



Our partners

French rare diseases networks



Our partners



Our institutional partners



French and international patient associations (non-exhaustive list)



Contact us

For more information and to access the data:

Email: genida@igbmc.fr

Website: <https://genida.unistra.fr/>

Genida International Project

GenIDA_project

Genida project

Scan this QR code to discover GenIDA



GenIDA

Platform for participatory clinical research on genetic forms of intellectual disability, autism and epilepsy

<https://genida.unistra.fr/>



Intellectual disability (ID) with or without manifestations of autism spectrum disorders (ASD) and/or epilepsy affects 1-2% of the population, and it is estimated that more than 50% of these cases have a single genetic cause (mutation in a gene, chromosomal abnormality or copy number variation / CNV).

Our understanding of the genetic causes of neurodevelopmental disorders (NDDs) has improved tremendously over the past decade, especially with the use of high-throughput sequencing since 2012.

More than 1000 genes and recurrent chromosomal abnormalities are involved in these genetic forms of ID or ASD, which often remain insufficiently described in terms of clinical spectrum, associated medical problems and natural history, due to their rarity and the often limited number of patients observed.



For instance, it required 10 years of diagnostic and phenotypic study of families affected by Fragile X syndrome worldwide (1991-2001)¹ to establish, from only 5 initial cases, that male carriers of the Fragile X premutation can present with a late-onset neurodegenerative disorder², and a further 3 years were needed to collect data to estimate the age-related penetrance of the disease³.

In such a context, how can patient cohorts be efficiently constructed to obtain sufficient and complete data to identify information of medical interest to families and professionals?



Professionals (medical and paramedical), create your profile and select your cohort(s) of interest in order to participate in the GenIDA study and access the medical data collected.



Access to the visualization of the results of our statistical analyses, according to your cohorts of interest.



More complete access to anonymised patient records upon request.



Secure data storage and exchange in compliance with the GDPR and French regulations.

Use GenIDA to:

- generate new and medically significant knowledge that can be translated into improved patient management (scientific publications, recommendations / guidelines, etc.);
- submit additional specific questions to subsets of patients;
- recruit patients for ethically approved research projects or clinical studies (subject to approval by GenIDA's Scientific Advisory Board).

Bibliography:

- Coutelle, Boedec, [...], Burger & Mandel (2022). The impact of lockdown on young people with genetic neurodevelopmental disabilities: A study with the international participatory database GenIDA. **BMC Psychiatry**, doi.org/10.1186/s12888-022-04213-6
- Burger, Colin, [...], Koolen & Mandel (2023). GenIDA: an international participatory database to gain knowledge on health issues related to genetic forms of neurodevelopmental disorders. **Journal of Neural Transmission**, doi.org/10.1007/s00702-022-02569-3
- Colin*, Burger*, [...], Koolen & Mandel (2023) GenIDA, an international participatory study of medical and natural history data in genetic forms of neurodevelopmental disorders: novel observations in a large cohort of patients with Koolen-de Vries syndrome. **Genetics in Medicine Open**, doi.org/10.1016/j.gimo.2023.100817
- Forbes, Morison, [...], Burger, [...], Mandel, [...], Morgan (2024) Speech and Language in DDX3X - neurodevelopmental disorder: a call for early augmentative and alternative communication intervention. **American Journal of Medical Genetics B**, doi.org/10.1002/ajmg.b.32971

- Access to all our publications from our website -

¹Rousseau et al. 1991, ²Hagerman et al. 2001, ³Jacquemont et al. 2004

Families, professionals, you want to participate?



It is easy > go to <https://genida.unistra.fr/> from a computer or a tablet

The registration procedure and access to the GenIDA study are presented in this video: <https://youtu.be/-8eJD9Chbe4>



Families, create your personal space and read the consent form to be able to participate in the GenIDA study by providing and regularly updating medical and quality of life information about the patient.

Consent to be given at the time of registration.
Your data are anonymised and hosted securely.

Project declared to the French Commission on Information Technology and Liberties (n°1907912) and approved by the Ethical Evaluation Committee of INSERM - CEEI-IRB (n°16-338).

What is the benefit to the participating families?

