



CELPHEDIA, a French Research Infrastructure, a reference center for animal research on rare diseases

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The best animal model to accelerate the comprehension of our genome and of human diseases

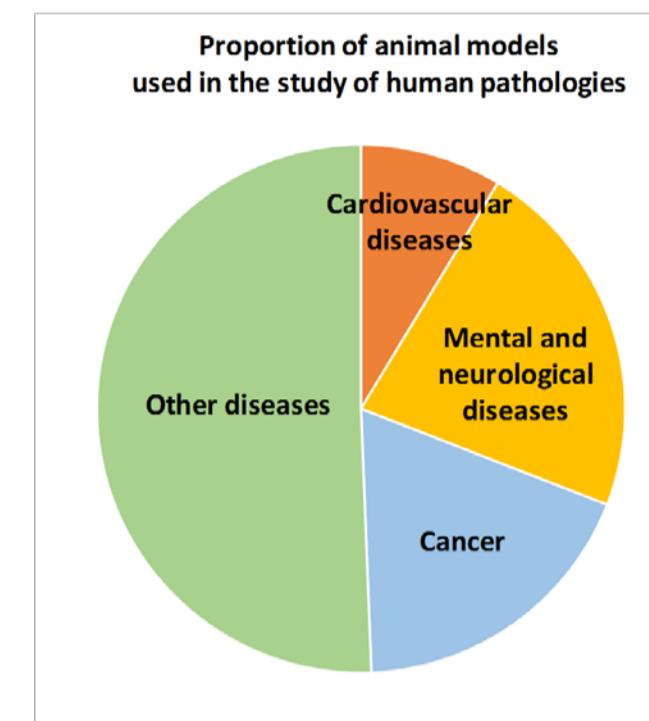
With 15 centers distributed over France, CELPHEDIA has developed innovative, standardized and massively parallel technological approaches:

- to accelerate the comprehension of the genome
- to generate human disease models
- to promote therapeutic innovations through validation of molecular targets

An unique access to 3 big families of model organisms to chose the best model to answer the questions of modern biology.

15 distributed centers, involved in 2 French infrastructures
phenomin & tefor
and in a European infrastructure INFRAFRONTIER

- 3 big families of organism models close to the users
5 ● Non mammals (TEFOR)
6 ● Mouse - Rat (PHENOMIN+3)
1 ● Other rodents
3 ● Non human primates



