

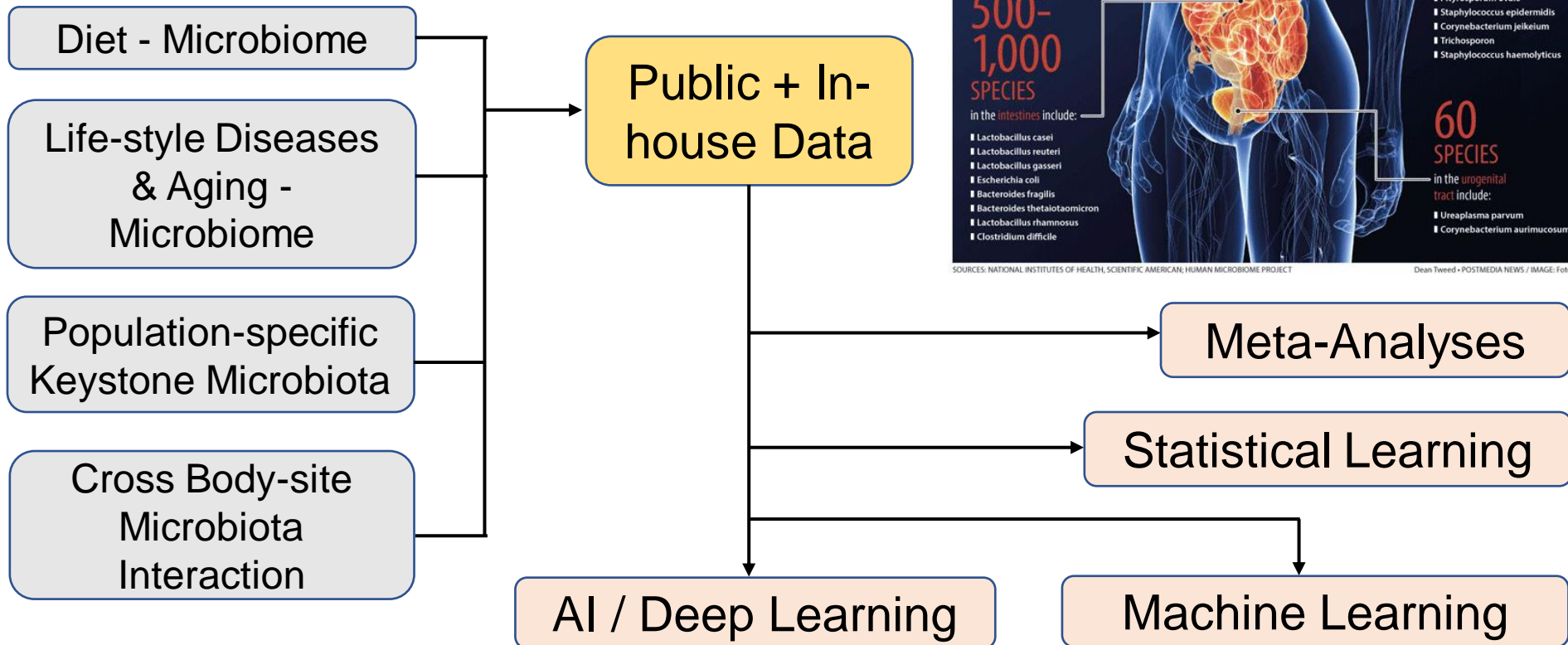
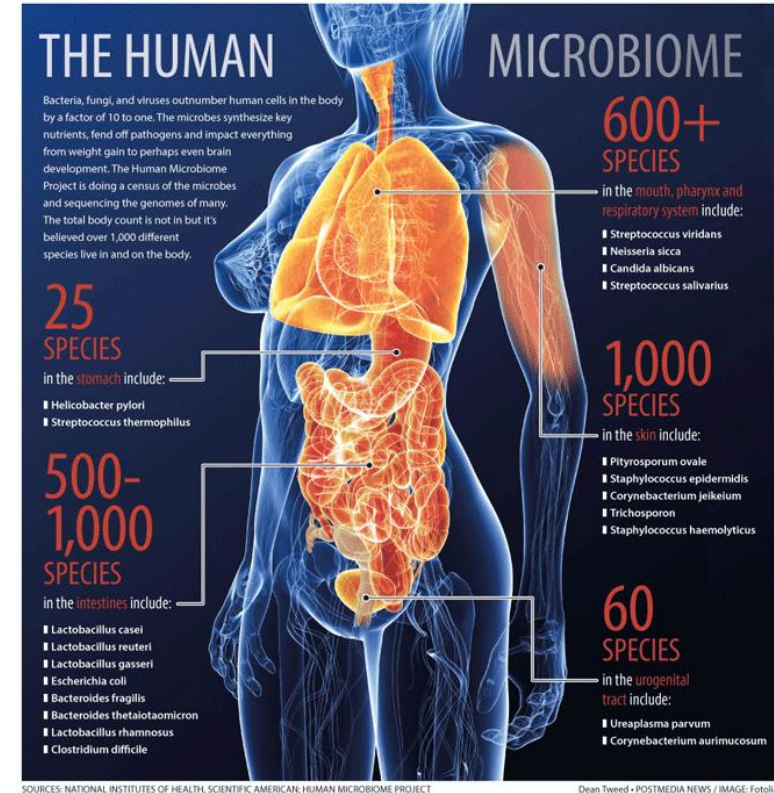
# Microbiome Revisited

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# About Myself ..

