

PREVENTING CHRONIC DISEASES VIA GUT MICROBIOTA:

THE USE OF PREBIOTIC FIBERS, FROM SCIENTIFIC DEMONSTRATION TO MARKETING COMMUNICATION

CAROLINE PERREAU & CLÉMENTINE THABUIS

ROQUETTE NUTRITION AND HEALTH R&D

27TH OF SEPTEMBER 2022

ADEBIOTECH

ROQUETTE: A LEADER WITH A PURPOSE

Ambition

Unlocking the potential of nature to offer the best ingredients for food, nutrition and health markets.

8,000+
Employees

25

Industrial sites

90
Years of industrial and operational excellence

100+
Countries served by a global commercial network

€3.9bn

Turnover

300

R&D workforce







AGENDA

- 1. Chronic diseases and gut microbiota
- 2. NUTRIOSE®, a good candidate to increase insulin sensitivity and thus to prevent diabetes
- 3. Gluconic acid to boost immune response and decrease antibiotics use in farms
- 4. How to put a new ingredient on the market and communicate on its health benefits?
- Novel food process
- Communication on the health benefits



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THE GUT MICROBIOTA: A MARKER OF HOST HEALTH

frontiers in CELLULAR AND INFECTION MICROBIOLOGY



The function of our microbiota: who is out there and what do they do?

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SCIENCE AND POLITICS OF NUTRITION

Role of the gut microbiota in nutrition and health

Ana M Valdes and colleagues discuss strategies for modulating the gut microbiota through diet and probiotics

icrobiome refers to the collective genomes of the microorganisms in a particular environment, and microbiota is the community of microorganisms themselves (box 1). Approximately 100 trillion micro-organisms (most of them bacteria, but also viruses, fungi, and protozoa) exist in the human gastrointestinal tract1 ²—the microbiome is now best thought of as a virtual organ of the body. The human genome consists of about 23 000 genes, whereas the microbiome encodes over three million genes producing thousands of metabolites, which replace many of the functions of the host,13 consequently influencing the host's fitness, phenotype, and health.

Box 1: Glossary

- Microbiome—the collective genomes of the micro-organisms in a particular environment
- Microbiota—the community of micro-organisms themselves
- Microbiota diversity—a measure of how many different species and, dependent on
 the diversity indices, how evenly distributed they are in the community. Lower
 diversity is considered a marker of dysbiosis (microbial imbalance) in the gut and
 has been found in autoimmune diseases and obesity and cardiometabolic conditions,
 as well as in elderly people
- Operational taxonomic unit—a definition used to classify groups of closely related organisms. DNA sequences can be clustered according to their similarity to one another, and operational taxonomic units are defined based on the similarity threshold (usually 97% similarity) set by the researcher
- Colonocytes-epithelial cells of the colon
- Germ-free animals—animals that have no micro-organisms living in or on them
- Short chain fatty acids—fatty acids with two to six carbon atoms that are produced by bacterial fermentation of dietary fibres

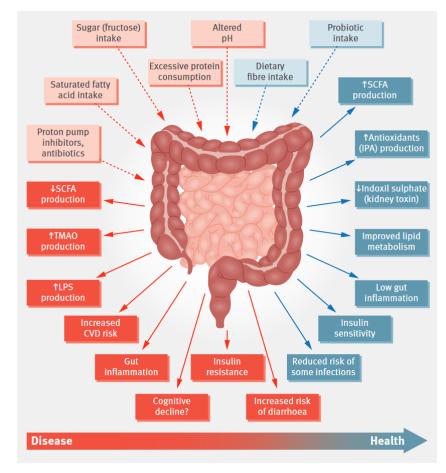


Fig 1 | Schematic representation of the role of the gut microbiota in health and disease giving some examples of inputs and outputs. CVD=cardiovascular disease; IPA=indolepropionic acid; LPS=lipopolysaccharide; SCFA=short chain fatty acids; TMAO=trimethylamine N-oxide



THE GUT MICROBIOTA & BLOOD GLUCOSE MANAGEMENT



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Impact of Gut Microbiota on Host **Glycemic Control**

Céline Gérard and Hubert Vidal*

Univ Lyon, CarMeN Laboratory, INSERM U1060, INRA U1397, INSA Lyon, Université Claude Bernard Lyon1, Oullins, France

