



FAKULTAS
KEDOKTERAN



Designing Next-Generation Probiotics and Prebiotics for Improved Human Health

Rina Agustina

Department of Nutrition

Faculty of Medicine, Universitas
Indonesia; Dr. Cipto Mangunkusumo
Hospital

Human Nutrition Research Center, IMERI
Jakarta, Indonesia

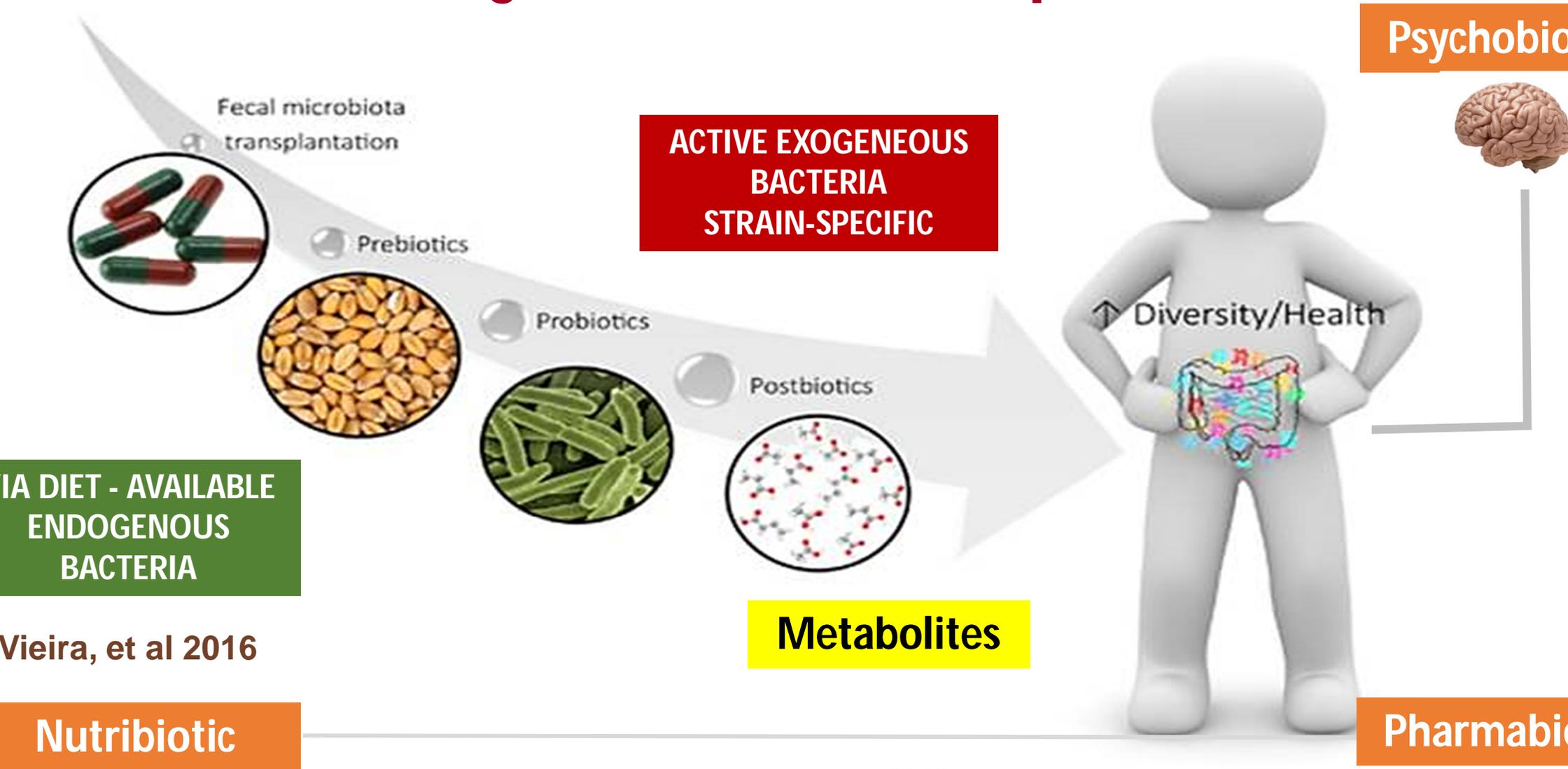
ILSI INDIA CONFERENCE DECEMBER 6, 2018



OUTLINE

- Potential strategies on modulating gut microbiota composition
- 'Current Generation' of Probiotics
- Advances Metagenomics
- Next Generation of Probiotics versus Life Bacterio-therapeutic product
- Novel Prebiotics
- Technical, Research and Regulatory challenge
- Conclusion

Potential strategies aimed at modulation of the gut microbiota composition



Vieira, et al 2016

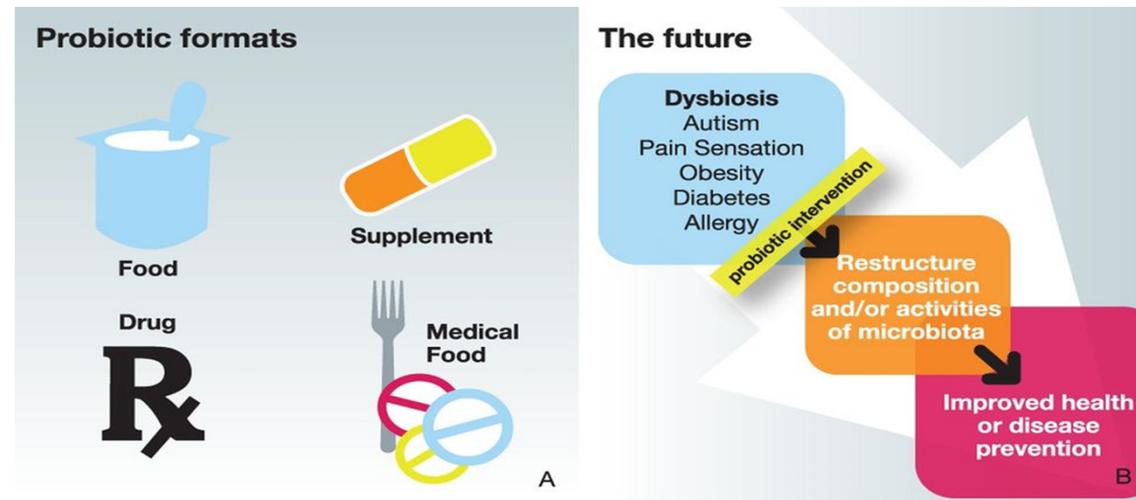
CURRENT PROBIOTIC EFFICACY

Some probiotics and prebiotics are suggested to play roles in alleviating disease symptoms, protection against infections, inflammatory bowel diseases, and atopic diseases, modulating the immune system, obesity and neurological problem by improving the beneficial gut microbiota colonization.