- Computational Microbiologist
- Human Microbiome
- Data-driven investigation



# The human gut microbiome by numbers

$10^{13} - 10^{14}$   
microbial cells

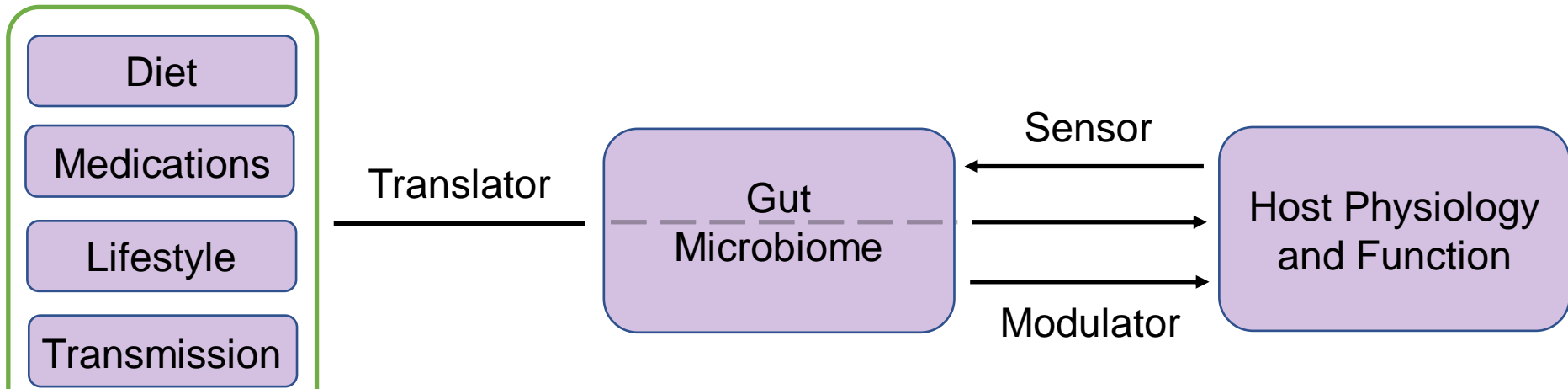


90% of the  
diseases  
associated with  
changes in the gut  
microbiome


Functional  
diversity of the  
microbiome to the  
human genome:  
150 : 1





Microbiome can  
be therapeutically  
modifiable

# The three facets of the gut microbiome



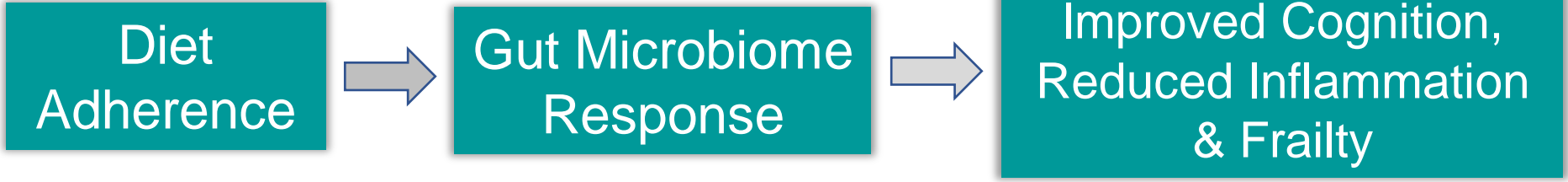
Nutrition  
Original research

Mediterranean diet intervention alters the gut microbiome in older people reducing frailty and improving health status: the NU-AGE 1-year dietary intervention across five European countries 

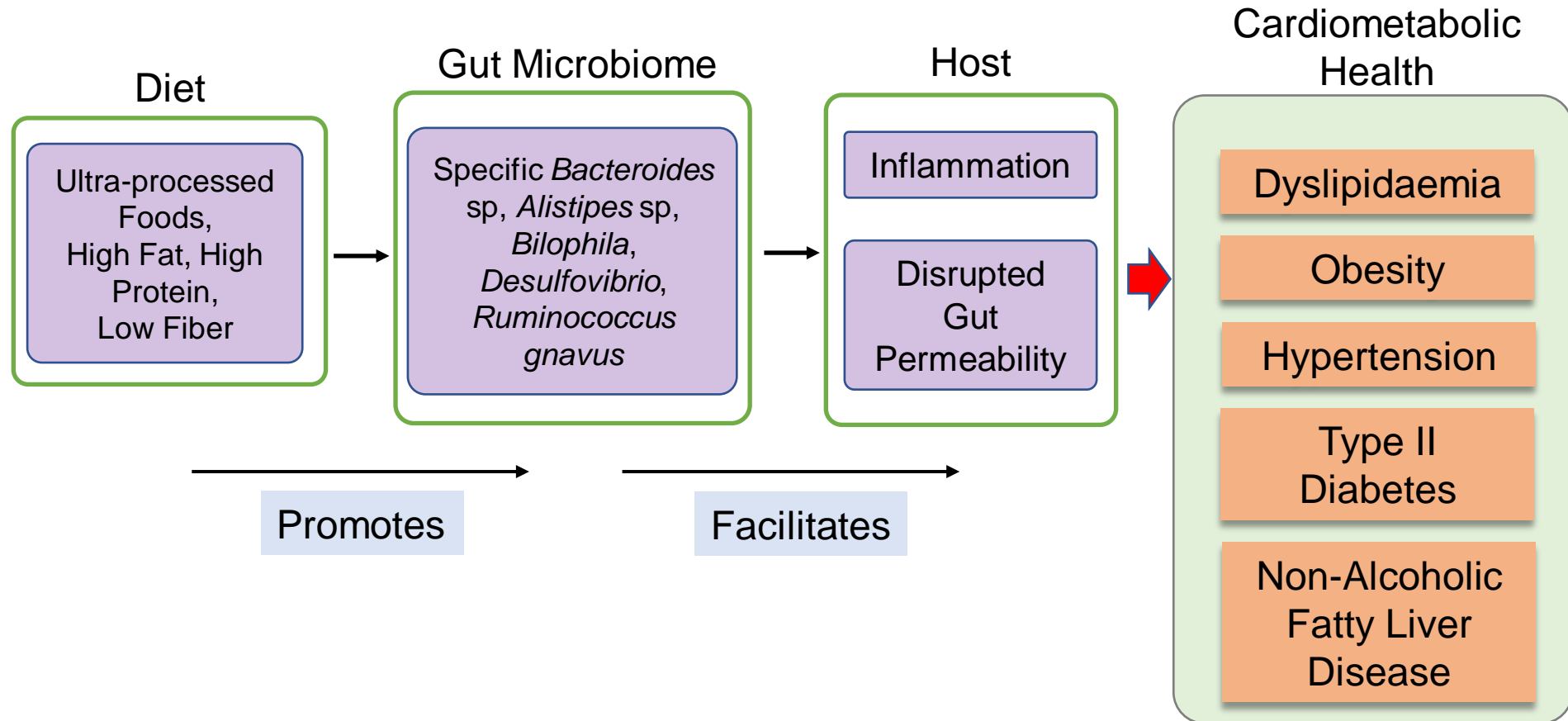
Tarini Shankar Ghosh <sup>1, 2</sup>, Simone Rampelli <sup>3</sup>, Ian B Jeffery <sup>1, 2</sup>, Aurelia Santoro <sup>4, 5</sup>, Marta Neto <sup>1, 2</sup>, Miriam Capri <sup>3</sup>, Enrico Giampieri <sup>4</sup>,  Amy Jennings <sup>6</sup>, Marco Candela <sup>3</sup>, Silvia Turrone <sup>3</sup>, Erwin G Zoetendal <sup>7</sup>,  Gerben D A Hermes <sup>7</sup>, Caumon Elodie <sup>8</sup>, Nathalie Meunier <sup>8</sup>, Corinne Malpuech Brugere <sup>9</sup>, Estelle Pujos-Guillot <sup>10</sup>, Agnes M Berendsen <sup>11</sup>, Lisette C P G M De Groot <sup>11</sup>, Edith J M Feskens <sup>11</sup>,  Joanna Kaluza <sup>12</sup>,  Barbara Pietruszka <sup>12</sup>, Marta Jeruszka Bielak <sup>12</sup>, Blandine Comte <sup>10</sup>, Monica Maijo-Ferre <sup>13</sup>, Claudio Nicoletti <sup>13, 14</sup>, Willem M De Vos <sup>7, 15</sup>, Susan Fairweather-Tait <sup>16</sup>, Aedin Cassidy <sup>17</sup>, Patrizia

