

# Ricerca clinica e nuove prospettive terapeutiche per la malattia di Huntington

Torino, 08/11/2019

**Dr Lorenzo Nanetti**  
UO Genetica Medica e Neurogenetica



Fondazione I.R.C.C.S.  
Istituto Neurologico Carlo Besta

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Sistema Socio Sanitario

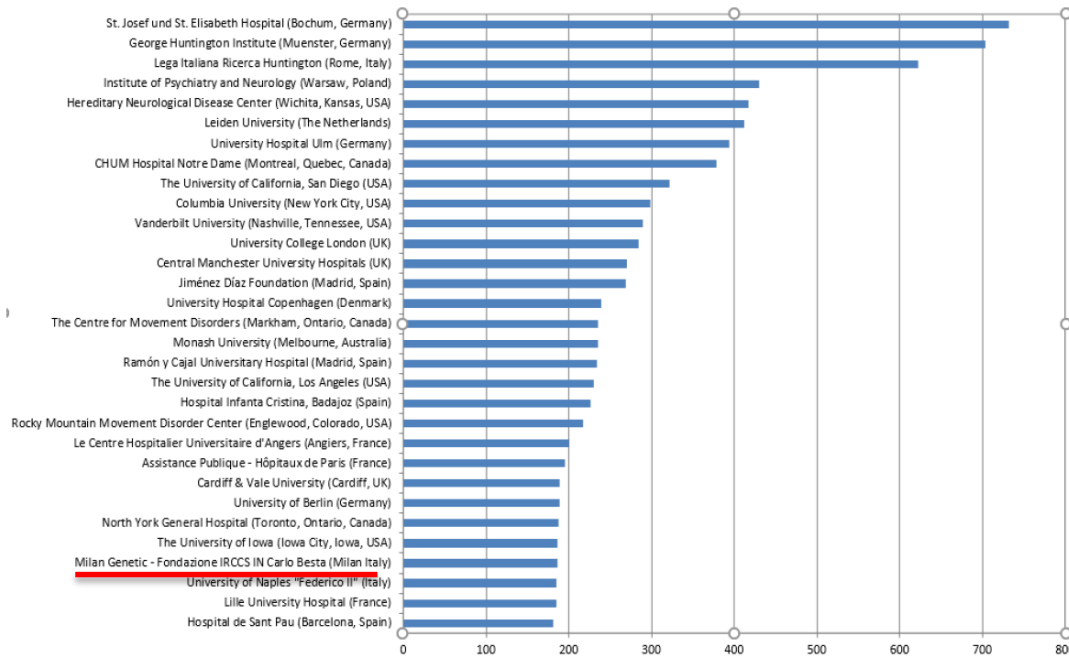


Regione  
Lombardia

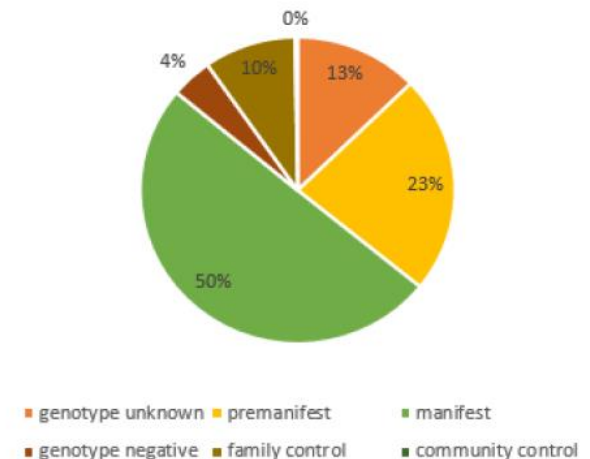
# Studi osservazionali per caratterizzare la storia naturale della malattia di Huntington



Top Enrolling Sites - Since Study Start (Active Participants)



>20000 soggetti arruolati nel mondo



# European Reference Network for Rare Neurological Diseases



PUBLIC HEALTH

European Commission > DG Health and Food Safety > Public health > European Reference Networks > Networks implementation > Call

## EUROPEAN REFERENCE NETWORKS

- **25 Centres from 14 EU countries**

Innsbruck, Milan, Brescia, Siena, Roma, Nijmegen, Amsterdam, Groningen, Paris, Bordeaux, Brussels, Budapest, Pecs, Athens, Tübingen, Bonn, Ulm, Lübeck, Munich, Prague, London, Aarhus, Oslo, Ljubljana, Barcelona

- **Disease focus on rare movement and cognitive disorders**

- Ataxias/Hereditary Spastic Paraplegias
- Frontotemporal degeneration
- Dystonia / Paroxysmal (non-epileptic)/ NBIA
- Leukodystrophies and neurometabolic movement disorders
- **Huntington and Chorea**s
- Rare and atypical Parkinson (including MSA, PSP)

Web site: <http://et-agentur.de/ern-rnd/>

The screenshot shows the website [et-agentur.de/ern-rnd/disease-resources.php](http://et-agentur.de/ern-rnd/disease-resources.php). The header features the ERN RND logo and the text "Initiative for European Reference Network for Rare Neurological Diseases". The navigation menu includes "Network", "Disease resources", "Dissemination, Training & Education", "Services", and "Internal". The "Disease Resources" section is active, displaying a list of nine disease categories, each with a circular icon and a right-pointing arrow:

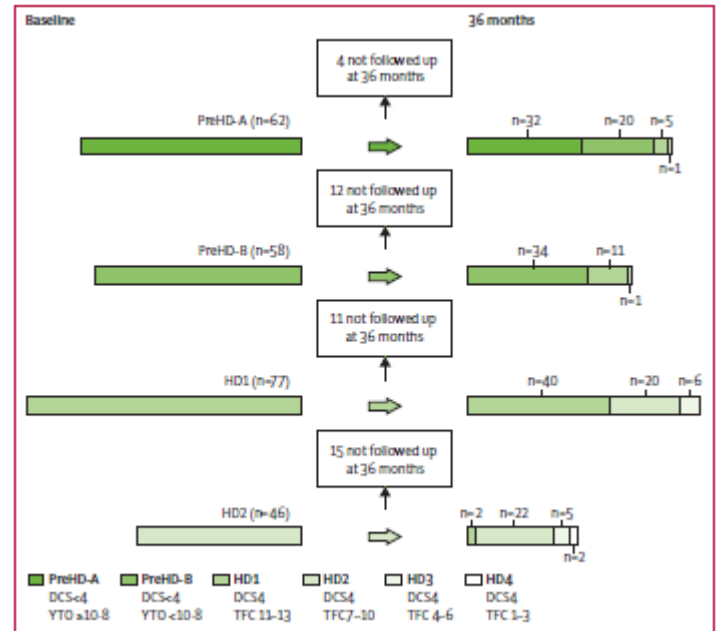
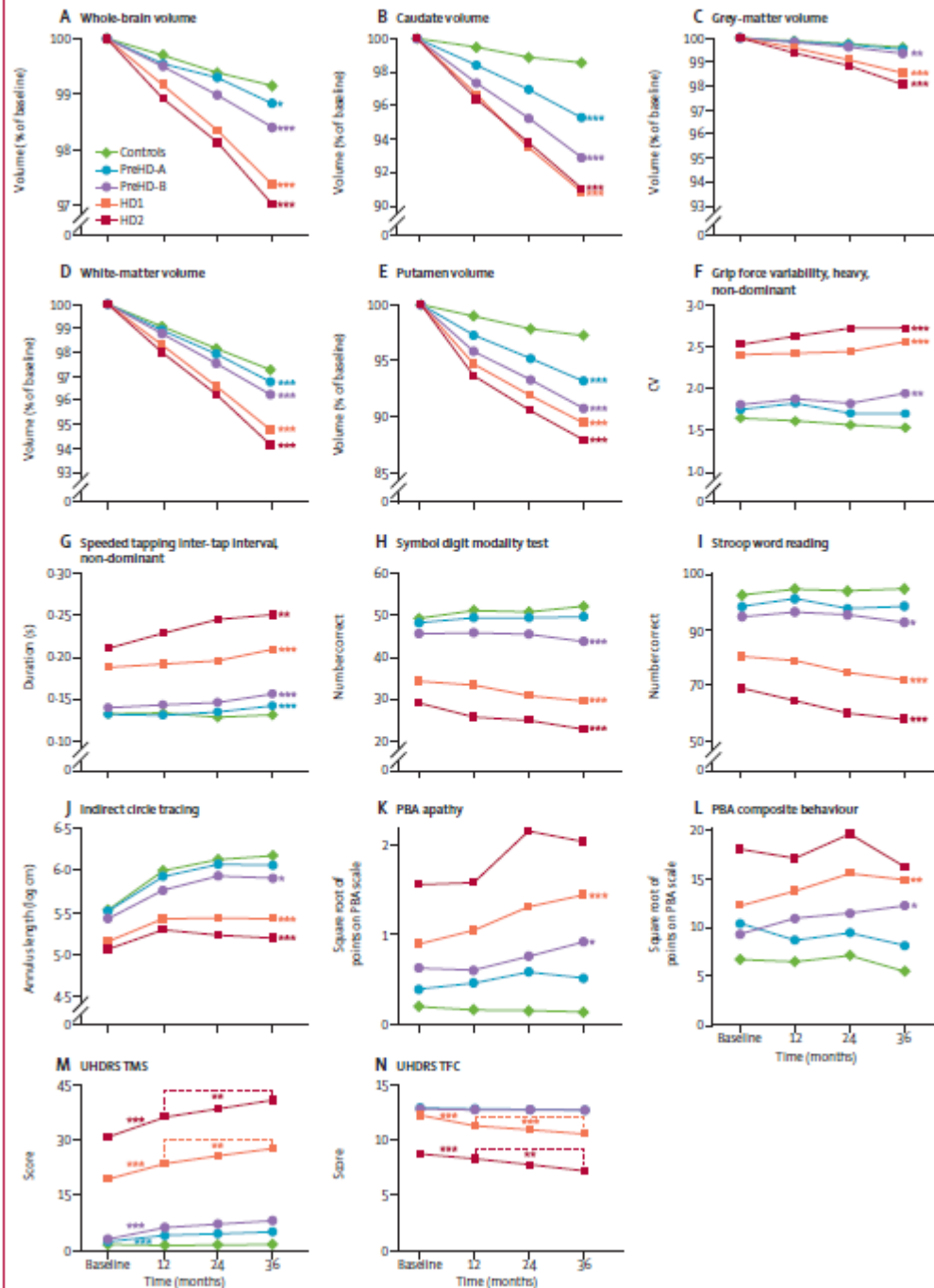
- Ataxias
- Chorea
- Dystonia
- FTD – Frontotemporal Dementia
- HSP – Hereditary Spastic Paraplegia
- Leukodystrophy
- MSA – Multiple System Atrophy
- NBIA – Neurodegeneration with Brain Iron Accumulation
- PSP – Progressive Supranuclear Palsy

