



THE ITALIAN MICROBIOME INITIATIVE FOR IMPROVED HUMAN HEALTH AND AGRI-FOOD PRODUCTION

Giovedì 11 luglio 2019 - ore 14,30



THE ITALIAN MICROBIOME INITIATIVE
For Improved Human Health and Agri-Food Production

Sala Monumentale Presidenza del Consiglio – Largo Chigi, 19 - Roma

THE ITALIAN MICROBIOME INITIATIVE FOR IMPROVED HUMAN HEALTH AND AGRI-FOOD PRODUCTION

Il Microbioma Umano dalla ricerca alla clinica attraverso i big data

University of Rome “Tor Vergata”
Department of Systems Medicine



MASSIMO FEDERICI

Center for Atherosclerosis
“Tor Vergata University Hospital”



THE HUMAN

MICROBIOME

Bacteria, fungi, and viruses outnumber human cells in the body by a factor of 10 to one. The microbes synthesize key nutrients, fend off pathogens and impact everything from weight gain to perhaps even brain development. The Human Microbiome Project is doing a census of the microbes and sequencing the genomes of many. The total body count is not in but it's believed over 1,000 different species live in and on the body.

25
SPECIES

in the stomach include:

- *Helicobacter pylori*
- *Streptococcus thermophilus*

500-
1,000
SPECIES

in the intestines include:

- *Lactobacillus casei*
- *Lactobacillus reuteri*
- *Lactobacillus gasseri*
- *Escherichia coli*
- *Bacteroides fragilis*
- *Bacteroides thetaiotaomicron*
- *Lactobacillus rhamnosus*
- *Clostridium difficile*

600+
SPECIES

in the mouth, pharynx and respiratory system include:

- *Streptococcus viridans*
- *Neisseria sicca*
- *Candida albicans*
- *Streptococcus salivarius*

1,000
SPECIES

in the skin include:

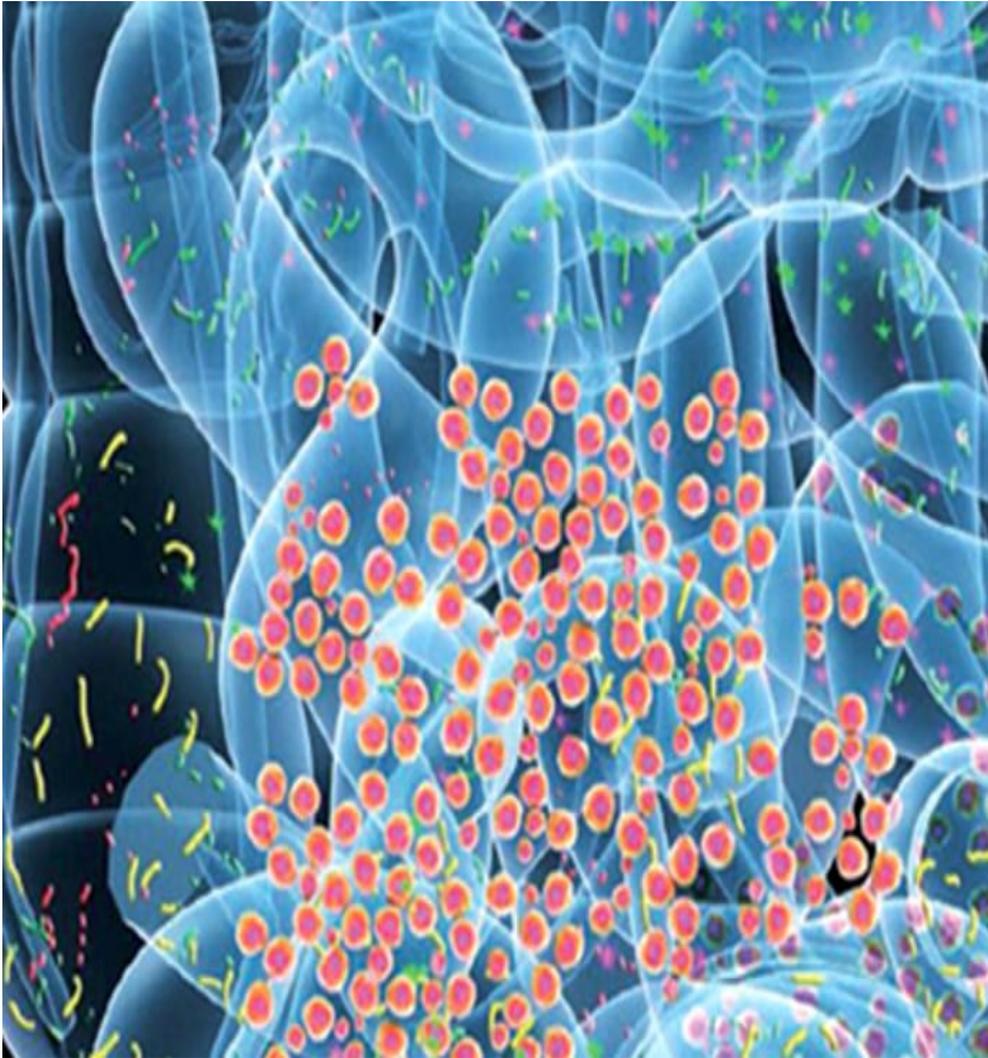
- *Pityrosporum ovale*
- *Staphylococcus epidermidis*
- *Corynebacterium jeikeium*
- *Trichosporon*
- *Staphylococcus haemolyticus*

60
SPECIES

in the urogenital tract include:

- *Ureaplasma parvum*
- *Corynebacterium aurimucosum*

Microbiota e Microbioma



COMPOSIZIONE

Ogni individuo ha uno specifico
«*microbial fingerprint*»

FUNZIONI

- Strutturale (mantenimento integrità della barriera intestinale);
- Assorbimento dei nutrienti;
- Metabolismo;
- Difesa;
- Sviluppo e regolazione dell'immunità

MICROBIOME DISTRIBUTION IN HUMAN GUT SYSTEM

Stomach 10^2

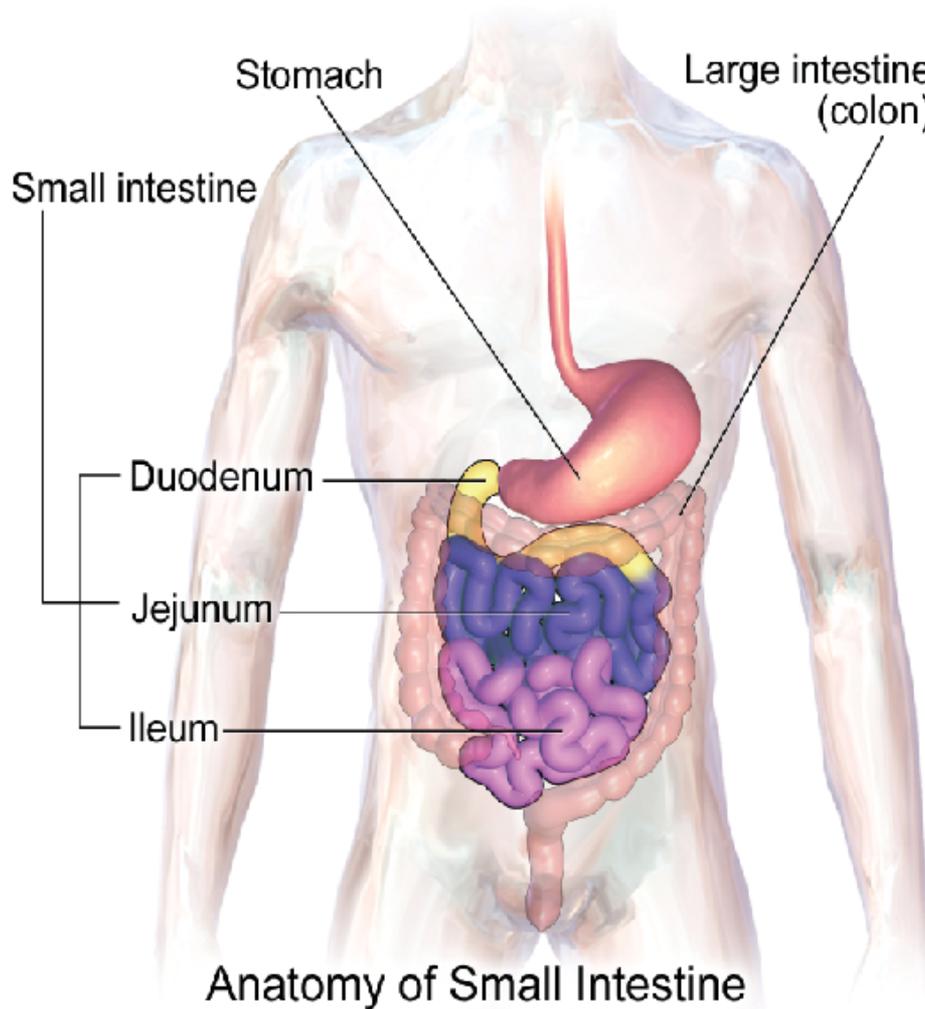
Lactobacillus
Candida
Streptococcus
Helicobacter pylori
Peptostreptococcus