**BMJ** Journals

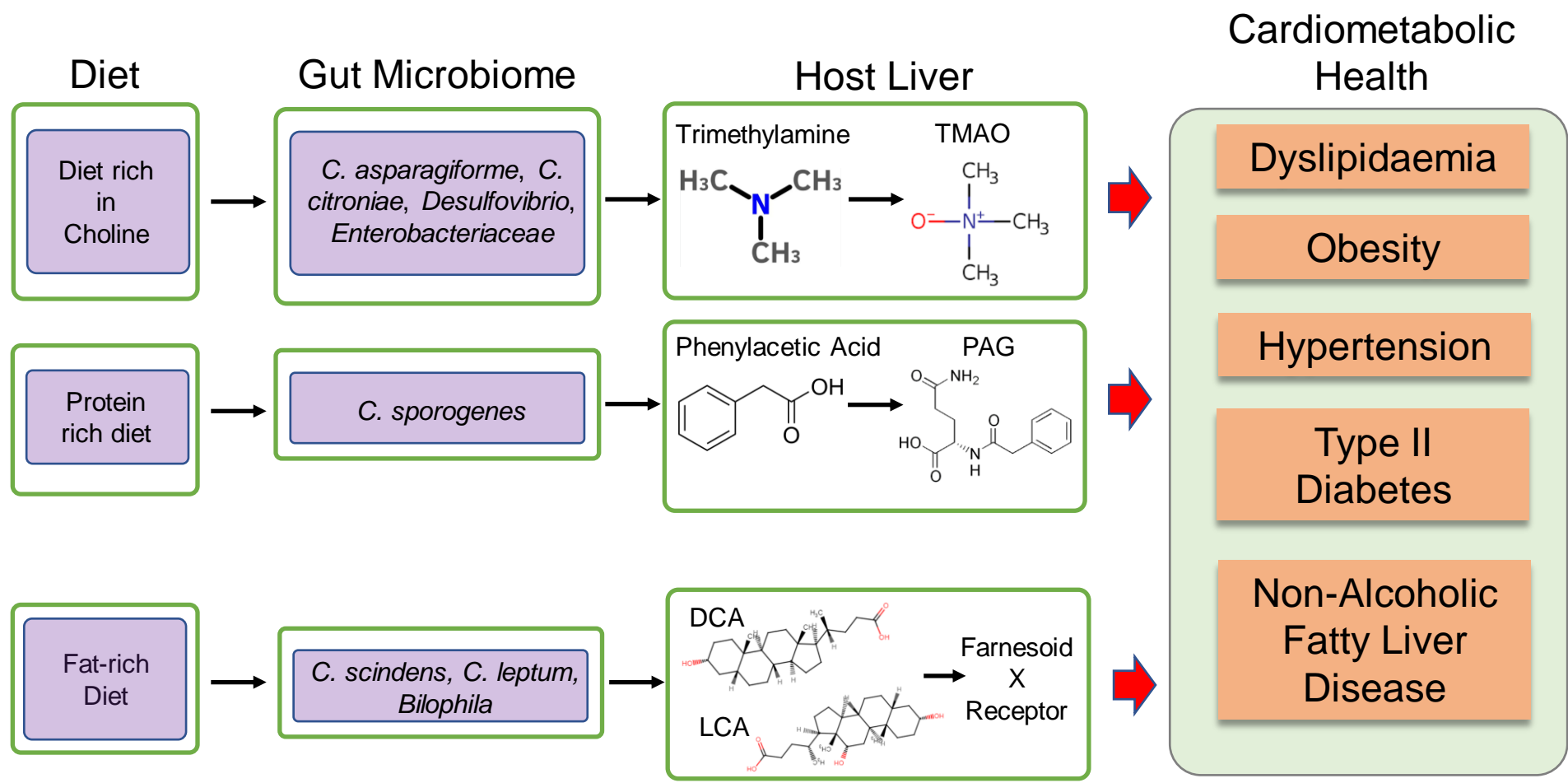
**Gut**



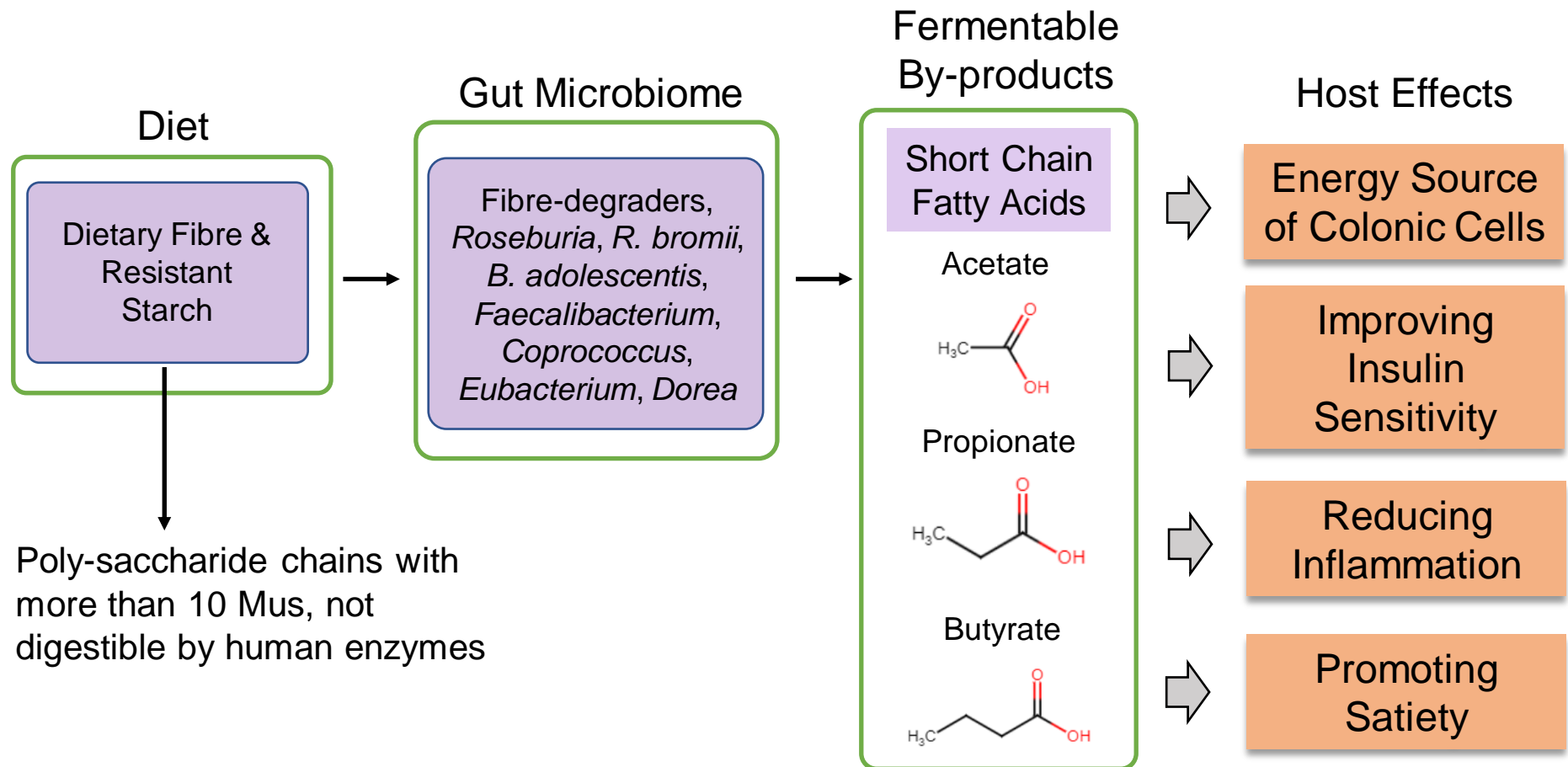
# Some examples of detrimental effects of bad diet on