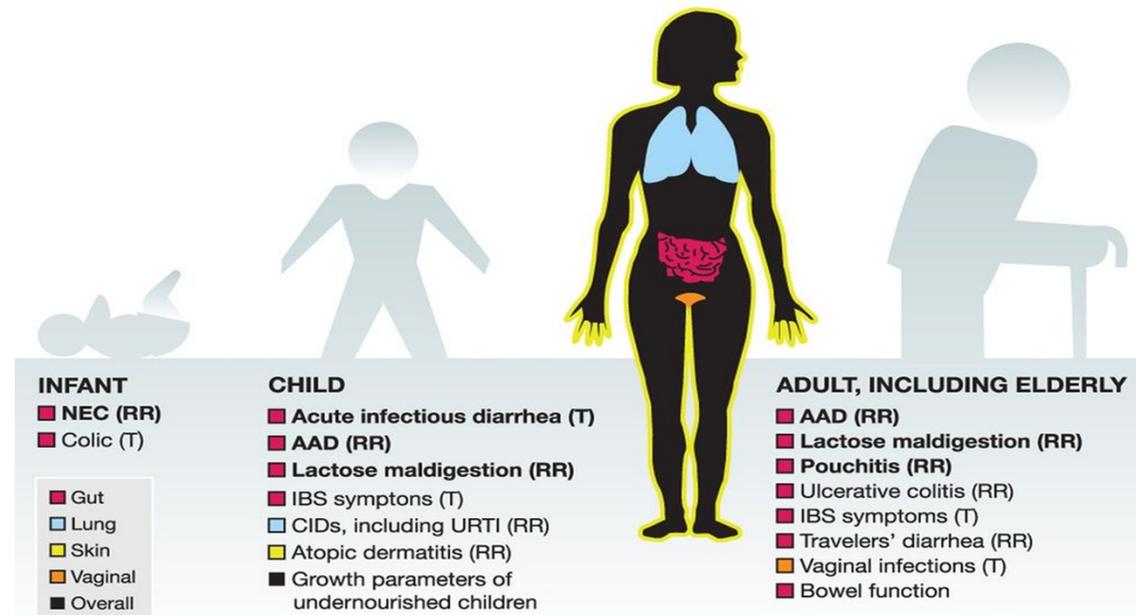
The best documented clinical areas for probiotics:
 Treatment of acute gastroenteritis in children
 Prevention of antibiotic-associated diarrhea both in children and in adults

However, the main existing probiotics to consumers are generally produced from a limited range of organisms (genera *Lactobacillus* – *Bifidobacterium*).

Rina Agustina - 2018



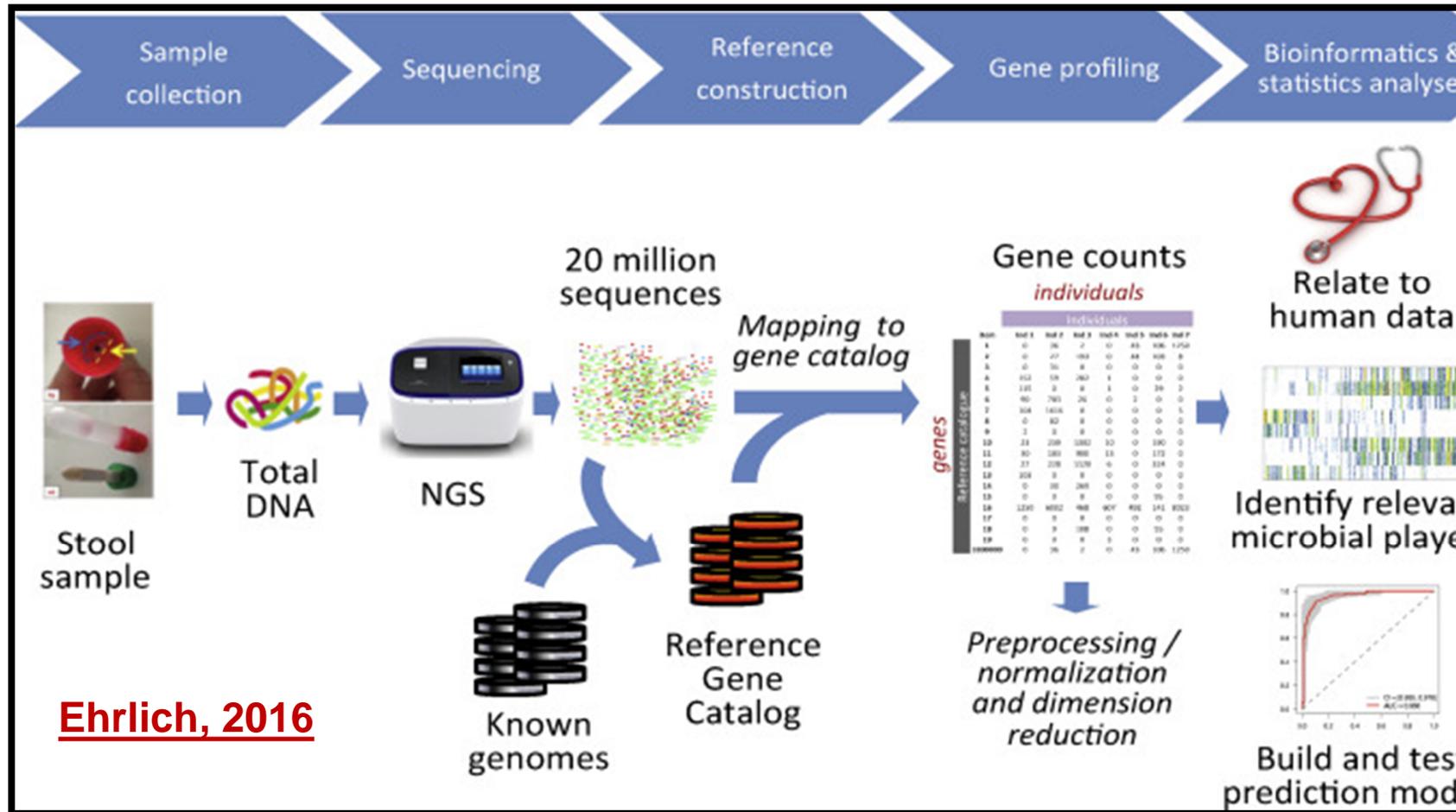
Health and clinical targets addressed by studies on orally administered probiotics conducted in human subjects of different ages



