

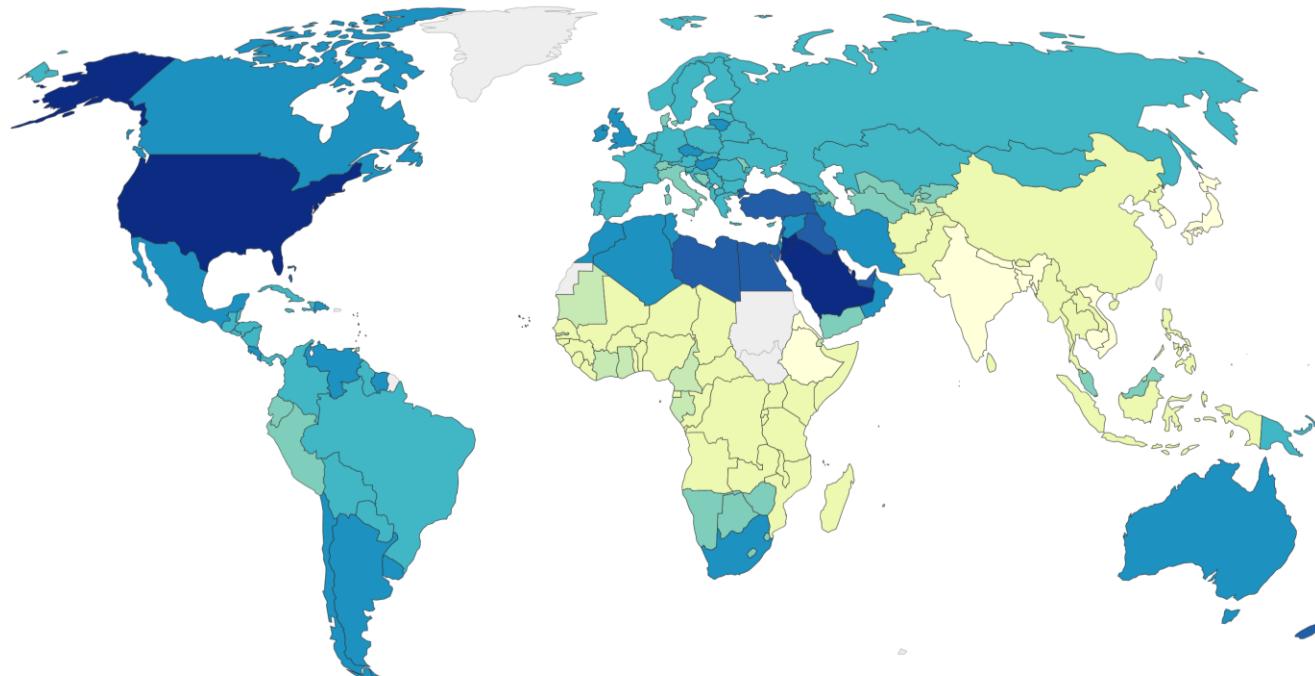
Obesitas, microbiota en inflammatie

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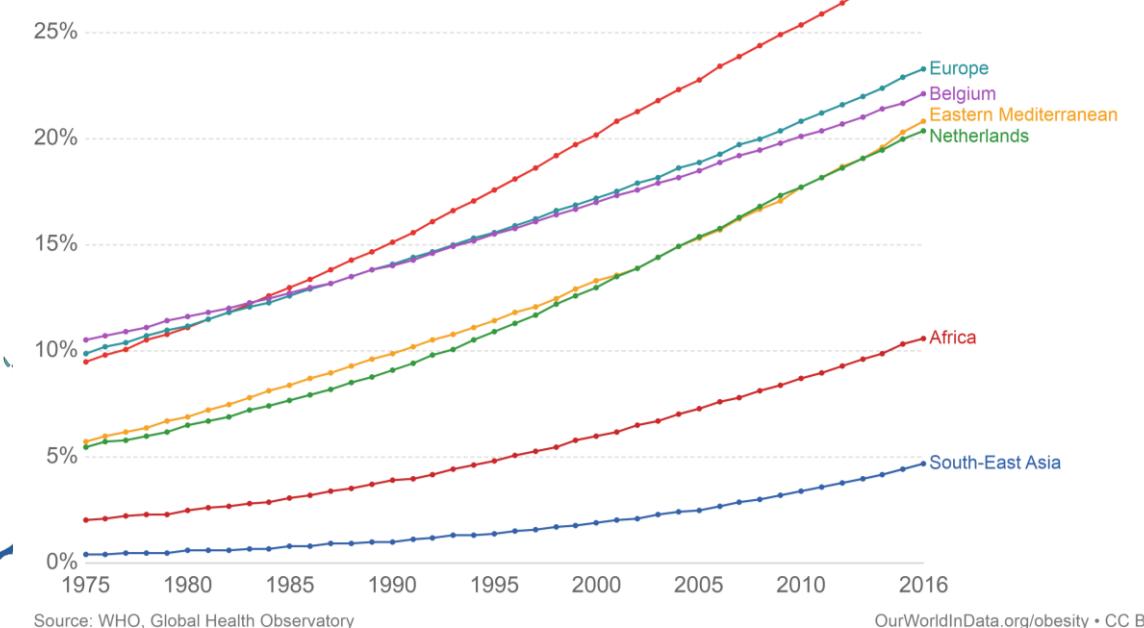


The obesity epidemic



Source: WHO, Global Health Observatory

OurWorldInData.org/obesity • CC BY



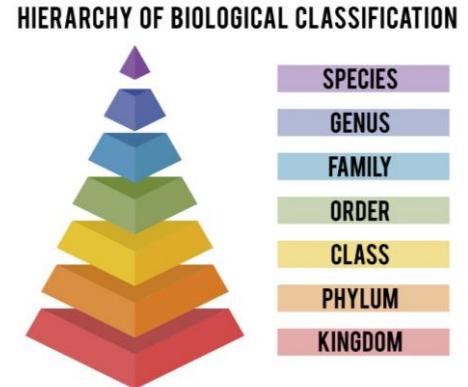
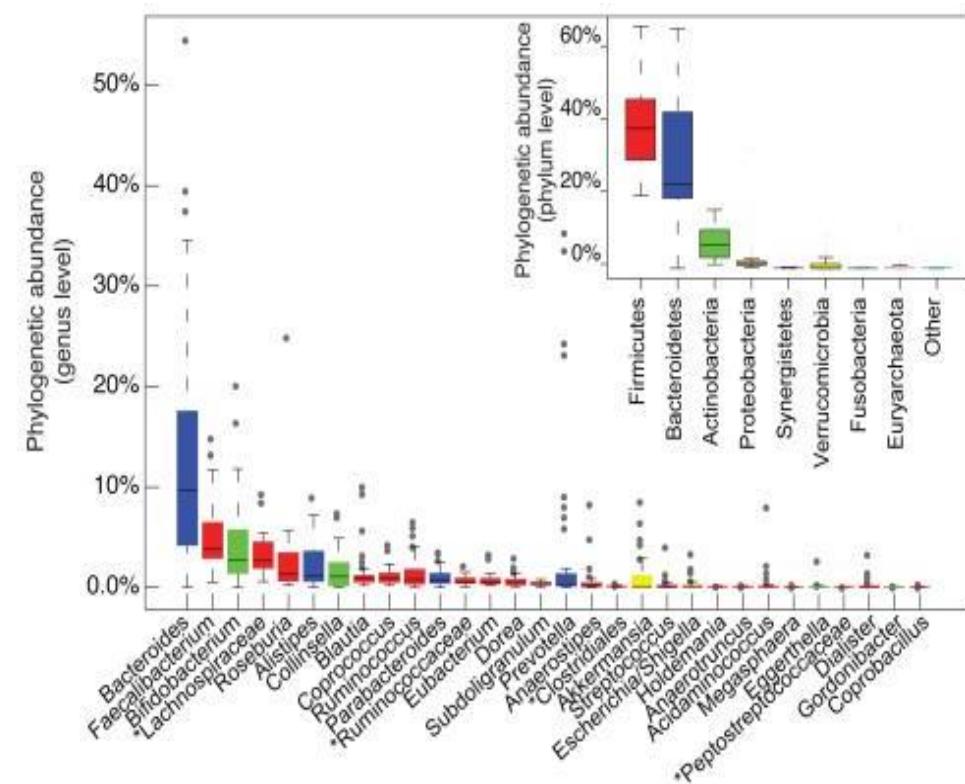
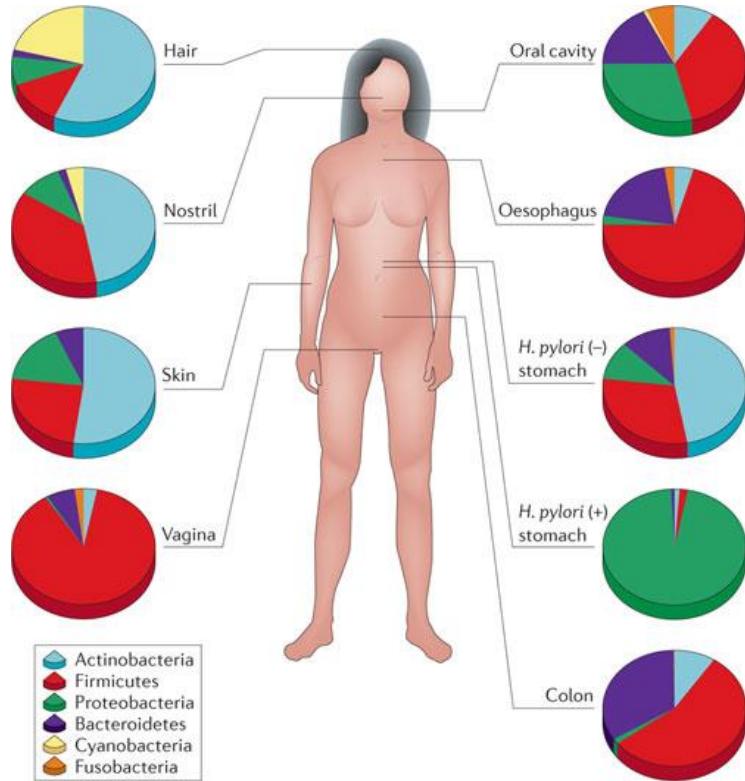
Source: WHO, Global Health Observatory