- [European Network for the Study of Dystonia Syndromes](#),
- [Leukotreat](#),
- [SPATAX Network](#): Clinical and Genetic Analysis of Spastic paraplegia and Ataxia,
- [European MSA Study Group](#)
- [Brain-Team](#) - French RD network focused on brain diseases,
- [Ataxia Study Group](#),
- [European Huntington Disease Network, EHDN](#)
- [German Ataxia / HSP Network](#) /
- [Genetic FTD Initiative](#),
- [Progressive supranuclear palsy \(PSP\) study group](#)
- [NBIA network \(TIRCON\)](#)

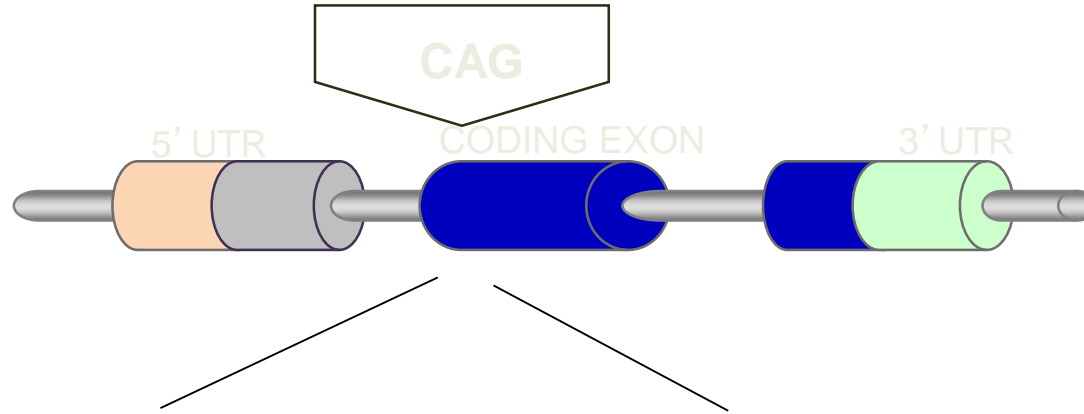
# Predictors of phenotypic progression and disease onset in premanifest and early-stage Huntington's disease in the TRACK-HD study: analysis of 36-month observational data

Sarah J Tabriz, Rachael I Sachill, Gail Owen, Alexandra Durr, Blair R Leavitt, Raymund A Roos, Beth Borowsky, Bernhard Landwehrmeyer, Chris Frost, Hans Johnson, David Craufurd, Ralf Reilmann, Julie C Stout, Douglas R Langbehn, and the TRACK-HD Investigators\*

Lancet Neurol 2013



# Huntington Disease: Expansion of triplet repeat CAG



DNA: Normal allele:  $\leq 26$  CAG

Mutated Allele:  $\geq 40$  CAG expansion

GTTTTC **CAG...CAG** CAACAG

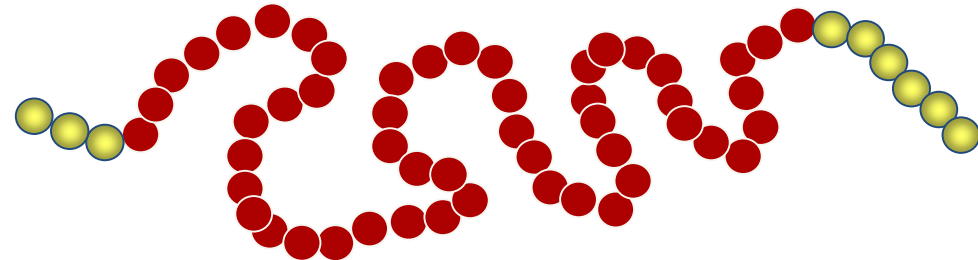
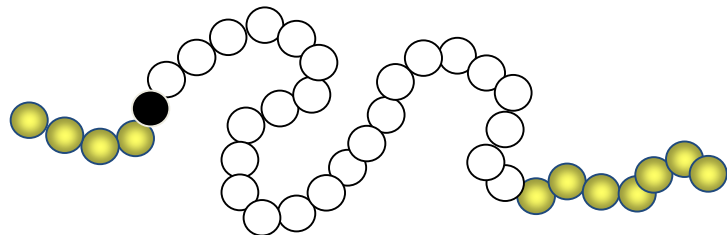
GTTTTC **CAGCAGCAG....CAGCAG** CAACAG