Sanders, BMJ, 2012

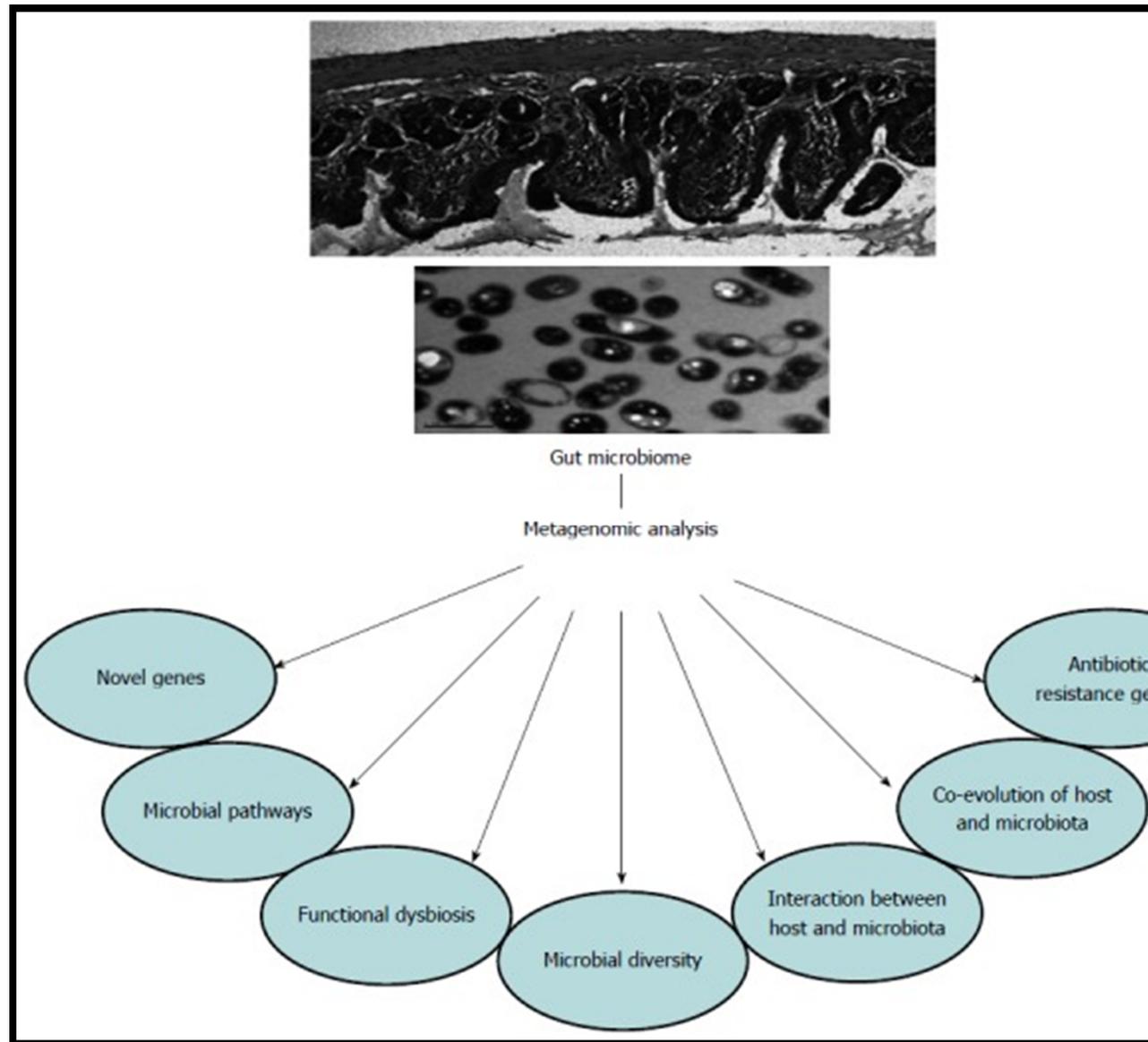
Metagenomics approach and bioinformatics analyses

- have provided the opportunity to study complex microbial communities and
- help ascertain general patterns that regulate human microbial ecosystems.



Metagenomics approach and bioinformatics analyses

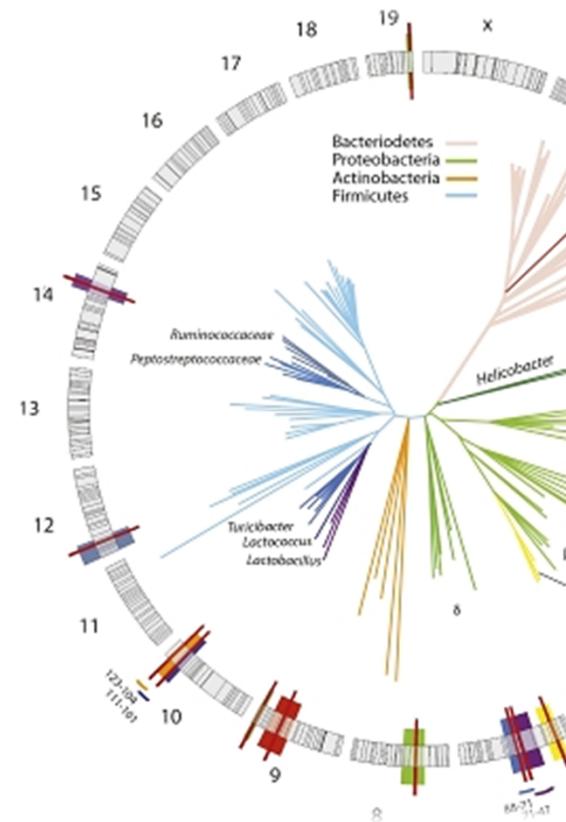
With this advancement, the discovery of new-generation probiotics and prebiotics will continue to evolve in modulating the intestinal and extra-intestinal health.





GUT MICROBIOTA "A COMPLEX COMMUNITY"

high complexity and (interpersonal) diversity



human cells and microbes, human and microbial genome

encode diverse functions and pathways

together affecting human metabolic and inflammatory features

In the population

- 1100 species
- 10 mil genes

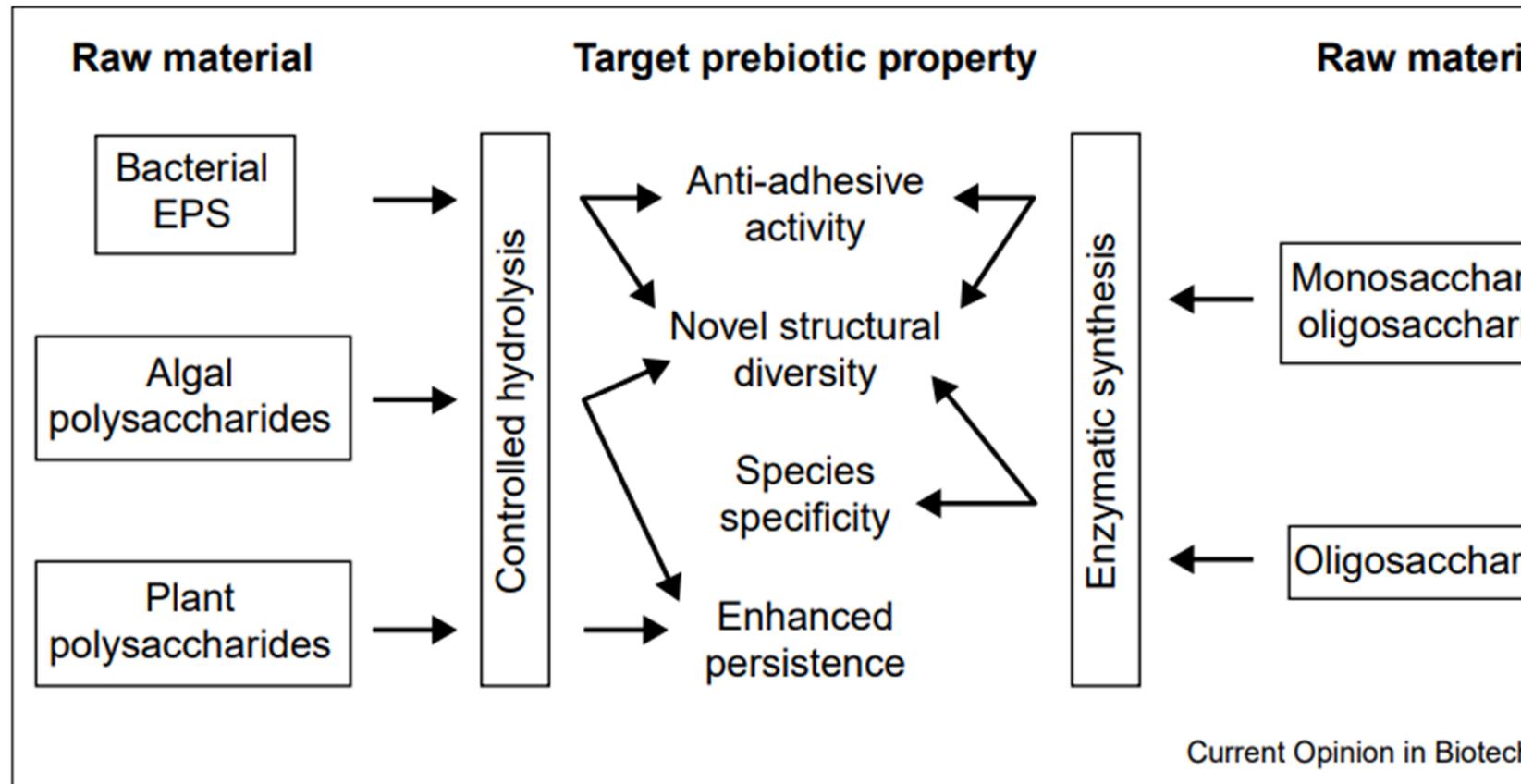
■ In each individual

- At least 160 species
- Approx 540k genes – we share almost 40% of these genes with any other person

