# VLDL1, VLDL2, and LDL apolipoprotein B-100 kinetics in patients with familial hypercholesterolemia of unknown origin (FH4)

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To study \*in vivo\* lipid and apolipoprotein metabolism in patients with FH4, and to compare the results with healthy family controls and dyslipidemic patients with mutations in established lipid genes(i.e. FH1 \* FH3). This will ultimately help to...

Ethical review Approved WMO

**Status** Recruitment stopped

**Health condition type** Lipid metabolism disorders

**Study type** Interventional

# **Summary**

### ID

NL-OMON45448

### **Source**

**ToetsingOnline** 

### **Brief title**

**APPRECIATION** 

### **Condition**

Lipid metabolism disorders

### Synonym

Familial hypercholesterolemia

### Research involving

Human

# **Sponsors and support**

**Primary sponsor:** Academisch Medisch Centrum

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**Source(s) of monetary or material Support:** Vidi grant [016.156.445] from the Netherlands Organisation for Scientific Research (NWO)

### Intervention

Keyword: apolipoprotien, hypercholesterolemia, kinetic, LDL

### **Outcome measures**

### **Primary outcome**

Tracer/tracee ratio, pool size, fractional catabolic rate, and production rate in FH4 subjects compared with (family) controls and/or dyslipidemic patients with mutations in established lipid genes.

### **Secondary outcome**

n.v.t.

# **Study description**

### **Background summary**

Familial hypercholesterolemia (FH) is characterized by increased low density lipoprotein (LDL) cholesterol and increased cardiovascular risk. There are 3 known genes (LDLR, ApoB, PCSK9) in which mutations can lead to the FH phenotype (FH1 to 3 respectively). However, in approximately 5-10% of patients such a mutation cannot be found, despite family-based linkage studies (the so called FH4 group). Therefore, a more elaborate approach is deemed necessary, where data derived from transcriptome, proteome, and metabolome studies are combined to find novel genes and metabolic pathways in lipid metabolism. We aim to acquire in depth phenotyping in selected FH4 patients by extensive lipoprotein kinetic flux studies yielding insights into the pathophysiological mechanism of FH4.

### **Study objective**

To study \*in vivo\* lipid and apolipoprotein metabolism in patients with FH4, and to compare the results with healthy family controls and dyslipidemic patients with mutations in established lipid genes(i.e. FH1 \* FH3). This will ultimately help to answer the fundamental question as to whether FH4 is a result of overproduction or a decreased catabolism of ApoB containing

lipoproteins.

### Study design

Matched case-control study

### Intervention

n.v.t.

### Study burden and risks

Participants will visit the AMC for 1 or 2 full day(s) (7.30 am till 6.00 pm). Metabolic fluxes will be measured with stable isotope tracers that behave like their natural substrates and are therefore not harmful. At 21 timepoints blood samples will be collected, in total 279 ml. In the week prior to the admission participiants will be asked to keep track of their daily diet. On the day after the study day a last blood sample will be collected at the patients home at 8.00 am.

Patiënts with FH4 will be asked to participate in 2 study days, 1 after a 4 week wash-out period of lipid lowering therapy.

We state that the scientific insight of our findings will outweigh the minimal burdens and risks in this study.

# **Contacts**

### **Public**

Academisch Medisch Centrum

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### Scientific

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# **Trial sites**

### **Listed location countries**

**Netherlands** 

# **Eligibility criteria**

### Age

Adults (18-64 years) Elderly (65 years and older)

### Inclusion criteria

- Diagnosis of familial hypercholesterolemia of unknown origin (FH4) based on Dutch Lipid Clinic Network criteria (Nordestgaard et al. 2013) and negative DNA-testing (mutations in LDLR, ApoB, PCSK9).
- Untreated LDL-cholesterol levels of > 95th percentile for affected members
- 18-65 years of age; Controls:
- Unaffected family member with untreated LDL-cholesterol between 20-60th percentile
- 18-65 years of age

Or:

- Persons with mutations in LDLR, APOB, or PCSK9 gene or other genes involved in lipid metabolism.
- 18-65 years of age

### **Exclusion criteria**

- Heavy alcohol use
- · Dysthyroidism
- · Pregnancy, breastfeeding
- · Renal insufficiency (creatinine >150 µmol/L)
- · Diabetes mellitus

# Study design

# **Design**

Study type: Interventional

Intervention model: Other

Allocation: Non-randomized controlled trial

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Masking: Open (masking not used)

Primary purpose: Basic science

### Recruitment

NL

Recruitment status: Recruitment stopped

Start date (anticipated): 22-05-2017

Enrollment: 60

Type: Actual

# **Ethics review**

Approved WMO

Date: 31-03-2017

Application type: First submission

Review commission: METC Amsterdam UMC

Approved WMO

Date: 31-07-2017

Application type: Amendment

Review commission: METC Amsterdam UMC

Approved WMO

Date: 10-10-2017

Application type: Amendment

Review commission: METC Amsterdam UMC

# **Study registrations**

# Followed up by the following (possibly more current) registration

No registrations found.

# Other (possibly less up-to-date) registrations in this register

ID: 26254

Source: Nationaal Trial Register

Title:

# In other registers

Register ID

CCMO NL61012.018.17 OMON NL-OMON26254

# **Study results**

Date completed: 20-04-2020

Actual enrolment: 2

## **Summary results**

Trial ended prematurely