mRNA: Normal messenger RNA

mRNA: mutated messenger RNA

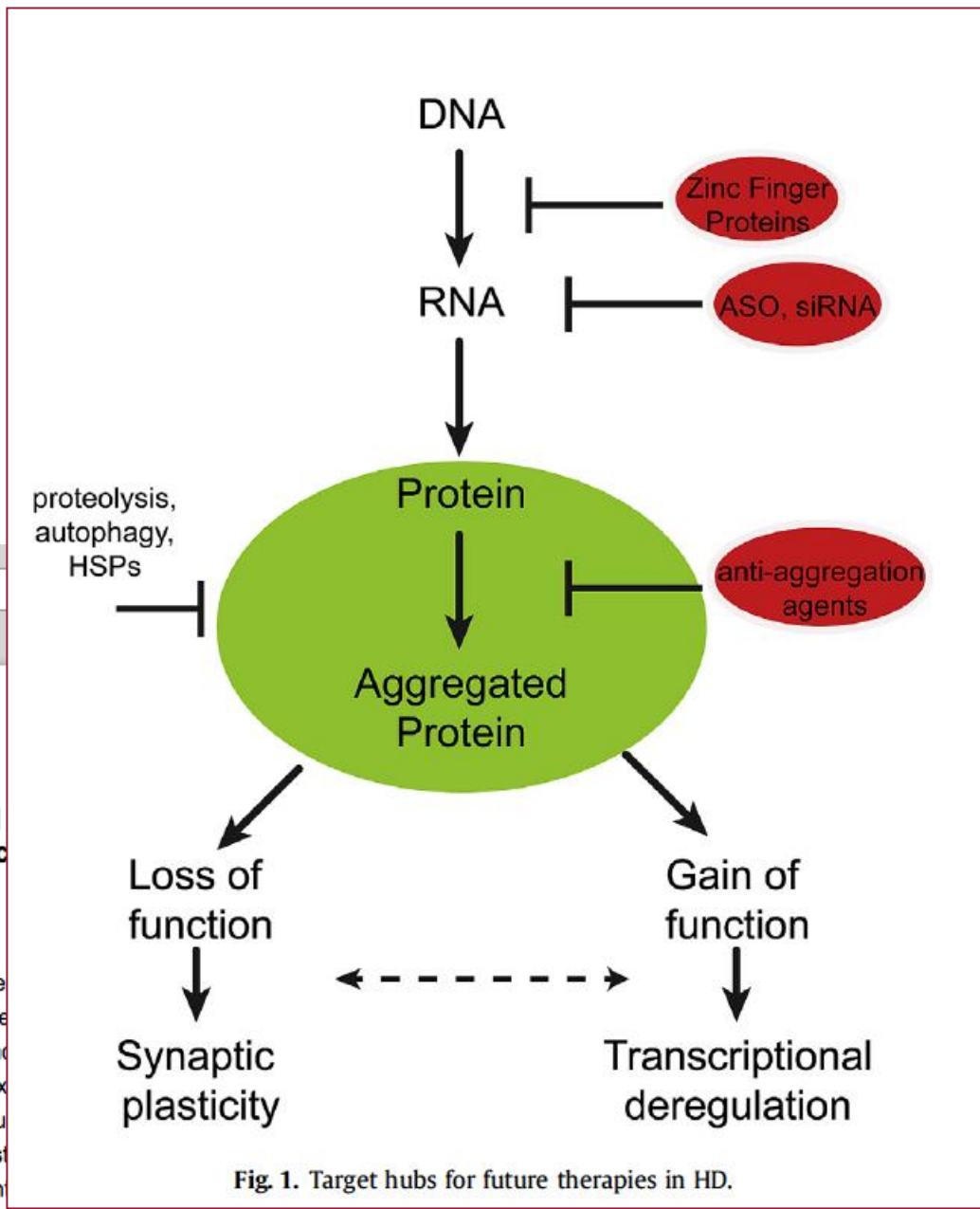
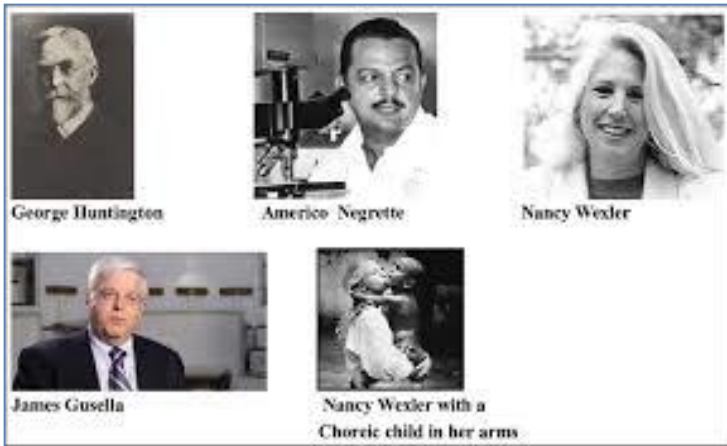
Normal (wild type) protein

Mutated protein



$\leq 26$  glutamine (Q)