Benson, 2010; Buccigrossi & Sommer & Bäckhed, 2012

Biotechnological manufacture of enhanced prebiotics

- At the same time, new manufacturing biotechnologies for prebiotics, and better understanding on the oligosaccharides metabolism by probiotics
- are facilitating the development of prebiotics and synbiotics for specific functional properties and health outcomes.



Why Next-generation Probiotic and Prebiotic?

Probiotics

- Limited list of genera, which mainly include *Lactobacillus* spp. and *Bifidobacterium* spp.
- The main existing probiotics to consumers are generally produced from a limited range of organisms, mainly sourced from the gut or from traditional fermented foods, such as pickles, yoghurts and kefir grains.

Prebiotics

- Several commercial prebiotics (or as synbiotics) products worldwide such as soybean oligosaccharides raffinose and stachyose, glycanases, Arabinogalacto-oligosaccharides, Pectic oligosaccharides Gentio-oligosaccharides or other, need to be thoroughly tested, that can be potential for development of novel prebiotics

Timeline of selected milestones in the history of probiotics and next-generation probiotics

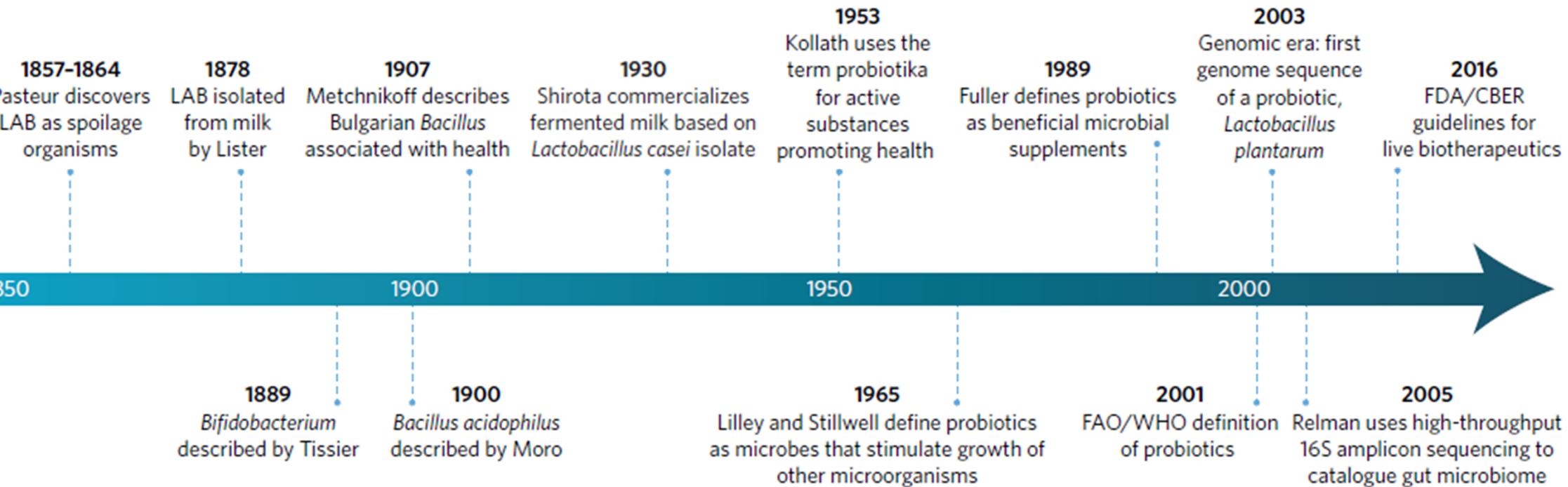


Figure 1 | Timeline of selected milestones in the history of probiotics and next-generation probiotics. LAB, lactic acid bacteria.

O'Toole, P., Marchesi, J., & Hill, C. (2017). Next-generation probiotics: the spectrum from probiotics to live biotherapeutics. *Nature Microbiology*, 2(5). doi: 10.1038/nmicrobiol.2017.57

Probiotics

FAO and WHO definition In 2001

Probiotics: *“Live microorganisms which when administered in adequate amounts, confer a health benefit on the host”*

