MICROBIOM

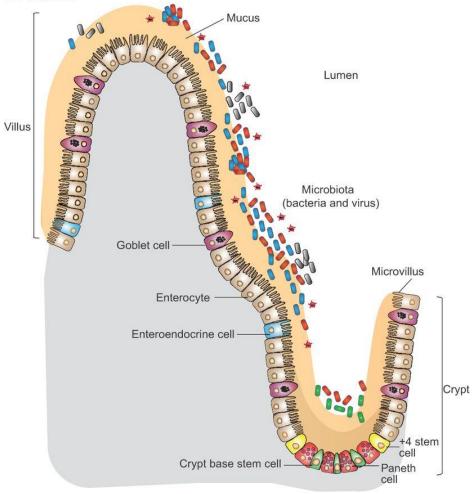
AND OBESITY

HEINZ GYAKY ® 2018 BUDAPEST

HUMAN MICROBIOM

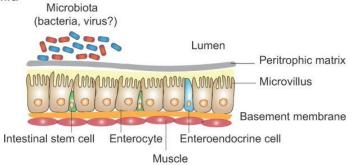
- 10 Billion bacterias are building a 1,5 2 kg heavy human microbiom
- It is located mainly in the human gut
- There is a intestinal controlled flora everywhere in our body
- Skin, mouth, stomach, blood, bladder, umbilical cord and placenta, vagina, lungs, ...
- Human microbiom is like a fingerprint, special and unique
- The bacterias have a system of communication and a controlled growing

A Human

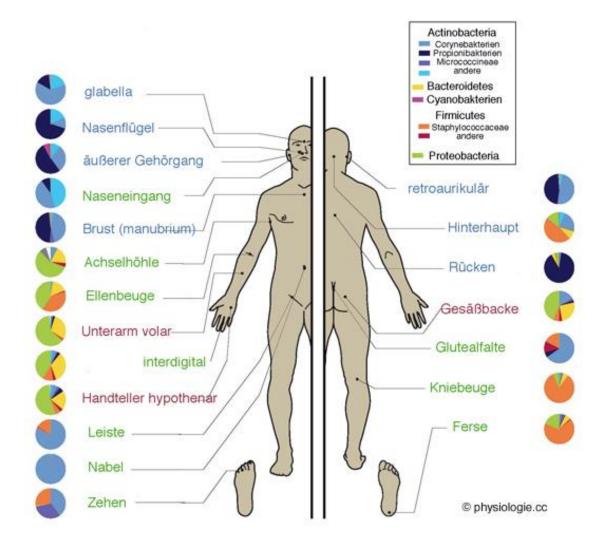


MICROBIOM AND GUT

B Drosophila



HUMAN MICROBIOM



TASKS OF OUR MICROBIOM

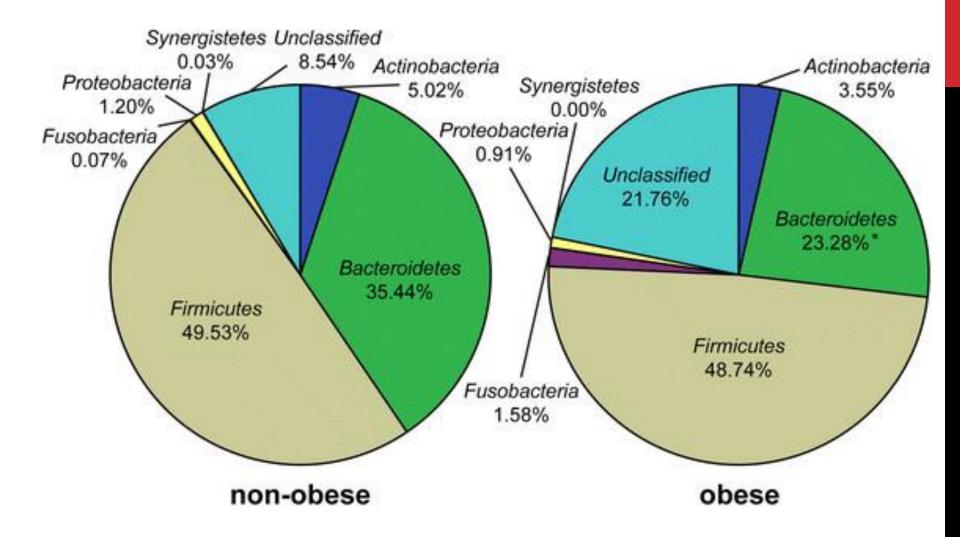
- Controlls 70-80 % of the immunity
- Controlls 100% of our contact with the environment
- Involved in our digestion
- Produces vitamines, hormons and enzyms
- Comunication with our brain, liver, adrenal gland
- The Comunication is always bidirectional