gut microbiome on cardiometabolic health



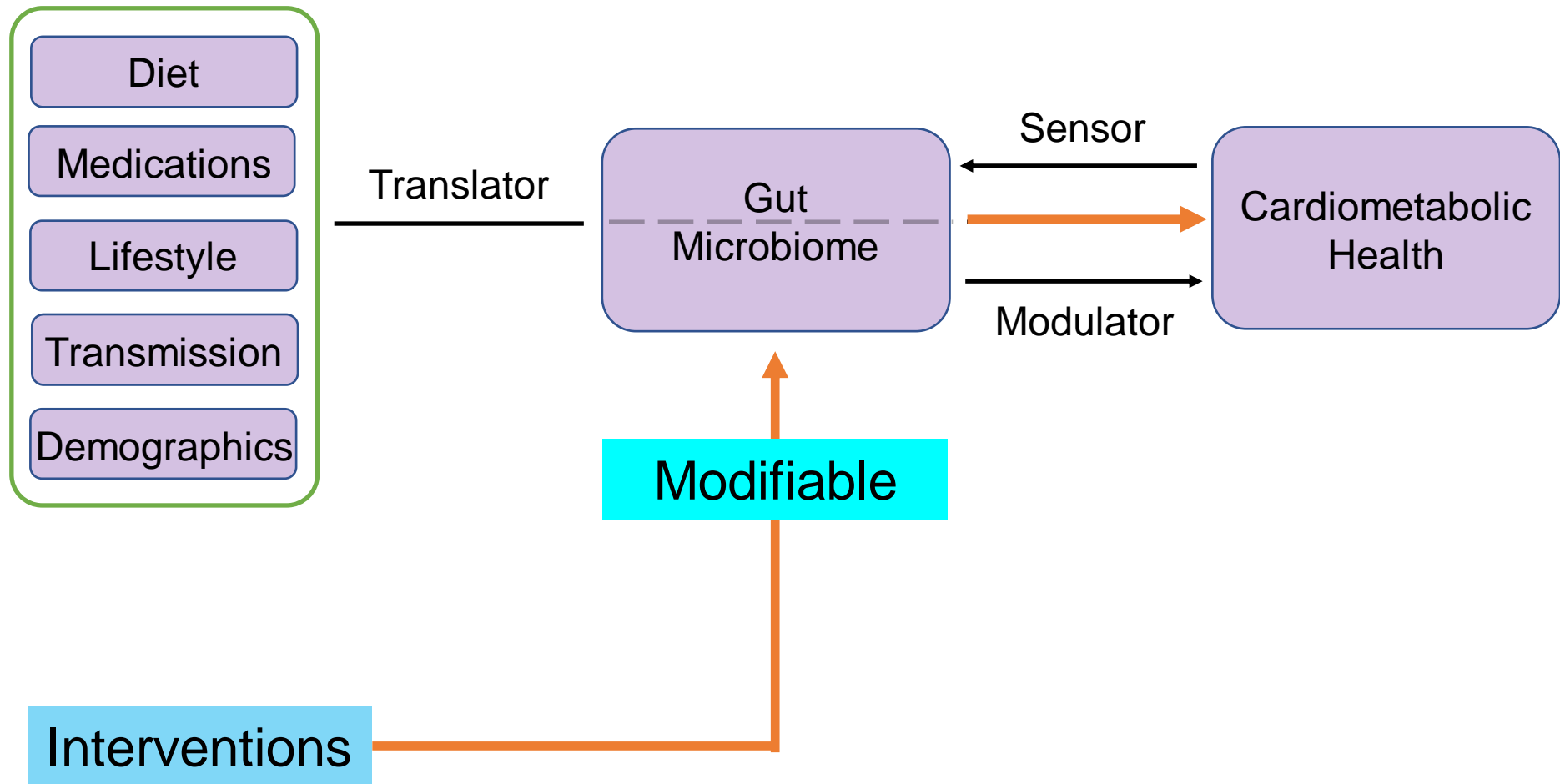
# Gut microbiota as a detrimental modulator/translator of cardiometabolic health



