

# What are the causes of individual differences between children in terms of aggressive behavior and associated behavioural problems?

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<b>Registration date</b> 01/10/2019	<b>Overall study status</b> Completed	<input type="checkbox"/> Protocol
<b>Last Edited</b> 01/10/2019	<b>Condition category</b> Mental and Behavioural Disorders	<input type="checkbox"/> Statistical analysis plan
		<input type="checkbox"/> Results
		<input type="checkbox"/> Individual participant data
		<input type="checkbox"/> Record updated in last year

## Plain English summary of protocol

### Background and study aims

Childhood aggression and its resulting social impairment inflict a huge personal and financial burden on affected children, their relatives, peers and society as a whole. The prevalence of clinical aggression in children ranges from 2-16%, and early-onset childhood aggression continues into adolescence and adulthood in a substantial proportion of children. There are large differences between children in aggression levels and this study asks "what are the causes of individual differences between in children for aggressive behaviour and associated behavioural problems?". We look at genetic, epigenetic and metabolomics markers, in different cohorts that collected these data in birth cohorts and population-based registries. We combine information from cohorts in a series of meta-analyses.

### Who can participate?

Children below age 18 years for whom information on aggression and attention problems is available and who have been genotyped on a genome-wide SNP (Single nucleotide polymorphism) arrays; participants of any age with information on aggressive and attention problems and genome-wide epigenetic data; children for whom information on aggressive and attention problems is available and for whom urine samples for metabolomics were collected.

### What does the study involve?

Estimating the associations between the phenotypes (aggression and attention problems) and the biomarker data. Primary analyses are carried out by each cohort and results are combined in meta-analyses.

What are the possible benefits and risks of participating? This project generates information on associations of behaviour with biomarkers. There are no individual-level risks or benefits to participants.

### Where is the study run from?

The cohort studies run in their own countries (USA, UK, Europe, Australia and New Zealand). The meta-analyses is carried out in the Netherlands.

When is the study starting and how long is it expected to run for?  
The cohort studies run in their own countries and typically are longitudinal projects. The meta-analyses will be finished and submitted for publication in 2019.

Who is funding the study?  
FP7-EU 602768

Who is the main contact?  
Dr. Boomsma  
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**Study website**  
<http://www.action-euproject.eu/>

## Contact information

**Type(s)**  
Scientific

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## Additional identifiers

**EudraCT/CTIS number**  
Nil known

**IRAS number**

**ClinicalTrials.gov number**  
Nil known

**Secondary identifying numbers**  
Nil known

## Study information

**Scientific Title**

**ACTION: Aggression in Children: Unraveling gene-environment interplay to inform Treatment and InterventiON strategies**

## **Acronym**

ACTION

## **Study objectives**

Improve the understanding of the genetic and non-genetic etiology of aggression in children to inform the development of novel prevention and treatment strategies by unraveling in large twin and genotyped cohorts the causes of variation in aggression, disentangling (epi)genetic and environmental effects and their interplay with a focus on critical developmental periods, gender, and comorbid disorders; by investigating metabolomic profiles of aggressive behavior to establish direction of causation for existing and new biomarkers and gain insight regarding the predictive power of pediatric aggression for adult outcome variables.

## **Ethics approval required**

Old ethics approval format

## **Ethics approval(s)**

Meta-analysis of existing cohort studies does not require ethical approval (confirmed by Central Ethics Committee on Research Involving Human Subjects of the VU University Medical Centre; BS7 Kamer H-565I; +31 (0)20 44 45585; metc@vumc.nl), ref: 2014.252

## **Study design**

Epidemiological meta-analysis

## **Primary study design**

Observational

## **Secondary study design**

Epidemiological study

## **Study setting(s)**

Community

## **Study type(s)**

Other

## **Participant information sheet**

<http://www.action-euproject.eu/content/data-protocols>

## **Health condition(s) or problem(s) studied**

Childhood behavioral problems, with a focus on aggression and attention problems

## **Interventions**

The researchers are doing a series of large meta-analyses, which either take cross-sectional or longitudinal data and test associations of trait outcomes (aggression and attention problems) with SNP data, epigenetics and metabolomics markers. In this work there is no follow-up (though the data mainly derive from ongoing population-based cohorts).

