

Role of bacterioPhages In gut Microbiome composition and glucose metabolism in Metabolic Syndrome subjects

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We speculate that the virome composition of the lean donor and the subsequent interaction between phages and bacteria in the recipients are crucial determinants of stability and diversity of gut bacterial composition and glucose metabolism in humans...

Ethische beoordeling	Positief advies
Status	Werving gestart
Type aandoening	-
Onderzoekstype	Interventie onderzoek

Samenvatting

ID

NL-OMON26916

Bron

NTR

Verkorte titel

PIMMS

Aandoening

Metabolic Syndrome; insulin resistance

Ondersteuning

Primaire sponsor: Investigator initiated

Overige ondersteuning: Diabetes II Breakthrough grant (Diabetes Fonds & ZonMW)

Onderzoeksproduct en/of interventie

Uitkomstmaten

Primaire uitkomstmaten

To assess and compare the effect of lean-donor phage transplantation with placebo treatment (saline) on glucose tolerance (via an OGTT) in obese, insulin resistant subjects at risk to develop diabetes.

Toelichting onderzoek

Achtergrond van het onderzoek

Rationale

Alterations in gut microbiota composition and bacterial metabolites have been increasingly recognized to affect host metabolism and are at the basis of metabolic diseases such as obesity and type 2 diabetes (T2DM). As shown by our group, modulation of the gut microbial composition by faecal microbiota transplantation (FMT) improves insulin sensitivity in insulin resistant subjects at risk to develop T2DM. Importantly, however, the effect of FMT is transient and effect size differs strongly between subjects. We have shown that subjects with lower diversity at baseline are more likely to benefit from donor FMT than subjects with higher bacterial diversity. In addition, the level of engraftment of the transplant (i.e., presence of donor bacterial composition in gut of recipient) might determine in part the efficacy of this procedure. Factors that determine gut microbiota diversity and engraftment after FMT and subsequent effects on glucose metabolism are largely unexplored.

Bacteriophages (phages) are viruses that exclusively infect and eliminate bacteria. There is substantial evidence for a role of phages in shaping microbial communities in many ecosystems. However, insight in and attention for the role of phages in the human gut microbial community is limited. The effect of interventions that have significant consequences for both bacteria and phage communities, such as FMT, have never been assessed.

Since one gram of faeces contains ~10⁹ phage-containing particles and ~10⁹ bacterial cells, a vast number of phages are co-transplanted during FMT. The contribution of phages to FMT success (improved glucose metabolism) might therefore be quite substantial but has thus far never been studied. We therefore here propose a study in which we will transfer faecal bacteriophages from lean, healthy subjects to the gut of obese, insulin resistant subjects and assess the effect on engraftment and glucose metabolism.

Hypothesis

We speculate that the virome composition of the lean donor and the subsequent interaction between phages and bacteria in the recipients are crucial determinants of stability and diversity of gut bacterial composition and glucose metabolism in humans after FMT. We hypothesize that lean donor phage transplantation will increase gut microbial diversity and

improves glucose handling in insulin resistant recipients at risk to develop T2DM.

Objective

- Primary objective: to assess and compare the effect of lean-donor phage transplantation with placebo treatment (saline) on glucose tolerance in obese, insulin resistant subjects at risk to develop diabetes.
- Secondary objective: to assess and compare changes in gut microbiota and phage composition following phage transplantation and placebo.

Study design

Prospective, double-blinded, randomised, single-center intervention study.

Study population

Male overweight, metabolic syndrome (MetSyn) subjects (N=24, ≥ 18 years old, BMI ≥ 25 kg/m², treatment naive.)

Intervention

Twelve MetSyn subjects will receive saline (placebo) whereas another group of twelve MetSyn subjects will receive a faecal filtrate transplant (FFT), which lacks bacteria and mainly consists of phages.

Main study parameters/endpoints

- An approximation of the potential of lean-donor phage transplantation to modify gut microbiota composition and improve glucose metabolism in individuals at risk to develop T2DM
- An overview of the dynamic changes in gut microbiota and virome population following phage transplantation compared to placebo.
- A comparison between phage composition in individuals at risk to develop T2DM at baseline with phage composition in lean, healthy subjects (donors).