# Fibre and Short Chain Fatty Acids



# Microbiome modifiable using interventions





# Microbiome associated interventions

**Prebiotics:** Substrates that are specifically utilized by 'beneficial' host microbes to promote health: beta-Glucan, Fructans, Inulin, Lactulose

**Dietary Interventions:**  
Mediterranean Diet,  
Increased Fibre Intake, Fruits  
and Vegetables Intake,  
Reduced Processed Foods

**Combinatorial  
therapies:** Dietary  
Supplementations  
(Diet + Pro/Pre-  
biotic)

**Probiotics:** Live organisms  
in viable forms that are  
thought to promote host  
health, either directly or  
through cross-feeding:  
Lactobacilli,  
Bifidobacteria, etc



**Fecal microbiome  
transplantation:** Transfer of  
fecal matter from a healthy  
donor to the GI tract of a  
patient

**Synbiotics:** Combination of probiotics  
and prebiotics

**Postbiotics:** Preparations of inanimate  
organisms or their components

# Lack of consistencies across studies poses challenges: An example of fibre supplementation

- Only 38% of the studies reported a significant link between dietary fibre and SCFAs

- Dependent on dose and fibre structure

- Sometimes the effect can be negative if the modulatory bacteria are absent



Systematic Review

## Effects of Dietary Fibers on Short-Chain Fatty Acids and Gut Microbiota Composition in Healthy Adults: A Systematic Review

Valentina Vinelli <sup>1</sup>, Paola Biscotti <sup>1</sup>, Daniela Martini <sup>1</sup>, Cristian Del Bo' <sup>1</sup>, Mirko Marino <sup>1</sup>, Tomás Meroño <sup>2,3</sup>, Olga Nikoloudaki <sup>4</sup>, Francesco Maria Calabrese <sup>5</sup>, Silvia Turroni <sup>6</sup>, Valentina Taverniti <sup>1</sup>, Andrea Unión Caballero <sup>2</sup>, Cristina Andrés-Lacueva <sup>2,3</sup>, Marisa Porrini <sup>1</sup>, Marco Gobetti <sup>4</sup>, Maria De Angelis <sup>5</sup>, Patrizia Brigidi <sup>7</sup>, Mariona Pinart <sup>8</sup>, Katharina Nimptsch <sup>8</sup>, Simone Guglielmetti <sup>1</sup> and Patrizia Riso <sup>1,\*</sup>

Gastroenterology 2023;164:228–240

## INFLAMMATORY BOWEL DISEASE

### Unfermented $\beta$ -fructan Fibers Fuel Inflammation in Select Inflammatory Bowel Disease Patients



Heather K. Armstrong, <sup>1,2,3</sup> Michael Bording-Jorgensen, <sup>1,2</sup> Deanna M. Santer, <sup>4</sup> Zhengxiao Zhang, <sup>1,5,6</sup> Rosica Valcheva, <sup>1,5</sup> Aja M. Rieger, <sup>7</sup> Justin Sung-Ho Kim, <sup>8,9</sup> Stephanie I. Dijk, <sup>1,10</sup> Ramsha Mahmood, <sup>3</sup> Olamide Ogungbola, <sup>4</sup> Juan Jovel, <sup>1</sup> France Moreau, <sup>11</sup> Hayley Gorman, <sup>11</sup> Robyn Dickner, <sup>1,2</sup> Jeremy Jerasi, <sup>1,2</sup> Inderdeep K. Mander, <sup>1,2</sup> Dawson Lafleur, <sup>1,2</sup> Christopher Cheng, <sup>1,2</sup> Alexandra Petrova, <sup>2</sup> Terri-Lyn Jeanson, <sup>3</sup> Andrew Mason, <sup>1,5</sup> Consolato M. Sergi, <sup>12</sup> Arie Levine, <sup>13</sup> Kris Chadee, <sup>11</sup> David Armstrong, <sup>8</sup> Sarah Rauscher, <sup>8,9,14</sup> Charles N. Bernstein, <sup>3</sup> Matthew W. Carroll, <sup>2</sup> Hien Q. Huynh, <sup>2</sup> Jens Walter, <sup>1,15</sup> Karen L. Madsen, <sup>1,5</sup> Levinus A. Dieleman, <sup>1,5</sup> and Eytan Wine <sup>1,2,10</sup>

There is to survey the evidence for clinical interventions targeting the gut microbiome in cardiometabolic disease

# The premise of the current study

214 human clinical intervention studies targeting the gut microbiome and cardiometabolic diseases