CELPHEDIA, a combination of expertise and skills

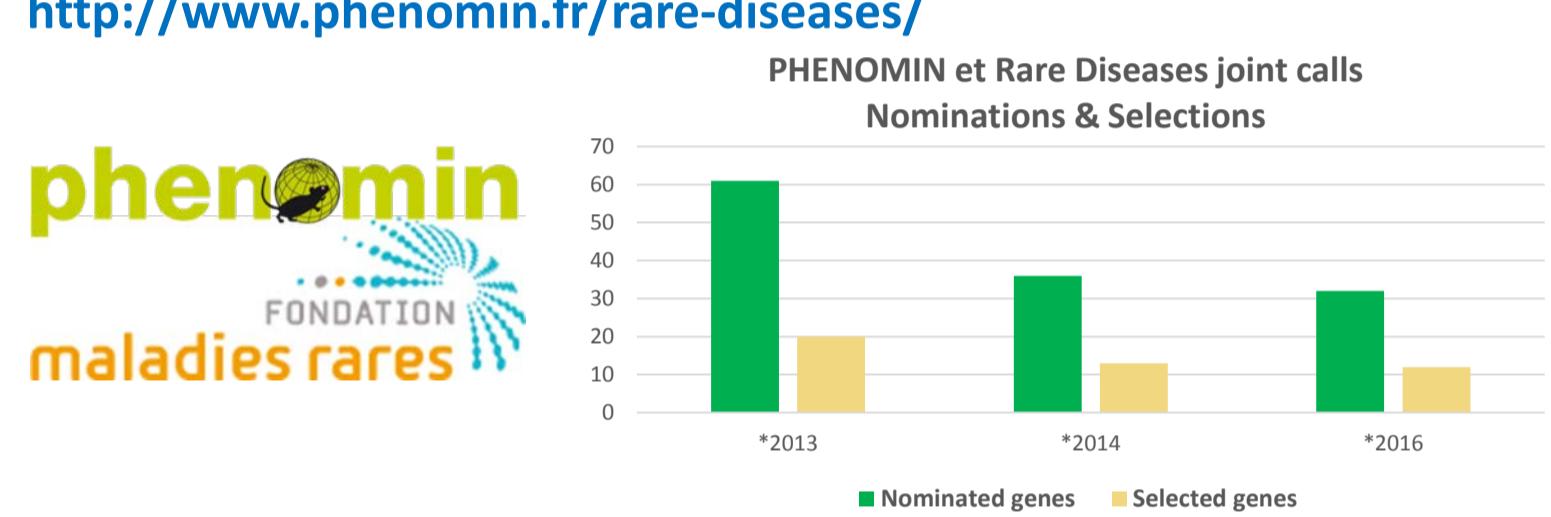


A remarkable palette of expertise, skills and knowledge unique in Europe. This unique access enables:

- scientific and technological multi-model approaches
- integrated comparative functional analyzes
- better cross-functionality of the results from a model organism to another.

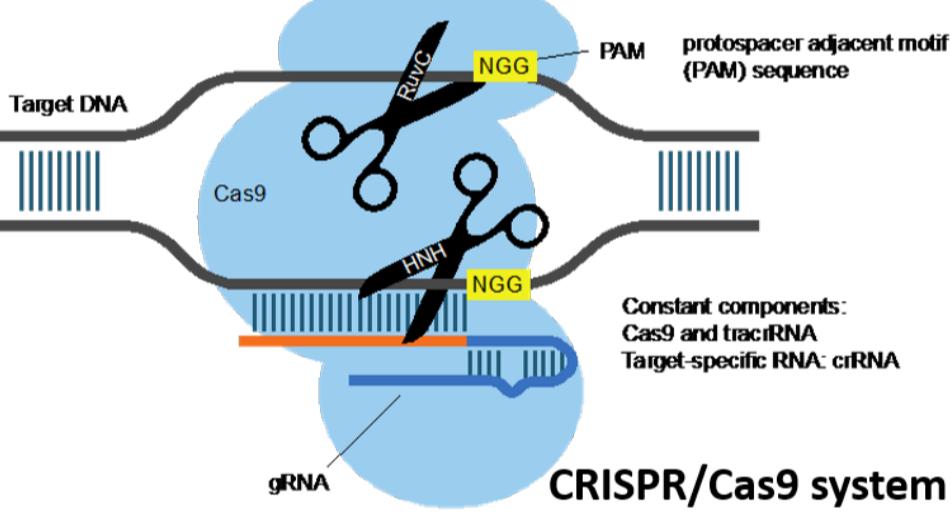
Mouse models and Rare Diseases

Rare Disease Foundation and PHENOMIN have launched 3 calls for joint research projects since 2013.
<http://www.phenomin.fr/rare-diseases/>