Duodenum 10^2

Streptococcus
Lactobacillus

Jejunum 10^2

Streptococcus
Lactobacillus



Proximal ileum 10^2

Streptococcus
Lactobacillus

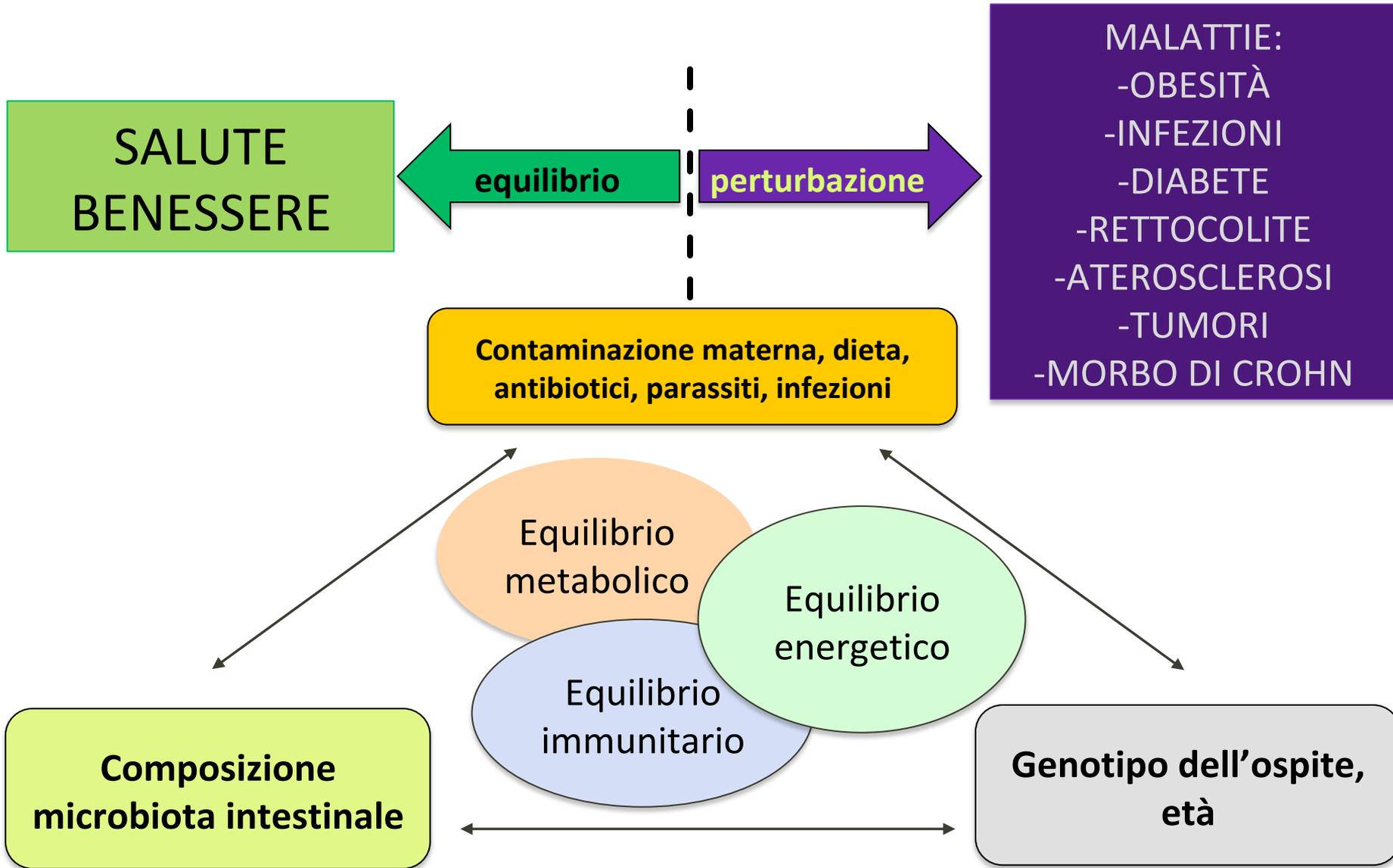
Distal ileum 10^8

Clostridium
Streptococcus
Bacteroides
Actinomycinae
Corneibacteria

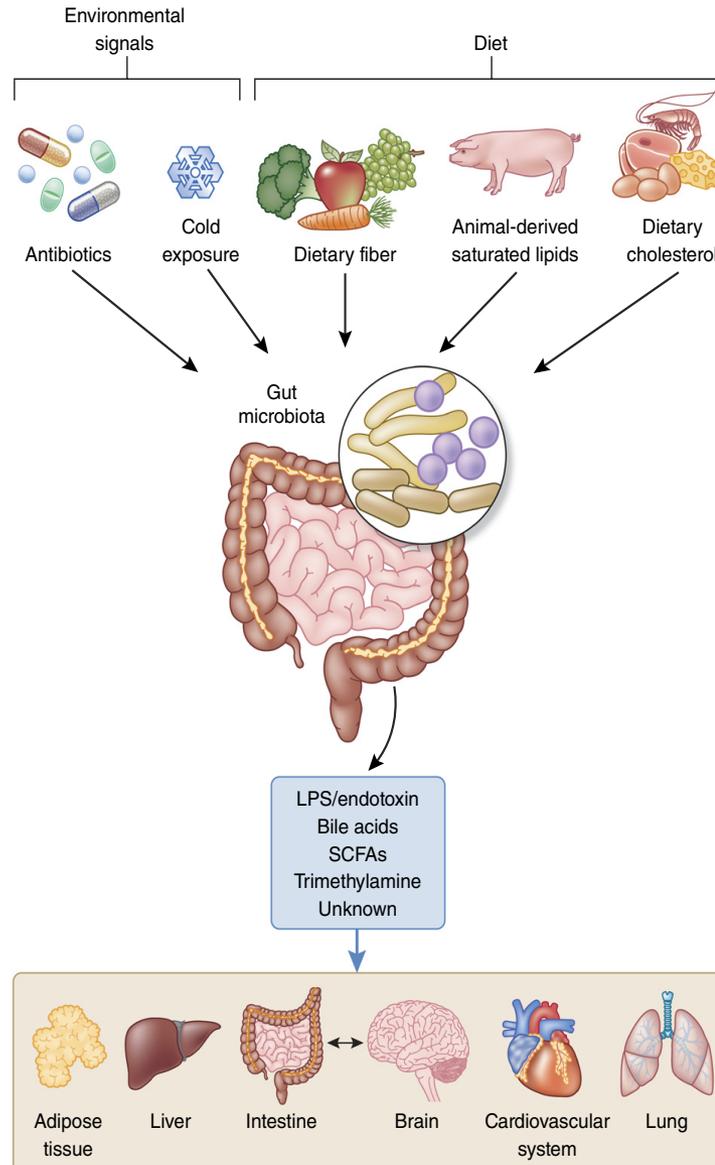