Diabetologia (2018) 61:810-820 https://doi.org/10.1007/s00125-018-4550-1

ARTICLE



Aberrant intestinal microbiota in individuals with prediabetes

Kristine H. Allin^{1,2} · Valentina Tremaroli^{3,4} · Robert Caesar^{3,4} · Benjamin A. H. Jensen⁵ · Mads T. F. Damgaard⁵ · Martin I. Bahl⁶ · Tine R. Licht⁶ · Tue H. Hansen¹ · Trine Nielsen¹ · Thomas M. Dantoft⁷ · Allan Linneberg^{7,8} · Torben Jørgensen 7,9,10 • Henrik Vestergaard 1,11 • Karsten Kristiansen • Paul W. Franks 12,13,14 • the IMI-DIRECT consortium · Torben Hansen 1,15 · Fredrik Bäckhed 3,4,16 · Oluf Pedersen 1,17

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ARTICLE

doi:10.1038/nature11450

A metagenome-wide association study of gut microbiota in type 2 diabetes

Junjie Qin¹*, Yingrui Li¹*, Zhiming Cai²*, Shenghui Li¹*, Jianfeng Zhu¹*, Fan Zhang²*, Suisha Liang¹, Wenwei Zhang¹, Yuanlin Guan¹, Dongqian Shen¹, Yangqinp Feng¹, Dongya Zhang¹, Zhuye Jie¹, Wenxian Wu¹, Youwen Qin¹, Wenbin Xue¹, Junhua Li¹, Lingehuan Han², Donghui Lu¹, Peixian Wu¹, Yali Dai², Xiaojuan Sun², Zesong Li², Aifa Tang², Shilong Zhong², Junnua Li', Lingenuan Han', Donghui Lu', Petikan Wu, Tan Dai, Alaquan Sun, Assonia Li, Ana Tang, Sindong Calong, Kiaoping Li', Weineng Chen', Ran Xu', Mingbang Wang', Qiang Feng', Meihua Gong', Jing Yu', Yanyan Zhang', Ming Zhang', Torben Hansen', Gaston Sanchez', Jeroen Raes^{2,8}, Gwen Falony', Shujiro Okuda', Mathieu Almeida', Emmanuelle LeChatelier', Pierre Renault', Nicolas Pons', Jean-Michel Batto', Zhaoxi Zhang', Hua Chen', Ruifu Yang^{1,10}, Pierre Renault', Nicolas Pons', Jean-Michel Batto', Zhaoxi Zhang', Hua Chen', Ruifu Yang^{1,10}, Pierre Renault', Nicolas Pons', Jean-Michel Batto', Zhaoxi Zhang', Hua Chen', Ruifu Yang^{1,10}, Pierre Renault', Nicolas Pons', Jean-Michel Batto', Zhaoxi Zhang', Hua Chen', Ruifu Yang^{1,10}, Pierre Renault', Nicolas Pons', Jean-Michel Batto', Zhaoxi Zhang', Hua Chen', Ruifu Yang^{1,10}, Pierre Renault', Nicolas Pons', Jean-Michel Batto', Zhaoxi Zhang', Hua Chen', Ruifu Yang^{1,10}, Pierre Renault', Nicolas Pons', Jean-Michel Batto', Zhaoxi Zhang', Hua Chen', Ruifu Yang^{1,10}, Pierre Renault', Nicolas Pons', Jean-Michel Batto', Zhaoxi Zhang', Hua Chen', Ruifu Yang^{1,10}, Pierre Renault', Nicolas Pons', Jean-Michel Batto', Zhaoxi Zhang', Hua Chen', Ruifu Yang^{1,10}, Pierre Ruifu Yang', Pierre Weimou Zheng¹, Songgang Li¹, Huanming Yang¹, Jian Wang¹, S. Dusko Ehrlich⁹, Rasmus Nielsen⁶, Oluf Pedersen⁵ Karsten Kristiansen^{1,13} & Jun Wang^{1,5,13}

Prebiotic/probiotic intervention

Gut microbiota



Incretin secretion

- · GLP-1 secretion
- Number of enteroendocrine cells
- · Enteroendocrine cell differentiation



SCFA production

- · Intestinal gluconeogenesis
- · Gut wall integrity
- GLP-1 secretion
- ß cell function and insulin secretion