Genome modification and creation of models

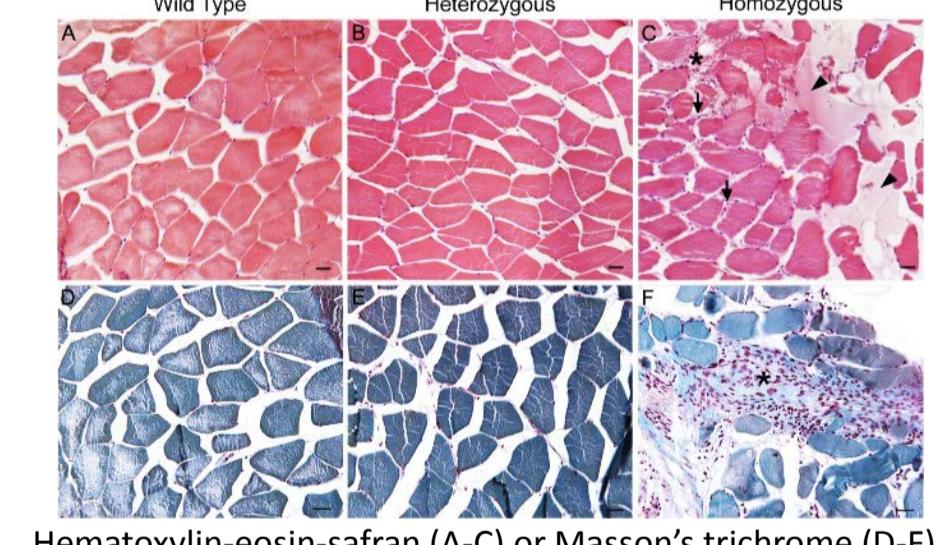
Design and production of genetically modified models, humanized mouse models, human pathology models, immunodeficient mice and rats :



The zebrafish to model human rare diseases

Bethlem myopathy : an incurable human collagen VI-related disease
The zebrafish line reproduces a human mutation within an essential splice donor site of *col6a1* gene using TALE nucleases, provoking an in-frame skipping of exon 14.

- Progressive disorganization of the muscle
- Co-dominantly inherited abnormal myofibers
- Enlarged sarcoplasmic reticulum
- Altered mitochondria
- Misaligned sarcomeres
- Development of fibrosis (* on C & F)
- Hypoxia-response behavior (locomotion tests)

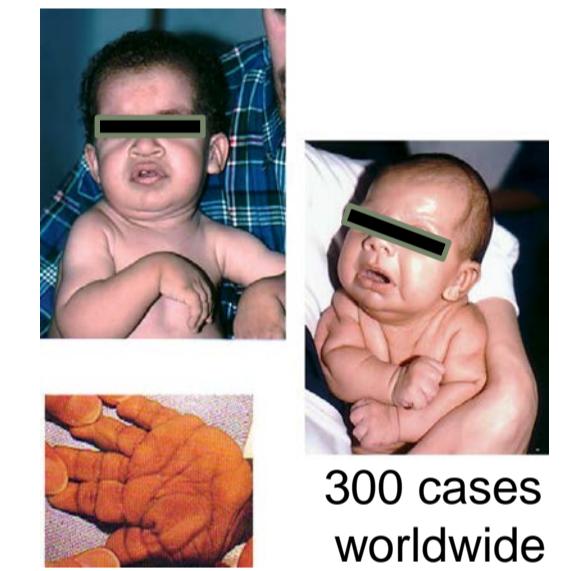


Radev Z, et al. PLoS One. 2015 - F. Sohm's team

The mouse to model human rare diseases

Costello syndrome: mouse model HRAS G12S reproduces most of the phenotypic characteristics observed in patients

CLINICAL SIGNS	« COSTELLO » PATIENTS	« COSTELLO » MICE
POST-NATAL GROWTH DELAY		?
RHABDOMYOSARCOME		?
HAIR/FUR ABNORMALITIES		?
SKIN ABNORMALITIES		?
FACIAL FEATURES/ MACROCEPHALY		
HEART ABNORMALITIES		
REDUCED MUSCLE STRENGTH		
REDUCED LOCOMOTOR ACTIVITY		
REDUCED COORDINATION		
LEARNING		



T. Sorg's team EXCELLENCE IN MOUSE PHENOMICS

RTHα genetic disease: mutations in the THRA gene coding the thyroid hormone receptor TRα1 generating various phenotypes