MICROBIOTA AND THE RELATIONSHIP TO DIET UND OBESITY

- The bacterial composition of lean and obese humans
- The impact of diet on the gut microbiota
- Can we manipulate the gut microbiota
- Mechanisms by which the gut microbiota can impact on weight gain

THE COMPOSITION OF LEAN AND OBESE HUMANS

- We are watching two groups of bacterias: Phylum bacteriodetes und Phylum firmicutes
- Obese humans have an increase of Firmicutes and the Firmicutes/bacteriodetes-ratio changes
- High Firmicutes populationes can be found in all obese humans and in their family members
- High "western" diet and maybe the microbiom transmission in the family are causes



THE IMPACT OF DIET ON THE GUT MICROBIOTA

The population of Bacteriodetes is increasing during a diet with fat restriction or carbohydrate restriction or calorie restriction and exercise

High dietary fiber causes an increase of Faecalbacterium prausnitzii

Low F. prausnitzii populations are seen in gut diseases with high inflamatory (Crohn, ...)

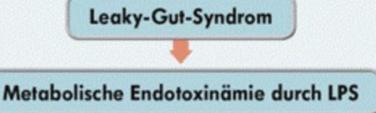
F. prausnitzii is responsible for the production of propionic acid and butyric acid, the main energy supplier of the tight junctions

MECHANISMS BY WHICH THE GUT MICROBIOTA CAN IMPACT ON WEIGHT GAIN

- Firmicutes increase the calorie intake by 20% and allowe the fermentation of fiber into glykogen and not into short chain fatty acides
- The absence of F. prausnitzii and the higher Firmicutes population causes a inflammatory in the gut
- The intestinal barrier gets leaky, endotoxin, a part of the gramnegativ bacterias, overload liver and the fatty tissue. A silent inflammatory starts.
- We measure high zonulin in faeces and high LPS endotoxine (lipopolysaccaride) in Blood as a marker for silent inflamation.
- Obesity is also associated with a decrease in the overall microbial diversity of the gut, although the total amount of microbes remains the same

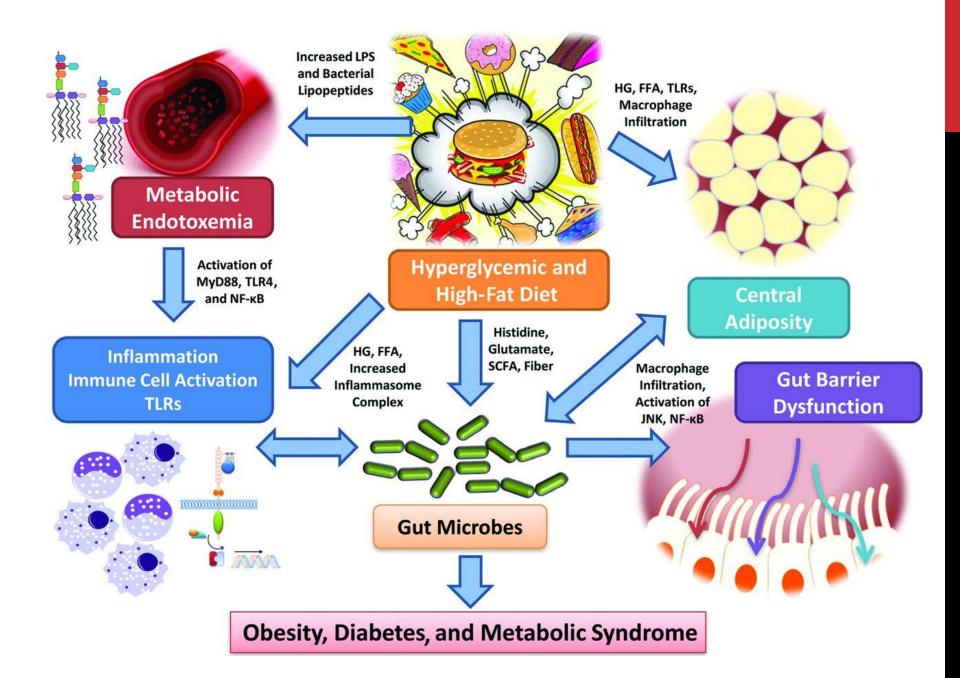
MECHANISMS BY WHICH THE GUT MICROBIOTA CAN IMPACT ON WEIGHT GAIN

Störungen der saccharolytischen und mukonutritiven Flora – Wegbereiter für das metabolische Syndrom Propion-/Buttersäure erniedrigt



Silent Inflammation

Metabolisches Syndrom und Diabetes mellitus Typ 2



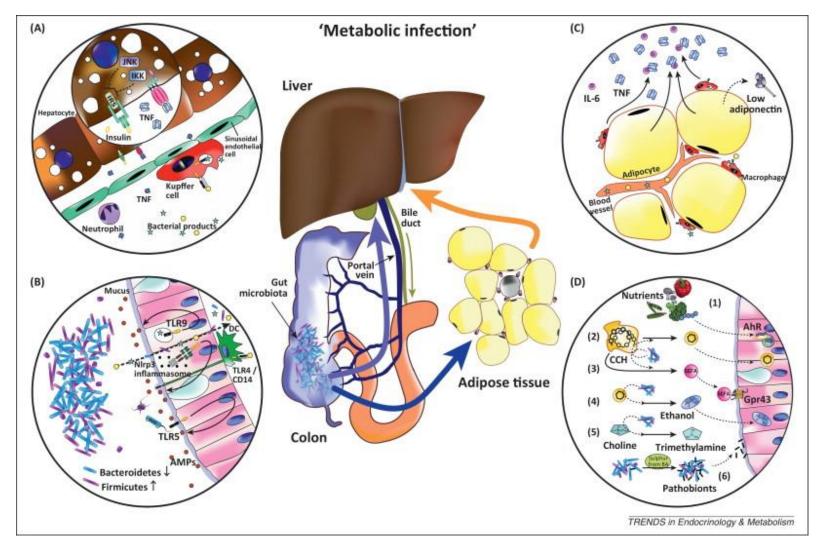
OVERVIEW ABOUT STUDIES AND MECHANISMS

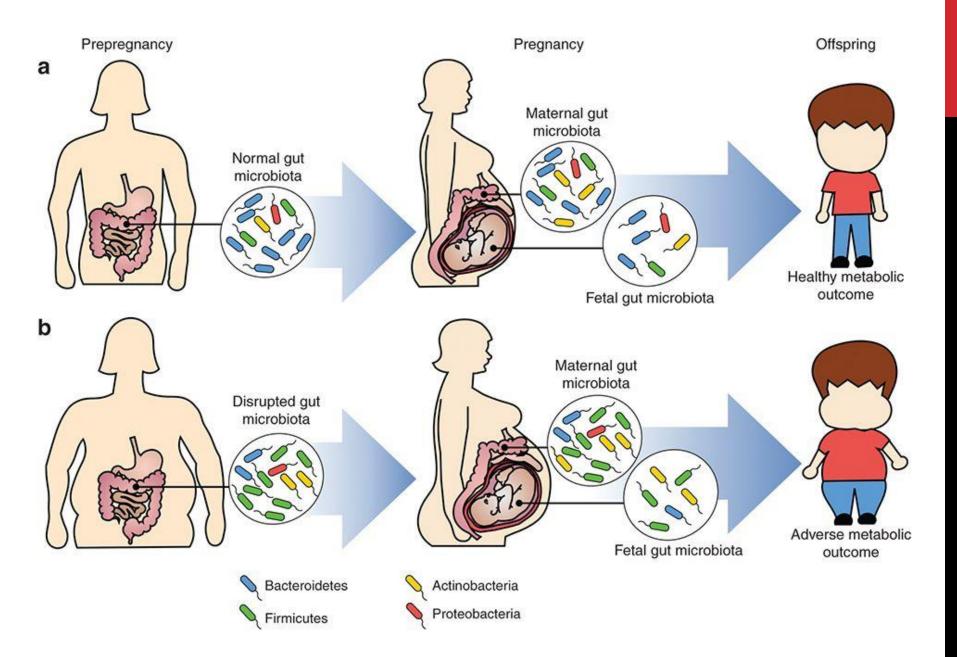
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Gut Microbes 3:3, 186-202; May/June 2012; © 2012 Landes Bioscience			
The gut microbiota and its relationship to diet and obesity New insights			
			eera, ¹ Paul R. Ross, ^{1,3} Fergus Shanahan, ^{3,5} Paul W. O'Toole ^{2,3} I D. Cotter ^{1,3,*}
			/ Department; University College Cork; Cork, Ireland; ³ Alimentary Pharmabiotic Centre; , Ireland; ⁵ Department of Medicine; University College Cork; Cork, Ireland
Keywords: gut microbiota, intervent	ion, prebiotic, probiotic, diet and obesity		
Obesity develops from a prolonged imbalance of energy	Gut Microbiota of Lean and Obese Animals		
intake and energy expenditure. However, the relatively recent discovery that the composition and function of the gut microbiota impacts on obesity has lead to an explosion of interest in what is now a distinct research field. Here, research	Mouse models are frequently employed by researchers investigating obesity and the role of the gut microbiota in obesity. The for		
relating to the links between the gut microbiota, diet and	lowing sub-sections will focus on the various mouse models th have been employed and the outcome from studies performed		