Expanded Poly-Q



Abstract

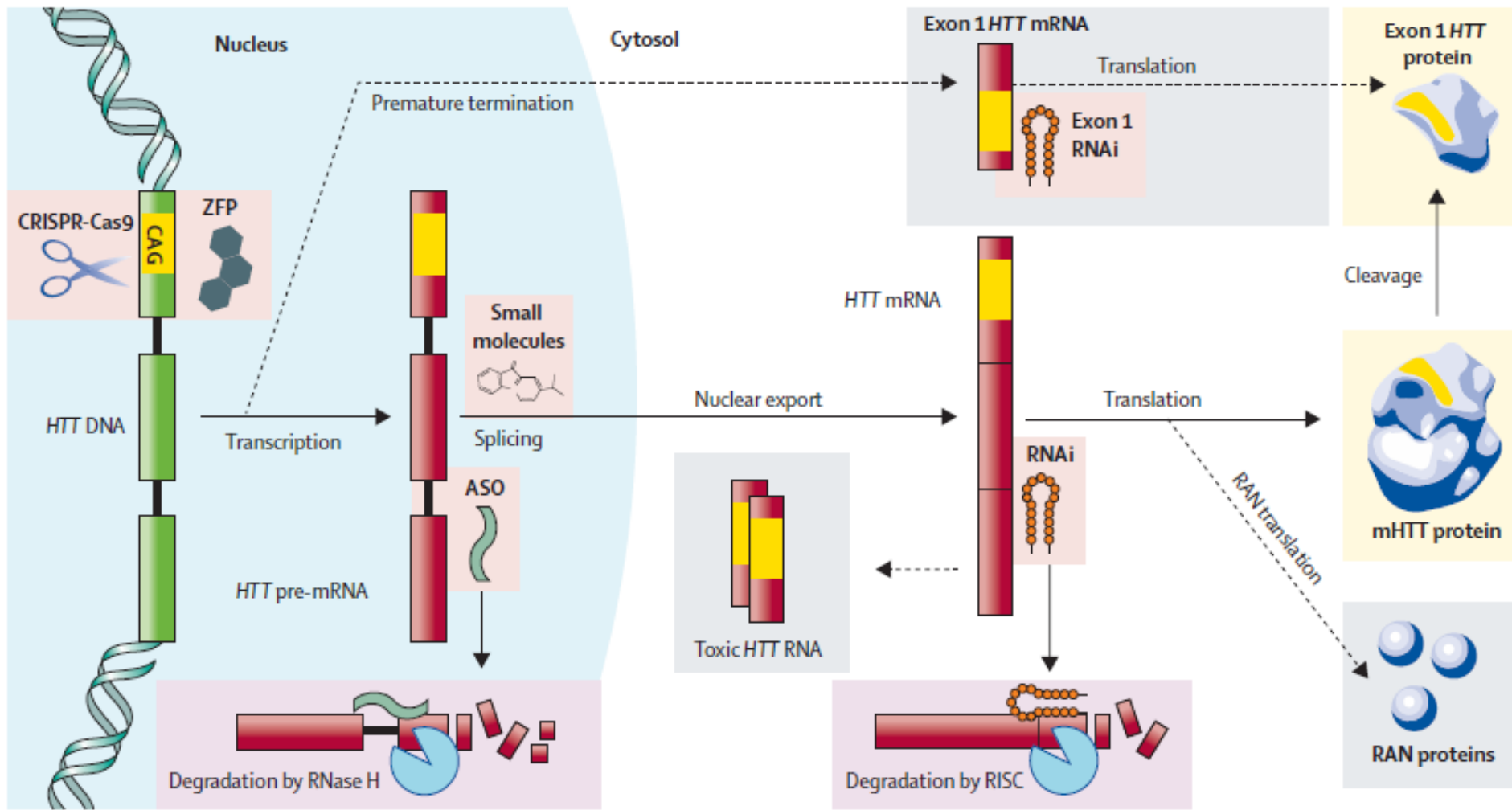
Cell. 1993 Mar 26;72(6):971-83.

**A novel gene containing a trinucleotide repeat that is expanded chromosomes. The Huntington's Disease Collaborative Research**

[No authors listed]

**Abstract**

The Huntington's disease (HD) gene has been mapped in 4p16.3 but has eluded identification. A small segment of 4p16.3 as the likely location of the defect from the target area contains a polymorphic trinucleotide repeat that is expanded and the normal range was observed on HD chromosomes from all 75 disease families examined. The (CAG)<sub>n</sub> repeat appears to be located within the coding sequence of an expressed but unrelated to any known gene. Thus, the HD mutation involves an unstable syndrome, spino-bulbar muscular atrophy, and myotonic dystrophy, acting in the con-







## The Nobel Prize in Physiology or Medicine 2006

"for their discovery of RNA interference - gene silencing by double-stranded RNA"



Photo: L. Cicero/Stanford

Andrew Z. Fire



Photo: R. Carlin/UMMAS

Craig C. Mello

The NEW ENGLAND JOURNAL of MEDICINE

ORIGINAL ARTICLE

# Nusinersen versus Sham Control in Infantile-Onset Spinal Muscular Atrophy

R.S. Finkel, E. Mercuri, B.T. Darras, A.M. Connolly, N.L. Kuntz, J. Kirschner, C.A. Chiriboga, K. Saito, L. Servais, E. Tizzano, H. Topaloglu, M. Tulinius, J. Montes, A.M. Glanzman, K. Bishop, Z.J. Zhong, S. Gheuens, C.F. Bennett, E. Schneider, W. Farwell, and D.C. De Vivo, for the ENDEAR Study Group\*