F. Flamant's team, UMS3444/US8 SFR BioSciences, Lyon

Generation of genetically modified mice at helix 12 of the receptor by CRISPR/Cas9

human & mouse wt THRA
NHRKHNIPHFWPKLILMKVTDLRMIAGACHASRFLHMVKCPTELFPPLFLEVQED
Helix 11
Helix 12

human mutations in C-term :
E403X NHRKHNIPHFWPKLILMKVTDLRMIAGACHASRFLHMVKCPTELFPPLFLEVQED
E403K NHRKHNIPHFWPKLILMKVTDLRMIAGACHASRFLHMVKCPTELFPPLFLEVQED
F397fs406X NHRKHNIPHFWPKLILMKVTDLRMIAGACHASRFLHMVKCPTELFPPLFLEVQED
P398R NHRKHNIPHFWPKLILMKVTDLRMIAGACHASRFLHMVKCPTELFPPLFLEVQED

CELPHEDIA at an international level

Involvement in European and international projects

Celphedia

Distributed French infrastructure
Reference center for animal research

phenomin

French infrastructure of phenogenomic

INFRAFRONTIER

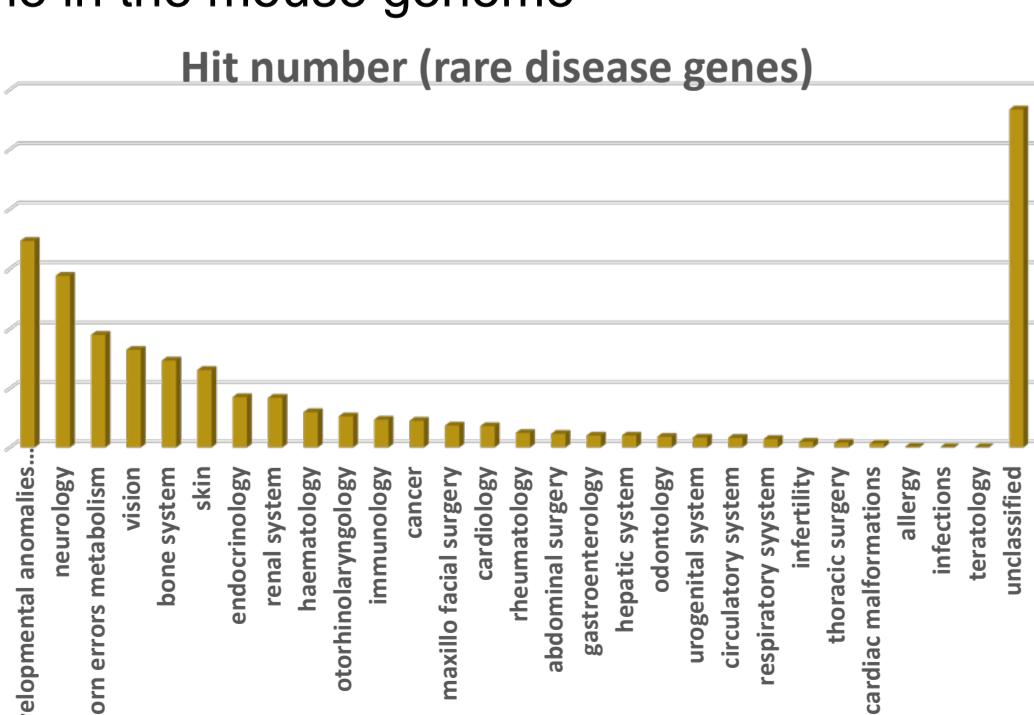
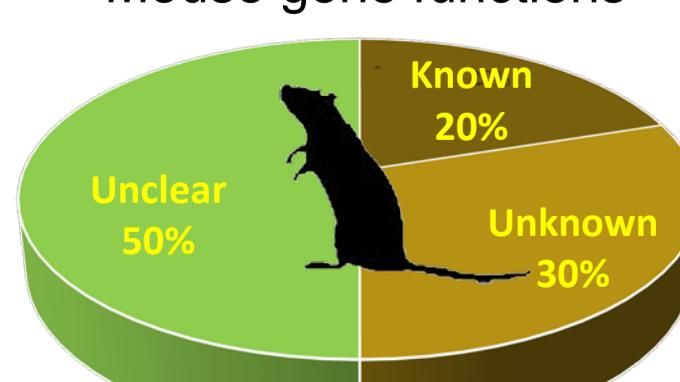
European network of phenotyping, archiving and distribution of mouse models

