# Genetic characteristics of Primary Ciliary Dyskinesia.

No registrations found.

**Ethical review** Not applicable

**Status** Pending

Health condition type -

**Study type** Observational non invasive

# **Summary**

#### ID

NL-OMON24319

**Source** 

NTR

**Health condition** 

Primary Ciliary Dyskinesia

## **Sponsors and support**

**Primary sponsor:** VU University Medical Center

Source(s) of monetary or material Support: VU University Medical Center

Fonds NutsOhra

## Intervention

#### **Outcome measures**

#### **Primary outcome**

- 1. Accuracy of MPS in detecting PCD mutation (validation);
- 2. Possible pathogenic mutations causing PCD.

## **Secondary outcome**

N/A

# **Study description**

#### **Background summary**

Primary Ciliary Dyskinesia (PCD) is an autosomal recessive hereditary disorder that causes dysfunction of cilia. Patients suffer from frequent respiratory infections and often develop bronchiectasis. Diagnosing PCD is difficult as a single gold standard is lacking. The diagnosis is usually based on a combination of clinical symptoms, abnormal movement of cilia on microscopic evaluation of respiratory epithelial biopsies and epithelial cell cultures, and/or identification of an ultra structural defect in the cilia by electron microscopy. Genetic testing is time consuming and very costly as there are many large genes involved. However, recent developments enable rapid DNA sequencing of many fragments in parallel.

We aim to validate Massive Parallel Sequencing (MPS) in Primary Ciliary Dyskinesia and identify novel disease causing mutations in our Dutch PCD population.

We aim to include all children and adults with PCD visiting the VU University Medical Center, Amsterdam.

## Study objective

We aim to validate Massive Parallel Sequencing in Primary Ciliary Dyskinesia (PCD) and identify novel disease causing mutations in the Dutch PCD population.

## Study design

N/A

#### Intervention

N/A

## **Contacts**

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# **Eligibility criteria**

#### Inclusion criteria

Primary Ciliary Dyskinesia.

## **Exclusion criteria**

Other recessive hereditary disorders.

# Study design

## **Design**

Study type: Observational non invasive

Intervention model: Parallel

Allocation: Non controlled trial

Masking: Open (masking not used)

Control: N/A, unknown

## **Recruitment**

NL

Recruitment status: Pending

Start date (anticipated): 01-01-2012

Enrollment: 83

Type: Anticipated

# **Ethics review**

Not applicable

Application type: Not applicable

# **Study registrations**

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

No registrations found.

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

No registrations found.

## In other registers

Register ID

NTR-new NL2990 NTR-old NTR3138

Other CWO VUmc : pro 11/68

ISRCTN wordt niet meer aangevraagd.

# **Study results**

## **Summary results**

N/A