2017

## Potent and specific genetic interference by double-stranded RNA in *Caenorhabditis elegans*

Andrew Fire<sup>\*</sup>, SiQun Xu<sup>\*</sup>, Mary K. Montgomery<sup>\*</sup>, Steven A. Kostas<sup>\*†</sup>, Samuel E. Driver<sup>‡</sup> & Craig C. Mello<sup>‡</sup>

<sup>\*</sup> Carnegie Institution of Washington, Department of Embryology, 115 West University Parkway, Baltimore, Maryland 21210, USA

<sup>†</sup> Biology Graduate Program, Johns Hopkins University, 3400 North Charles Street, Baltimore, Maryland 21218, USA

<sup>‡</sup> Program in Molecular Medicine, Department of Cell Biology, University of Massachusetts Cancer Center, Two Biotech Suite 213, 373 Plantation Street, Worcester, Massachusetts 01605, USA

Experimental introduction of RNA into cells can be used in certain biological systems to interfere with the function of an endogenous gene<sup>1,2</sup>. Such effects have been proposed to result from a simple antisense mechanism that depends on hybridization between the injected RNA and endogenous messenger RNA transcripts. RNA interference has been used in the nematode *Caenorhabditis elegans* to manipulate gene expression<sup>3,4</sup>. Here we investigate the requirements for structure and delivery of the interfering RNA. To our surprise, we found that double-stranded RNA was substantially more effective at producing interference than was either strand individually. After injection into adult animals, purified single strands had at most a modest effect, whereas double-stranded mixtures caused potent and specific interference. The effects of this interference were evident in both the injected animals and their progeny. Only a few molecules of injected double-stranded RNA were required per affected cell, arguing against stoichiometric interference with endogenous

an Publishers Ltd 1998

NATURE | VOL 391 | 19 FEBRUARY 1998

	Allele selectivity	Delivery	Vector	Sponsor	Key advantages	Key disadvantages	References
<b>Antisense oligonucleotides</b>							
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Pre-mRNA degradation	SNP-targeted	Intrathecal	None	Wave Life Sciences (Cambridge, MA, USA)	Selective silencing of mutant allele	Several drugs required to treat most patients; SNP targeting limits choice of RNA-binding sequences	Hersch, 2017; <sup>28</sup> Butler, 2015 <sup>29</sup>
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mRNA degradation	None	Intracranial	AAV2	Spark (Philadelphia, PA, USA)	Single treatment provides sustained HTT reduction	Invasive delivery; limited treatment volume; cannot be reversed if adverse events occur	Harper, 2005; <sup>15</sup> Franich, 2008; <sup>16</sup> McBride, 2011 <sup>17</sup>
mRNA degradation	None	Intracranial	AAV1	Voyager (Cambridge MA, USA)	As above	As above	Stanek, 2015 <sup>31</sup>
mRNA degradation	None	Intracranial	AAV5	UniQure NV (Amsterdam, Netherlands)	As above	As above	Miniarikova, 2016; <sup>32</sup> Samaranch, 2017 <sup>33</sup>
<b>Small molecules</b>							
Screening programme*	Unknown	Potentially oral	None	CHDI Foundation (New York, NY, USA)	Potentially highly accessible route of delivery; potentially readily reversible	More difficult to achieve selectivity for HTT than with nucleotide approaches	Doherty, 2017 <sup>34</sup>

SNP= single nucleotide polymorphism. AAV1, 2, and 5= adeno-associated virus 1, 2, and 5. HTT= huntingtin protein. \*The mechanisms of action, route of delivery, and advantages and disadvantages of the small molecules being investigated by CHDI Foundation remain to be determined because the programme is currently at the phenotypic screen stage.