As soon as the number of participants in a cohort is sufficient, we make the anonymised results of our analyses available.

The larger the number of participants, the better the quality and usefulness of the results for families and professionals (doctors, researchers, etc.).

You can download a single PDF containing all your answers, which can be added to the patient's medical file.

GenIDA

GenIDA* is an **international participatory database** initiated in 2016, with the aim of **better characterising the clinical manifestations and natural histories of these genetic forms of ID and/or ASD**.

The aim of this project is to **accelerate knowledge about these rare diseases** by **strengthening the participation of affected individuals, their families and the associations concerned** in order to **create international cohorts of patients of sufficient size** so that doctors, researchers and other professionals can **extract new medically significant data** to improve the care of affected individuals.

GenIDA also enables longitudinal studies to be carried out, in particular to monitor the evolution of cognitive skills and possible behavioural problems in people with the disease.



Collection of health and behavioural data



Statistical analysis of data and synthesis of information



Sharing data with patients/families & professionals



Promoting interaction between patients / families & professionals



Generating new and medically meaningful data

In order to do this, patients and their families are asked to answer a structured questionnaire currently available in 8 languages covering physical, medical, cognitive and behavioural aspects of the disease.

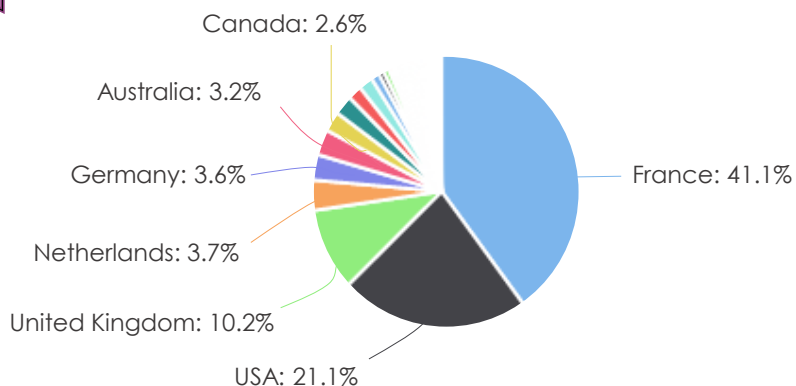
GenIDA, July 2024



> 1980 completed patient records
& > 250 professionals (clinicians, geneticists, etc.) registered



> 60 nationalities represented:



Selected cohorts in GenIDA:

COHORT	GENE / GENETIC DEFECT	NB OF PARTICIPANTS
Koolen-de Vries syndrome	<i>KANSL1</i> & 17q21.31 deletion	276
Kleefstra syndrome	<i>EHMT1</i> & 9q34.3 deletion	222
DDX3X	<i>DDX3X</i>	69
Rasopathies	<i>PTPN11</i> , <i>BRAF</i> , <i>KRAS</i> , etc.	68
KBG syndrome	<i>ANKRD11</i>	57
MED13L	<i>MED13L</i>	50
SETD5	<i>SETD5</i>	37
DYRK1A syndrome	<i>DYRK1A</i>	35
White-Sutton syndrome	<i>POGZ</i>	35

Collaborations

GenIDA was initially developed in collaboration with Radboud University Medical Centre, Nijmegen, The Netherlands (Pr Tjitske Kleefstra, Dr David A. Koolen, Dr Charlotte Ockeloen, Pr Bert B. A. de Vries).

GenIDA collaborates with many professionals in order to develop new cohorts and to study the data thus collected in order to bring out new information of medical interest, allowing for better patient care (non-exhaustive list):

- Murdoch Children's Research Institute, Melbourne, Australia (Pr. A. T. Morgan);
- Royal Manchester Children's Hospital, Manchester, UK (Pr. B. Kerr);
- St Georges Hospital Medical School, London, UK (Pr. M. Patton);
- The Goldschleger Eye Institute, Dept of Ophthalmology, Sheba Medical Center, Tel Aviv University, Israel (Dr. D. Landau Prat);
- ErasmusMC Rotterdam, The Netherlands (Pr T. Kleefstra);
- Dijon University Hospital (Pr L. Faivre-Olivier);
- Lille University Hospital (Pr J. Ghomid, Dr T. Smol, Dr R. Caumes);
- Hospices Civils de Lyon (Dr N. Chatron);
- Montpellier University Hospital (Pr D. Geneviève, Dr V. Ruault);
- APHP, Paris (Pr A. Verloes);
- Strasbourg University Hospital (Dr A. Piton, Dr R. Coutelle, Dr A. de Saint-Martin, Dr E. Schaefer, Dr B. Durand, etc.).