IMPC

International consortium of mouse phenotyping
9 countries
16 mouse clinics

- To undertake broad based primary phenotyping of about 20,000 mutants
- To determine the function of every gene in the mouse genome

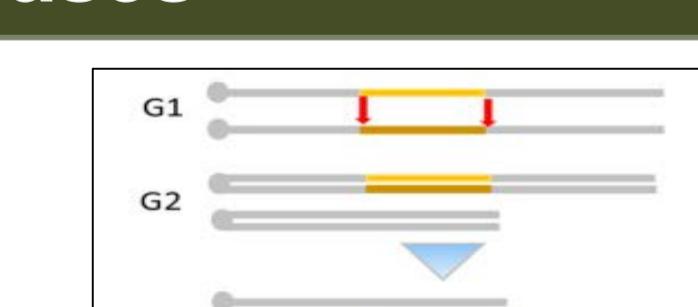
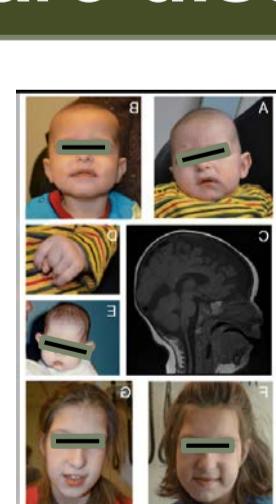
Mouse gene functions



The rat to model human rare diseases

MRD7 syndrome

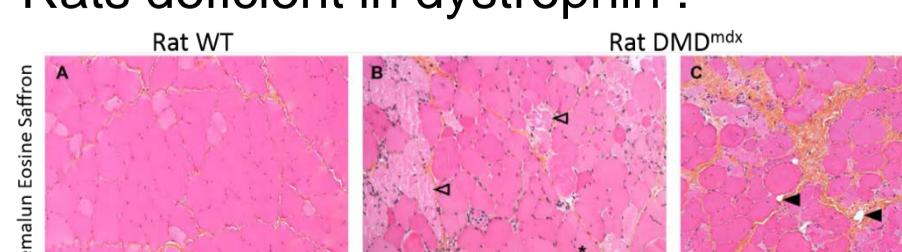
Clinical signs identified from 7 months to 7 years: microcephaly, growth delay, skeletal abnormalities, difficulty with nutrition, language delay, intellectual deficit, anxiety, aggressiveness, autism...



Rat model MRD7: Inactivation of a *Dyrk1a* allele with deletion of a gene sequence by CRISPR/Cas9

Duchenne myopathy

Rats deficient in dystrophin :



Hemalum Eosin Saffron
Rat WT
Rat DMD^{mdx}
A 3 month
B 7 month
← centers of small regenerating centronucleated fibers;
→ fibrous necrosis; ★ fibrosis; ▲ adipose tissue

I. Anegon's team

	DMD patient	GRMD dog	DMD pig	HFMD cat	Mdx mouse	Dmd ^{mdx} rat
Revertant fibers	1 to 3%	<1%	ND	ND	5%	>5%
Necrotic fibers	0.5 to 3.5%	2%	absent	present	5%	10%
Regeneration	ND	15%	ND	present	10%	10%
Calification	mild	mild to marked	ND	severe	mild	absent
Fibrosis	marked	present	present	mainly diaphragm	late & mainly absent	marked
Lipomatosis	severe	absent	absent	absent	mild	absent
Cardiomyopathy	marked, major cause of death	mild	absent	present	absent or late	marked
Strength reduction	marked	marked	ND	ND	mild	marked
Locomotion	severely impaired	impaired	impaired	ND	normal	impaired

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