Location of the trials

Demographics

Cohort Size

Duration

Different Intervention Types

Diet

Probiotic

Prebiotic

Synbiotics

Diet + Others

Clinical Targets

Obesity

Cardiovascular Disease

T2D/Insulin Resistance/GDM

Metabolic Syndrome & General CMD Risk

NAFLD & Variants

Intervention Efficacy

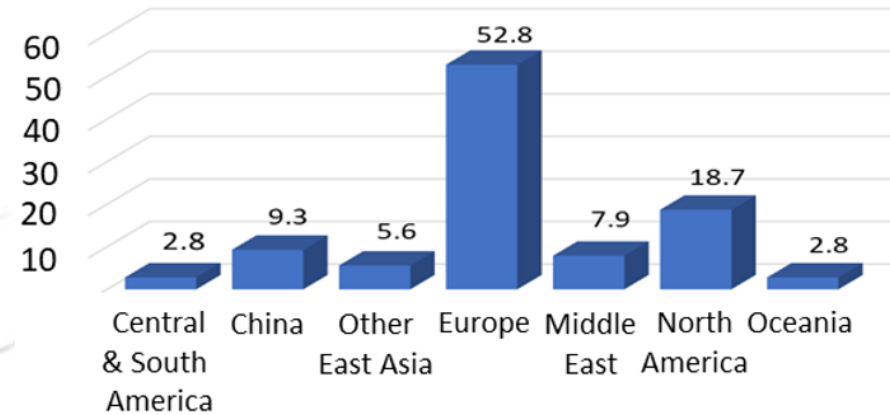
Host Effect

Microbiome Effect

Microbiome Mediated Effect

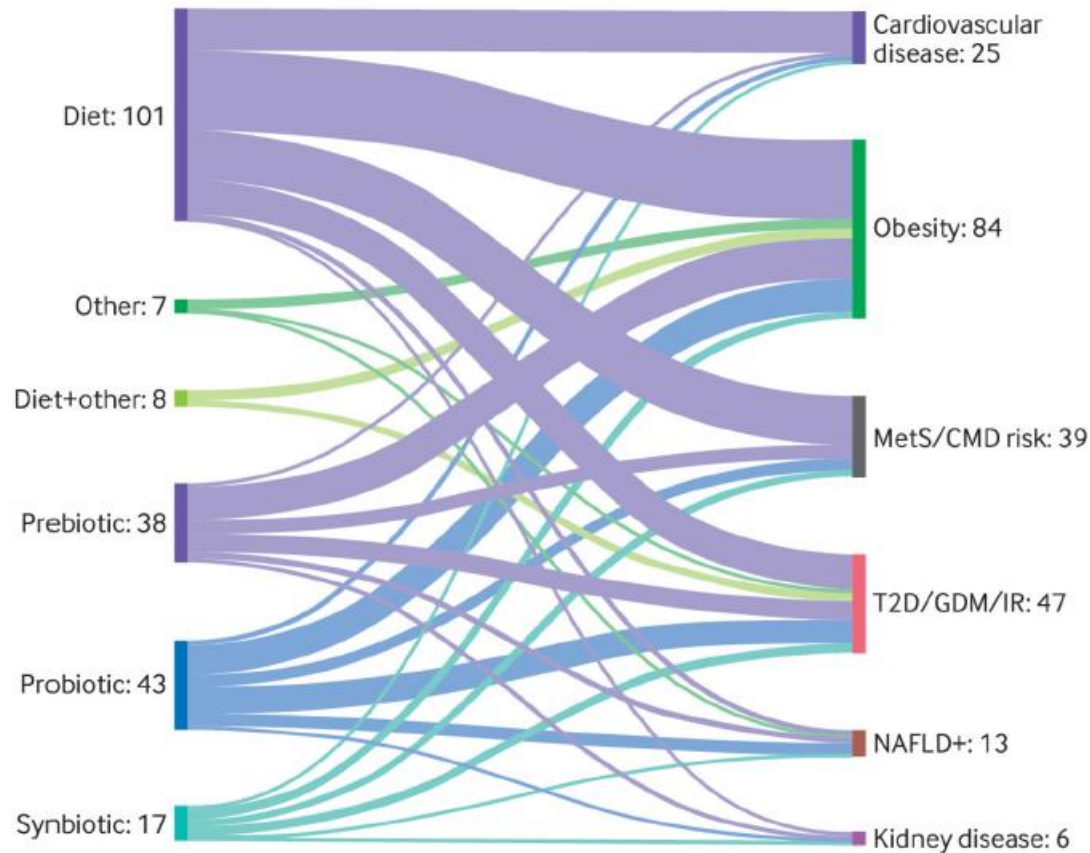
# Large-scale geographical trials in trial locations

Countries where trials have been performed    Region-specific distribution of trials



- Location of trials predominantly in EU > North America > China
- No representation from South Asia and Africa !!

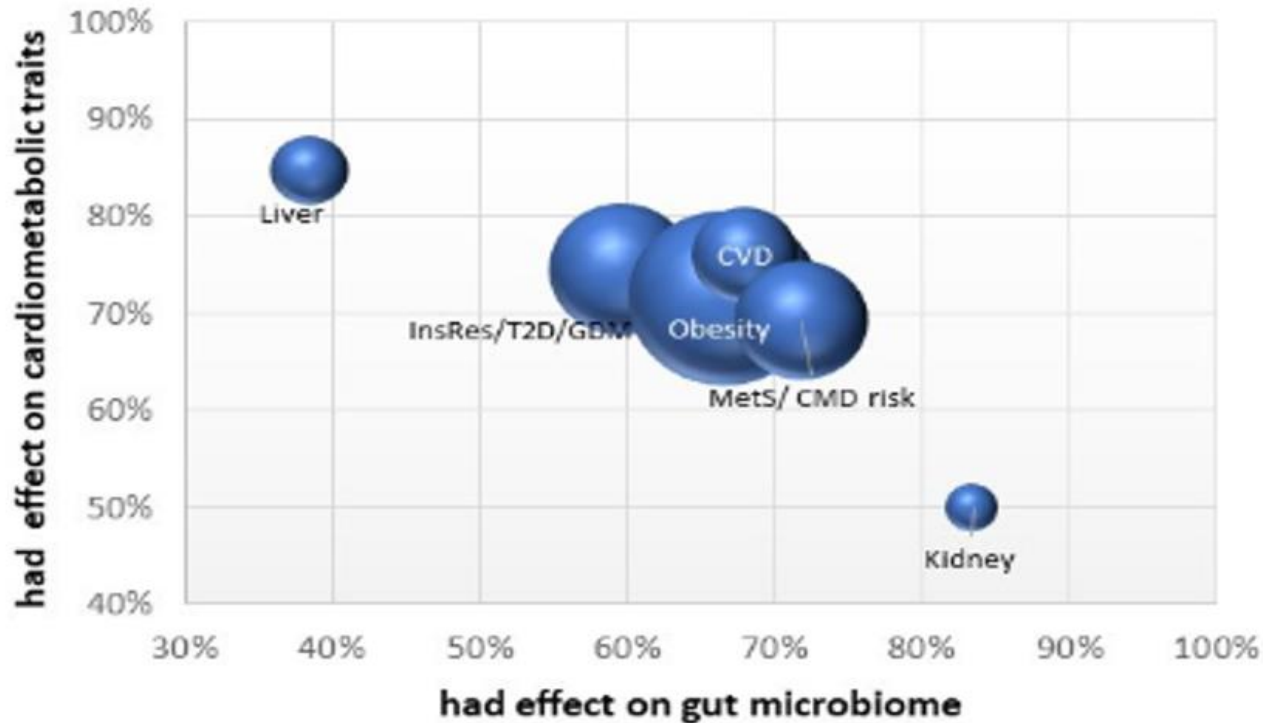
# Majority (51%) involved dietary interventions and were focused on Obesity



Interventions are mostly still seen as extensions of nutrition and “healthy-eating” regimes, not as medical interventions for a given clinical condition

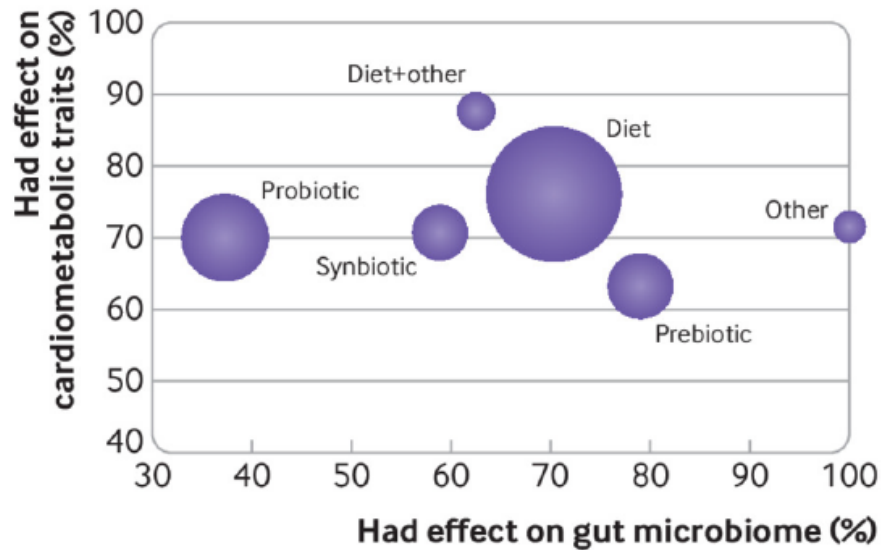
Gut microbiome is primarily a read-out in many interventions

72.5% had efficacy with respect to host clinical outcomes and 63.% had significant effects on the gut microbiome



The pattern was invariant across targeted conditions, with minor exceptions of Liver and Kidney Diseases

# There were variations in effect depending on the intervention type



## Host-specific Effect

Prebiotics < Probiotic/Synbiotic < Diet +

Difference of efficacy between interventions with highest (Diet +) and lowest efficacy (Prebiotics) was marginal ( $P < 0.094$ )

## Gut Microbiome Effect

Probiotics < Diet / Diet + < Prebiotic

Probiotics had significantly lower effect on the microbiome when compared to Diet & Prebiotics ( $P < 0.034$ )

No significant difference between single and multi-strain probiotic interventions



# Probiotic colonization and action have been previously reported to be driven by baseline gut microbiome composition

## Personalized Gut Mucosal Colonization Resistance to Empiric Probiotics Is Associated with Unique Host and Microbiome Features

Cell

Niv Zmora,<sup>1,2,11</sup> Gili Zilberman-Schapira,<sup>1,11</sup> Jotham Suez,<sup>1,11</sup> Uria Mor,<sup>1,11</sup> Mally Dori-Bachash,<sup>1</sup> Stavros Bashiardes,<sup>1</sup> Eran Kotler,<sup>3,4</sup> Maya Zur,<sup>1</sup> Dana Regev-Lehavi,<sup>1</sup> Rotem Ben-Zeev Brik,<sup>1</sup> Sara Federici,<sup>1</sup> Yotam Cohen,<sup>1</sup> Raquel Linevsky,<sup>1</sup> Daphna Rothschild,<sup>3,4</sup> Andreas E. Moor,<sup>3</sup> Shani Ben-Moshe,<sup>3</sup> Alon Harmelin,<sup>5</sup> Shalev Itzkovitz,<sup>3</sup> Nitsan Maharshak,<sup>6,7,8</sup> Oren Shibolet,<sup>6,7,8</sup> Hagit Shapiro,<sup>1</sup> Meirav Pevsner-Fischer,<sup>1</sup> Itai Sharon,<sup>9,10</sup> Zamir Halpern,<sup>6,7,8,12,\*</sup> Eran Segal,<sup>3,4,12,\*</sup> and Eran Elinav<sup>1,12,13,\*</sup>

Metagenomic analysis reveals distinct patterns of gut lactobacillus prevalence, abundance, and geographical variation in health and disease

gut  
microbes

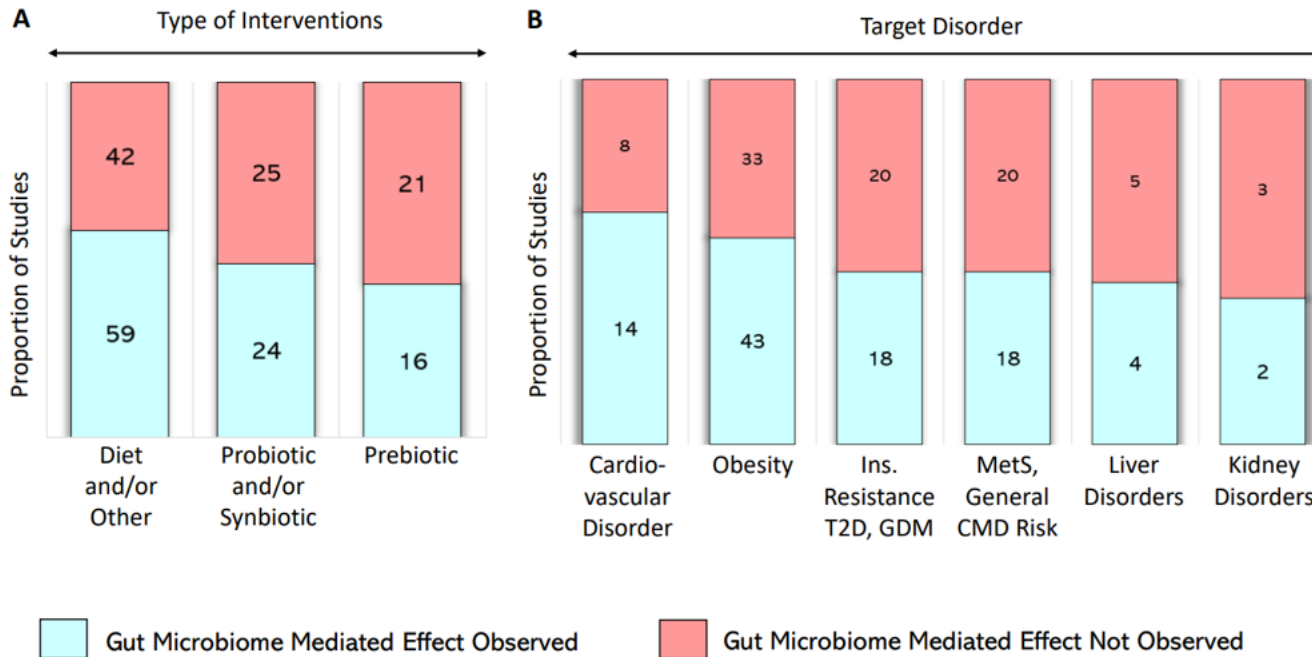
Tarini Shankar Ghosh, Jerome Arnoux & Paul W. O'Toole



Taylor & Francis  
Taylor & Francis

# Gut microbial changes mediating effects of cardiometabolic health

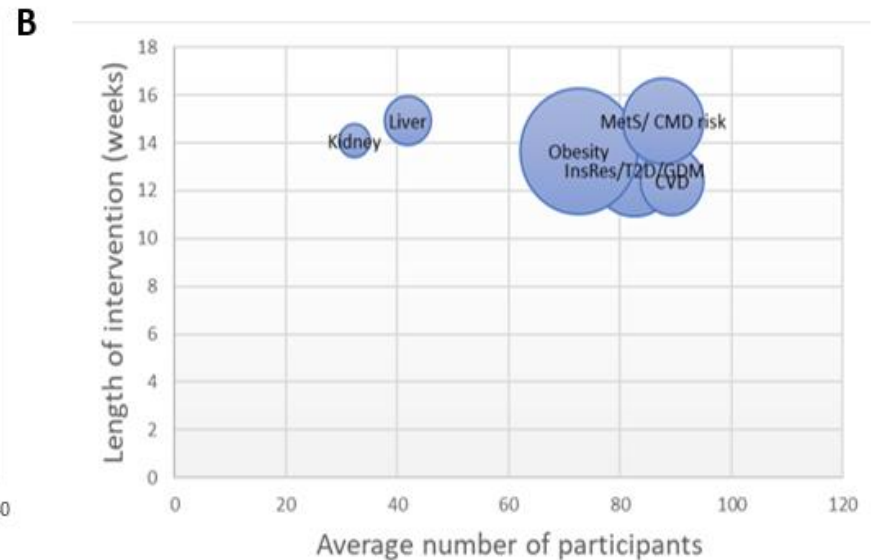
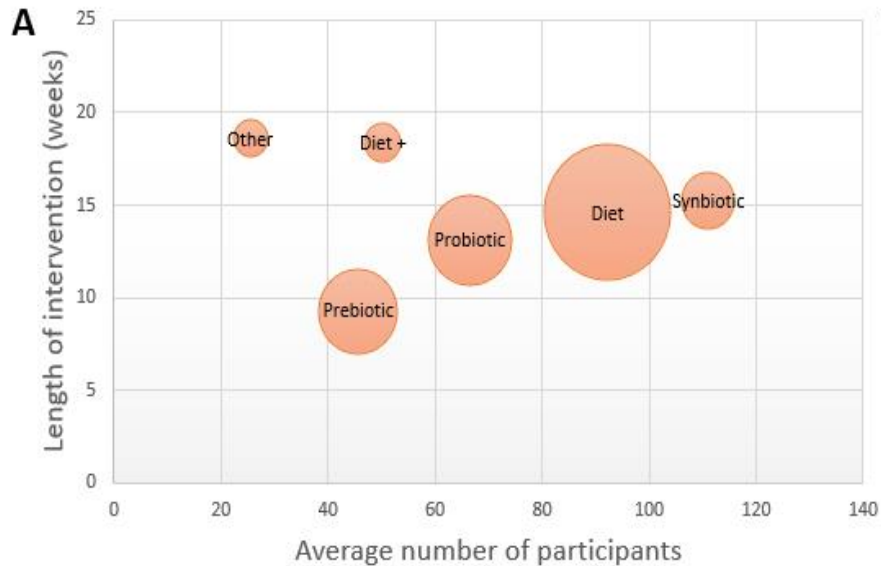
- In studies reporting significant changes in host phenotype, 63% reported these (or additional) effects to be mediated by gut microbiome changes



- Minor variations across intervention types and clinical conditions

- No uniform strategy to measure these effects, ranging from associations to mediation models to Machine learning models

# There were also variations in the cohort size and duration of the interventions



Prebiotic interventions had noticeably lower mean length of interventions and smaller cohort size

May result in lesser power and effect of intervention trials

# Summary of findings

- 72.4% of studies reporting significant improvements in one or more cardiometabolic traits
- Interventions tackling cardiovascular or organ specific diseases (such as atherosclerosis, hypertension, and non-alcoholic fatty liver disease), not just obesity, is fairly modest in proportion.
- Large geographic disparity exists in the location of these trials
- Probiotic interventions are less effective in improving the gut microbiome than broader interventions like diet and prebiotic that target multiple resident gut microbiome members simultaneously
- Lack of unified cohort, study design and effect measurement protocols

# Future Directions

- Better candidate probiotics derived from resident gut microflora, e.g. Akkermansia and Faecalibacterium
- Mechanistic Multi-OMIC level investigations to generate a robust microbial metabolite markers of cardiometabolic risk
- Well-designed large studies involving functional investigation and integration of multiple host phenotypic data
- Formulation of efficient integrative data-investigation strategies to probe host-diet-microbiota interactions

# Acknowledgements



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