Essential: Microbial, Viable, Beneficial to health and **Strain-specific**

The active bacteria are mostly *Lactobacillus* and *Bifidobacteria*

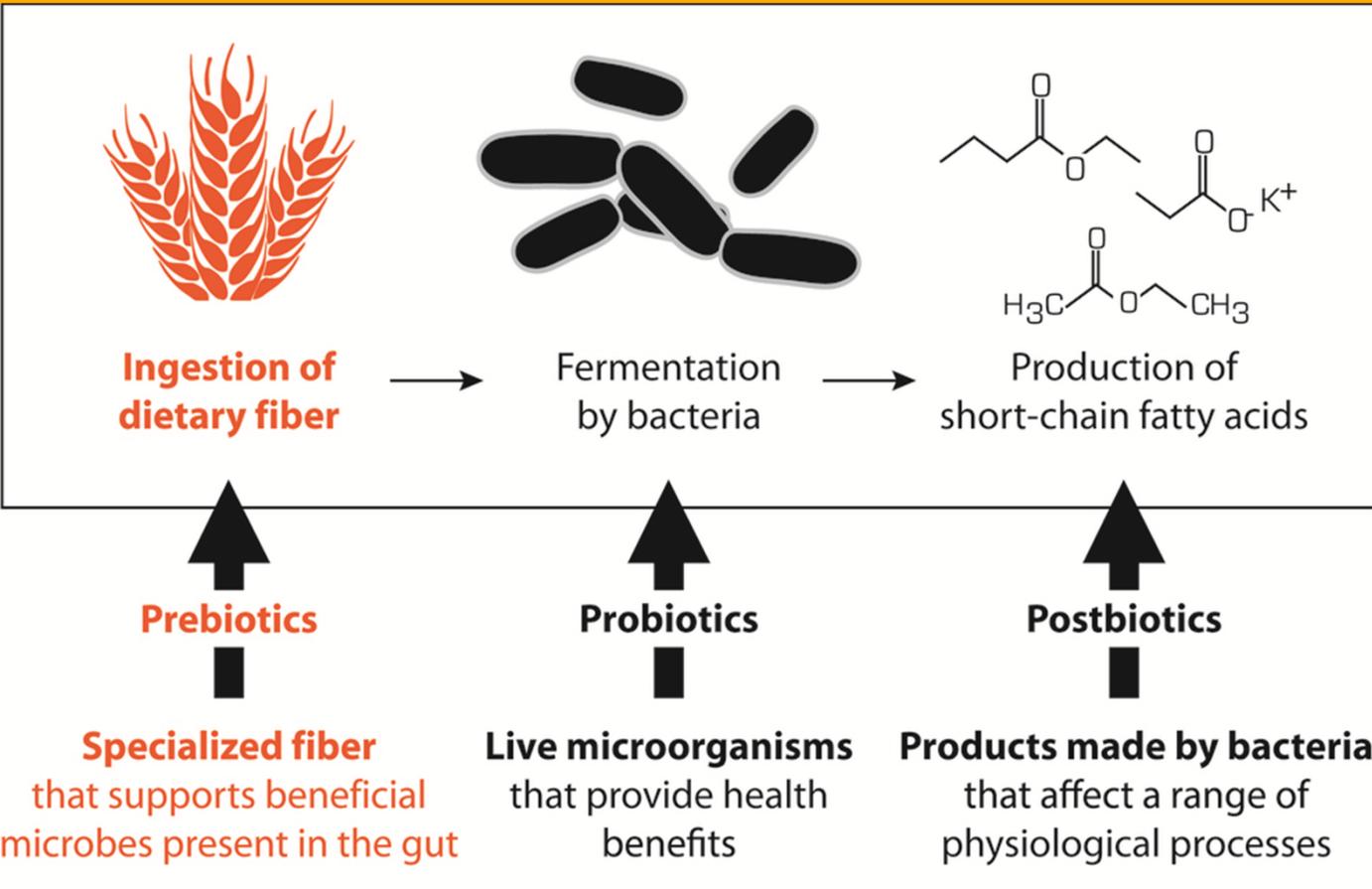
Rina Agustina - 2018



The global probiotics market is projected to reach a turnover value of US\$46.55 billion by 2020

(<http://www.marketsandmarkets.com/PressReleases/probiotics.asp>)

Prebiotics



"nondigestible oligosaccharides, such as fructooligosaccharides, galactooligosaccharides, lactulose, and inulin, which have the potential to stimulate growth of selective and beneficial gut bacteria, particularly lactobacilli and bifidobacteria"

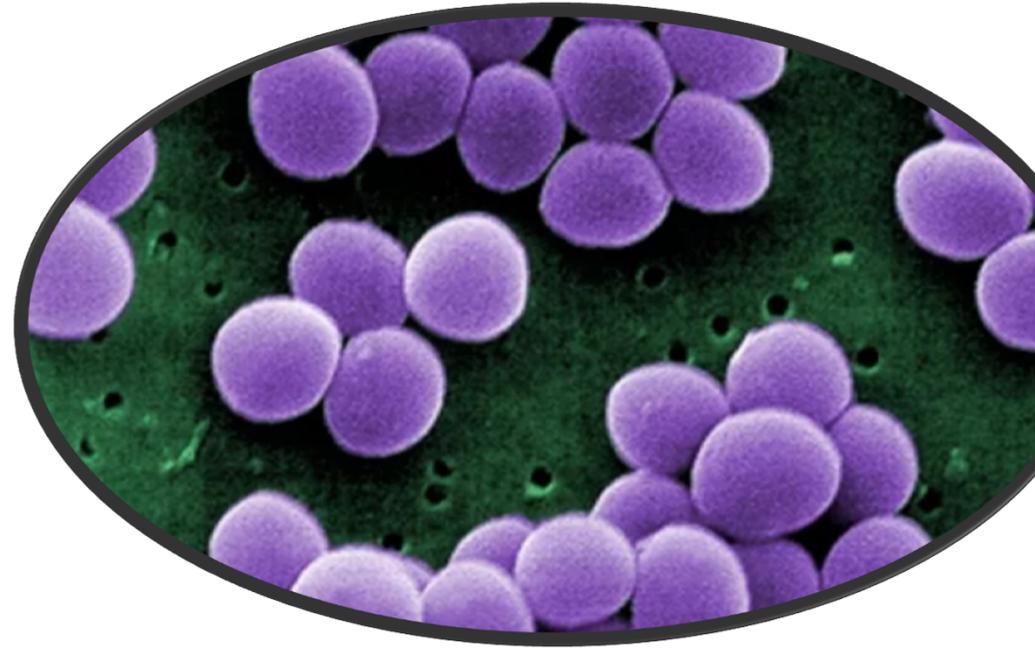
Gibson and Roberfroid, 1995

fructooligosaccharides, galactooligosaccharides, lactulose, and inulin

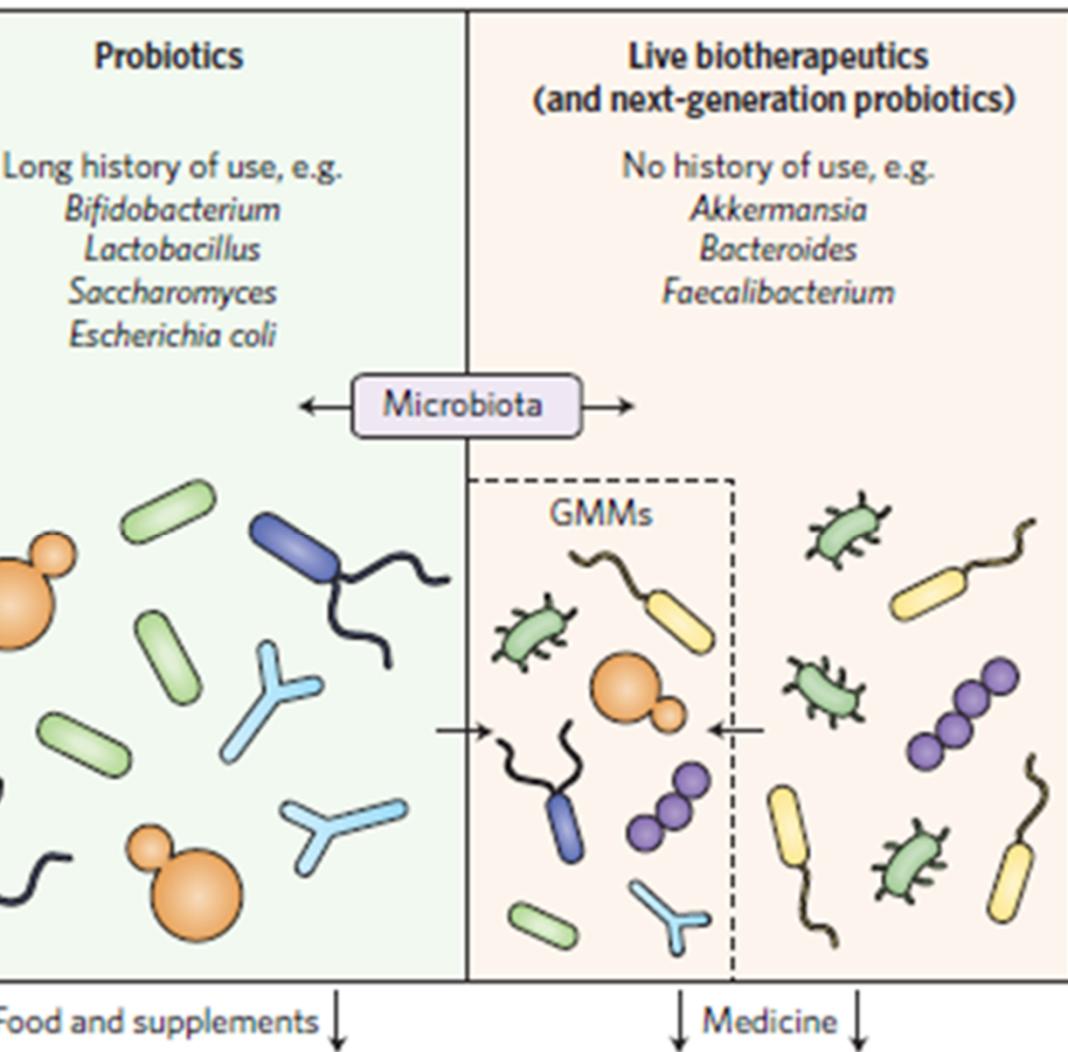
Next-generation Probiotics (NPGs)

