CANDY MULTIPLEX STUDY

Published: 12-01-2022 Last updated: 13-06-2024

1) Assess the whole genome sequence (WGS) and clinical profile of the MPX families to identify how rare and common genetic variants interact to produce shared versus distinct NDDs and clinical features in individuals, or different combinations of...

Ethical review	Approved WMO
Status	Recruiting
Health condition type	Mental impairment disorders
Study type	Observational invasive

Summary

ID

NL-OMON52192

Source ToetsingOnline

Brief title Multiplex

Condition

- Mental impairment disorders
- Developmental disorders NEC

Synonym Neurodevelopmental Disorder

Research involving Human

Sponsors and support

Primary sponsor: Radboud Universitair Medisch Centrum **Source(s) of monetary or material Support:** Horizon2020 programme of the EU (project [CANDY]);grant number 847818

Intervention

Keyword: ADHD, Autism Spectrum Disorder (ASD), Epilepsy, Intellectual Disability

Outcome measures

Primary outcome

1) Complete genetic profile of all participants (including rare and common genes).

2) Refined phenoptyping of all participants by means of questionnaires and interviews in terms of: ASD symptoms and traits, Cognitive and neurocognitive level, Developmental level, Somatic comorbidities (clinical epilepsy, allergies, immune disease, gastro-intestinal problems) and Psychiatric comorbidities.

A subsample of the participants will be invited for a nested follow-up study with assessment of EEG/ERP, eye-tracking and cognition; for this, we will submit an amendment at due time.

Secondary outcome

n.a.

Study description

Background summary

Neurodevelopmental disorders (NDDs) are frequent and significantly impact peoples* lives. According to the DSM-5, NDDs include Autism Spectrum Disorder (ASD), Attention Deficit Hyperactivity Disorder (ADHD), Intellectual Disability (ID), specific learning disorders, language disorders and motor disorders.

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ADHD, ID and epilepsy co-occur more often within an individual with ASD than in the general population. Further, NDDs are highly heritable and cluster in partly the same families. Most studies investigate families with a single affected member but few studies have investigated the other family members who are likely to be affected by NDDs or epilepsy. The relationship between genotype and phenotype is only partially understood, especially in families in which several individuals are affected by heterogeneous forms of NDDs. In the current study, we will investigate the shared and distinct genetic and biologic mechanisms involved in those NDDs by recruiting so-called multiplex (MPX) families in which there are at least two patients, one proband with ASD, and at least one other first-degree or second-degree relative with epilepsy, ASD, ID or ADHD.

Study objective

1) Assess the whole genome sequence (WGS) and clinical profile of the MPX families to identify how rare and common genetic variants interact to produce shared versus distinct NDDs and clinical features in individuals, or different combinations of NDDs and/or epilepsy in an individual.

Nota Bene: We will not perform whole genome sequencing analysis in participants younger than 16 years, but only assess variation in "common genes".

2) Identify the molecular pathways that are involved in the clinical phenotype and cut across NDD in relationship with the genetics variants by combining different technique used to identify (i) mutated genes; (ii) common genetic variants; (iii) the genetic background; and (iv) the clinical profile of the individual.

Study design

The CANDY multiplex study is an observational multi-centre study conducted at five sites: Robert-Debré Hospital (Paris), King*s College London (London), Radboudumc (Nijmegen), Ghent University and Karolinska Institutet (Stockholm). The study will be conducted over 36 months (time between the first and last visit).

Study burden and risks

We will collect biosamples by means of venapuncture in all participants for genetic studies and immune profiling, and collect data by questionnaires and interviews. In addition to adults and adolescents, children of 3 years and older will be included in this study, as we are also investigating the development of and emergence of NDDs over a wide age range. A study of very young and young children through to adults is vital to understanding how the clinical and genetic/ biomarker profile of NDDs changes at different time points. As this is an explorative study, the hypotheses that are generated as a result of these experiments will stimulate and inform future NDD research. Overall, this work will lead to new insights into the common and distinct genetic and biologic mechanisms that underpin ASD, ADHD, ID and epilepsy. This will lead to better counseling of families with NDDs and development of targeted treatments.

Contacts

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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)

Inclusion criteria

Multiplex (MPX) families will be enrolled when there are at least two diagnosed individuals, one with ASD, and at least one other first or second-order member

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with epilepsy, ASD, ID or ADHD (age 3 years and older). All unaffected first-degree relatives of an MPX family will be invited to participate as well.

Exclusion criteria

None of the biological parents of the ASD proband is available / willing to participate

Study design

Design

Study type: Observational invasive		
Masking:	Open (masking not used)	
Control:	Uncontrolled	
Primary purpose:	Basic science	

Recruitment

NL	
Recruitment status:	Recruiting
Start date (anticipated):	05-06-2023
Enrollment:	100
Туре:	Actual

Ethics review

12-01-2022
First submission
CMO regio Arnhem-Nijmegen (Nijmegen)
18-05-2022
Amendment
CMO regio Arnhem-Nijmegen (Nijmegen)

Date:	06-04-2023
Application type:	Amendment
Review commission:	CMO regio Arnhem-Nijmegen (Nijmegen)
Approved WMO Date:	01-08-2023
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 CCMO

ID NL76487.091.21