Bile acid metabolism

- Chemical diversity of bile acid pool
- FXR and TGR5 metabolic signaling
- FGF19 secretion



Adipose tissue regulation

- LPS-induced inflammation
- · White adipose tissue browning

Improved host glycemic control





THE GUT MICROBIOTA & LOW-GRADE INFLAMMATION





Down Regulation of NF-kB

Revie

Probiotics and Prebiotics for the Amelioration of Type 1 Diabetes: Present and Future Perspectives

Sidharth Mishra ^{1,3}, Shaohua Wang ¹, Ravinder Nagpal ¹, Brandi Miller ¹, Ria Singh ¹, Subhash Taraphder ³ and Hariom Yadav ^{1,2},*

- Department of Internal Medicine-Molecular Medicine, Centre for Diabetes, Obesity and Metabolism, Wake Forest School of Medicine, Winston-Salem, NC 27101, USA; spmishra@wakehealth.edu (S.M.); shaowang@wakehealth.edu (S.W.); rnagpal@wakehealth.edu (R.N.); bcmiller@wakehealth.edu (B.M.); ria.singh@gmail.com (R.S.)
- Department of Microbiology and Immunology, Centre for Diabetes, Obesity and Metabolism, Wake Forest School of Medicine, Winston-Salem, NC 27101, USA
- Department of Animal Genetics and Breeding, West Bengal University of Animal and Fishery Science, Kolkata 700 037, India; subhash.taraphder@gmail.com
- * Correspondence: hyadav@wakehealth.edu; Tel.: +1-336-713-5049

Inflammation

Maintaining Luminal pH Intestinal Lumen Enhanced Intestinal Barrier Function B Cell Humoral Immune Response Plasm Cell Intestinal Epithelium Mucus Damaged Tissue Stimulation of Mucus production repair Acetate Maintaining the gut microbiota Carbohydrate Butyrate Activated Akt Anti-inflammatory Cytokines Gut Microbiota Expression of Low grade intestinal

Increasing Intestinal Integrity

Gut microbiota

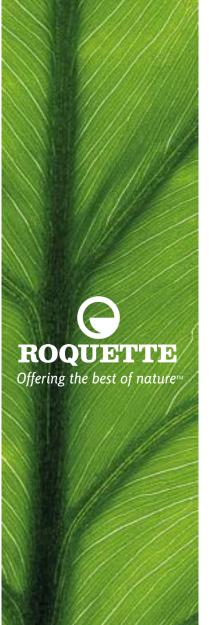


Original research

Gut microbiota composition reflects disease severity and dysfunctional immune responses in patients with COVID-19

Yun Kit Yeoh , 1,2 Tao Zuo , 2,3,4 Grace Chung-Yan Lui, 5,5 Fen Zhang, 2,3,4 Qin Liu, 2,3,4 Amy YL Li, Arthur CK Chung, Chung, Chun Pan Cheung, 2,3,4 Eugene YK Tso, Kitty SC Fung, Veronica Chan, Lowell Ling, Gavin Joynt, David Shu-Cheong Hui, 5,5 Kai Ming Chow , Susanna So Shan Ng, 5,5 Timothy Chun-Man Li, 5,5 Rita WY Ng, Terry CF Yip, 4,6 Grace Lai-Hung Wong , 3,4 Francis KL Chan , 2,3,4 Chun Kwok Wong, Paul KS Chan, 1,2,10 Siew C Ng , 2,3,4