Defined as is with the WHO-FAO definition of a probiotic, with an extension of understanding to primarily referring to those microorganisms that have not currently been used as agents to promote health especially in a pharmaceutical product, than a food delivery route

Non-traditional microorganisms

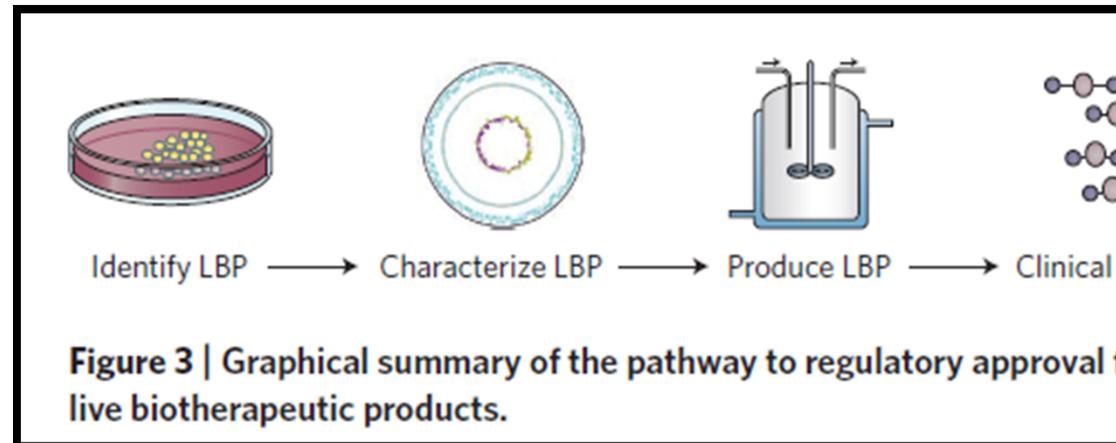


Next - Generation Probiotics (NGPs) versus Live Biotherapeutic Products (LBPs)



The US Food and Drug Administration (FDA) definition of a LBP: “a biological product that:

1. Contains live organisms, such as bacteria
2. is applicable to the prevention, treatment, cure of a disease or condition of human beings
3. is not a vaccine”.



Selected examples of next-generation probiotics

Organism	Type	Disease target	Level of evidence	Study type
<i>Bacteroides xylanisolvens</i> DSM 23694	Natural (human)	Cancer	Medium: safety in humans has been established while levels of TFA-specific IgM have been shown to be elevated in humans	Human
<i>Bacteroides ovatus</i> D-6	Natural (human)	Cancer	Low to medium: increases levels of murine TFA-specific IgM and IgG	Preclinical in mice
<i>Bacteroides ovatus</i> V975	GMO (originally from human gut samples) expressing KGF-2	Intestinal inflammation	Medium: shows abrogation of symptoms of DSS induced in murine colitis model	Preclinical in mice
<i>Bacteroides ovatus</i> V975	GMO expressing TGF- β 1	Intestinal inflammation	Medium: shows abrogation of symptoms of DSS induced in murine colitis model	Preclinical in mice
<i>Bacteroides dorei</i> D8	Natural (human)	Heart disease	Low: depletion of cholesterol <i>in vitro</i>	Preclinical <i>in vitro</i>

(O'Toole, P., Marchesi, J., & Hill, C., 2017).

Selected examples of next-generation probiotics

Organism	Type	Disease target	Level of evidence	Study type
<i>Bacteroides fragilis</i> ZY-312	Natural (human)	Clearance of infectious agents	Low: data only <i>in vitro</i>	Preclinical <i>in vitro</i>
<i>Bacteroides acidifaciens</i> JCM 10556(T)	Natural (mouse)	Clearance of infectious agents	Low to medium: increases IgA levels in the large intestine of gnotobiotic mice	Preclinical in mice
<i>Clostridium butyricum</i> MIYAIRI 588	Natural (human)	Multiple targets including cancer, inflammation and infectious agents	Low to medium: evidence gathered for claims in human and animals trials	Human
<i>Faecalibacterium prausnitzii</i>	Natural (human)	Mainly IBD but also asthma, eczema and type 2 diabetes	Low to medium: mainly focused animal models of colitis and in associative studies	Preclinical in mice and <i>in vitro</i>
<i>Lactococcus lactis</i> ::elafin	GMO (host isolated from food)	Mainly inflammatory diseases such as IBD	Medium: good evidence from animal models of IBD	Preclinical in mice
<i>Lactococcus lactis</i> ::trefoil factor 1 or IL-10	GMO (host isolated from food)	Allergen sensitivity and autoimmune diseases — type 1 diabetes	Medium: mainly animal-based efficacy	Human, phase 1 trial

(O'Toole, P., Marchesi, J., & Hill, C., 2017).

Next Generation Probiotic: *Akkermansia muciniphila* as a Therapeutic candidate?

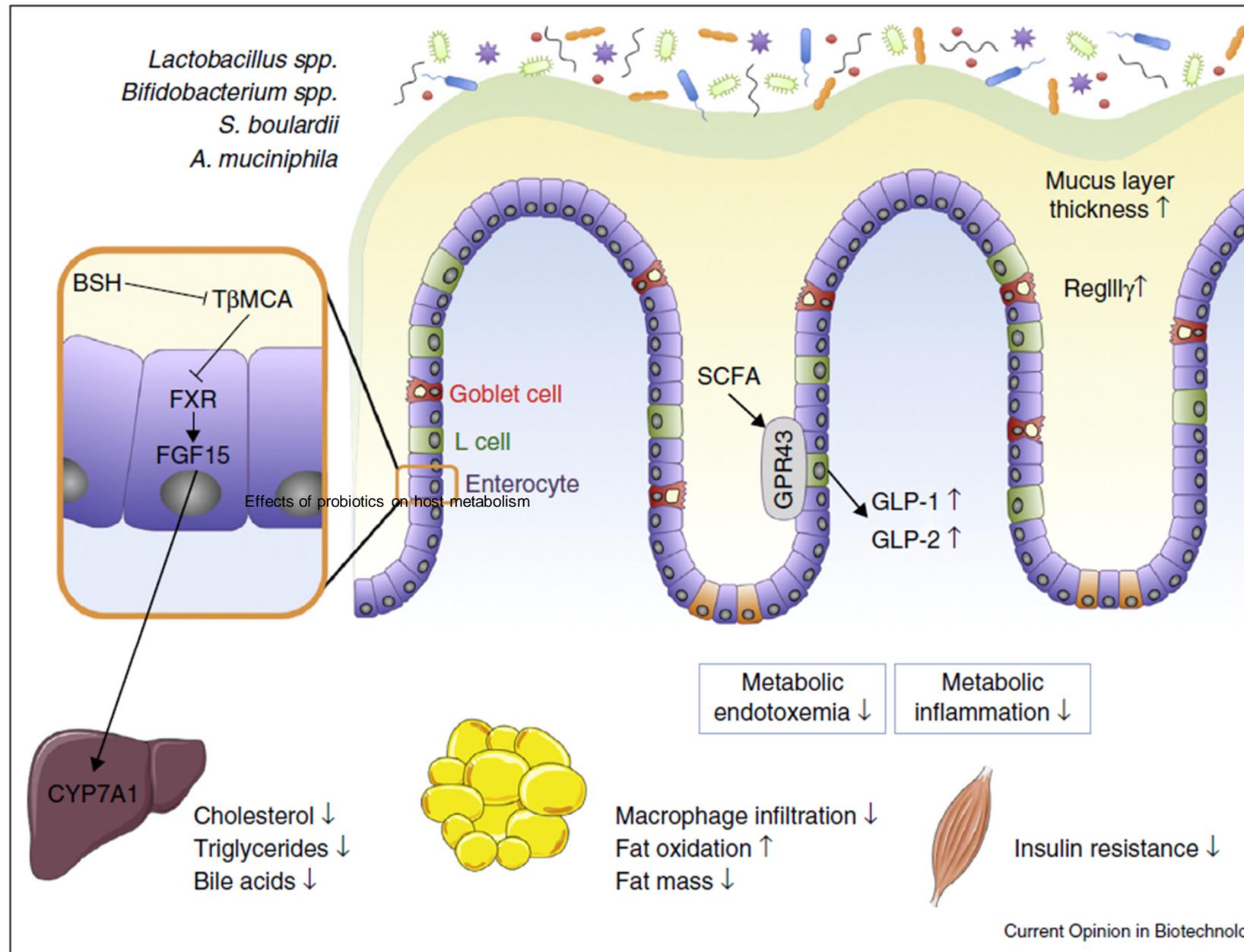
(Sani, P., & Van Hul, M.
2018).