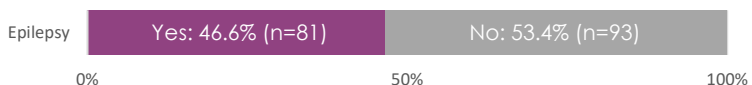


7th GenIDA Scientific Advisory Board, ESHG Berlin, June 2024.



KdVS / Epilepsy (study conducted in collaboration with Dr Nicole Collot, expert physician in pharmacovigilance):

The frequency, type of epilepsy and age of onset are consistent with previously published data⁸:



Study of the frequency of use of different anti-epileptic drugs, their perceived efficacy and associated side effects:

Anti-epileptic drugs	Drug use report	Good efficiency		Secondary effects	
		n/total	%	n/total	%
Levetiracetam	30	20/30	67%	14/30*	47%
Valproate	24	20/24	83%	10/24	42%
Oxcarbazepine	13	11/13	85%	1/13	8%
Topiramate	6	3/6	50%	2/6	27%
Carbamazepine	5	2/5	40%	3/5*	60%
Lamotrigine	5	4/5	80%	0/5	
Zonisamide	4	4/4		0/4	
Phenobarbital	3	2/3		0/3	
Lacosamide	2	2/2		0/2	
Phenytoin	2	2/2		0/2	
ACTH	1	0/1		1/1	
Vigabatrin	1	1/1		0/1	
Benzodiazepines					
Clobazam	7	6/7	86%	4/7	57%
Diazepam	4	4/4		1/4*	
Midazolam	3	3/3		0/3	
Clonazepam	2	1/2		1/2*	
Lorazepam	2	2/2		1/2	
Others	6	2/6		0/6	

* Major adverse effects reported

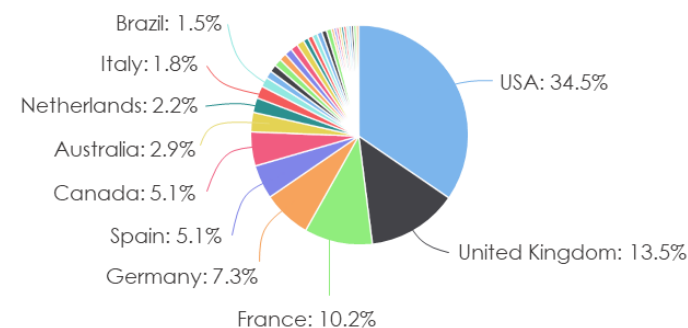
The two most commonly used anti-epileptic drugs are Levetiracetam and Valproate, with a trend (to be verified) towards better efficacy and lower adverse effects for Valproate. Oxcarbazepine, although less used, seems to have a good profile.