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Oorzaken van obesitas

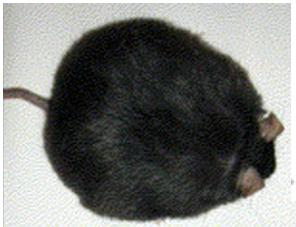


De mens is een superorganisme

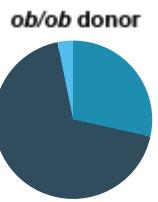


Obese muizen

obese donor (*ob/ob*)



ob/ob donor



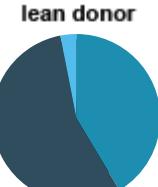
ob/ob recipients



lean donor



lean donor



lean recipients



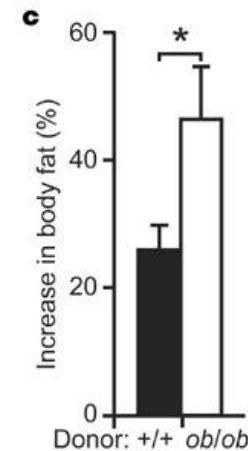
lean recipients

Firmicutes

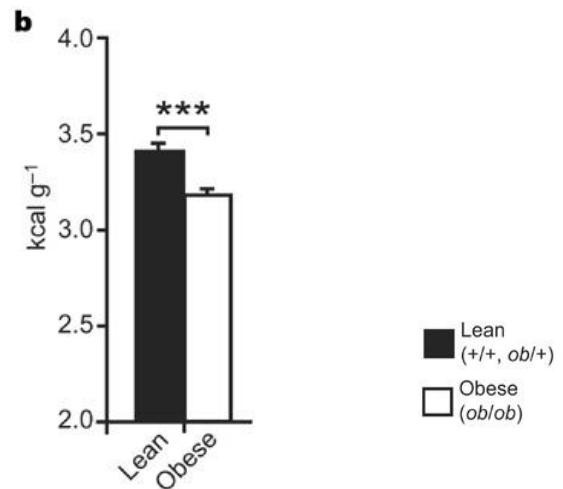
Bacteroidetes

Other

Increase in body weight



Energy in feces

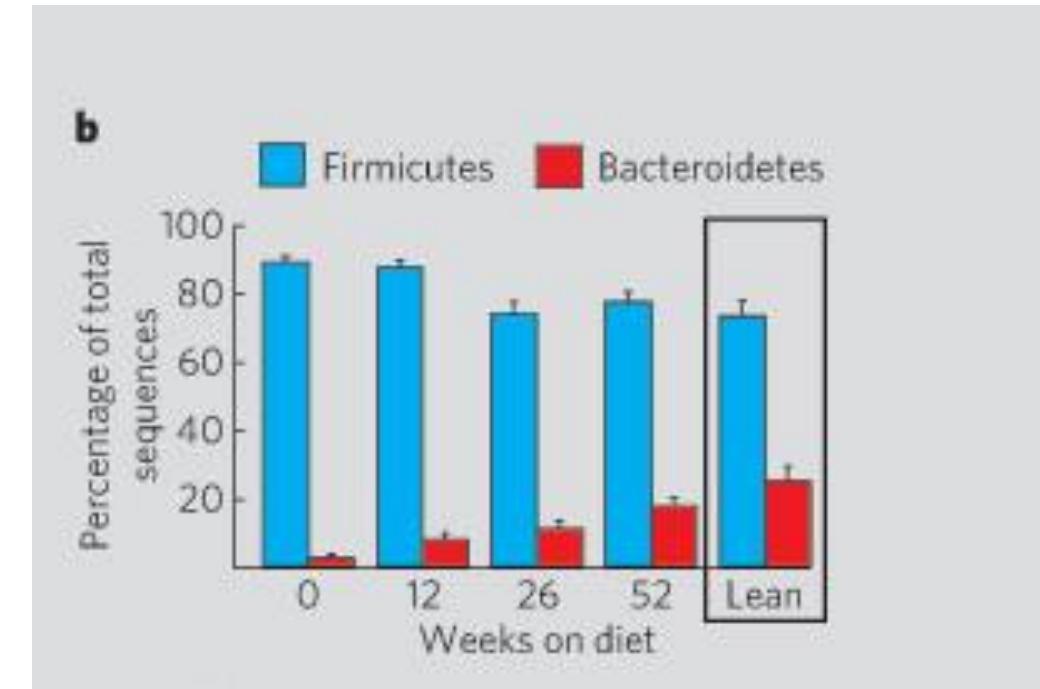


obese microbiome possesses metabolic pathways that are highly efficient at extracting energy from food

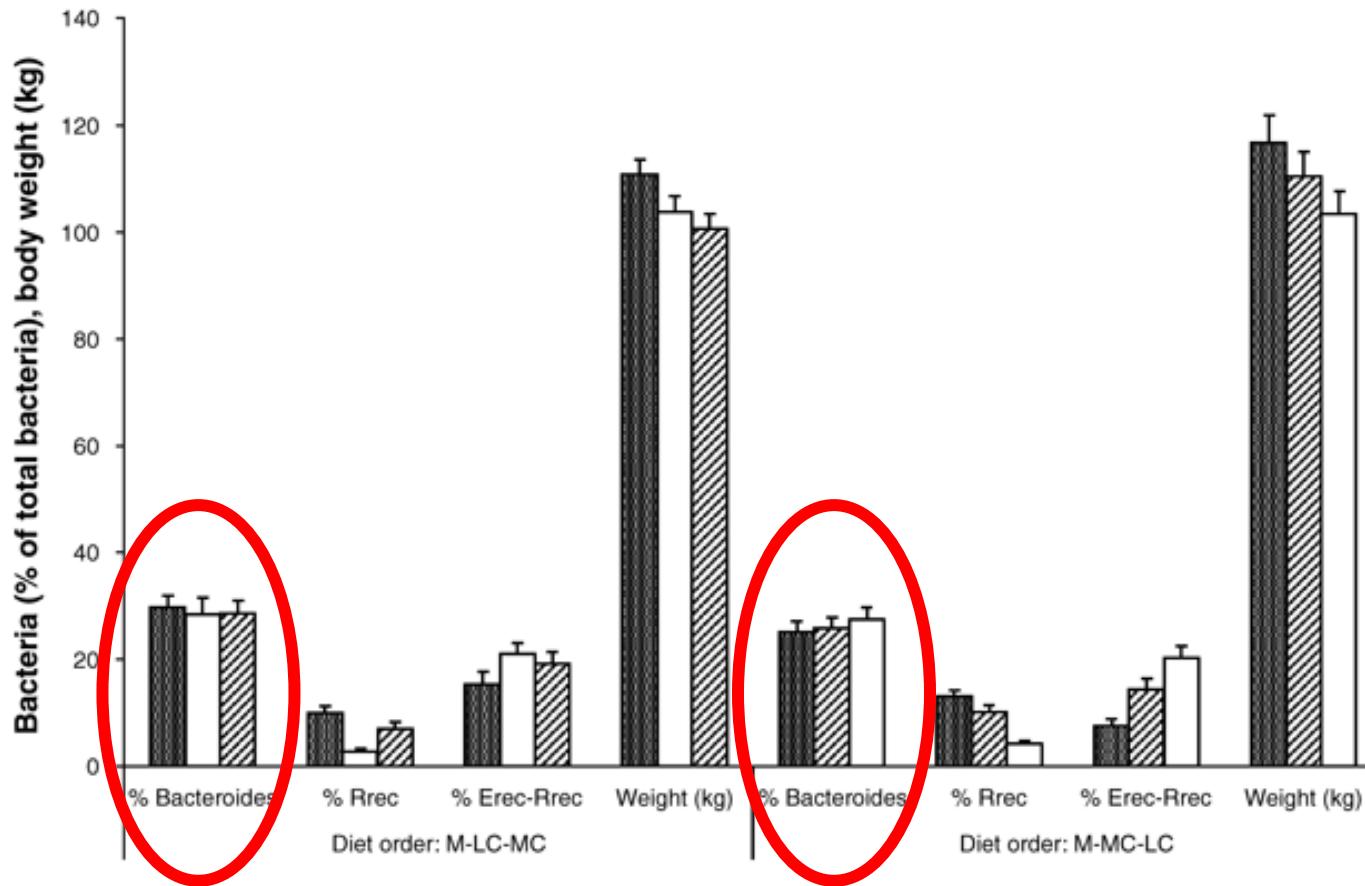
Microbiota samenstelling in obese personen



12 obese personen
↓
Laag calorie dieet



Others do not confirm



No effect on % Bacteroides
after weight loss

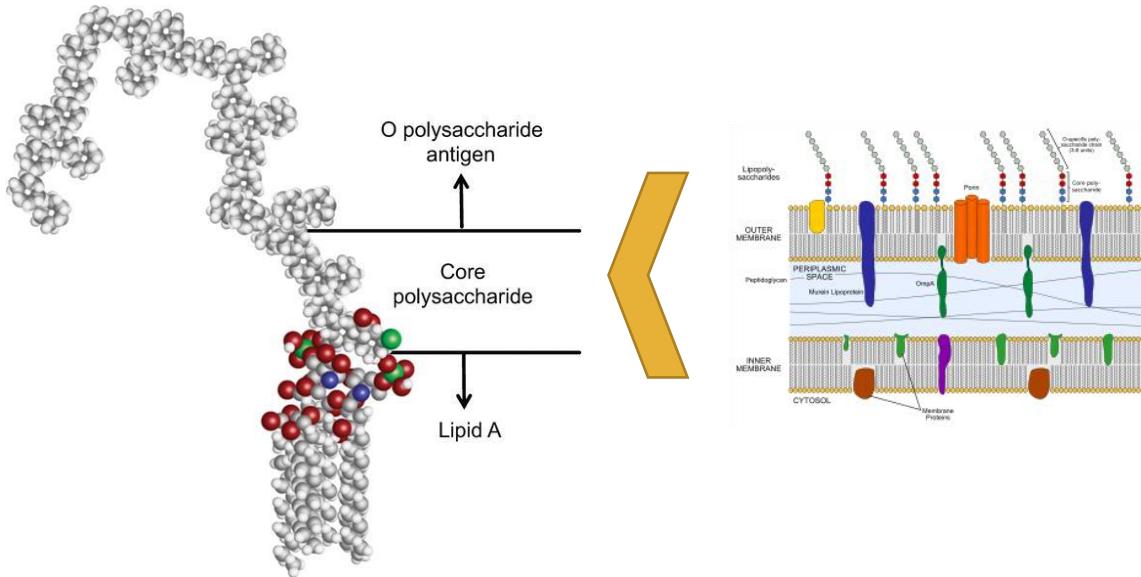
Gut microbiota and obesity

- Phylumwide changes currently not considered as biomarker for obesity
- Changes at lower taxonomic level may better define dysbiosis
- Several genera increased or decreased in obesity – species may follow opposite trend
 - Lactobacilli ↗
 - Bifidobacteria ↘
 - *Faecalibacterium prausnitzii* ↘
 - *Akkermansia muciniphila* ↗
- Mainly association studies -> cause or consequence?

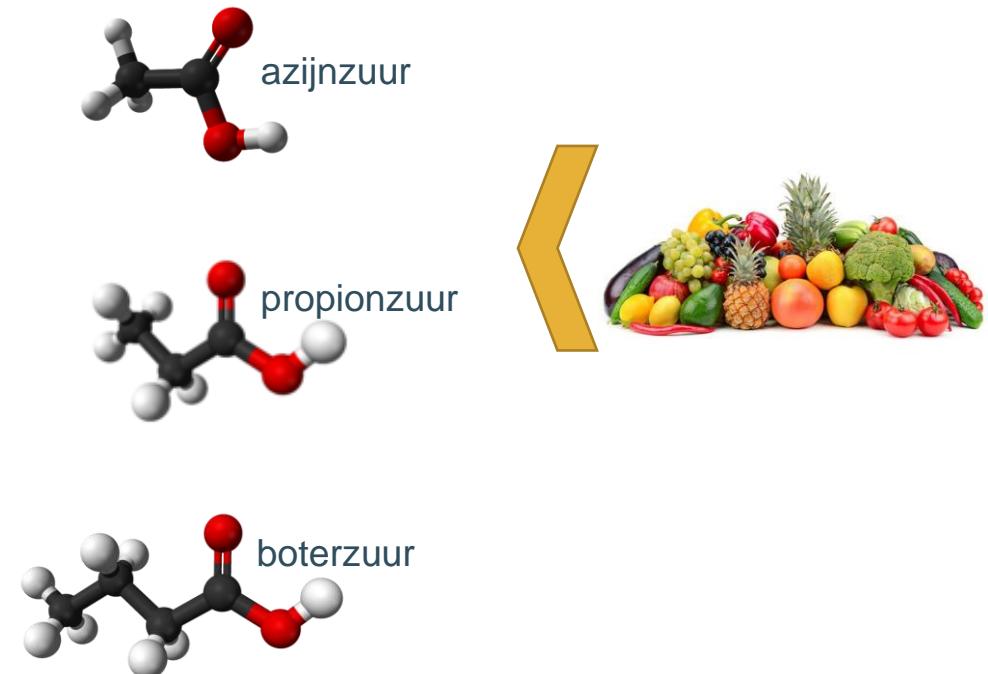
It is still impossible to define an obese microbiome

Microbial signals linked to obesity

Lipopolysaccharide (LPS)

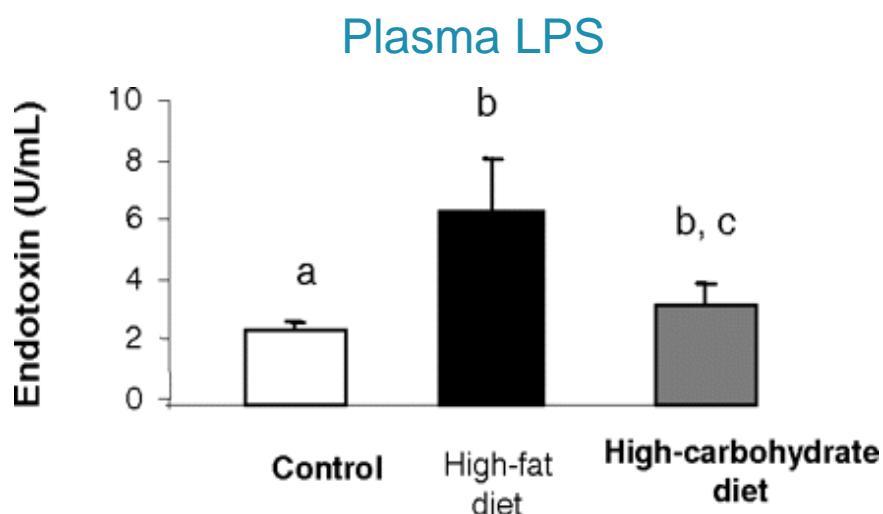


SCFA

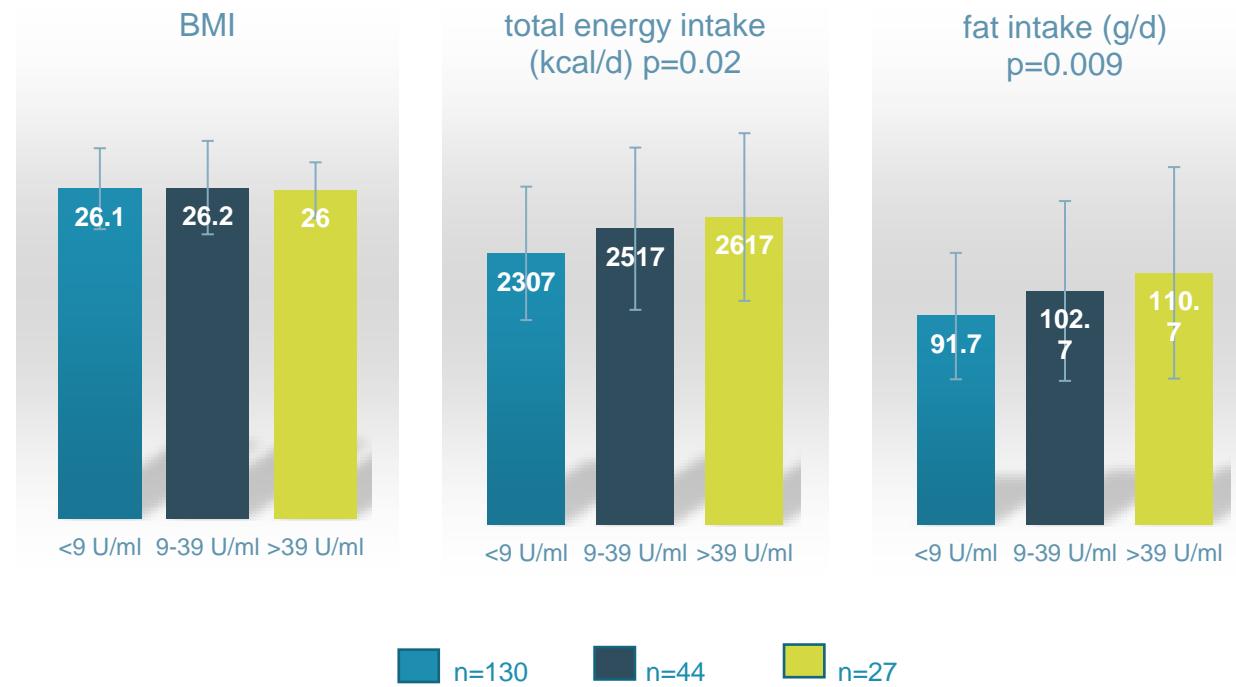


High fat intake increases plasma LPS

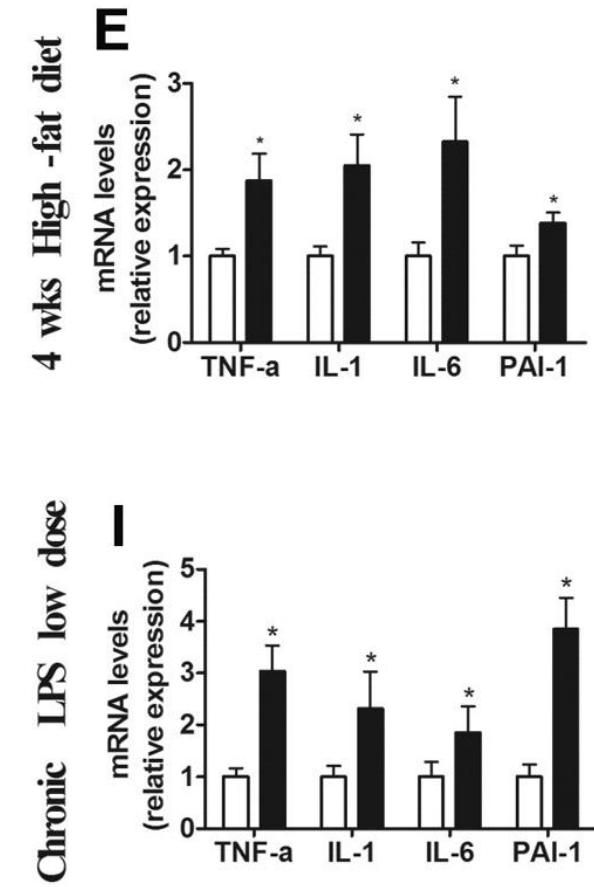
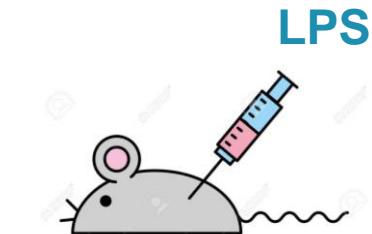
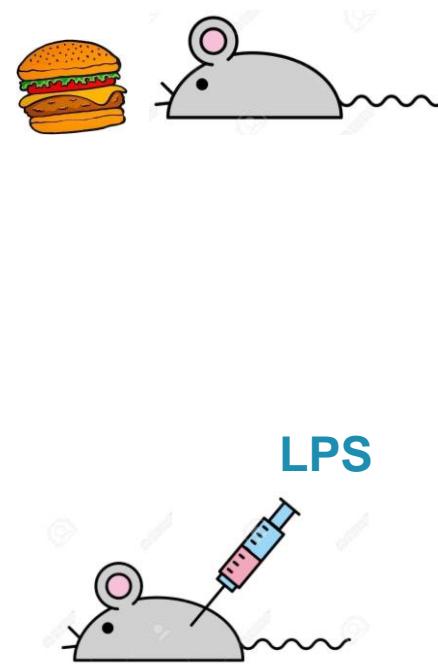
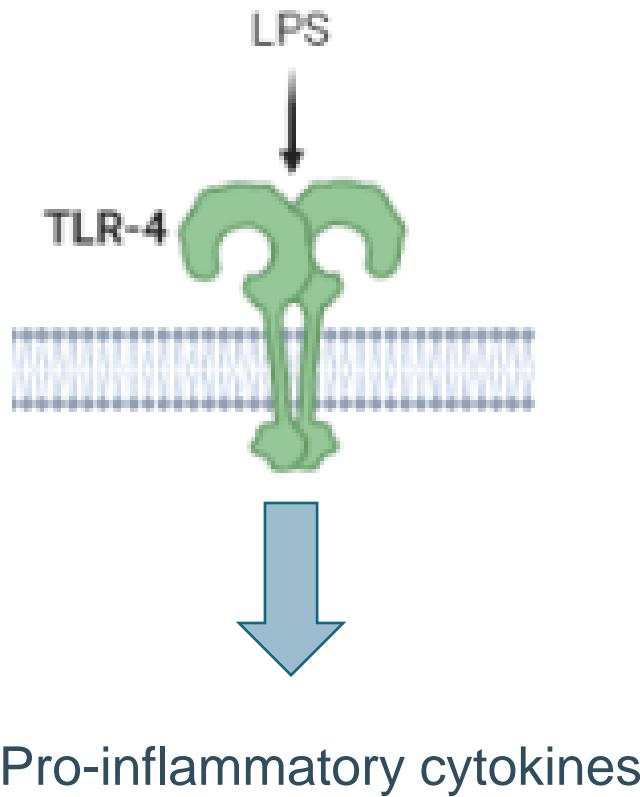
Mice
Intervention for 4 wks



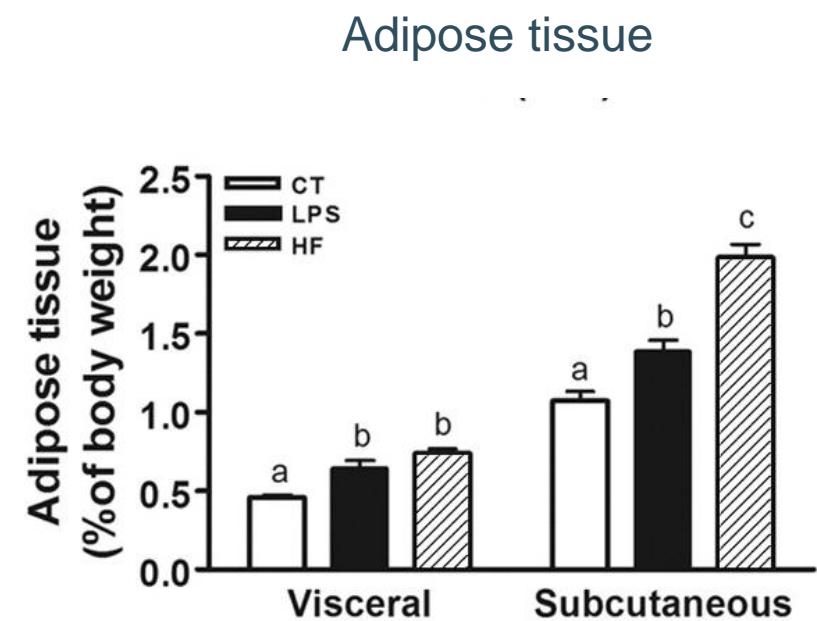
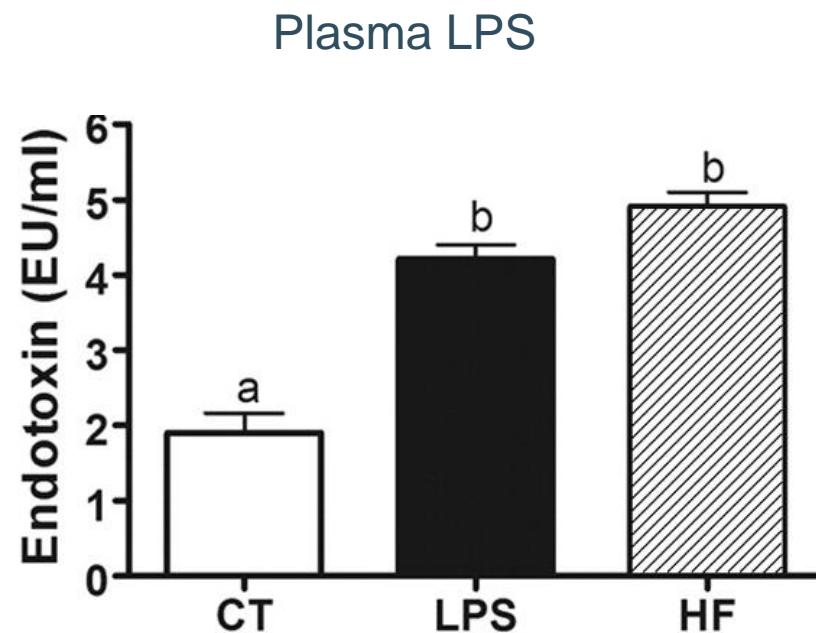
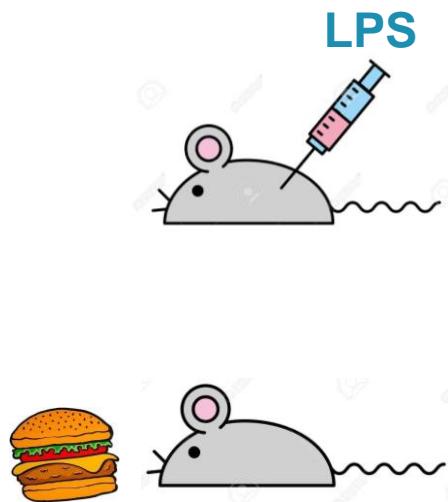
Humans
Cross-sectional analysis, n=201



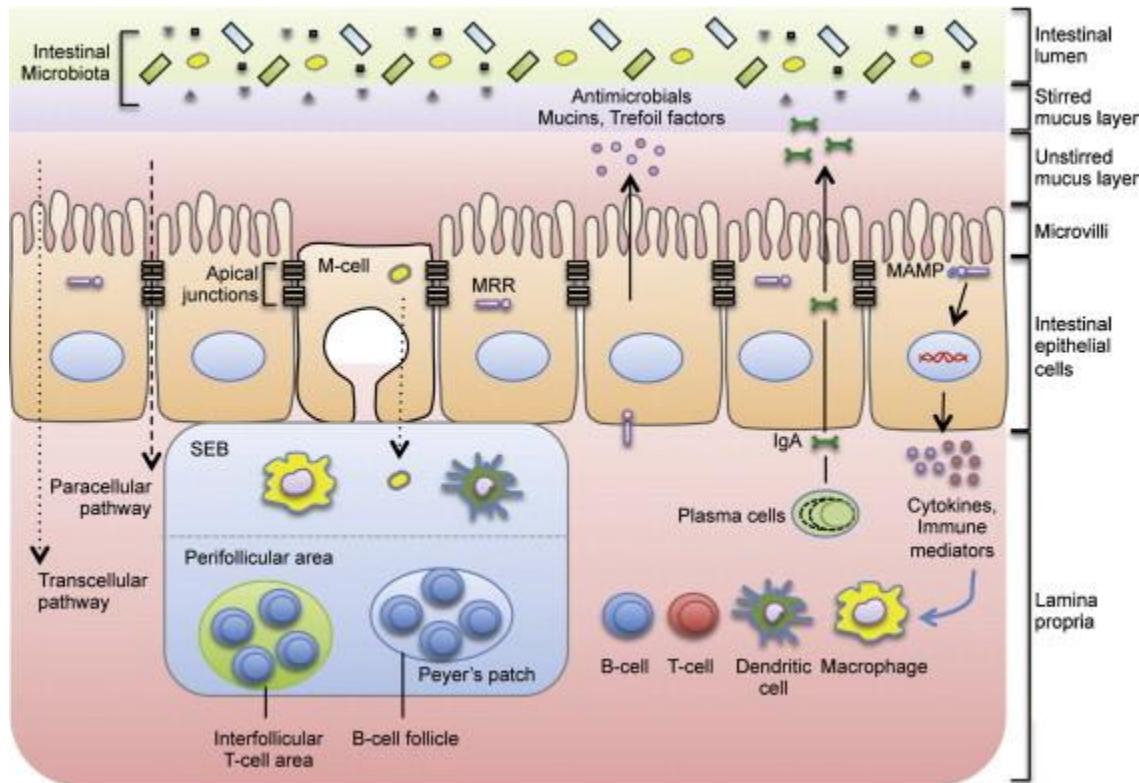
Increased plasma LPS induces inflammation



Increased plasma LPS induces obesity

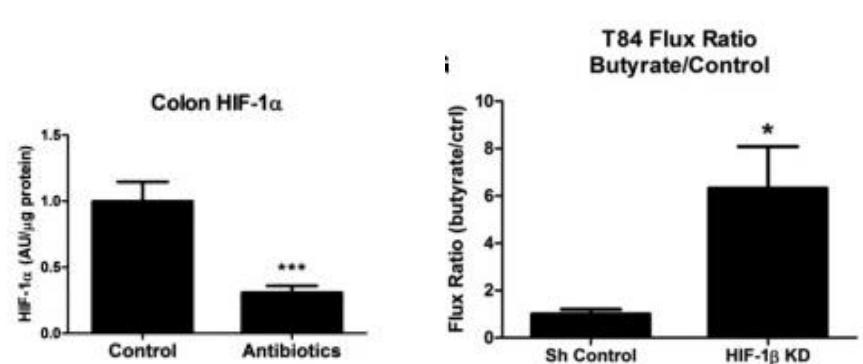
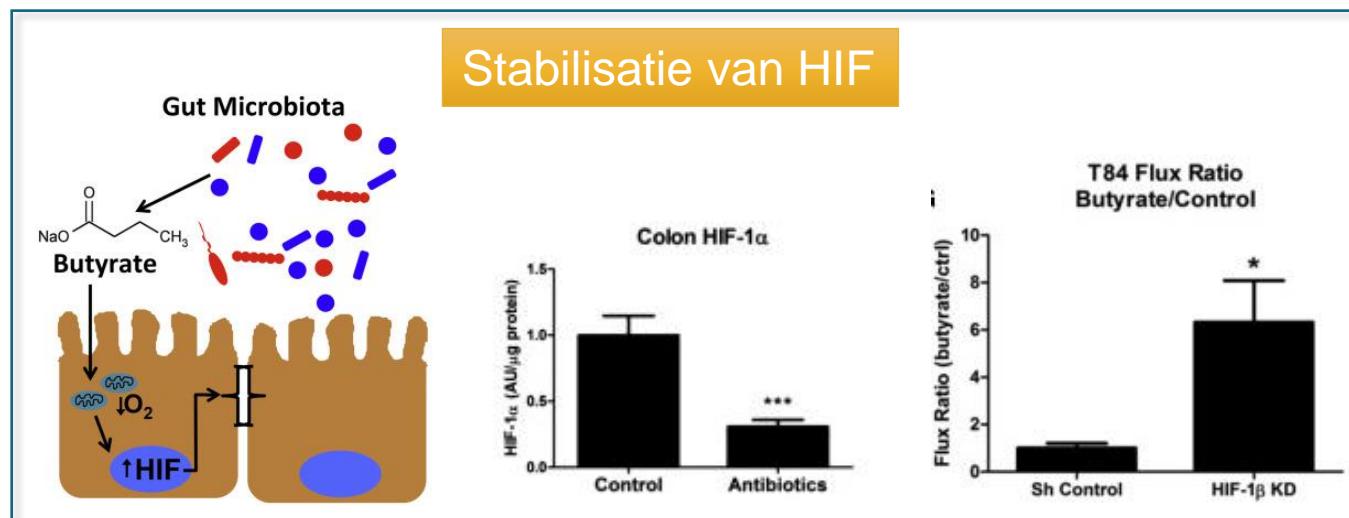
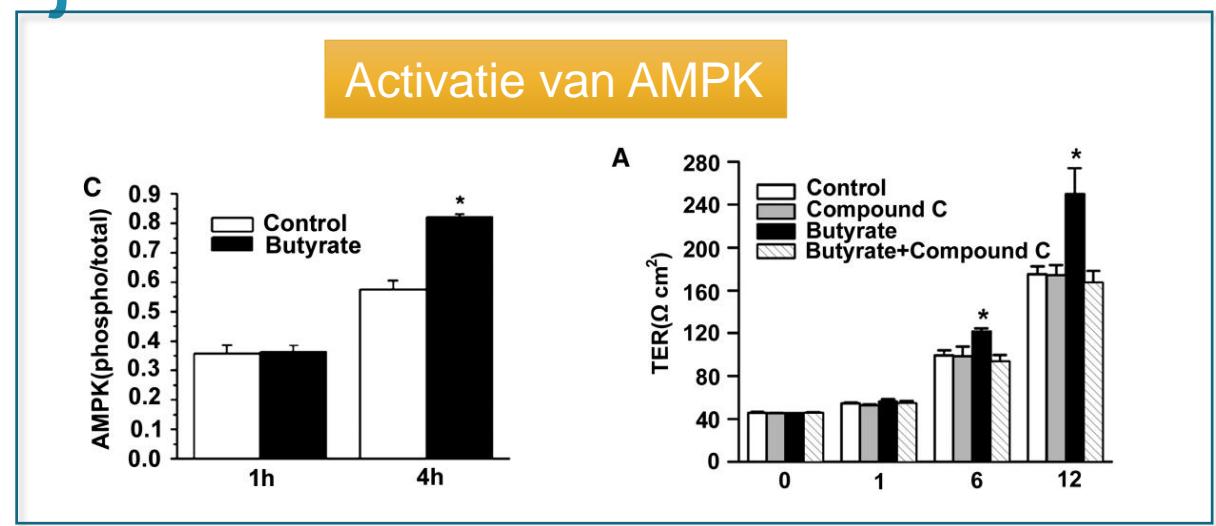
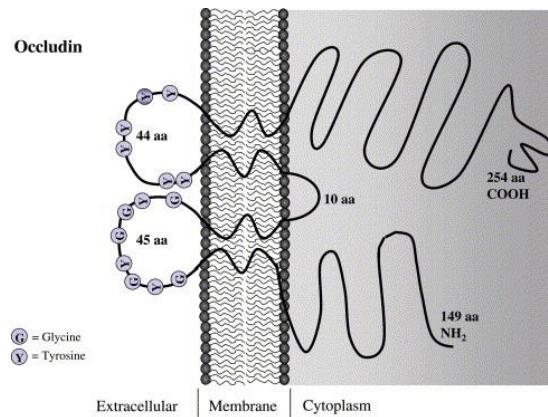
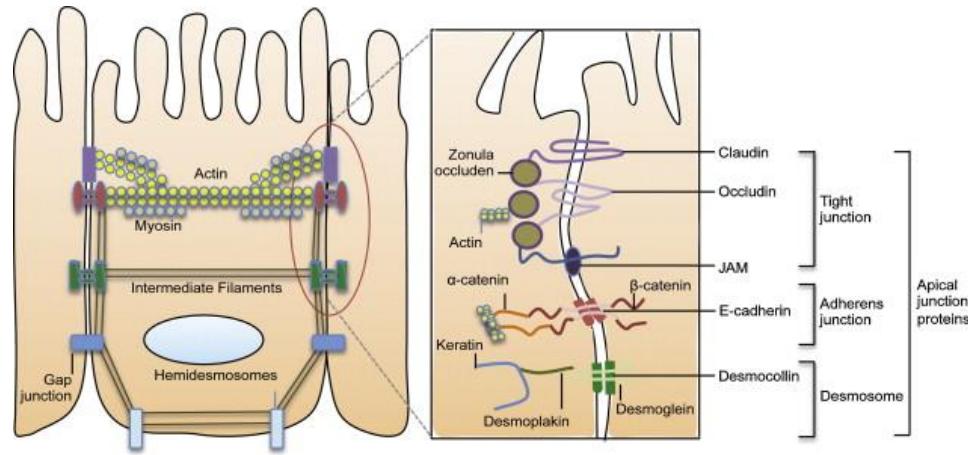


SCFA protect against inflammation

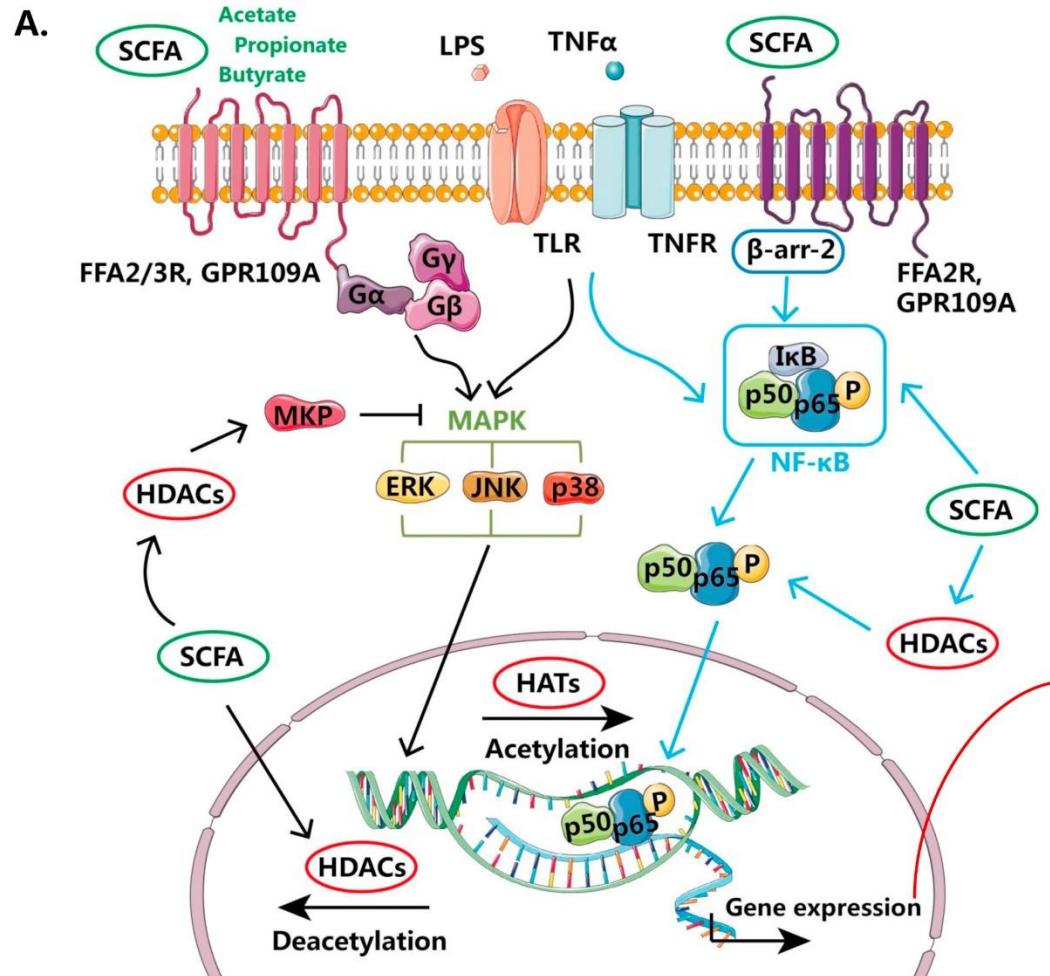


- Epithelial cells
- Mucus
- Immune system
- antimicrobials

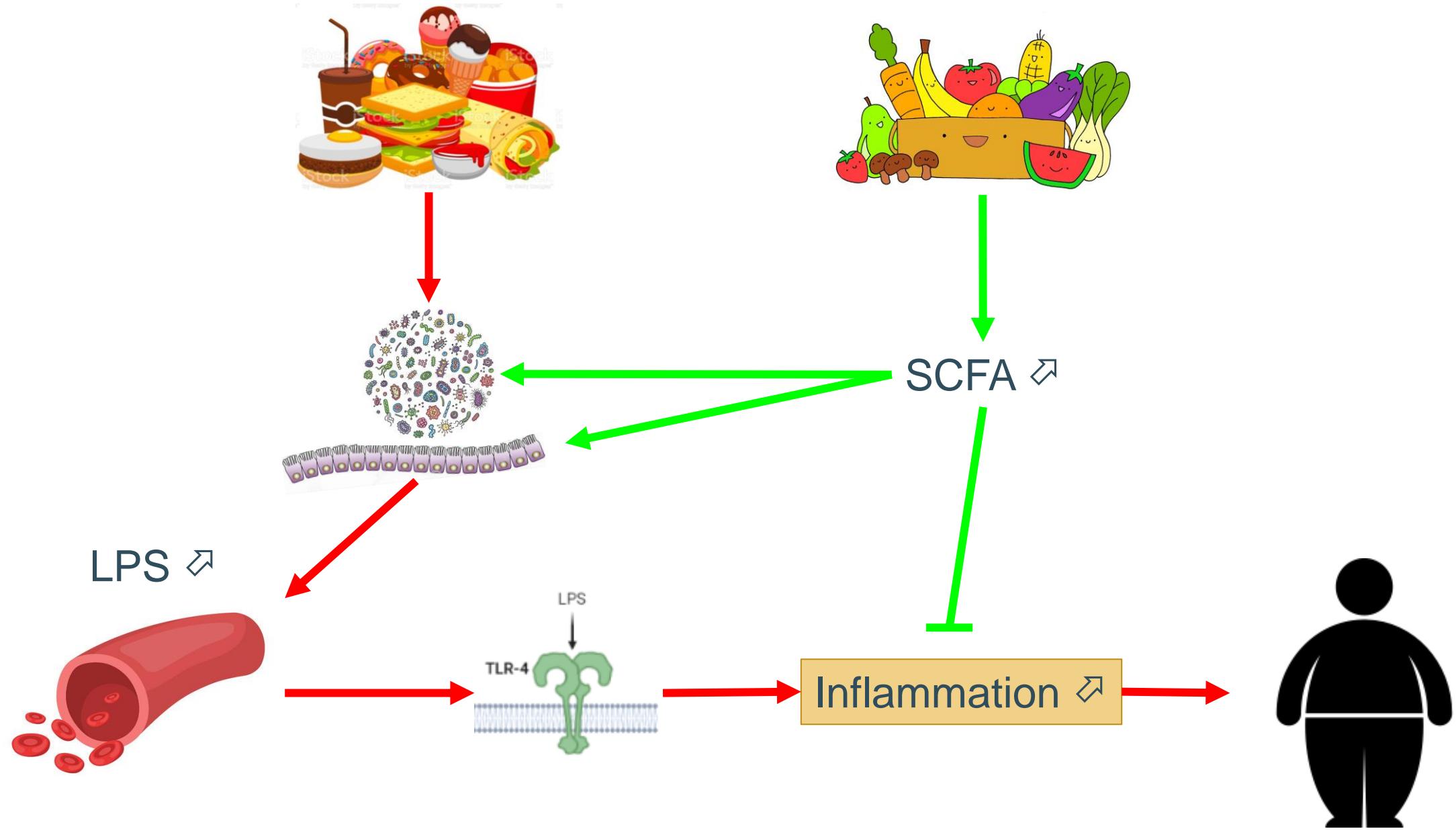
SCFA versterken de tight junctions



SCFA interfereren met het immuunsysteem

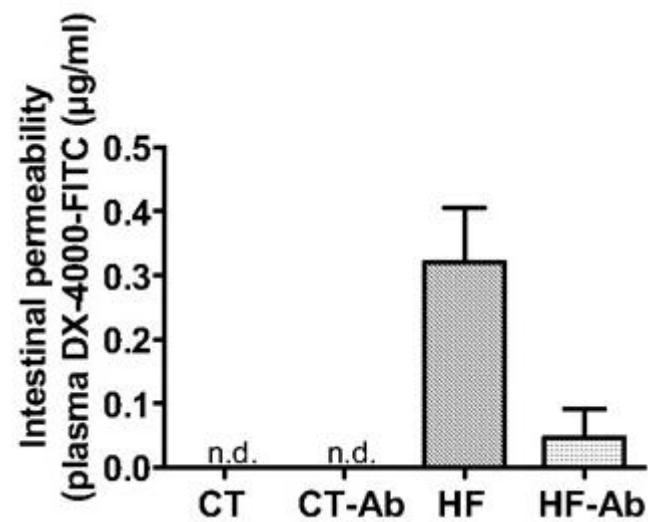


- Stimulatie van MAPK
- Inhibitie van NF- κ B
- Inhibitie van HDACs

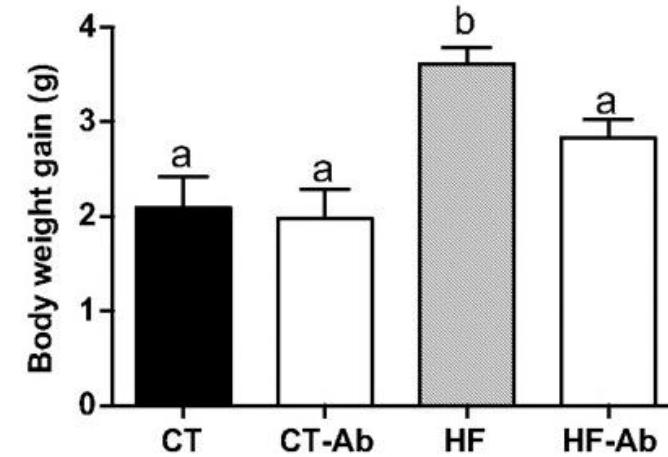


Obese mice have increased gut permeability

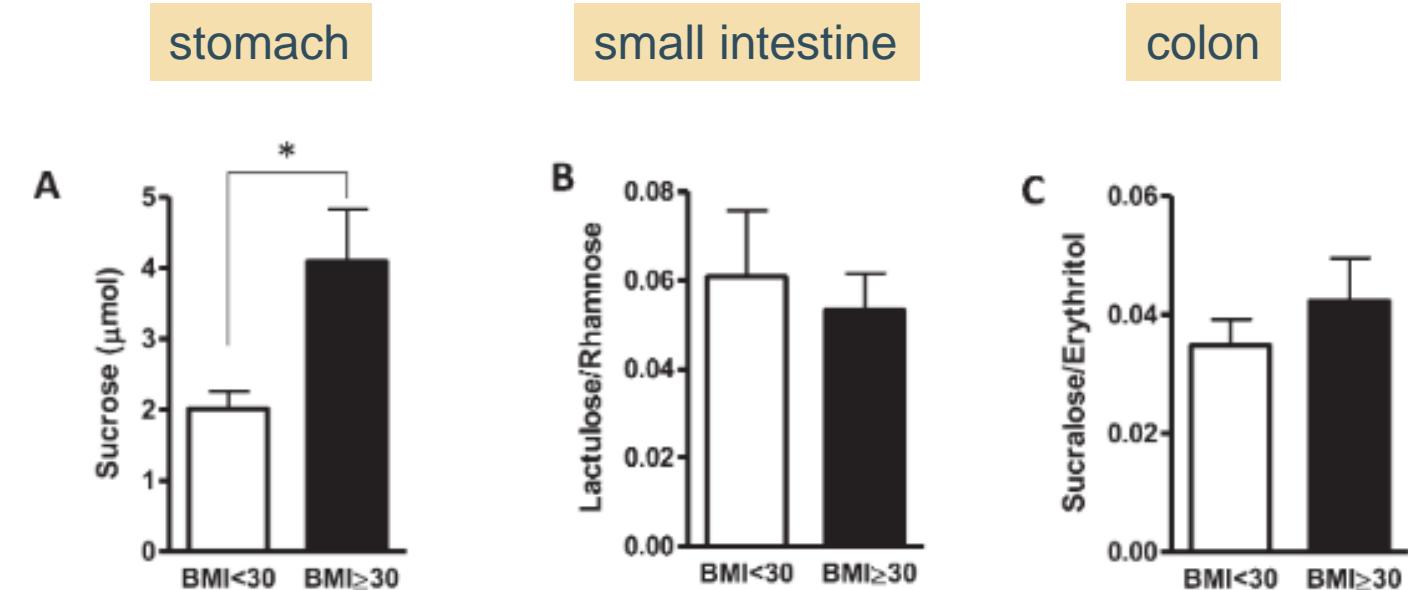
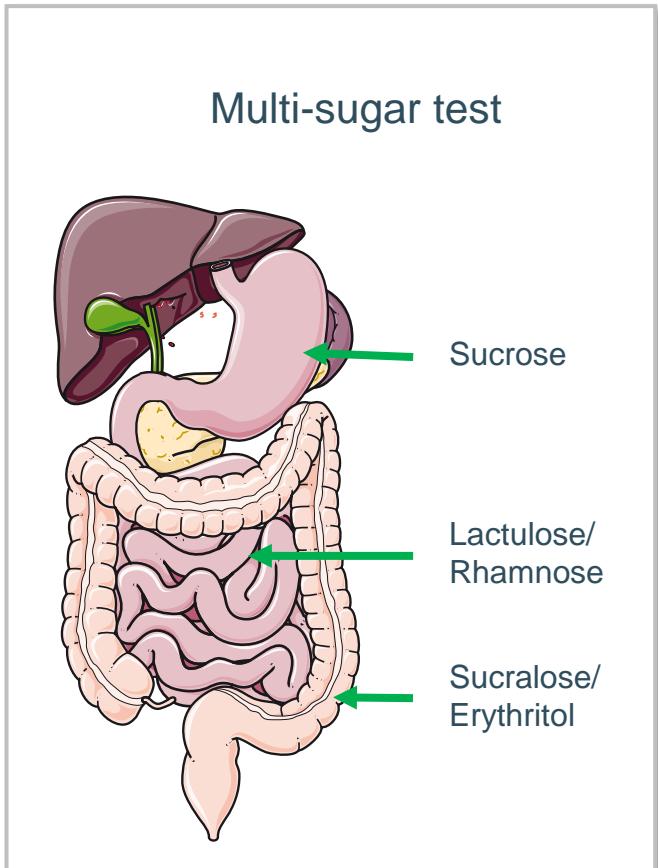
In vivo permeability



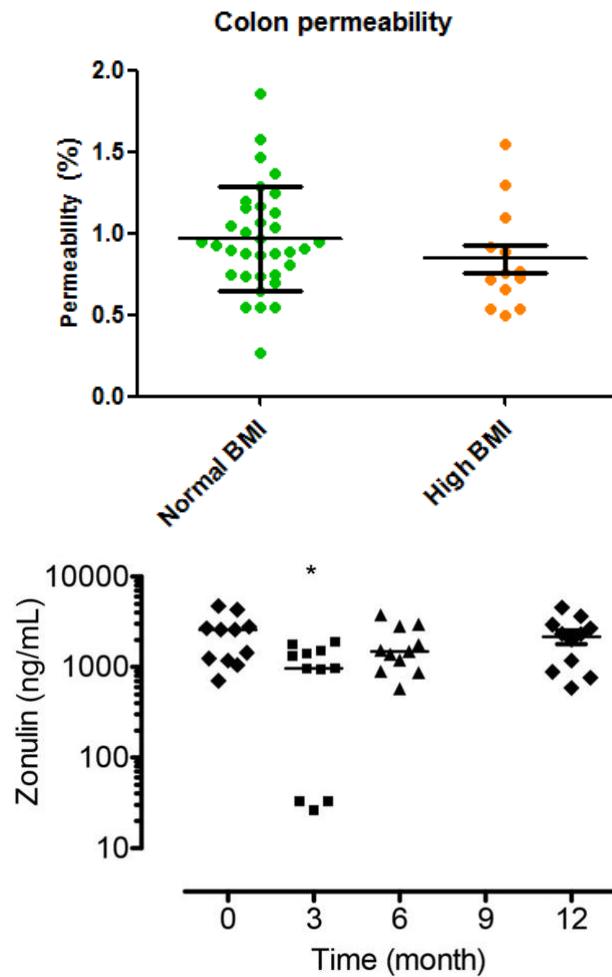
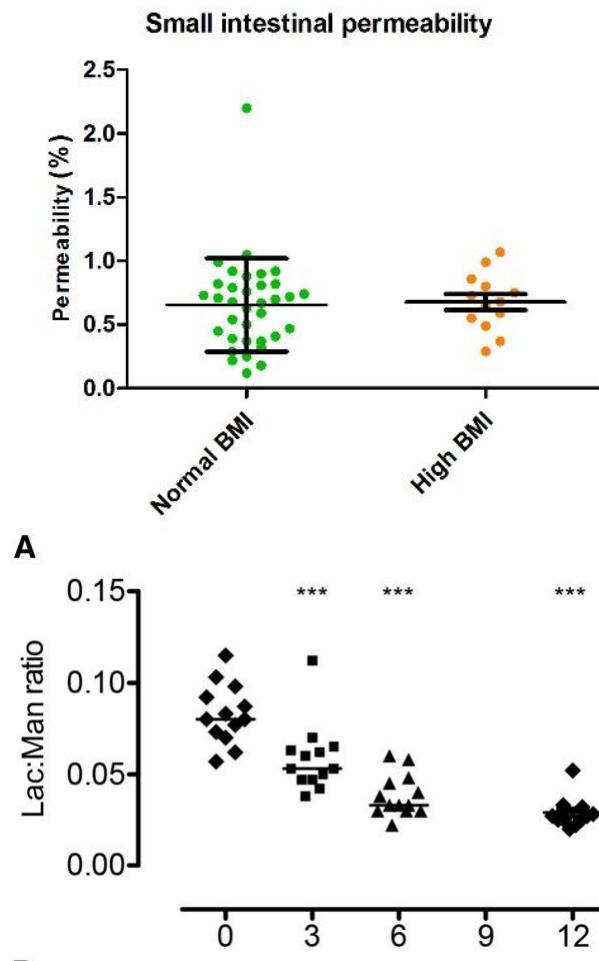
Body weight



Gut permeability in humans



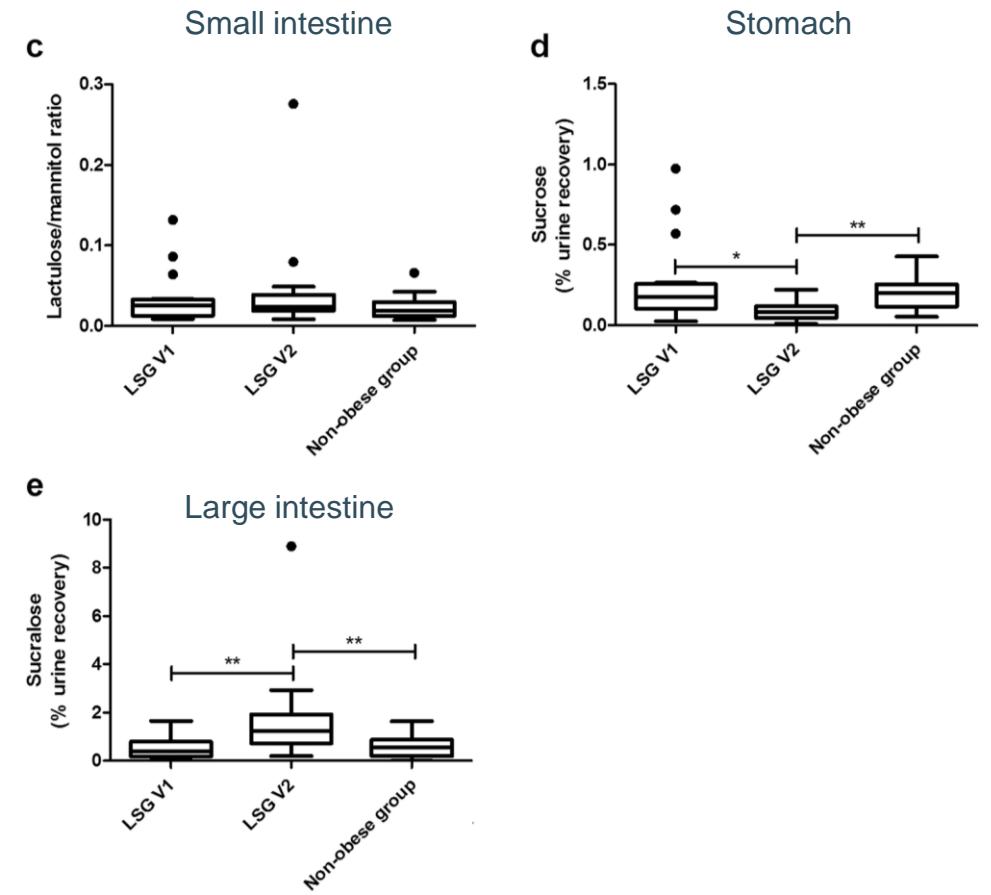
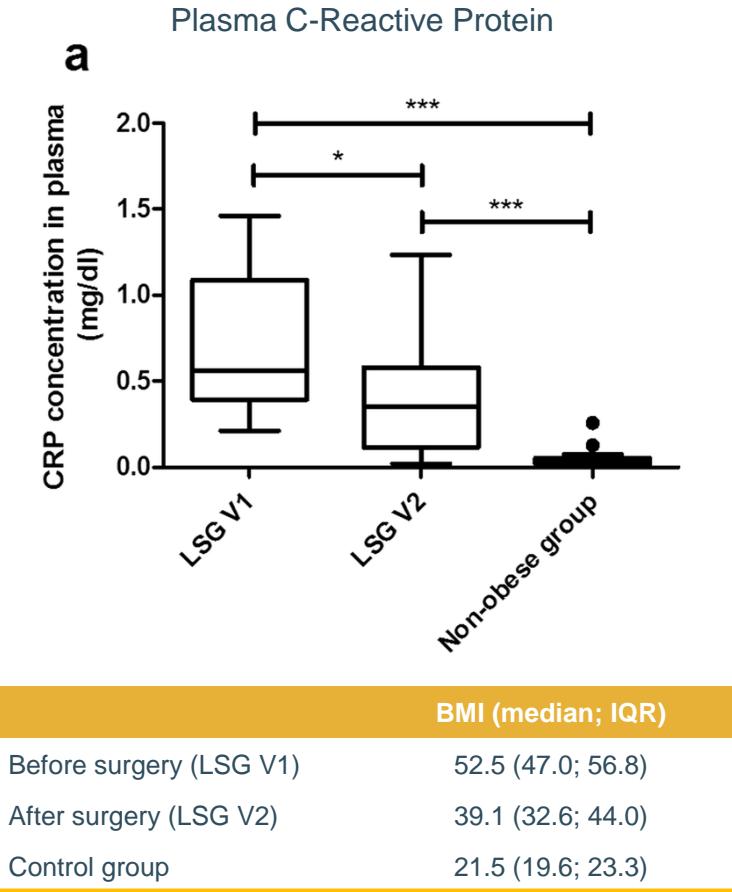
Gut permeability in humans



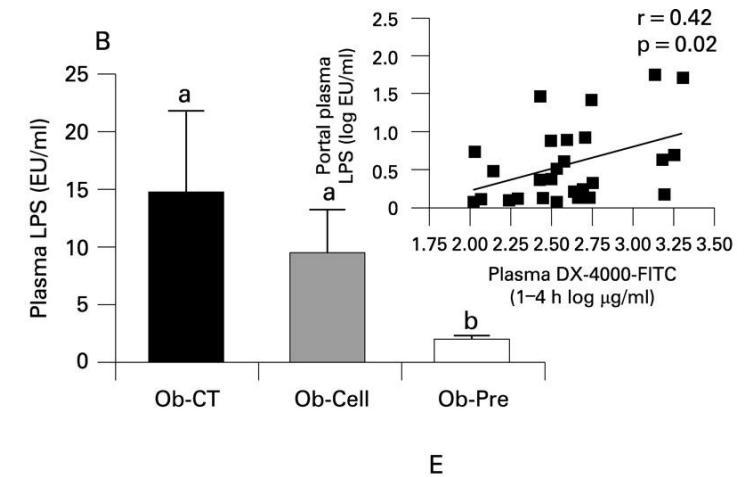
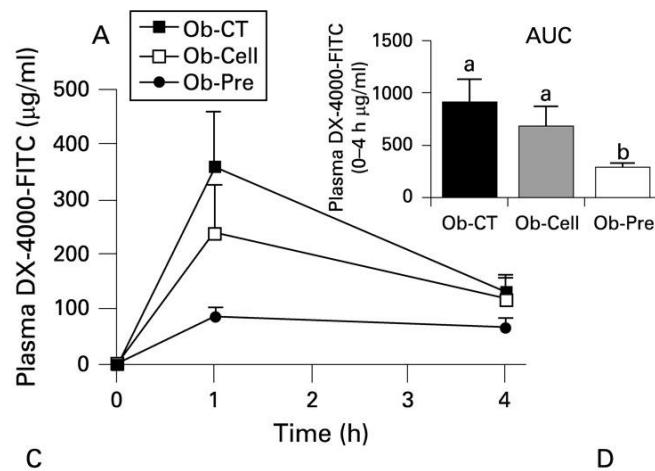
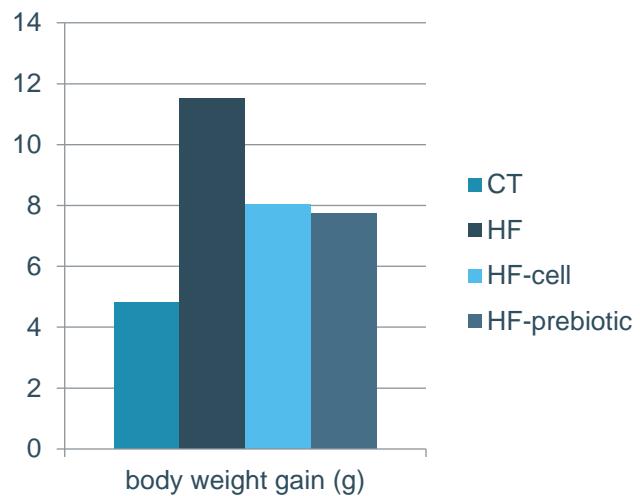
Obese subjects BMI: 32 (IQR 31-34)

Obese subjects BMI:
 t_{0m} : 43.7 ± 5.2
 t_{12m} : 36.4 ± 5.1

Gut permeability in humans

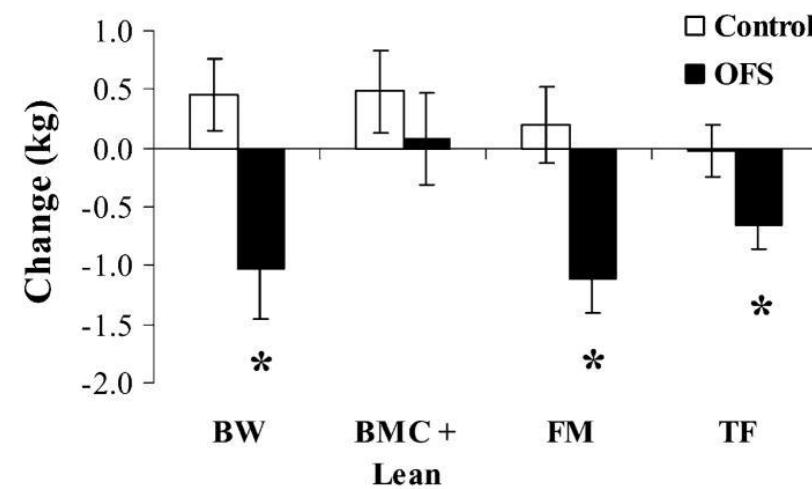
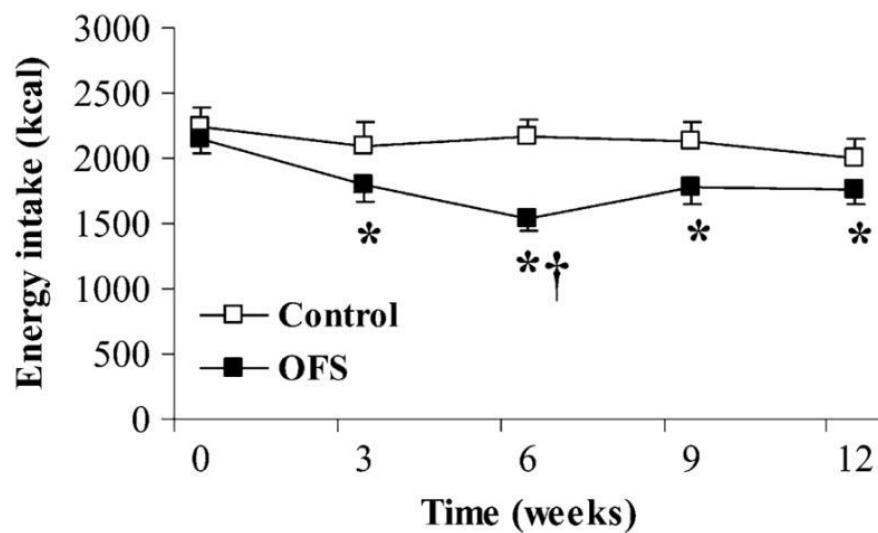


Prebiotics to treat obesity: mice studies



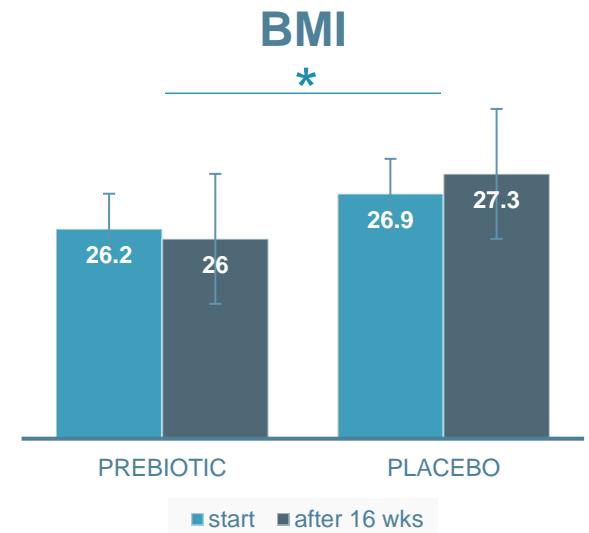
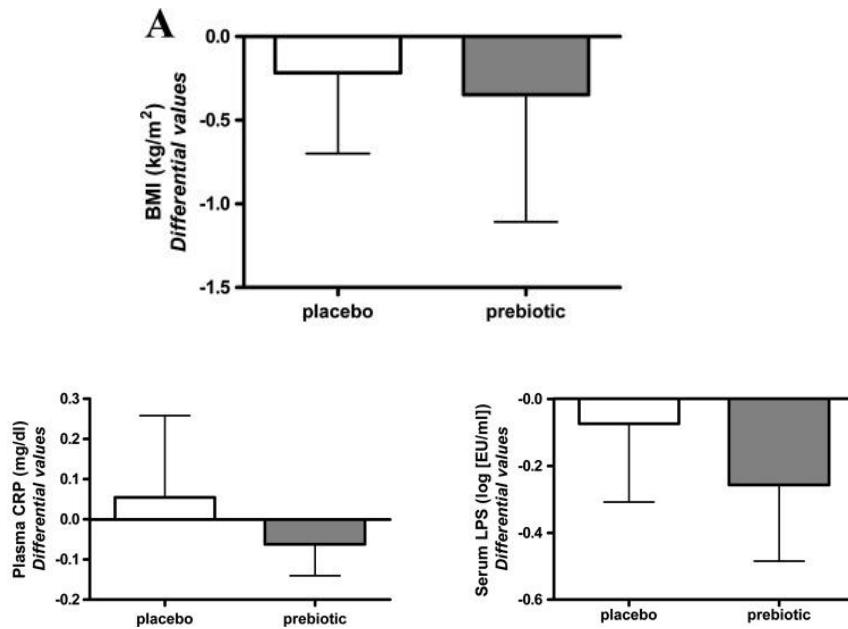
Prebiotics in humans

- 48 healthy adults with $\text{BMI} > 25 \text{ kg/m}^2$
- 21 g oligofructose or placebo for 12 weeks

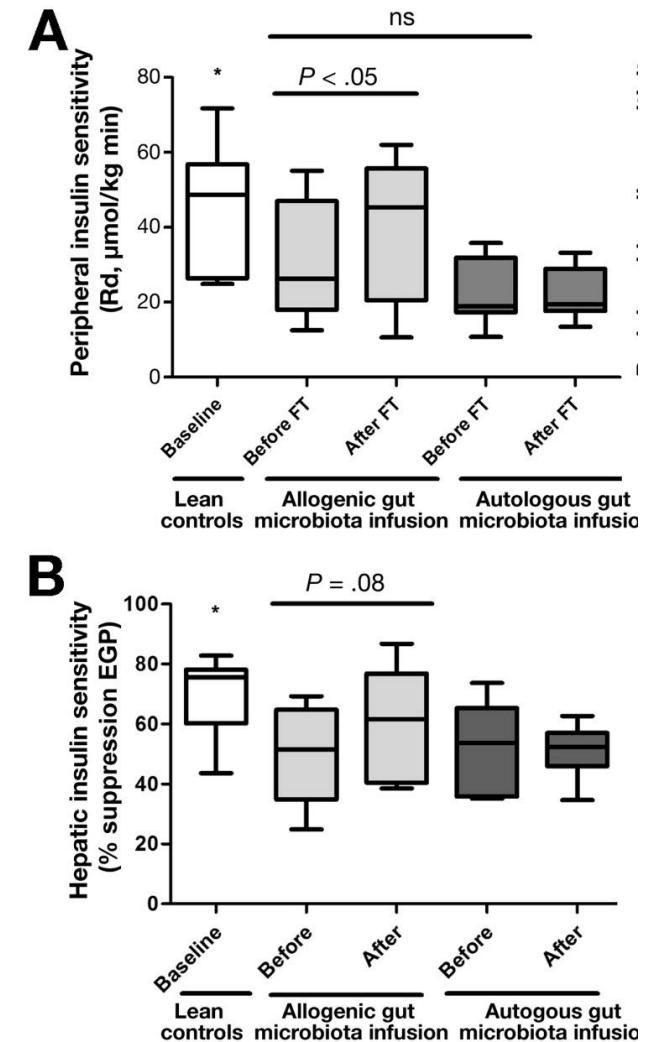
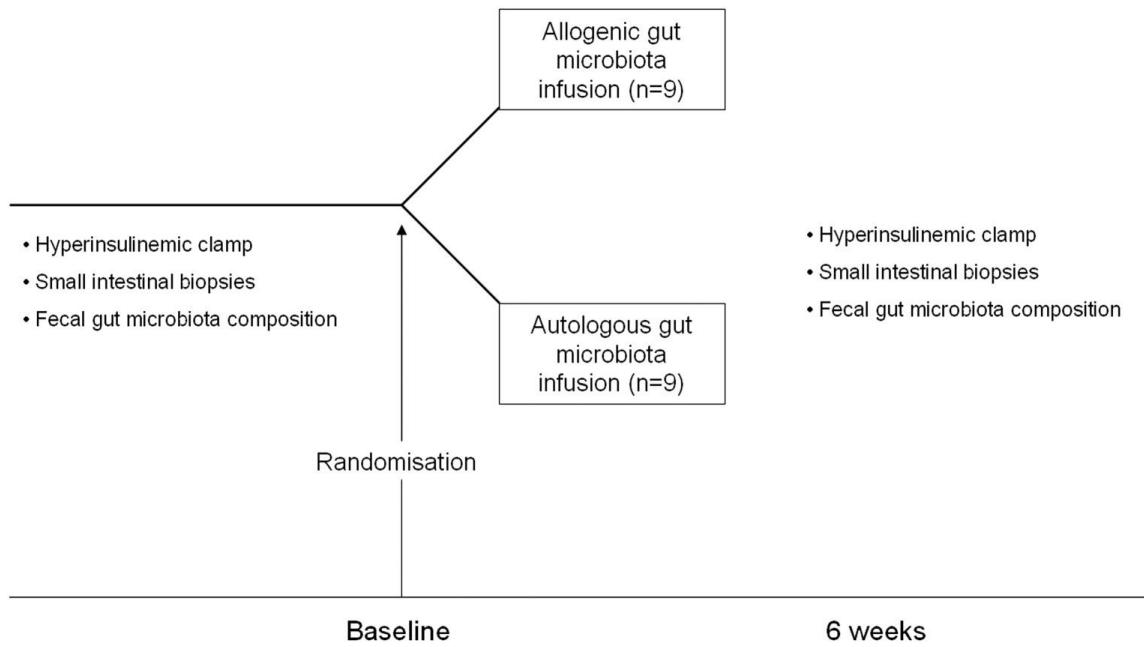


Prebiotics in humans

- 30 obese women
- 16 g oligofructose/inulin (50/50) or placebo for 3 months
- 106 obese people (m/f)
 - caloric restriction with 30% of energy intake
 - 16 g inulin or placebo for 3 months
- 42 obese children 7-12y
 - 8 g oligofructose/inulin (50/50) or placebo for 16 wks



Fecal microbiota transplantation



Take home messages

- Er zijn meer en meer aanwijzingen dat de microbiota betrokken is in het ontstaan van obesitas
- Microbiële componenten/metabolieten kunnen fungeren als signaalmoleculen
 - In combinatie met een hoog-vet dieet wordt meer LPS opgenomen dat obesitas en inflammatie induceert
 - SCFA verhinderen inflammatie
- De rol van darmpermeabiliteit bij de mens is niet duidelijk
- Interventies bij proefdieren die de microbiota moduleren leveren veelbelovende resultaten
- De effecten van dergelijke interventies bij de mens zijn eerder bescheiden

Bedankt voor uw aandacht!