Data were drawn from the following cohorts:

Amsterdam Born Children and their Development Study  
Avon Longitudinal Study of Parents and Children  
Brain dEvelopment and Air polluTion ultrafine particles in scHool childrEn  
Child and Adolescent Twin Study in Sweden  
Christchurch Health and Development Study  
Collaborative Studies on the Genetics of Alcoholism  
Copenhagen Prospective Studies on Asthma in Childhood 2010  
Dunedin Multidisciplinary Health and Development Study  
Environmental Risk Longitudinal Twin Study  
Finnish Twin Cohort  
Generation R Study  
German Infant study on the influence of Nutrition Intervention PLUS environmental and genetic influences on allergy development / The influence of Life-style factors on the development of the Immune System and Allergies in East and West Germany  
Great Smoky Mountains Study  
Institute for Behavioral Genetics  
Infancia y Medio Ambiente  
Impact of Neurodevelopmental disorders and School performance: genes and environment  
Minnesota Center for Twin and Family Research  
Norwegian Mother and Child Cohort Study  
Michigan State University Twin Register  
Mater University of Queensland Study of Pregnancy  
Northern Finland Birth Cohort 1986  
Netherlands Twin Register  
Queensland Institute of Medical Research  
Western Australian Pregnancy Cohort (Raine) Study  
Swedish Twin study of CHild and Adolescent Development  
Twin Early Development Study  
TRacking Adolescents' Individual Lives Survey  
Virginia Twin Study of Adolescent Behavioral Development  
Young Finns Study

## **Intervention Type**

Other

## **Primary outcome measure**

Measures of association (from regression analyses) for biomarker-outcome analyses. For each set of biomarkers (metabolomics, SNP and epigenetic data) an appropriate level of statistical significance is specified. Outcome data on problem behaviors were collected by birth and child cohorts by standardized surveys.

The most commonly employed instruments to assess childhood aggressive and attention problems came from the Achenbach System of Empirically based Assessment (ASEBA; Achenbach et al. 2017) and the Strengths and Difficulties Questionnaire (SDQ; Goodman 2001). These two instruments accounted for > 70% of the phenotype data.

-Achenbach TM, Ivanova MY, Rescorla LA (2017) Empirically based assessment and taxonomy of psychopathology for ages 1½–90+ years: Developmental, multi-informant, and multicultural

findings. Compr Psychiatry 79:4–18. doi: 10.1016/J.COMPPSYCH.2017.03.006  
- Goodman R (2001) Psychometric Properties of the Strengths and Difficulties Questionnaire. J Am Acad Child Adolesc Psychiatry 40:1337–1345. doi: 10.1097/00004583-200111000-00015

### **Secondary outcome measures**

Based on primary analyses, gene-based and network tests were carried out.

### **Overall study start date**

01/06/2014

### **Completion date**

01/06/2019

## **Eligibility**

### **Key inclusion criteria**

Biomarker- outcome associations are tested in child (2-18 years) cohorts and for epigenetics also in adults. Inclusions into the original population-based cohort studies were voluntary.

### **Participant type(s)**

Mixed

### **Age group**

All

### **Sex**

Both

### **Target number of participants**

For metabolomics the target number of participants was 1600. For genetics and epigenetics meta-analyses the target was the largest possible number of participants.

### **Key exclusion criteria**

For genetic studies, to avoid population stratification, ancestries that are different from EU are excluded

### **Date of first enrolment**

01/06/2014

### **Date of final enrolment**

01/06/2019

## **Locations**

### **Countries of recruitment**

Australia

Denmark

Finland

Germany

Netherlands

New Zealand

Norway

Sweden

Switzerland

United Kingdom

United States of America

**Study participating centre**

**Vrije Universiteit**

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## **Sponsor information**

**Organisation**

Vrije Universiteit - FGB

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University/education

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**ROR**

<https://ror.org/008xxew50>

# Funder(s)

## Funder type

Government

## Funder Name

Seventh Framework Programme

## Alternative Name(s)

EC Seventh Framework Programm, European Commission Seventh Framework Programme, EU Seventh Framework Programme, European Union Seventh Framework Programme, EU 7th Framework Programme, European Union 7th Framework Programme, Siebten Rahmenprogramm, Séptimo Programa Marco, Septième programme-cadre, Settimo programma quadro, 7th Framework Programme, Seventh EU Framework Programme, FP7

## Funding Body Type

Government organisation

## Funding Body Subtype

National government

## Location

# Results and Publications

## Publication and dissemination plan

Planned publication in a high-impact peer-reviewed journal

## Intention to publish date

01/12/2019

## Individual participant data (IPD) sharing plan

Over 30 cohorts are included in this effort, which each have their own repositories. The association results from all cohorts are uploaded to a central repository in the Netherlands. The full set of meta-analysis association results will be made available upon publication of the papers.

## IPD sharing plan summary

Stored in repository