

The relationship between age at menopause loci and the response of controlled ovarian hyperstimulation in women undergoing IVF: a candidate gene study.

No registrations found.

Ethical review	Positive opinion
Status	Pending
Health condition type	-
Study type	-

Summary

ID

NL-OMON28369

Source

Nationaal Trial Register

Brief title

RiSPONS

Health condition

Subfertile women who will undergo IVF treatment.

Sponsors and support

Primary sponsor: Ferring

Source(s) of monetary or material Support: Ferring

Intervention

Outcome measures

Primary outcome

The primary outcome parameter is the responsiveness of the ovaries to exogenous FSH stimulation during IVF treatment, i.e. number of retrieved oocytes.

Secondary outcome

Secondary study parameters are other outcomes of the IVF treatment, described by the number of follicles after stimulation, quality of retrieved oocytes, quality of embryo's as well as pregnancy rate and live-birth rate.

Study description

Background summary

Rationale: Over the last decade genome-wide association studies (GWAS) have identified hundreds of common genetic variants (Single Nucleotide Polymorphisms, SNPs) associated with age of natural menopause (ANM). The peri-menopausal stage is preceded by a decrease in fertility eventually leading to sterility in the 10 years before menopause. Poor response to controlled ovarian hyperstimulation (COH) during IVF treatment is a reasonable predictor of diminished ovarian reserve. Therefore, we hypothesize that presence of genetic variants associated with earlier age of natural menopause might also be involved in diminished responsiveness of the ovaries to exogenous FSH stimulation during IVF treatment. In other words, patients that carry more SNPs associated with early age of menopause might as well have a poor response to COH treatment. Consequently, if that is the case, these SNPs might be used as a predictor for ovarian response during IVF treatment using polygenic risk scores.

Objective: In this project, we hypothesize that presence of genetic variants associated with earlier age of natural menopause might be involved in diminished responsiveness of the ovaries to exogenous FSH stimulation during IVF treatment. Furthermore, we aim to investigate whether polygenic risk scores of genetic variants associated with age at natural menopause based on the large-scale GWAS are predictive of the response to IVF stimulation in patients starting their first IVF treatment cycle leading to a more patient-tailored care-setting.

Study design: This candidate gene study will be conducted in a retrospective as well as prospective cohort study.

Study population: Patients will be recruited at the IVF outpatient clinic of the Division of Reproductive Medicine at the Erasmus MC, University Medical Center, Rotterdam. All women with a regular menstrual cycle undergoing their first cycle of IVF because of male infertility or unexplained infertility will be asked to participate in this study.

Study protocol: Baseline characteristics, including medical history, age, BMI, smoking, SES, and endocrine parameters such as FSH, LH, progesterone, estradiol, AMH will be determined. Extra blood samples will be taken during standard blood sampling. DNA will be extracted

from peripheral blood, and genotyping shall be executed using Illumina arrays. Standard quality-control procedures are performed. SNPs were selected to be included in the Polygenic Risk Score calculation based on their p-values in the original GWAS. Patients will undergo hormonal stimulation according to a standard protocol in our clinic. Response to the IVF stimulation in terms of dose of exogenous FSH, number of follicles and number and quality of obtained oocytes as well as pregnancy will be monitored.

Main study parameters: The primary outcome parameter is the responsiveness of the ovaries to exogenous FSH stimulation during IVF treatment, i.e. number of retrieved oocytes. Secondary study parameters are other outcomes of the IVF treatment, described by the number of follicles after stimulation, quality of retrieved oocytes, quality of embryo's as well as pregnancy rate and live-birth rate.

Nature and extent of the burden and risks associated with participation, benefit and group relatedness: Single blood sampling is needed to execute this candidate gene study, although this will be combined with standard blood sampling women will need to visit the hospital one extra time.

Study objective

We hypothesize that presence of genetic variants associated with earlier age of natural menopause might be involved in diminished responsiveness of the ovaries to exogenous FSH stimulation during IVF treatment.

Study design

Before IVF treatment the extra blood samples will be taken.

Intervention

Extra blood samples will be taken during standard blood sampling.

Contacts

Public

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Scientific

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Eligibility criteria

Inclusion criteria

- A regular menstrual cycle
- Undergoing IVF because of male infertility or unexplained infertility
- First cycle IVF treatment
- Age between 18 and 45 years at the time of the IVF treatment

Exclusion criteria

- A history of ovarian surgery, chemotherapy or radiation therapy
- Non Northern European ethnicity

Study design

Design

Intervention model:	Other
Allocation:	Non controlled trial
Masking:	Open (masking not used)
Control:	N/A , unknown

Recruitment

NL	
Recruitment status:	Pending
Start date (anticipated):	01-01-2021
Enrollment:	500
Type:	Anticipated

IPD sharing statement

Plan to share IPD: No

Ethics review

Positive opinion

Date: 06-01-2021

Application type: First submission

Study registrations

Followed up by the following (possibly more current) registration

ID: 54409

Bron: ToetsingOnline

Titel:

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

No registrations found.

In other registers

Register	ID
NTR-new	NL9173
CCMO	NL75062.078.20
OMON	NL-OMON54409

Study results