Doel van het onderzoek

We speculate that the virome composition of the lean donor and the subsequent interaction between phages and bacteria in the recipients are crucial determinants of stability and diversity of gut bacterial composition and glucose metabolism in humans after FMT. We hypothesize that lean donor phage transplantation will increase gut microbial diversity and improves glucose handling in insulin resistant recipients at risk to develop T2DM.

Onderzoeksopzet

Visits to the AMC: screening and inclusion, visit 1 (day 0) and visit 2 (day 28);

Collection of fecal samples: day -7, 0, 2, 4, 7, 14, 28, 42

Continuous glucose monitoring: from day -7 to day 7

Dietary diary: from day -7 to day 7

OGTT: during visit 1 (day 0) and visit 2 (day 28)

FFT or placebo: during visit 1 (day 0)

Onderzoeksproduct en/of interventie

Twelve MetSyn subjects will receive saline (placebo) whereas another group of twelve MetSyn subjects will receive a faecal filtrate transplant (FFT), which lacks bacteria and mainly consists of phages.

Contactpersonen

Publiek

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Wetenschappelijk

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Deelname eisen

Belangrijkste voorwaarden om deel te mogen nemen (Inclusiecriteria)

- Caucasian male/female
- Age: ≥ 18 years old
- BMI: ≥ 25 kg/m²
- At least 3 of the following criteria:
 - o Fasting plasma glucose ≥ 5.6 mmol/L OR HOMA-IR index ≥ 2.5 (HOMA-IR is measured as (fasting insulin (pmol/L) x fasting glucose (mmol/L)) / 135)
 - o Waist-circumference ≥ 102 cm
 - o HDL-cholesterol ≤ 1.02 mmol/L
 - o Blood pressure $\geq 130/85$ mmHg
 - o Triglycerides ≥ 1.7 mmol/L
- Subjects should be able to give informed consent

Belangrijkste redenen om niet deel te kunnen nemen (Exclusiecriteria)

- A history of cardiovascular event (cerebrovascular accident (CVA), myocardial infarction (MI)) or pacemaker implantation
- Use of any medication including proton pump inhibitors, antibiotics and pro-/prebiotics in the past three months or during the study period
- (Expected) prolonged compromised immunity (due to recent cytotoxic chemotherapy or human immunodeficiency viruses (HIV) infection with a CD4 count < 240/mm³)
- Presence of overt type 1 diabetes mellitus (T1DM) or T2D
- History of chronic diarrhoea (≥ 3 stools/day for >4 weeks), chronic obstipation (<2 defecations/week for >3 months), Irritable Bowel Syndrome (IBS) (according to Rome IV criteria) or Inflammatory Bowel Disease (IBD).
- Smoking or illicit drug use (MDMA/amphetamine/cocaine/heroin/GHB) in the past three months or use during the study period
- Use of >5 units of alcohol daily on average in the past three months or use of >2 units of alcohol during the study period
- History of cholecystectomy

Onderzoeksopzet

Opzet

Type:	Interventie onderzoek
Onderzoeksmodel:	Parallel
Toewijzing:	Gerandomiseerd
Blindering:	Dubbelblind
Controle:	Placebo

Deelname

Nederland	
Status:	Werving gestart
(Verwachte) startdatum:	31-07-2019
Aantal proefpersonen:	24
Type:	Verwachte startdatum

Voornemen beschikbaar stellen Individuele Patiënten Data (IPD)

Wordt de data na het onderzoek gedeeld: Nog niet bepaald

Ethische beoordeling

Positief advies

Datum: 15-01-2020

Soort: Eerste indiening

Registraties

Opgevolgd door onderstaande (mogelijk meer actuele) registratie

Geen registraties gevonden.

Andere (mogelijk minder actuele) registraties in dit register

Geen registraties gevonden.

In overige registers

Register	ID
NTR-new	NL8289
Ander register	METC AMC : METC 2018_231, NL67136.018.18

Resultaten