Type of data collected

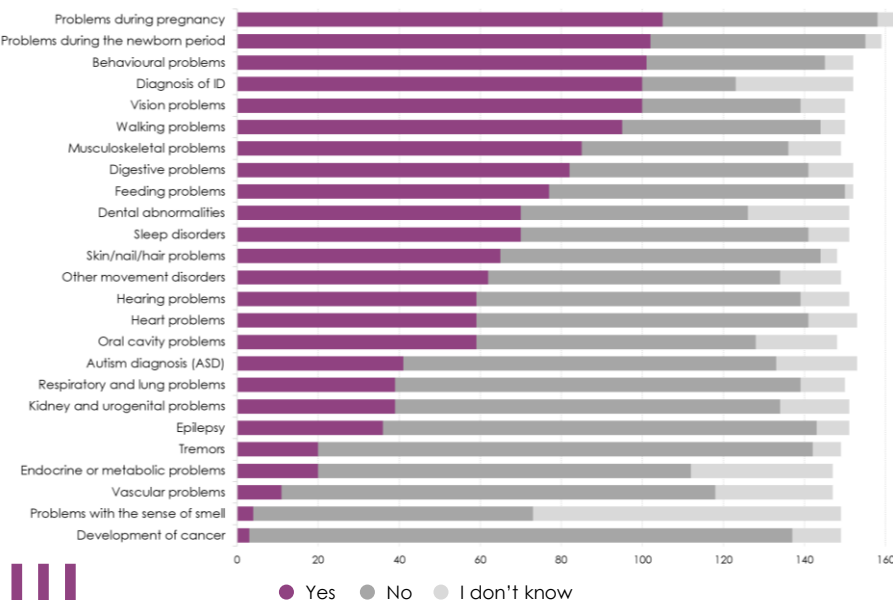
An overview of the data is automatically generated for each cohort (updated every 24 hours).

GenIDA - Kleefstra syndrome cohort (May 2022)

174 participants, including 90 women & 84 men



Overview:



⁸Myers et al. 2017

Validation of the GenIDA approach

The GenIDA data generally confirm the information reported in the literature, but also allow new observations.

Koolen-de Vries syndrome (KdVS) is caused by the 17q21.31 deletion or by a pathogenic variant of the *KANSL1* gene^{4,5,6}. The main features are intellectual disability, hypotonia, suggestive facial features, variable frequency of epilepsy, congenital anomalies and various neuromuscular and orthopaedic manifestations.

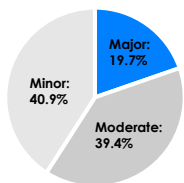
GenIDA - KdVS cohort (May 2022)

 235 participants, including  120 women &  115 men
195 patients (83%) have the 17q21.31 deletion, and 40 (17%) carry a pathogenic *KANSL1* variant.

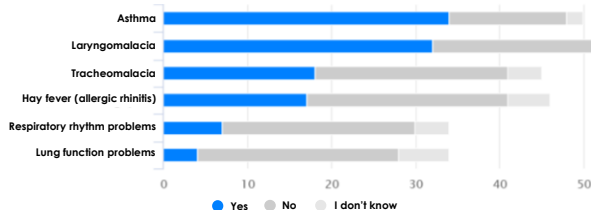
KdVS / Respiratory problems - New observations:

Respiratory problems briefly mentioned in Unique guidelines and one mention of pneumonia in Koolen et al. 2016⁴; no mention in OMIM, GeneReviews, or Zollino et al. 2015⁷.

Severity of respiratory problems reported in GenIDA



Respiratory problems (male & female)



Asthma and pneumonia or other respiratory infections are among the most reported comorbidities for KdVS in GenIDA and these problems are considered major by many families.

⁴Koolen et al. 2006, ⁵Koolen et al. 2012, ⁶Koolen et al. 2016, ⁷Zollino et al. 2015

Richness of open-ended answers from families:

f	10.0	Until the age of 6 she had several pneumonias and bronchial problems. She has recovered and is without problems now
m	8.0	Repetitive pneumonia
m	8.0	He had asthma triggered by respiratory infections from birth to around 9 years old. This subsided as he got older and is now completely gone.
m	11.0	Re-occurring pneumonia
f	4.5	Inflammation des bronches peut être due à un reflux
f	1.0	Classée comme asthme du nourrisson après 3 épisodes de bronchiolite. Traitement au flixotide d'octobre à mars
f	21.0	Système respiratoire fragile avec bronchite et laryngites
m	3.3	Broncopneumopatia cronica, ricoverata 10/12/2010 per 13 giorni in rianimazione per insufficienza respiratoria. Successivamente bronchiti ricorrenti [...].

Koolen-de Vries, Kleefstra (KS) and KBG (KBGS) syndromes - Behavioural problems

A lower frequency of behavioural problems was reported in GenIDA for KdVS (54%), compared to KS (70%) and KBGS (80%).

The KS had the highest score for repetitive/stereotypic behaviour, while the KBGS had the highest scores for attention deficit, impulsivity and aggressivity:

