

Research into the progression of Parkinson's disease in patients with a mutation in the GBA1 gene, based on medical history and genetic analyses

No registrations found.

| | |
|------------------------------|----------------------------|
| Ethical review | Positive opinion |
| Status | Recruiting |
| Health condition type | - |
| Study type | Observational non invasive |

Summary

ID

NL-OMON28601

Source

Nationaal Trial Register

Brief title

CHDR1838 / LTI-OBS-001

Health condition

Parkinson's disease and the GBA-PD subset of patients

Sponsors and support

Primary sponsor: Lysosomal Therapeutics Incorporated

Source(s) of monetary or material Support: Lysosomal Therapeutics Incorporated

Intervention

Outcome measures

Primary outcome

Based on multiple parameters, the following compound measurements are constructed:

- Parkinson's disease - Progression Rate Inventory (pD-pRI)
- GBA1 genotype and PD associated SNPs
- Parkinson's Risk Score (PRS)
- Rate of disease progression as rated by a clinical expert: Slow, Intermediate, Fast

The endpoints are the analyses of the associations between:

- genetic factors and phenotypic factor
- genetic factors and the rate of disease progression as rated by a Movement Disorder Neurologist
- genetic factors and a data-driven algorithm based on phenotypic characteristics

Secondary outcome

N.A.

Study description

Background summary

This protocol is a follow-up on the previous study CHDR1707 (Toetsing online number: NL61137.056.17), titled "Genetic screening in Parkinson's Disease in order to identify patients who can participate in clinical trials with new targeted therapies." In this previous study, Parkinson's patients throughout the Netherlands were genetically screened for presence of mutations in the GBA1 gene and LRRK2 gene. In approximately 15% of all screened patients, a mutation was found in the GBA1 gene. The current protocol aims to further characterize this subgroup of patients with a GBA1 mutation, based on phenotype, as assessed by medical history, and on genotype, as assessed by Parkinson's disease related Single Nucleotide Polymorphism (SNP) analysis. The goal of this study is to exploratively investigate whether clinical and genetic factors may contribute to the rate of clinical progression in patients with Parkinson's disease associated with a GBA1 mutation in the gene encoding GCase (GBA-PD)

Study objective

- To determine the relationship between phenotypic and genetic characteristics of GBA-PD patients. Phenotypic characteristics will be obtained by patient dossier review. Genetic characteristics of the GBA1 gene and a panel of SNP's (gene markers), previously associated with risk or progression of idiopathic Parkinson's disease, will be assessed.

- To determine the correlation between genetic characteristics (as described above) and the estimated disease progression rating (Fast, Intermediate, Slow) by a Movement Disorders Neurologist, based on retrospective phenotypic characteristics (as described above).
- o Inter-rater correlation between Movement Disorders Neurologists will be determined in their rating of disease progression.
- To determine the correlation between genetic characteristics (as described above) and a data-driven algorithm based on phenotypic characteristics (as described above).

Study design

N.A. Observational study of hospital records and genotype records of PD patients (of CHDR1707 study)

Contacts

Public

Centre for Human Drug Research
Geert Jan Groeneveld

+31 71 5246 400

Scientific

Centre for Human Drug Research
Geert Jan Groeneveld

+31 71 5246 400

Eligibility criteria

Inclusion criteria

- Signed informed consent prior to any study-mandated procedure.
- Minimum age of 18 years.
- Clinical diagnosis of Parkinson's disease at least 6 months prior to screening, confirmed by a Movement Disorder's Neurologist.
- Mutation(s) in the glucocerebrosidase GBA1 gene. Reference Appendix A for a list of GBA1 mutations.

Exclusion criteria

- N.A.

Study design

Design

| | |
|---------------------|----------------------------|
| Study type: | Observational non invasive |
| Intervention model: | Other |
| Allocation: | Non controlled trial |
| Masking: | Open (masking not used) |
| Control: | N/A , unknown |

Recruitment

| | |
|---------------------------|-------------|
| NL | |
| Recruitment status: | Recruiting |
| Start date (anticipated): | 15-10-2018 |
| Enrollment: | 350 |
| Type: | Anticipated |

IPD sharing statement

Plan to share IPD: No

Ethics review

| | |
|-------------------|------------------|
| Positive opinion | |
| Date: | 07-05-2019 |
| Application type: | First submission |

Study registrations

Followed up by the following (possibly more current) registration

ID: 46323
Bron: ToetsingOnline
Titel:

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

No registrations found.

In other registers

| Register | ID |
|----------|----------------|
| NTR-new | NL7717 |
| CCMO | NL67297.056.18 |
| OMON | NL-OMON46323 |

Study results