Identify novel opportunities for
next-generation
probiotics targeting
metabolic syndrome.

- Fei and Zhao reported that the *Enterobacter cloacae* B29 strain isolated from the gut microbiota of an obese human induced obesity in germ-free mice
- Conversely, *A. muciniphila*, a mucin-degrading bacterium that resides in the mucus layer, is present at lower levels in obese and type 2 diabetic subjects
- Prebiotic treatment (i.e., oligofructose) increased the abundance of *A. muciniphila* in obese mice → resulted in an improved metabolic profile.
- *A. muciniphila* treatment decreased metabolic endotoxemia and adipose tissue inflammation by improving intestinal mucosal barrier function at different levels.

Effects of Probiotics on host metabolism

Canani, P., & Van Hul, M. (2018).



Next-Generation Beneficial Microbes:

Akkermansia muciniphila on host metabolism

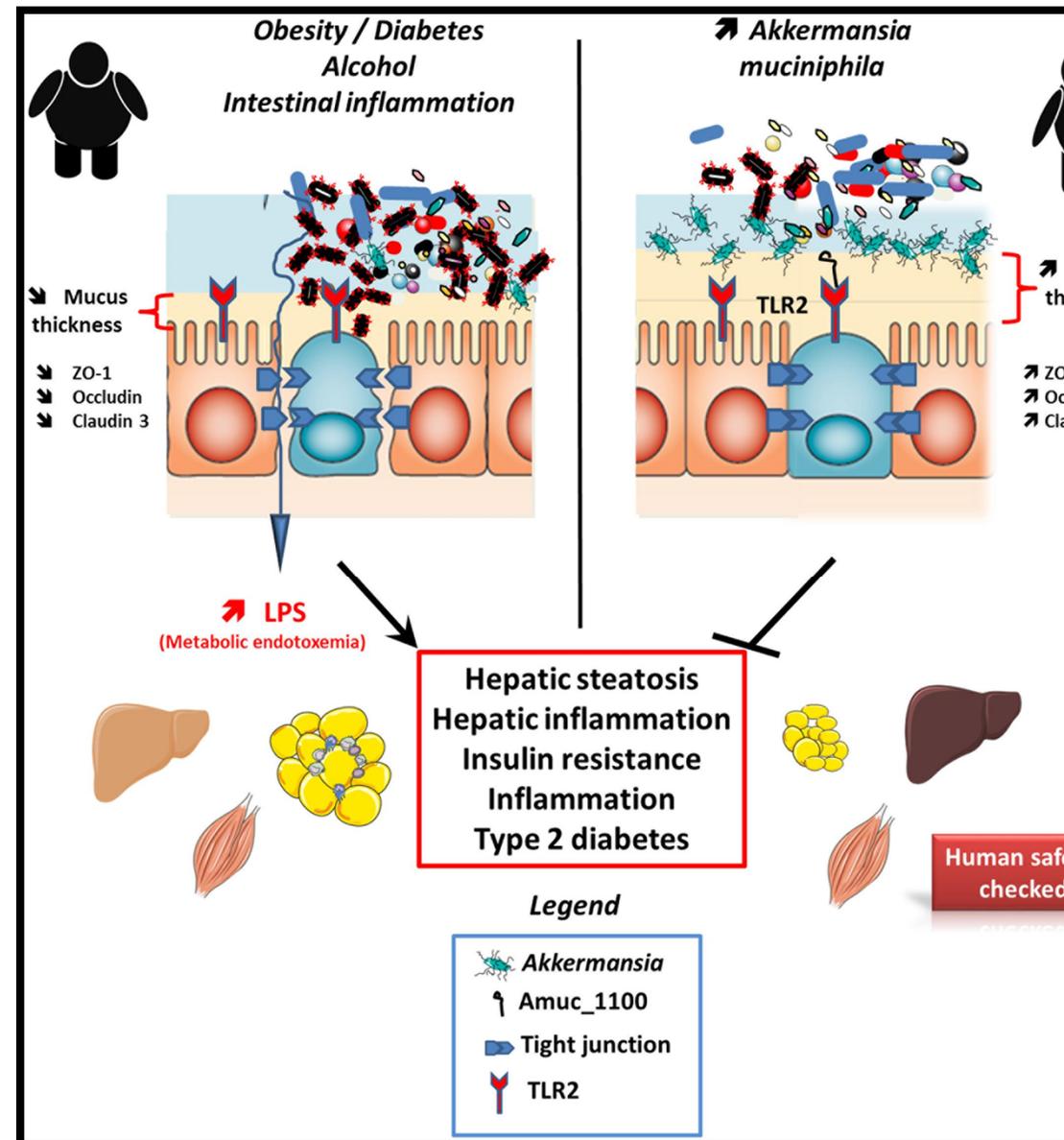
atrice D. Cani and Willem M. de Vos

A. muciniphila has been found to be lower in several conditions such as during obesity, diabetes, intestinal inflammation, liver diseases, or chronic alcohol consumption.

This is associated with an altered gut barrier function leading to an increased plasma LPS levels and eventually triggering low grade inflammation and metabolic disorders.

A. muciniphila alive or pasteurized as well as Amuc_1100 has been shown to restore gut barrier function likely by acting on TLR2 and restoring appropriate tight junction expression.

All these results are associated with an increased mucus layer thickness and an improvement of metabolic disorders. It is worth noting that an exploratory human investigation has shown that *A. muciniphila* is apparently safe.



Next Generation Probiotic: Novel strain?

