

Hereditary hearing impairment in the Netherlands: elucidation of genetic causes, and clinical characterization.

Published: 31-05-2011

Last updated: 04-05-2024

The goal of this study is to collect a large cohort of patients mainly with NSHI. In the study cohort, the genetic causes underlying the NSHI will be determined which can be either in known or novel deafness genes. Also, the clinical characteristics...

Ethical review	Approved WMO
Status	Recruiting
Health condition type	Ear and labyrinthine disorders congenital
Study type	Observational non invasive

Summary

ID

NL-OMON47077

Source

ToetsingOnline

Brief title

Hereditary hearing impairment in the Netherlands

Condition

- Ear and labyrinthine disorders congenital

Synonym

hereditary deafness, hereditary hearing impairment

Research involving

Human

Sponsors and support

Primary sponsor: KNO

Source(s) of monetary or material Support: NWO (ZONMW),Het Heinsius Houbolt fonds;Fonds NutsOHRA

Intervention

Keyword: genotype, hearing impairment, inheritance, phenotype

Outcome measures

Primary outcome

1, pathologic genetic variants that are known to or are likely to be causative for hearing impairment. 2, genotype-phenotype correlations for at least part of the genetic subtypes leading to improved counselling.

Secondary outcome

A strategy for DNA-diagnostics for hearing impairment in the Dutch population.

Study description

Background summary

Currently, many of the causative genes for syndromic hearing impairment are known. However, especially for non-syndromic hereditary hearing impairment (NSHI) the majority of the genes remain to be identified which means that in ~70% of NSHI patients, the genetic defect remains unknown in routine DNA-diagnostics. Knowledge of the genetic cause is important to patients in order to distinguish between genetic and non-genetic hearing impairment, to give them a more grounded genetic counselling and prognosis, to distinguish in an early stage between syndromic and non-syndromic hearing impairment, and to determine whether treatments such as cochlear implantation may be helpful to the patient.

Study objective

The goal of this study is to collect a large cohort of patients mainly with NSHI. In the study cohort, the genetic causes underlying the NSHI will be determined which can be either in known or novel deafness genes. Also, the clinical characteristics associated with the specific genetic defects will be determined. Collecting a large cohort of patients will help us to provide 1. accurate genetic counselling to patients, 2. a more reliable prognosis, 3. to discover new genes associated with hearing impairment, 4. to develop a strategy for future routine DNA diagnostics for hereditary hearing impairment and 5.

insight in inner ear (dys)function.

Study design

Patients will be informed about the study by their ENT specialist, clinical geneticist or audiologist. The usual diagnostic examinations for hearing impairment will be performed (otoscopy, tympanometry, pure tone audiometry, speech audiometry), and a blood sample will be drawn for usual DNA diagnostic screening for hereditary hearing impairment.

Patient will receive a questionnaire, with questions about their hearing, balance and origin of their grandparents. Besides, the patient will be asked to give consent on the further genetic research when no mutations in known deafness genes can be found.

A small number of patients will be invited for more psychophysical hearing tests, in order to get a more detailed phenotype.

Study burden and risks

The only burden for a patient could be the possible extra examinations (psychophysical testing, takes 1 hour). The other examinations would be performed anyway for diagnostic purposes.

Contacts

Public

Selecteer

Geert Grooteplein Zuid 10
Nijmegen 6525 GA
NL

Scientific

Selecteer

Geert Grooteplein Zuid 10
Nijmegen 6525 GA
NL

Trial sites

Listed location countries

Netherlands

Eligibility criteria

Age

Adolescents (12-15 years)
Adolescents (16-17 years)
Adults (18-64 years)
Children (2-11 years)
Elderly (65 years and older)

Inclusion criteria

patients with hearing impairment in whom common non-genetic causes for hearing impairment are excluded.

Exclusion criteria

patients with age-related hearing impairment

Study design

Design

Study type: Observational non invasive

Masking: Open (masking not used)

Control: Uncontrolled

Primary purpose: Basic science

Recruitment

NL

Recruitment status: Recruiting

Start date (anticipated): 15-06-2011

Enrollment: 2050

Type: Actual

Ethics review

Approved WMO

Date: 31-05-2011

Application type: First submission

Review commission: CMO regio Arnhem-Nijmegen (Nijmegen)

Approved WMO

Date: 14-01-2015

Application type: Amendment

Review commission: CMO regio Arnhem-Nijmegen (Nijmegen)

Approved WMO

Date: 29-07-2015

Application type: Amendment

Review commission: CMO regio Arnhem-Nijmegen (Nijmegen)

Approved WMO

Date: 30-12-2015

Application type: Amendment

Review commission: CMO regio Arnhem-Nijmegen (Nijmegen)

Approved WMO

Date: 27-07-2017

Application type: Amendment

Review commission: CMO regio Arnhem-Nijmegen (Nijmegen)

Approved WMO

Date: 28-12-2017

Application type: Amendment

Review commission: CMO regio Arnhem-Nijmegen (Nijmegen)

Approved WMO

Date: 02-05-2018

Application type: Amendment

Review commission: CMO regio Arnhem-Nijmegen (Nijmegen)

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
CCMO	NL33648.091.10