**Table 1: Huntington-lowering programmes targeting mRNA, by class and mechanism**

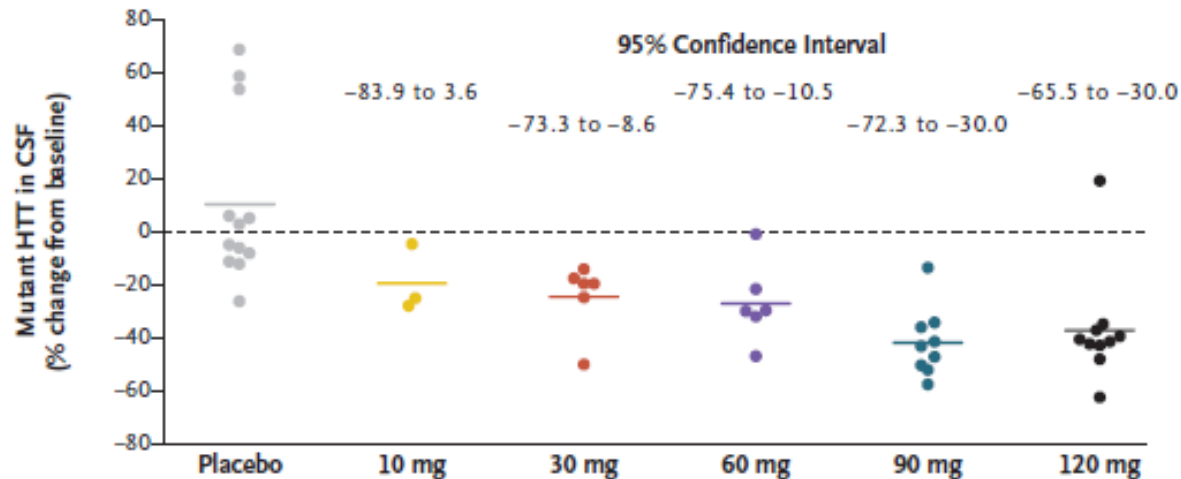
ORIGINAL ARTICLE

## Targeting Huntingtin Expression in Patients with Huntington's Disease

Sarah J. Tabrizi, M.B., Ch.B., Ph.D., Blair R. Leavitt, M.D., C.M., G. Bernhard Landwehrmeyer, M.D., Edward J. Wild, M.B., B.Chir., Ph.D., Carsten Saft, M.D., Roger A. Barker, M.R.C.P., Ph.D., Nick F. Blair, M.B., B.S.,\* David Craufurd, M.B., B.S., Josef Priller, M.D., Hugh Rickards, M.D., Anne Rosser, M.B., B.Chir., Ph.D., Holly B. Kordasiewicz, Ph.D., Christian Czech, Ph.D., Eric E. Swayze, Ph.D., Daniel A. Norris, Ph.D., Tiffany Baumann, B.S., Irene Gerlach, Ph.D., Scott A. Schobel, M.D., Erika Paz, B.S., Anne V. Smith, Ph.D., C. Frank Bennett, Ph.D., and Roger M. Lane, M.D., for the Phase 1–2a IONIS-HTT<sub>®</sub> Study Site Teams†

- ✓ **Fase I-II**
- ✓ **46 pazienti in doppio cieco (3:1)**
- ✓ **somministrazione intratecale**
- ✓ **4 dosi a cadenza mensile**

**B** Percentage Change in CSF Concentration of Mutant HTT, According to Dose Group



## Non-allele-selective ASO selected for clinical development

Data from RG6042 to date suggest the non-allele-specific approach is well tolerated and has the broadest patient eligibility

- Non-allele-specific approach preferentially developed due to:
  - broad eligibility for all HD patients irrespective of individual SNP
  - ability to screen the entire *HTT* gene to identify a highly potent ASO with favorable safety profile

### Preclinical safety

- Lowering of total HTT in the CNS with irreversible (e.g. siRNA) or reversible (e.g. ASO) approaches appear safe in normal animals<sup>1-3</sup>

### Preclinical efficacy

- Non-allele-specific ASOs have demonstrated efficacy in transgenic animal models, similar to allele selective approaches<sup>1,2,4,5</sup>

### Pharmacology

- RG6042 results in the dose-titratable, partial and reversible reduction of HTT
- Approach appears well tolerated in Phase I/IIa and OLE
- >200 doses of RG6042 have been administered in the OLE study to date

RG6042 (previously known as IONIS-HTTRx) is an investigational medicine and has not yet received regulatory approval in any country.

ASO, antisense oligonucleotide; HD, Huntington's disease; *HTT*, Huntingtin gene; HTT, Huntingtin protein; OLE, open-label extension; siRNA, small interfering RNA; SNP, single nucleotide polymorphism.

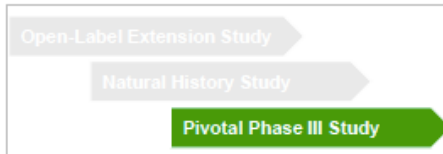
1. Kordasiewicz HB, et al. *Neuron* 2012; 74:1031–1044; 2. Drouot V, et al. *Ann Neurol*. 2009; 65:276–285; 3. Stiles DK, et al. *Exp Neurol*. 2012; 233:463–471; 4. Stanek LM, et al. *Hum Gene Ther*. 2014; 25:461–474; 5. Boudreau RL, et al. *Mol Ther*. 2009; 17:1053–1063. For further details see poster J03: Leavitt B, et al. Partial lowering of total huntingtin levels to treat adults with HD: Potential benefits and theoretical risks from human studies and animal models.

## GENERATION HD1 Study | Global sites



# GENERATION HD1 – RG6042 Pivotal Phase III study design

Objective: Evaluate efficacy and safety of intrathecally-administered RG6042 in adult patients with manifest HD