NON-ALCOHOLIC STEATOHEPATITIS: A MICROBIOTA-DRIVEN DISEASE

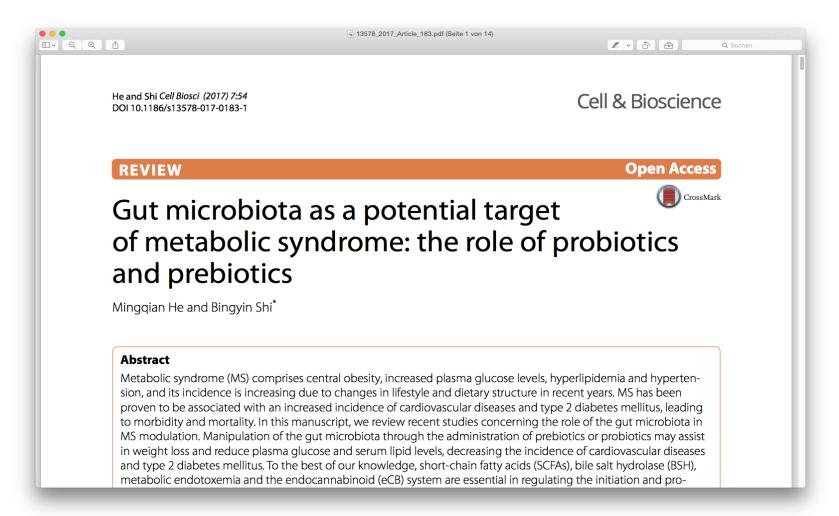
- Whereas overnutrition and obesity are crucially involved in the development of a simple fatty liver, it remains unclear why approximately 10% of all affected individuals develop the 'inflammatory' phenotype so-called nonalcoholic steatohepatitis (NASH).
- A link between the intestinal microbiota and the development of obesity and its metabolic consequences including NAFLD is becoming clearer.
- First clinical studies are suggesting that microbiotal factors are driving forces of hepatic steatosis and inflammation that involve Toll-like receptors and proinflammatory cytokines such as tumor necrosis factorα (TNFα).

NON-ALCOHOLIC STEATOHEPATITIS: A MICROBIOTA-DRIVEN DISEASE





CONCLUSION



CONCLUSION

Accumulating evidence suggests that gut microbiota plays a significant role in the initiation and progression of MS. The gut microbiota was proven to modulate plasma glucose, appetite, serum lipids and pro-inflammation.

In addition, prebiotics or probiotics, which are widely used to manipulate the microbiota, can reduce low-grade intestinal inflammation and improve gut barrier integrity to reduce plasma glucose and serum lipid levels, induce weight loss and decrease insulin resistance. Based on these current achievements, the gut microbiota may be a potential therapeutic target for MS.

However, clinical trials addressing the efficacy and efficiency of current or potential treatments on therapeutic applications in metabolic syndrome are needed.

TAKE HOME

- Reduce alcohol, carbohydrates and fat in your food !
- Raise salat, vegetables und resistant starch.
- Watch antibiotics, PPIs and heavy metals.
- Probiotics are usefull in the treatment of obesity, diabetes, irritable bowel syndrome, crohn disease, ...
- The treatment with probiotics lasts at least 6 months





TAKE HOME



Thank you for your attention