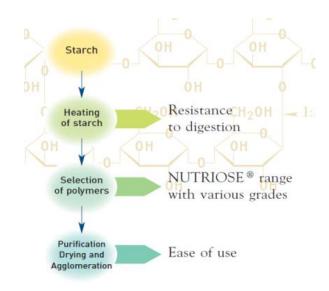




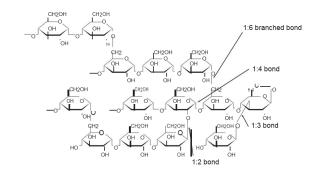
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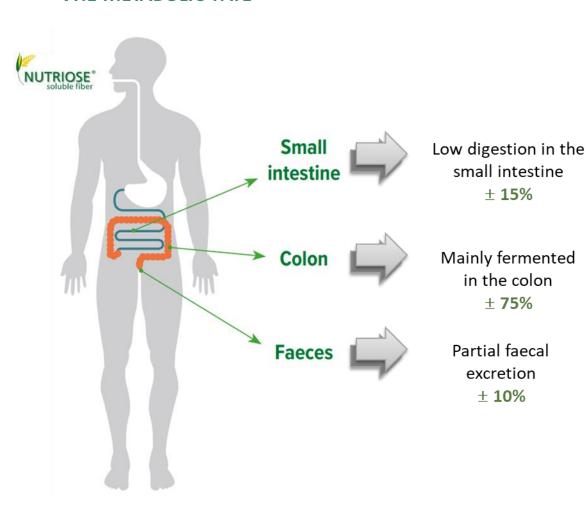
A SPECIFIC PROCESS FOR A SPECIFIC FIBER



OSIDIC LINKAGES NOT HYDROLYSED BY DIGESTIVE ENZYMES



THE METABOLIC FATE



4 HEALTH BENEFITS SUPPORTED BY CLINICAL STUDIES



BLOOD GLUCOSE MANAGEMENT

GUT HEALTH SUSTAINED ENERGY RELEASE

SATIETY







Glycaemic response Blood glucose control Tolerance Prebiotic - microbiota Intestinal balance

Colonic fermentations Physiologic/metabolic energy

Satiety Weight management



Investigate the effects of 14g/day NUTRIOSE® for 28 days in humans on satiety, metabolic parameters and microbiota

THE STUDY DESIGN

- Chronic, randomized, cross over study
- Setting up in the UK (University of Reading)

THE POPULATION

Volunteers

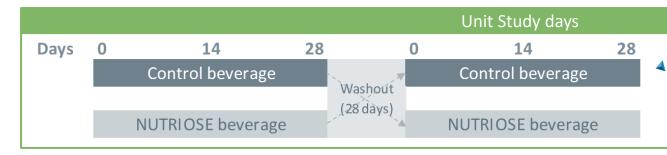


20 normal weight volunteers (21.0-24.9 kg/m²)

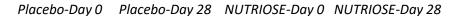


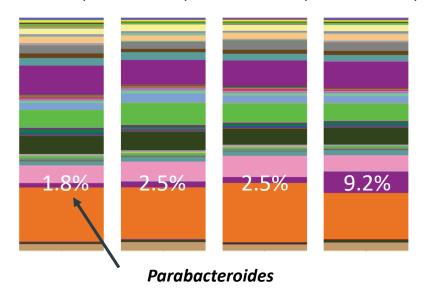
16 overweight volunteers $(24.9-30.0 \text{ kg/m}^2)$

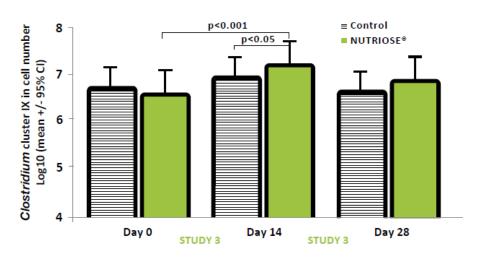
- Aged 22-55 y
- Usual daily dietary fibre intake: 16g/day



Outcomes: Appetite evaluation, Fecal microbiota, blood glucose measurements after an orange juice intake







FISH technics: Clostridium cluster IX (+)

= a Bacteroides subgroup of the phylum Bacteroidetes

Relative abundance using faecal 16SrRNA sequencing

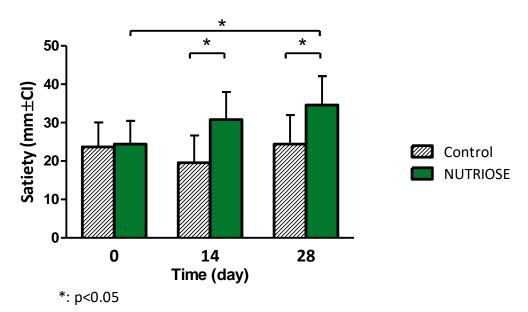
- Significant increase in the relative abundance of *Parabacteroides* genus, a saccharolytic genus
- Significant increase in *Clostridium cluster 9 group*, a propionate producing bacterial group



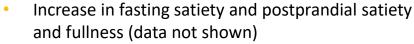
Nutriose® has a lowering effect on long term post prandial glucose and on caloric intake