Colon 10^{12}

Bacteroides
Clostridium
Bifidobacterium
Enterobacteriaceae

Un ecosistema in equilibrio

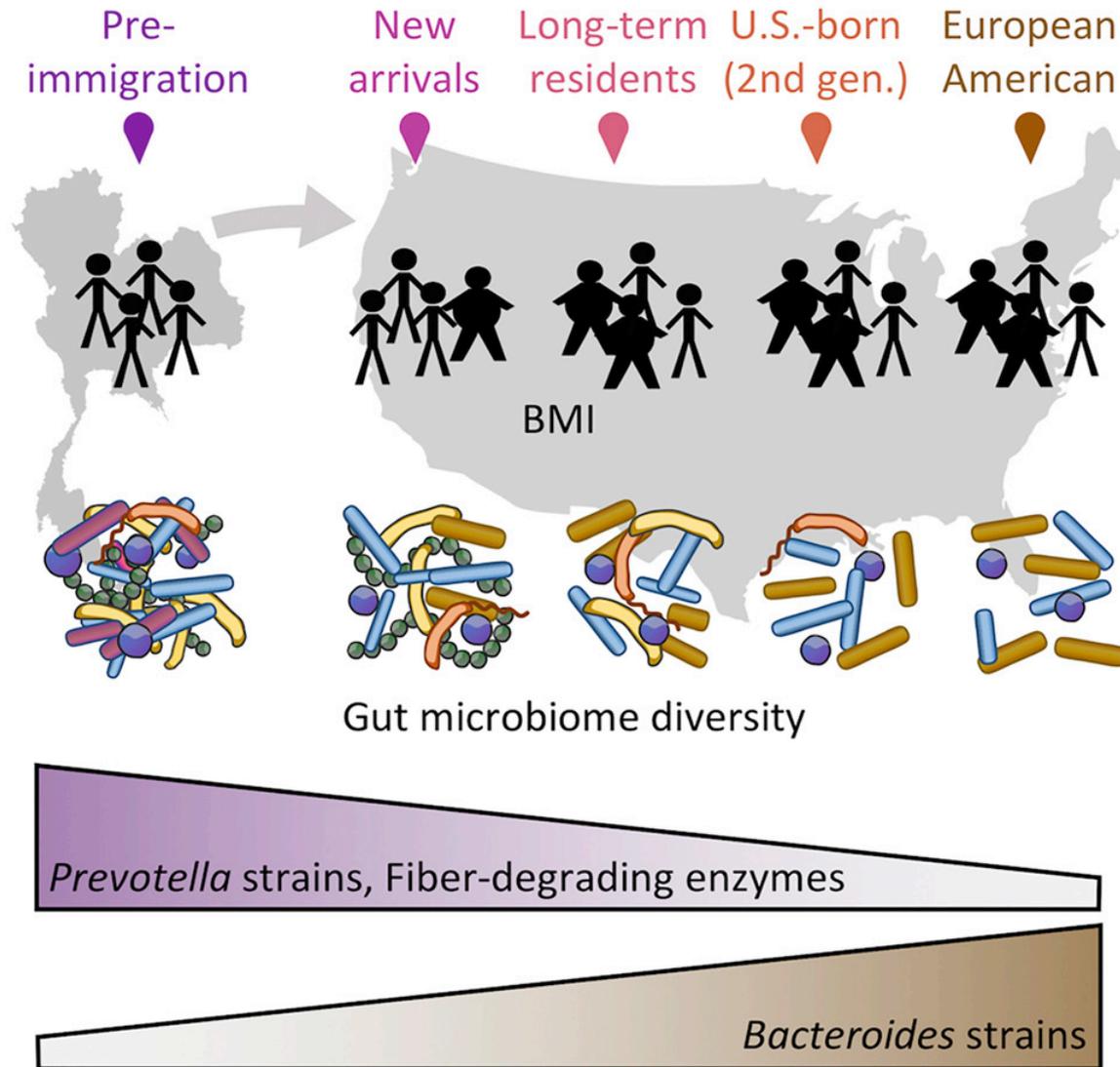


Ambiente e Microbiota



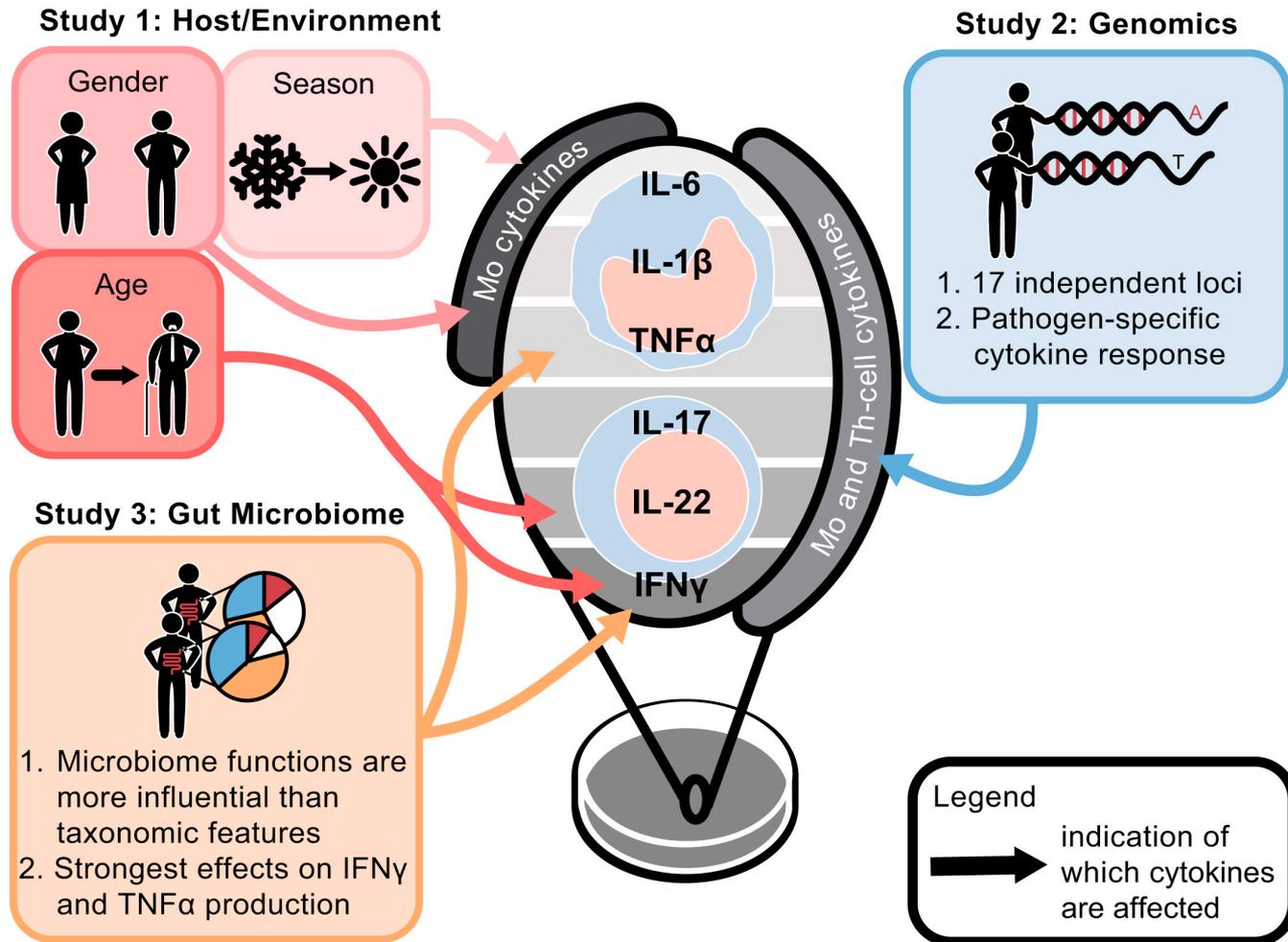
Debbie Maizeis/Nature Publishing Group

L'immigrazione occidentalizza il microbioma umano



MICROBIOMA INTESTINALE E INFIAMMAZIONE SISTEMICA

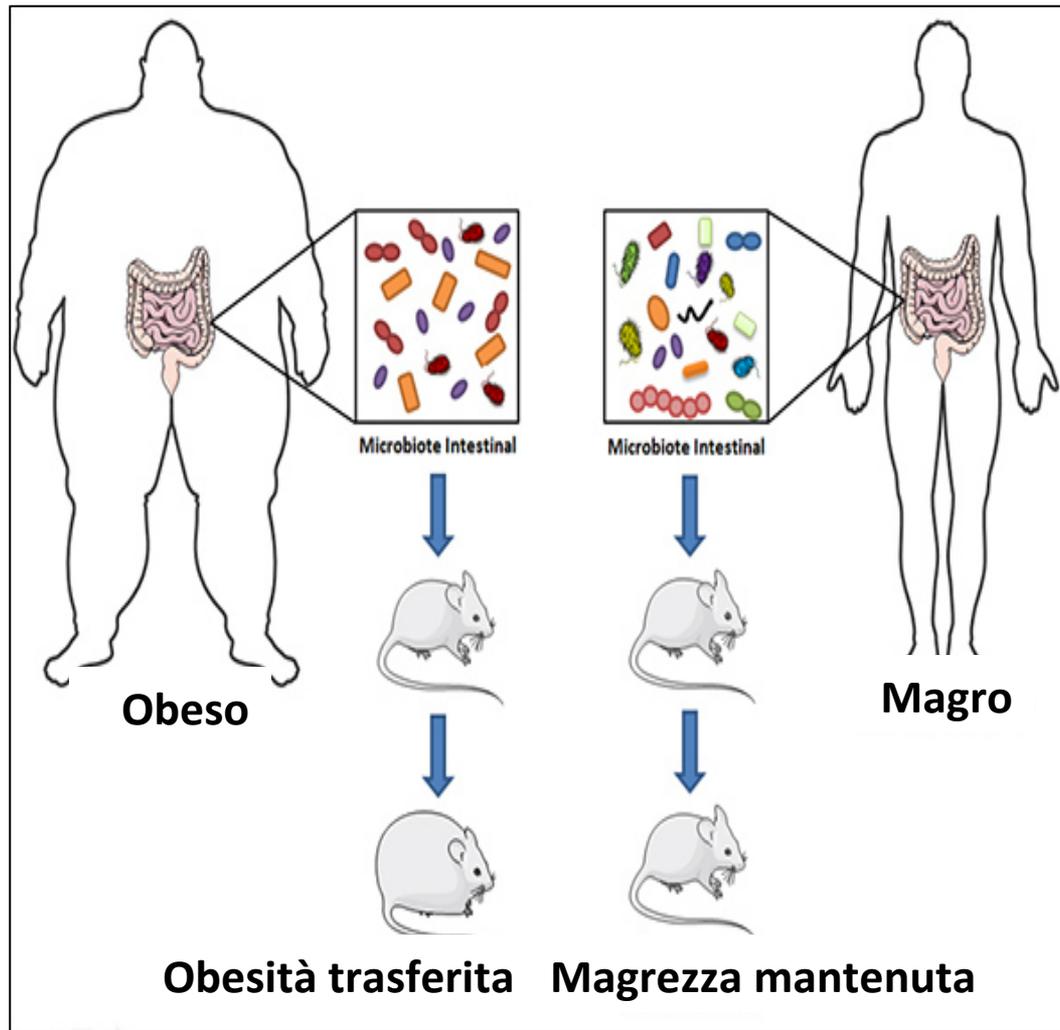
Funzioni del microbioma più influenti della tassonomia



Microbiota intestinale e Malattie

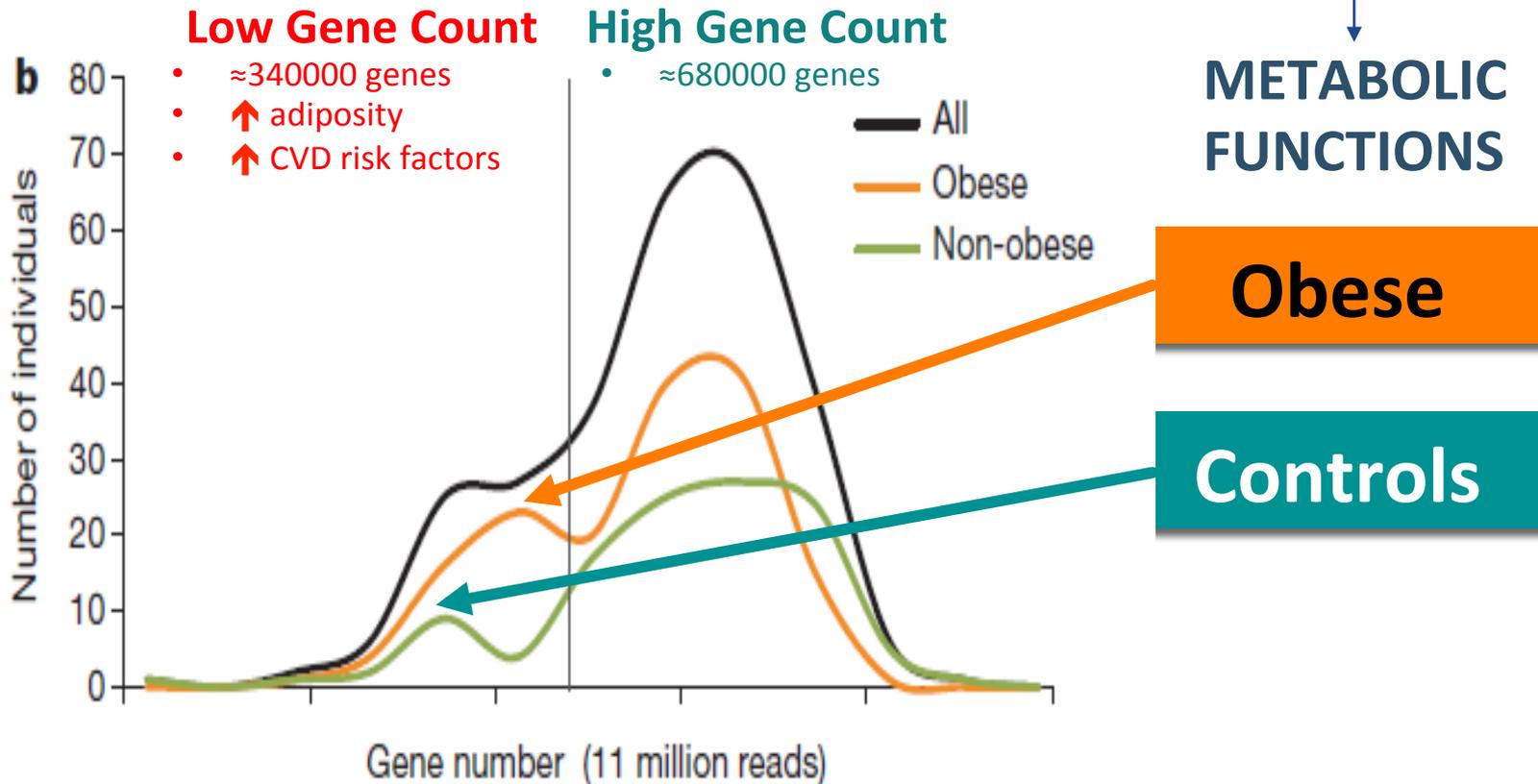
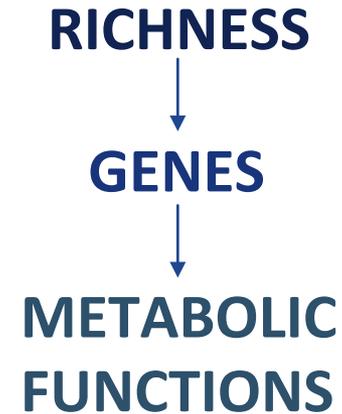


MICROBIOMA INTESTINALE ED OBESITA': EFFETTI CAUSALI

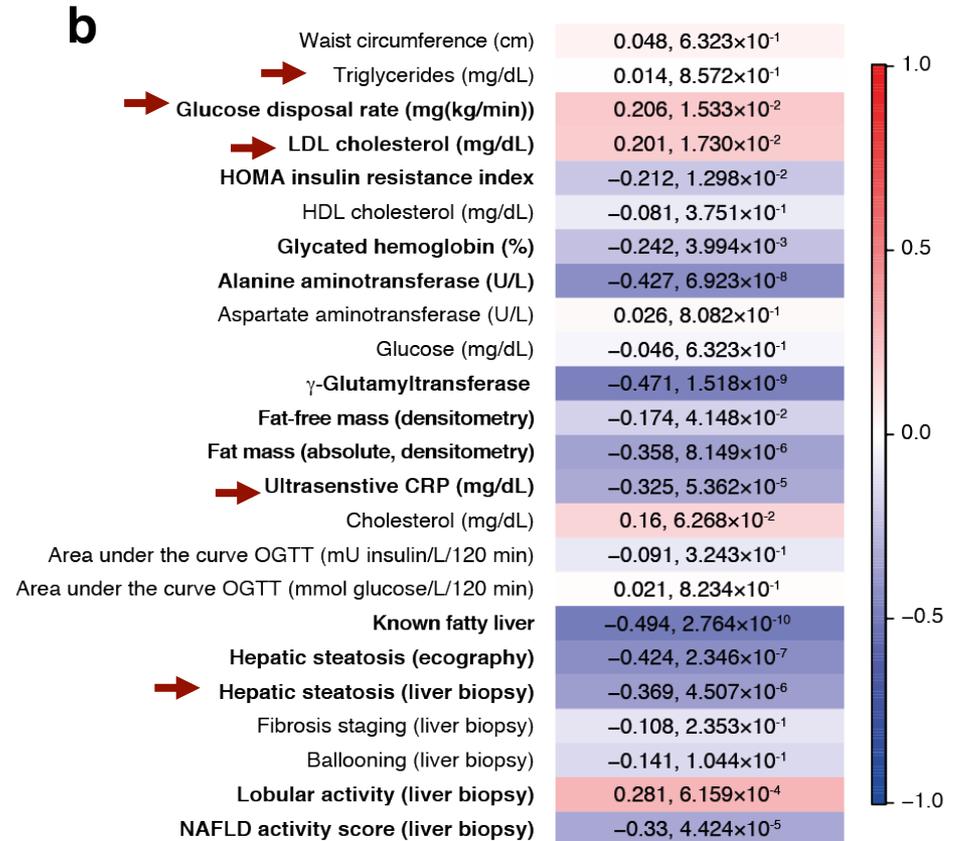
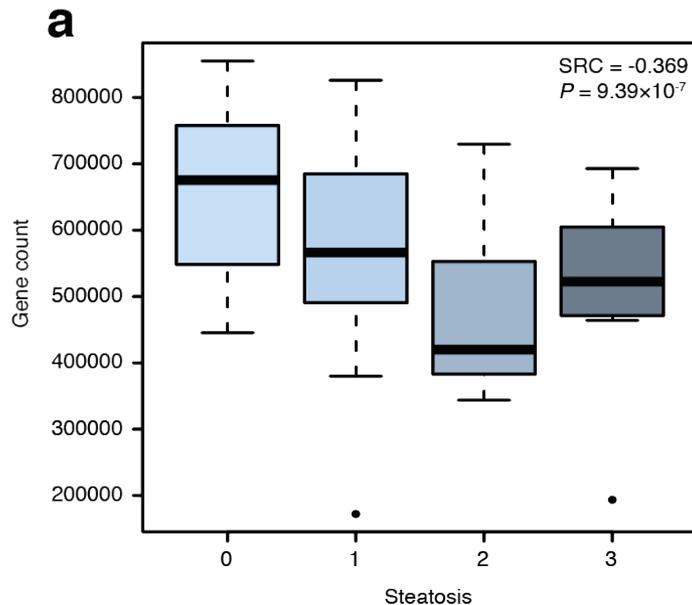


Richness of human gut microbiome correlates with metabolic markers

Emmanuelle Le Chatelier^{1*}, Trine Nielsen^{2*}, Junjie Qin^{3*}, Edi Prifti^{1*}, Falk Hildebrand^{4,5}, Gwen Falony^{4,5}, Mathieu Almeida¹, Manimozhiyan Arumugam^{2,3,6}, Jean-Michel Batto¹, Sean Kennedy¹, Pierre Leonard¹, Junhua Li^{3,7}, Kristoffer Burgdorf², Niels Grarup², Torben Jørgensen^{8,9,10}, Ivan Brandslund^{11,12}, Henrik Bjørn Nielsen¹³, Agnieszka S. Juncker¹³, Marcelo Bertalan¹³, Florence Levenez¹, Nicolas Pons¹, Simon Rasmussen¹³, Shinichi Sunagawa⁶, Julien Tap^{1,6}, Sebastian Tims¹⁴, Erwin G. Zoetendal¹⁴, Søren Brunak¹³, Karine Clément^{15,16,17}, Joël Doré^{1,18}, Michiel Kleerebezem¹⁴, Karsten Kristiansen¹⁹, Pierre Renault¹⁸, Thomas Sicheritz-Ponten¹³, Willem M. de Vos^{14,20}, Jean-Daniel Zucker^{15,16,21}, Jeroen Raes^{4,5}, Torben Hansen^{2,22}, MetaHIT consortium†, Peer Bork⁶, Jun Wang^{3,19,23,24,25}, S. Dusko Ehrlich¹ & Oluf Pedersen^{2,26,27,28}



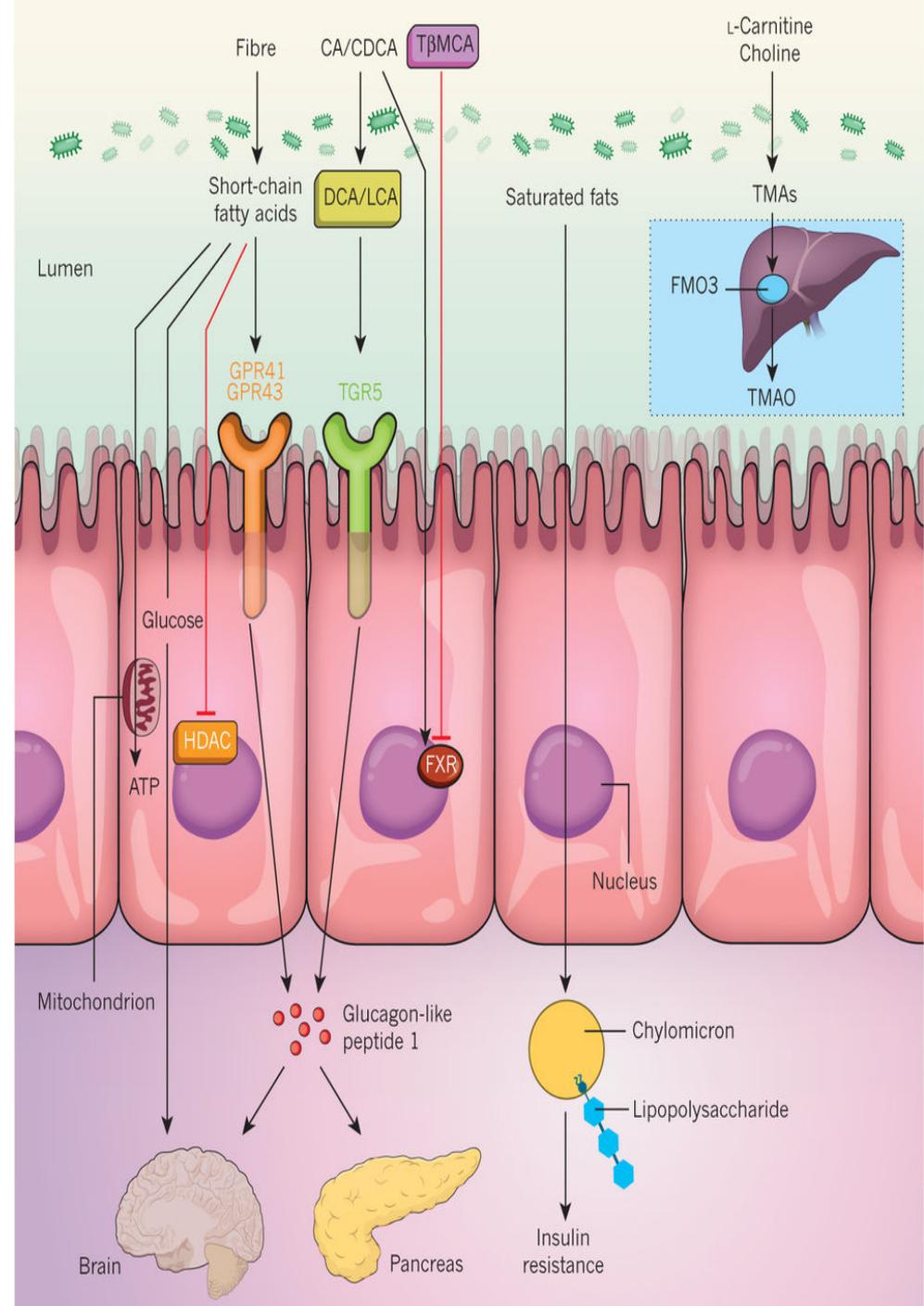
Association of Low Microbial Gene Richness with Steatosis (A) and metabolic variables (B)



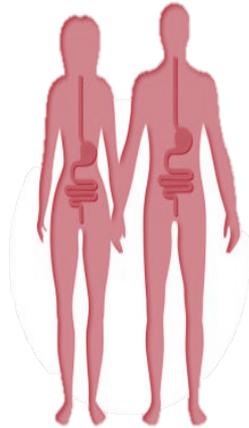
High microbial gene richness =
metabolic health: how?

ENDOCRINE SIGNALS

INVASIVE SIGNALS

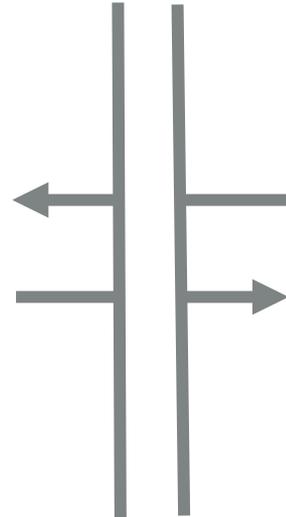


UNA VISIONE FUTURIBILE



OSPITE

**SISTEMA
IMMUNITARIO/
METABOLISMO**



**BARRIERA/
ASSORBIMENTO**

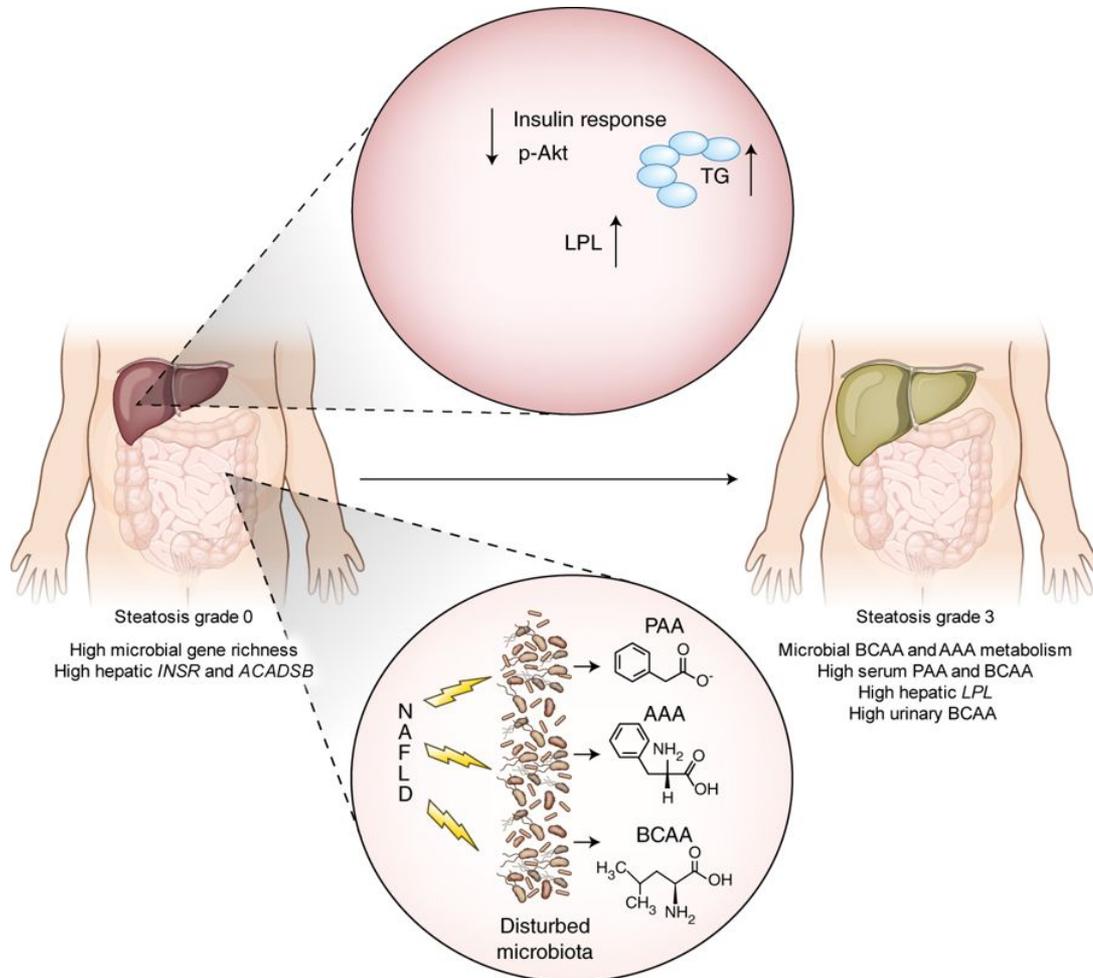


MICROBIOMA

**COMPOSIZIONE
/ FUNZIONE**

**DAL MICROBIOMA:
BIOMARCATORI E OBIETTIVI TERAPEUTICI**

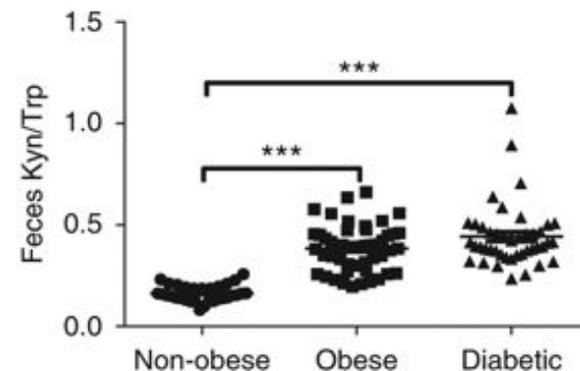
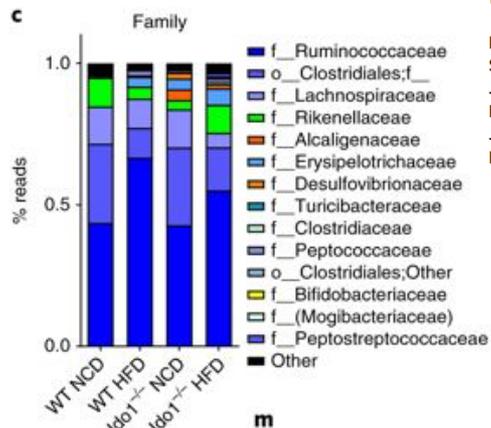
Il fenilacetato (PAA), un metabolita di origine microbica è un potenziale biomarcatore di NAFLD



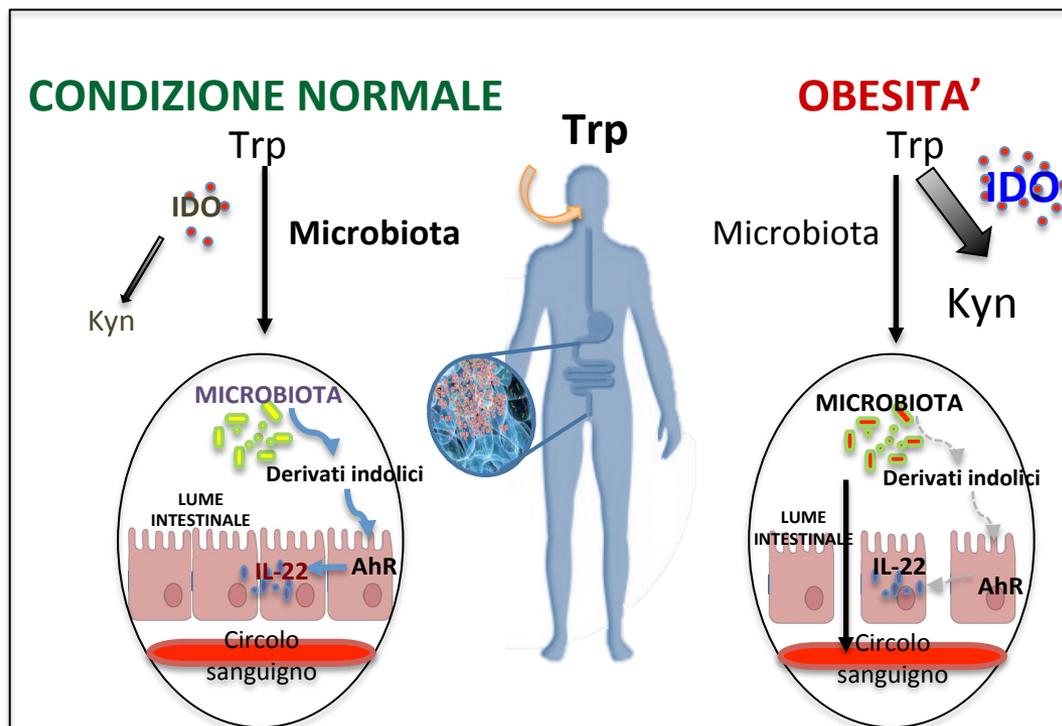
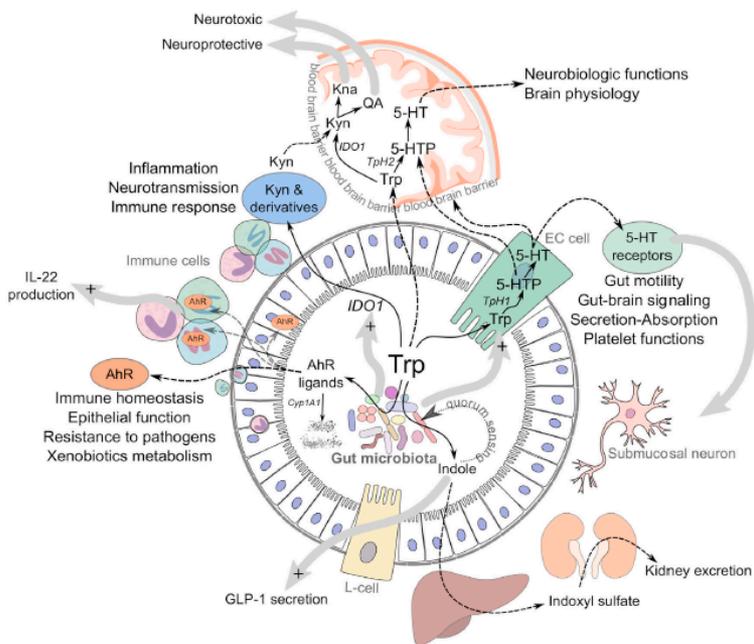


Genetic deficiency of indoleamine 2,3-dioxygenase promotes gut microbiota-mediated metabolic health

Ludivine Laurans¹, Nicolas Venteclef², Yacine Haddad¹, Mouna Chajadine¹, Fawaz Alzaid³, Sarvenaz Metghalchi¹, Bruno Sovran³, Raphael G. P. Denis⁴, Julien Dairou⁵, Marina Cardellini⁶, Jose-Maria Moreno-Navarrete^{7,8}, Marjolene Straub⁹, Sarah Jegou⁹, Claire McQuitty⁹, Thomas Viel¹, Bruno Esposito¹, Bertrand Tavitian¹, Jacques Callebert¹⁰, Serge H. Luquet⁴, Massimo Federici^{6,5}, José Manuel Fernandez-Real⁷, Remy Burcelin¹¹, Jean-Marie Launay¹⁰, Alain Tedgui¹, Ziad Mallat^{1,12}, Harry Sokol^{3,9,13} and Soraya Taleb^{1*}

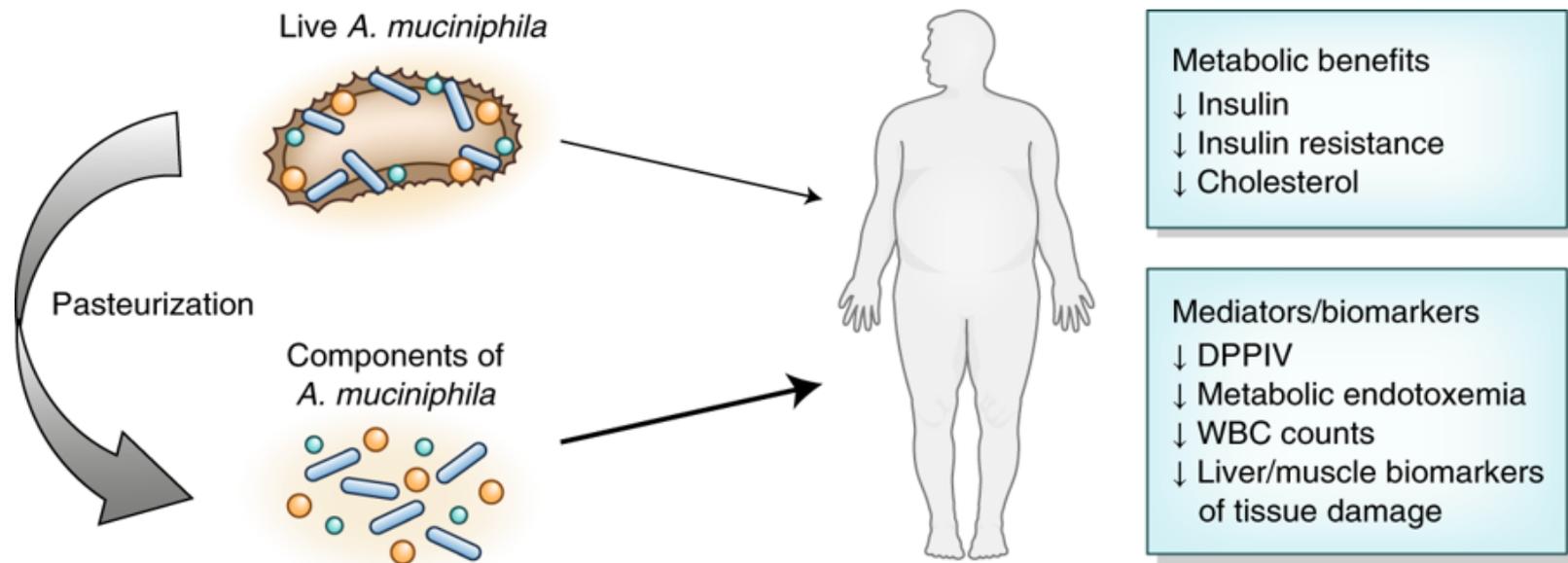


Indoleamina 2,3-Diossigenasi (IDO)



Supplementation with *Akkermansia muciniphila* in overweight and obese human volunteers: a proof-of-concept exploratory study

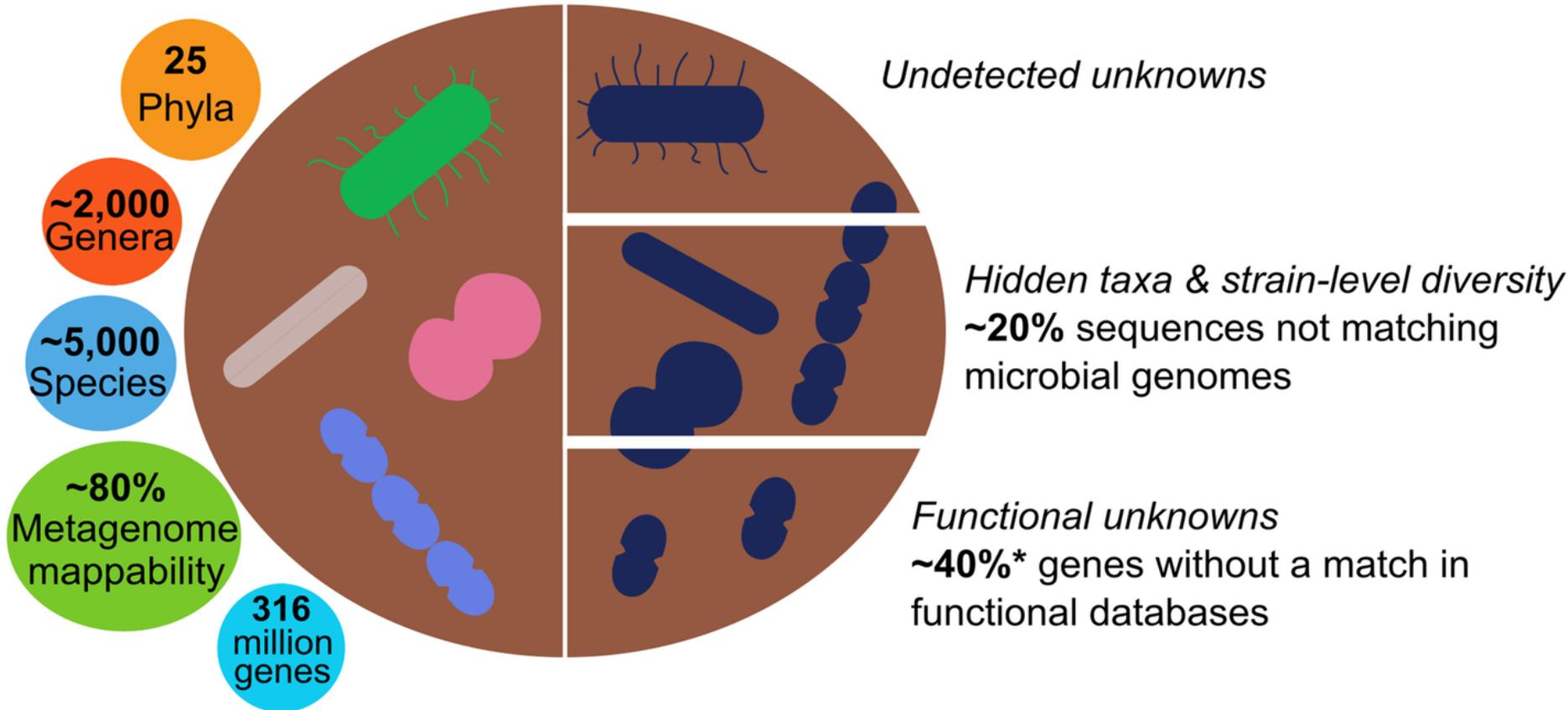
Clara Depommier^{1,9}, Amandine Everard^{1,9}, Céline Druart¹, Hubert Plovier¹, Matthias Van Hul¹, Sara Vieira-Silva^{2,3}, Gwen Falony^{2,3}, Jeroen Raes^{2,3}, Dominique Maiter^{4,5}, Nathalie M. Delzenne⁶, Marie de Barse^{4,5,10}, Audrey Loumaye^{4,5,10}, Michel P. Hermans^{4,5,10}, Jean-Paul Thissen^{4,5,10}, Willem M. de Vos^{7,8,10} and Patrice D. Cani^{1*}



