make results more reproducible, and that would facilitate the analysis and interpretation of data across multiple studies.

Studies based on characterizing genetic material, proteins or metabolites using high-throughput analyses will remain the norm for the foreseeable future. To produce useful results, however, researchers must adopt better data-sharing practices.



E' STATO COME AL PROCESSO DI NORIMBERGA Woody Allen





La pace è l'unica battaglia che vale la pena combattere. Albert Camus