AND WEIGHT MANAGEMENT IN HUMANS

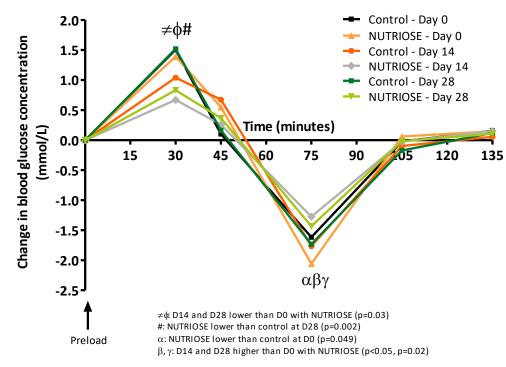
Fasting satiety values (before breakfast)



Fasting satiety ratings in the latter postprandial phase



No measured impact in caloric intake



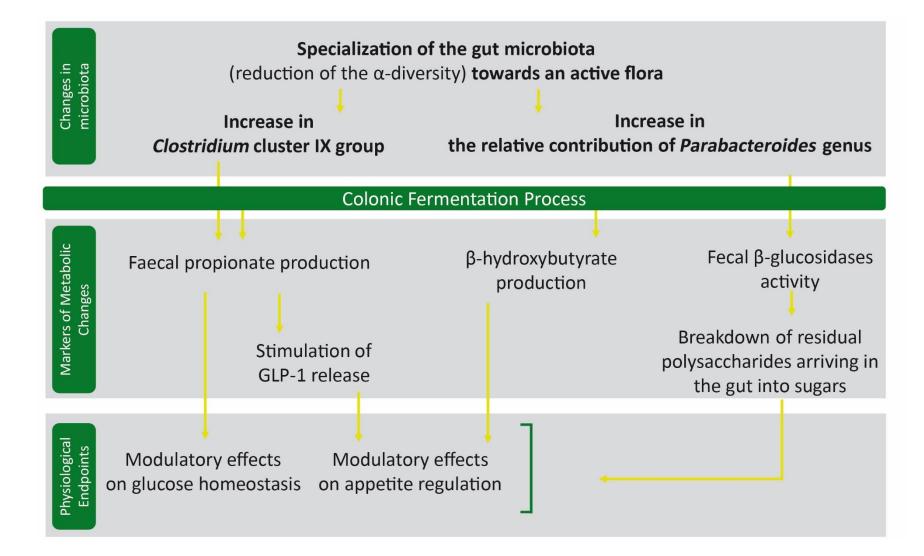
Changes in blood glucose concentrations following intake of a carbohydrate-preload

- Reduction of the overall postprandial glucose rise after a carbohydrate-preload administration
- Long term biological effects beyond a simple reduction in glycaemic carbohydrates





NUTRIOSE® INDUCES A POSITIVE IMPACT ON THE OVERALL GUT ECOSYSTEM IN HUMANS AND EXERTS POTENTIAL METABOLIC HEALTH BENEFITS





GUT HEALTH: GLUCONEOGENESIS, THE PUTATIVE MECHANISM OF ACTION



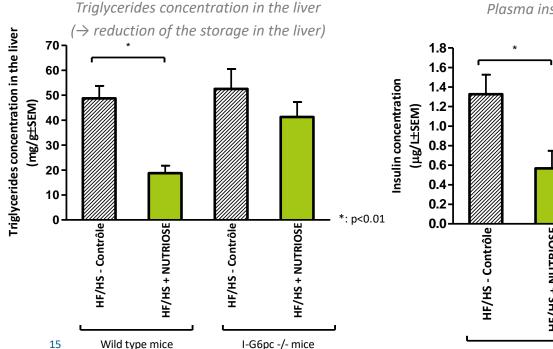
THE HEALTH BENEFITS OF NUTRIOSE® MAY BE EXPLAINED BY A CENTRAL ROLE OF INTESTINAL GLUCONEOGENESIS / THE GUT-BRAIN AXIS

METHODOLOGY:

- Wild type mice and I-G6pc -/- mice (no intestinal glucose production = no gluconeogenesis) France
- 3-week study period
- High fructose/High sucrose (HF/HS) diet supplemented with 10% NUTRIOSE®
- Blood parameters, Tissue weight, Fecal microbiota, SCFA

- Modulation of some parameters (triglycerides, insulin and other) and in gene expression (data not shown) in wild type mice only suggesting the role of intestinal gluconeogenesis
- Fecal microbiota modulation
- Increase in caecal butyrate (data not shown)

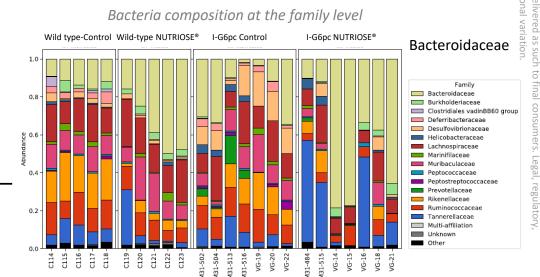
Intestinal gluconeogenesis may explain the clinical outcomes like satiety and blood glucose management

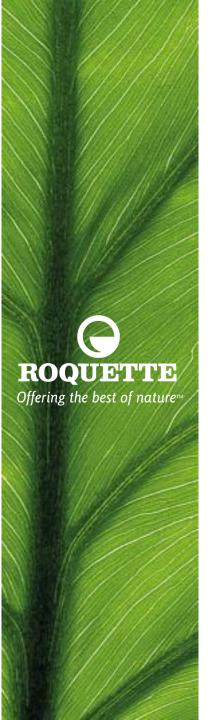


Plasma insulin concentration

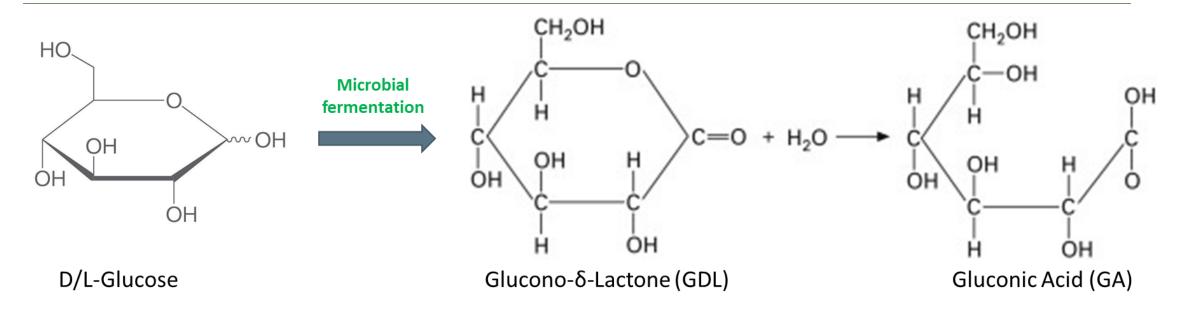
Wild type mice

I-G6pc -/- mice





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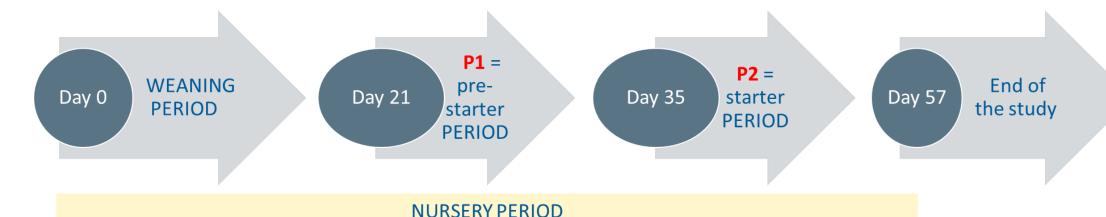
- Nutritional value can be added to glucose syrup using <u>microbial fermentation</u>
- GDL is used as a food stabilizer
- According to literature, GA is <u>poorly absorbed in the upper part of the GI</u> and can be converted <u>into butyrate by gut microbiota</u>
- Gluconic acid <u>prebiotic properties</u> were investigated in the present study







<u>Objective</u>: to evaluate the effect of a dietary Sodium Gluconate or Gluconic Acid supplementation on weaning performances and intestinal health in piglets during the nursery period.



Significant impact on the main parameters of **zootechnical performance**:

- feed intake at the end of the P1 & P2
- apparent digestibility at the highest dose

Positive impact on **immune status**:

- significant decrease in blood lymphocytes indicating **lower infection levels**
- lower antibiotic use

IMPACT on **gut microbiota**:

- impact on **specific bacteria** (*Lactobacillus, F. prausnitzii*...) → increase in **butyrate producers**
- higher butyric acid for both doses





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HOW TO PUT A NEW "PRODUCT" ON THE MARKET?

For European regulation -> within each company, an evaluation of a novel food dossier of an ingredient is usually done by a committee gathering marketing, scientific and regulatory experts



Evaluation of human consumption to significant degree within the Union of the ingredient or assessment of equivalence to another ingredient identified before 15 May 1997 and taking into account these specific points (not exhaustive):

- food with a new or intentionally modified molecular structure,
- isolated from or produced from a plant or a variety of the same species
- production process already used or not

NOVEL FOOD

Any food that was not used for human consumption to a significant degree within the Union before 15 May 1997, and that falls under at least one of the 10 categories.

https://food.ec.europa.eu/system/files_en?file=2016-10/novel-food_guidance_human-consumption_en.pdf

Novel food dossier



HOW TO COMMUNICATE ON SCIENCE IN EUROPE?

B2B communication

RECOMMANDATIONS

- Ensure the communication does not relate to a product intended for the final consumer
- Remain **proportionate** and do not be misleading
- Remain factual (e.g. clinical studies data, scientific literature...)
- As we speak to professional, the vocabulary must be scientific and technical
- Use of disclaimers* when the health benefits of the ingredients are not yet approved/evaluated by regulatory organizations

Scientific communication

SCIENTIFIC PEER-REVIEWED PAPER **POSTER** SCIENTIFIC PRESENTATION TECHNICAL MEETING WITH CUSTOMERS **WEBINARS**

Health claims

APPROVED HEAITH CLAIMS or GENERIC HFALH CLAIMS

Must refer to the exact approved health claim/wording and the conditions of use.

B2C communication

Roquette is NOT able to provide its customers with risk assessments on their final products. We can only support them.



RECIPE OF THE FINAL PRODUCT?

(Nutriose® = Just **ONE INGREDIENT AMONG OTHERS**)



LEVEL OF RISK THAT THE B2C **COMPANY IS** WILLING TO TAKE?





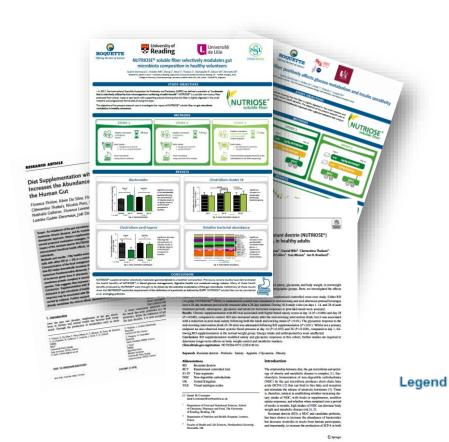
26/09/2022

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NUTRIOSE® – HEALTH BENEFITS AND CLAIMS



Scientific communication

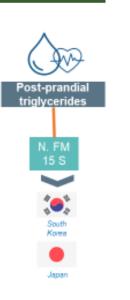


Health claims













Proven health benefits: Roquette proprietary clinical evidence

Accessible health-related claims by country

The examples given are suggestions, associated with more or less regulatory risk. A decision about whether a wording is acceptable or not may need to be made on a case-by-case basis. Front of pack claims are the responsibility of the responsible for the product marketing / placing on the market.

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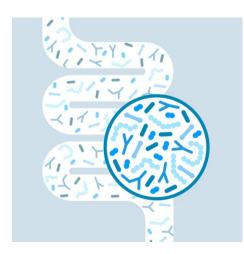
CONCLUSION



The gut microbiota is a marker of host health.

Microbiota dysbiosis <-> Diseases

- Thanks to its short- and long-term impact on blood glucose management,
 NUTRIOSE® can be a good candidate to increase insulin sensitivity and thus to prevent diabetes.
 - Microbiota modulation and gluconeogenesis -> putative mechanism of action
- Gluconic acid can boost immune response and decrease antibiotics use in farms
- Before putting a new product on the market -> evaluation of novel food dossier
- B2B communication allow **factual communication** between two professionals, before evaluation of B2C communication with approved health claims in Europe.





Offering the best of nature™