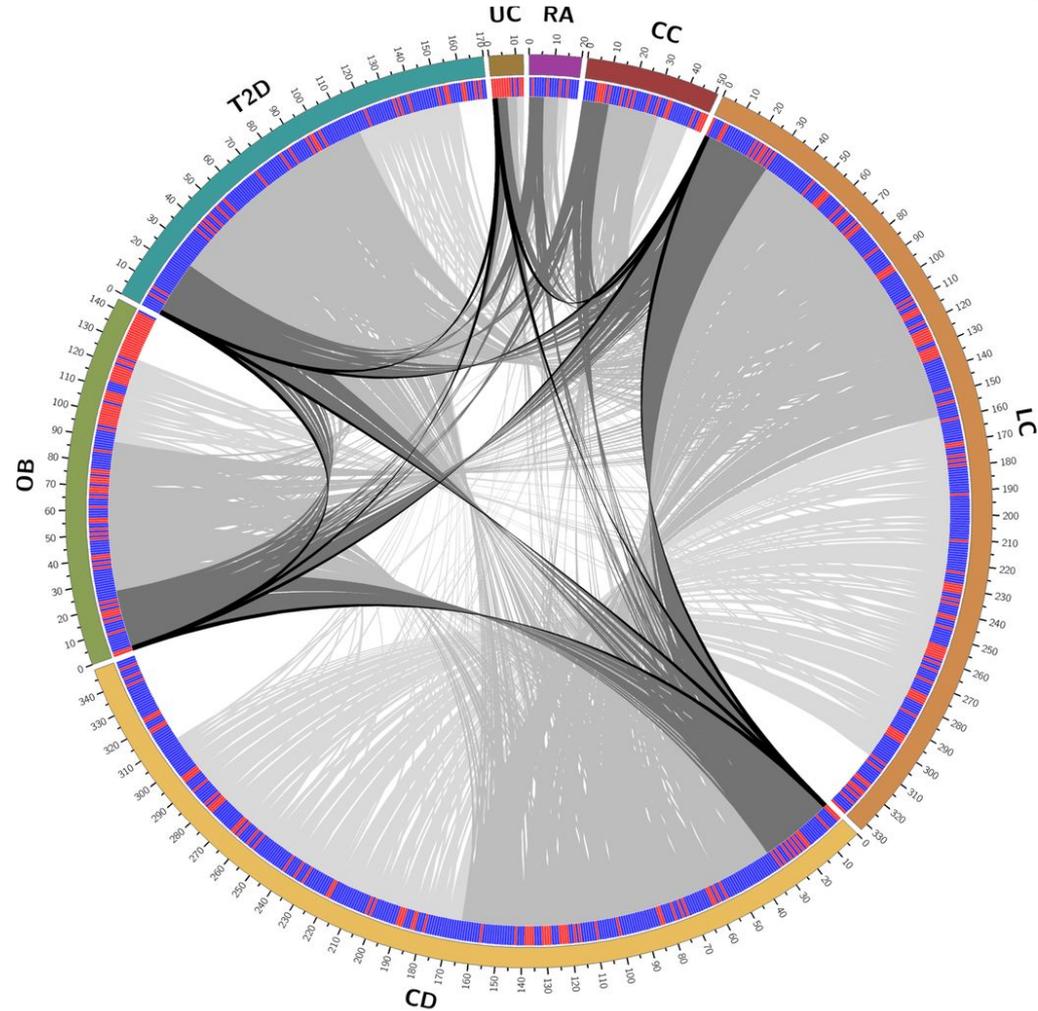
The human microbiome

What is known?

What is unknown?



FATTORI COMUNI E DISTINTI DEL MICROBIOMA UMANO IN MALATTIE CRONICHE NON TRASMISSIBILI (NCDs)



Outer ring colors

- Rheumatoid arthritis (RA)
- Colorectal cancer (CC)
- Liver cirrhosis (LC)
- Crohn's disease (CD)
- Obesity (OB)
- Type II diabetes (T2D)
- Ulcerative colitis (UC)

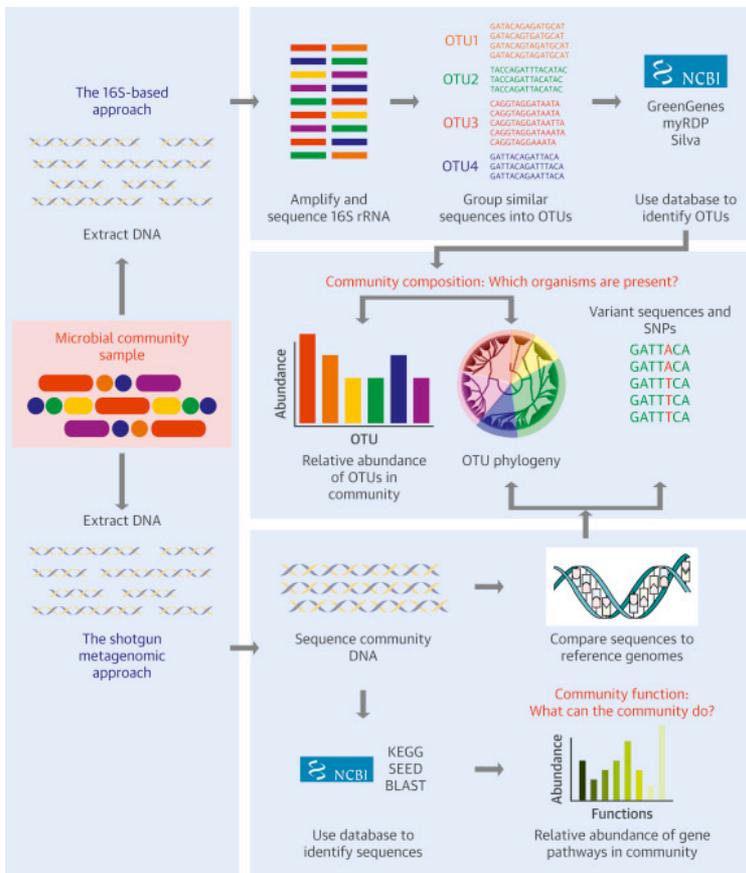
Inner ring colors

- Increased in cases
- Decreased in cases

Arc colors

- 5 diseases
- 4 diseases
- 3 diseases
- 2 diseases
- 1 disease

MICROBIOMA INTESTINALE E MALATTIE



PROCEDURE OPERATIVE STANDARDIZZATE:

- 16S RNA vs shotgun metagenomics
- Tassonomia vs Genomica Funzionale (basata sull'effetto fisiologico dei prodotti genici)
- Approccio Statistico (FDR, PCA, RF, diversity, etc)

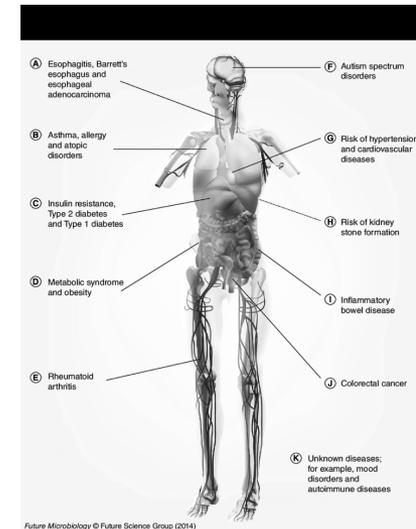
USO APPROPRIATO:

- Specie come obiettivi farmacologici?
- Dati Genomici per diagnosi e personalizzazione delle malattie?

DEFINIZIONE DEI CRITERI DI FISILOGIA/NORMALITA'

- Biobanca di soggetti italiani
- Confronto con bio-banche europee
- es. UK BioBank 500.000 soggetti

Schroeder and Backhed Nature Medicine 2016



Università di Roma Tor Vergata

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Nikolaus Marx

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Jurgen Adamski



EURHYTHDIA



dedicated to finding a cure