Faecalibacterium prausnitzii

- *F. prausnitzii* may possess in vivo and in vitro anti-inflammatory effects.
- Able to reduce Th1 and Th17 proinflammatory cytokines in Mesenteric Lymphatic Node (MLN) and colon tissues both DNBS and DSS colitis model.
- Can induce the Clostridium-specific IL-10-secreting regulatory T cell subset, present in several human colon cells.
- Its capacity for lowering IL-12 and IFN γ production indicates that the interaction between *F. prausnitzii* and the host shape and maintain the gut barrier immune function (Quévrain et al., 2016).

Breyner et al. (2017). Microbial anti-inflammatory molecule (MAM) from *Faecalibacterium prausnitzii* shows a protective effect on DNBS and DSS-induced colitis model in mice through inhibition of NF- κ B pathway.

Next Generation Probiotic: Novel train?

Bacteroides fragilis and *Bacteroides uniformis*

- *B. fragilis* produces polysaccharide A (PSA), an immunomodulatory molecule that activates the T-cell dependent immune responses (Troy and Kasper, 2010). It is involved in the development and homeostasis of the host immune system (*Troy and Kasper, 2010*).
- Oral administration of *Bacteroides uniformis* (*B. uniformis*) CECT 7771 in high fat diet-fed mice improved lipid profile, reduced glucose insulin and leptin levels, increased TNF- α production by dendritic cells (DCs) in response to LPS stimulation, and increased phagocytosis (*Gauffin Cano et al., 2012*).
- It can ameliorate metabolic disorder and immunological dysfunction related to intestinal dysbiosis in obese mice (*Gauffin Cano et al., 2012; Yang et al., 2016*).

Probiotics on central nervous system (CNS) disorders: *children with autism, anxiety and depression*

an association between
 shifts in the gut microbiota
 composition in children
 with neurodevelopmental
 disorders (e.g. autism
 spectrum disorder, anxiety
 and depressive behaviors)

temporal profile of
 neurodevelopmental
 sequences in relation to the
 age of onset of mental
 disorders and degree of
 microbiota
 stability/diversity
 throughout life

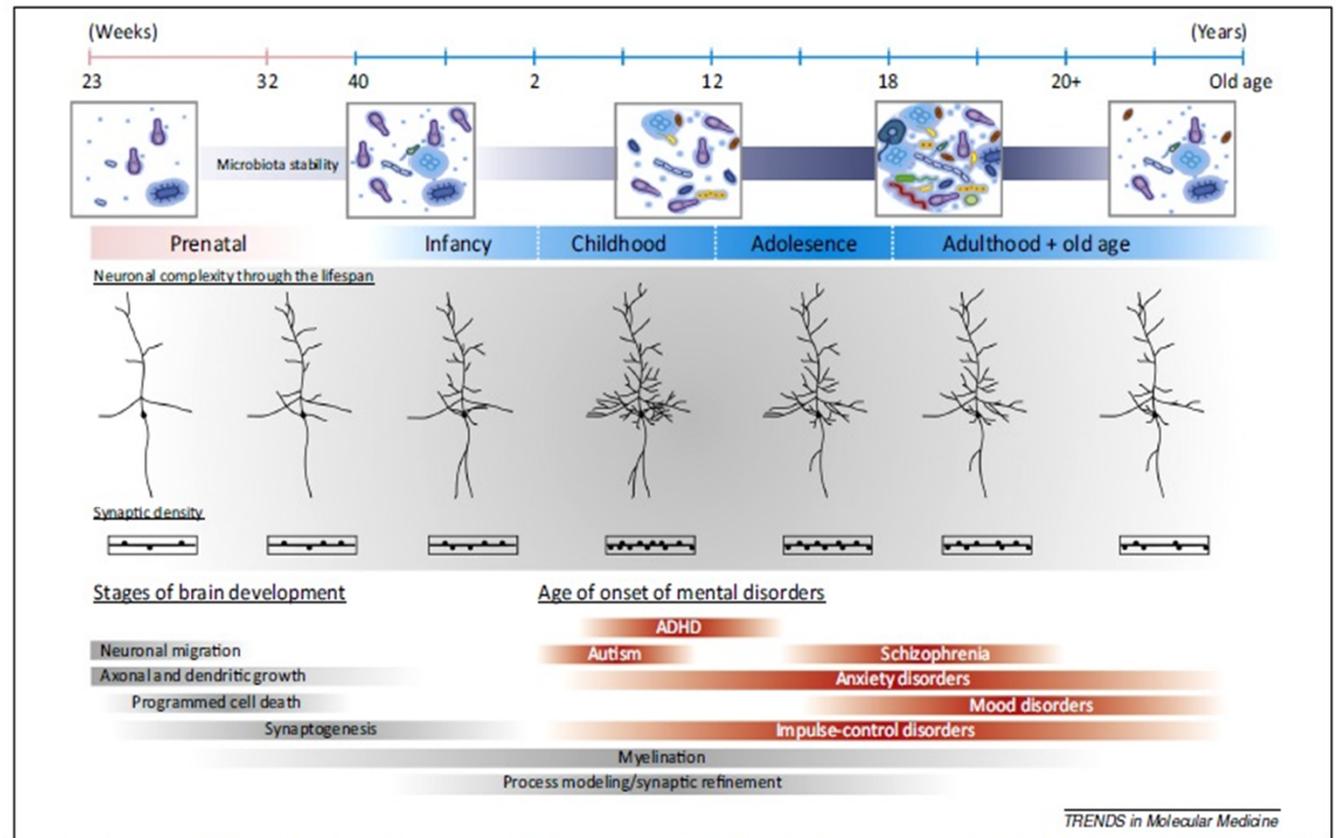


Figure 2. Temporal profile of neurodevelopmental sequences in relation to the age of onset of mental disorders and degree of microbiota stability/diversity throughout life. Gut microbiota is essential to human health and is a key player in the bidirectional communication between the gastrointestinal tract and the central nervous system. The microbiota dynamically changes throughout lifespan, establishing its symbiotic rapport with the host with critical windows during infancy, adolescence, and aging. During these windows, the organism is vulnerable to external stressors, which may result in mental disorders. Early life perturbations of the developing gut microbiota can impact neurodevelopment and potentially lead to adverse mental health outcomes later in life. The concept of parallel and interacting microbial–neural critical windows opens new avenues for developing novel microbiota-modulating based therapeutic interventions in early life to combat neurodevelopmental deficits and brain disorders. Abbreviation: ADHD, attention deficit hyperactivity disorder.

**The Lancet 2018: sequencing
 genes – 18 genes candidates**

Technical Challenges

- **Stability of the probiotic during the probiotic production** → The product manufacturing and storage processes may impact the viability of the bacterial strains, influencing probiotic stability and properties.
- **Viability of the probiotics after consumption** → Bacterial strains should remain viable at sufficient numbers through the gastrointestinal tract (GIT) passage → the selection of optimal culture medium and cell protectants is crucial

Research challenges and opportunities

- Strategy of how scientists may meet this challenge in designing the NPGs with new strains from the lessons learned working with traditional probiotics, and prebiotics.

Importantly, series of clinical trials (Phase 1 for safety and dosage ranges; Phase 2 will for the primary endpoint in a small groups; preferably the Phase 3 for evaluating the efficacy, side effects and relative benefits in a larger group) need to be determined in general use or for vulnerable populations.

Regulatory Challenges

The definition and classification of probiotics by regulatory agents throughout the world is different → the status of probiotic products is still uncertain. How the regulation of new product of NGPs will align with LBPs

Thus, reservations about probiotic products claims may arise among regulatory bodies, producers, and consumers.



THANK YOU