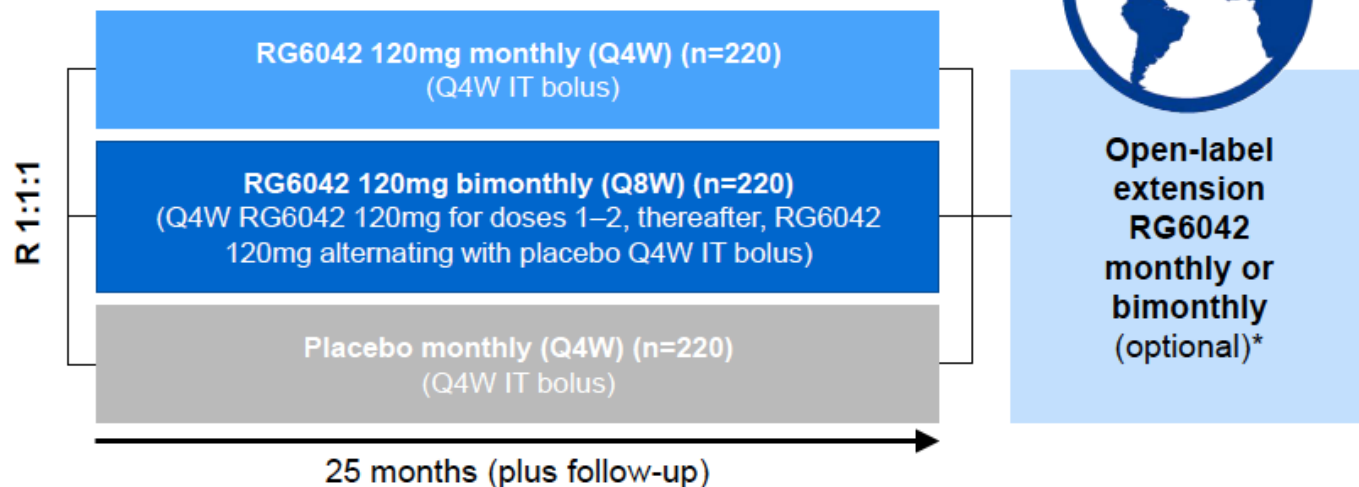
Study launch planned for end of 2018 with patients enrolling by early 2019  
Countries: ~15 countries worldwide (80–90 sites)

## Randomised, multicenter, double-blind, placebo-controlled study

### Key inclusion criteria

- Clinically diagnosed manifest HD (DCL=4)
- Aged 25–65 years
- CAP >400
- Independence scale > or equal to 70
- Ambulatory, verbal

**n=660**



**Inclusion criteria for pivotal study are broader than OLE and HD NHS studies<sup>†</sup>**

\*Provided participants meet eligibility criteria, the data for RG6042 support continued development and the study is approved by Authorities and Ethics Committees/Investigational Review Boards.

<sup>†</sup>Pivotal Phase III study protocol is pending approval by Health Authorities, Investigational Review Boards and Ethics Committees.

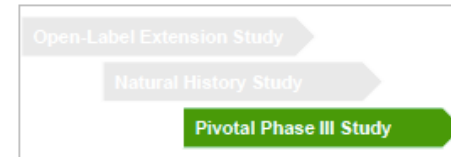
CAP, CAG-age product; DCL, diagnostic confidence level; GENERATION HD1, Global Evaluation of Efficacy and Safety of Roche/Genentech AnTIsense OligoNucleotide for Huntington's Disease; HD, Huntington's disease; IT, intrathecal; NHS, Natural History Study; OLE, open-label extension; Q4W, once-a-month.

## CRITERI DI INCLUSIONE

- Età compresa tra i 25-65 anni
- Segni manifesti di malattia (non soggetti presintomatici portatori della mutazione genetica)
- Discrete capacità funzionali in base ad una scala detta «Indipendence Scale», ovvero pazienti almeno in grado di vestirsi, lavarsi da soli, cucinare ed usare le posate
- Peso di almeno 40 kg
- Per le donne in età fertile necessaria adeguata contraccezione (assunzione pillola estroprogestinica ecc..)
- Presenza di altre malattie croniche (HIV, HCV, epatopatia, insufficienza renale ecc..)
- Assunzione di sostanze stupefacenti (cannabis ecc..)
- Consigliata la presenza di un caregiver (familiare/amico) disponibile ad accompagnare il paziente a tutte le visite ed eseguire alcuni compiti/questionari al domicilio (smart-phone)

## Decision on primary endpoint for global Phase III study

*UHDRS clinical measures are well positioned to demonstrate clinically meaningful efficacy across disease domains*



- **cUHDRS** will be the global primary endpoint
  - Best tracks multidomain decline
  - Related to biology and function
  - Supported by EMA
- **The TFC** will be the primary endpoint in US only
  - A component of the cUHDRS
  - Tracks unilateral functional decline well when measured over longer time periods, and consistency of decline is helped by CAP score
  - Required measure of daily function by FDA

**Consistency of effect anticipated between cUHDRS and TFC**

FDA requires the primary endpoint to measure daily functional abilities, so TFC will be primary endpoint in US only  
CAP, CAG-age product; cUHDRS, composite UHDRS; EMA, European Medicines Association; FDA, Food and Drug Administration; HD, Huntington's disease; HTT, Huntingtin protein; TFC, Total Functional Capacity; UHDRS, Unified Huntington's Disease Rating Scale.



# Dispositivi da utilizzare durante lo studio

## Orologio

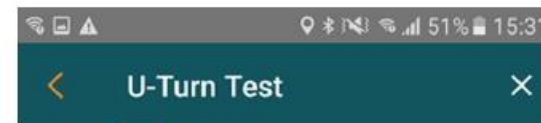
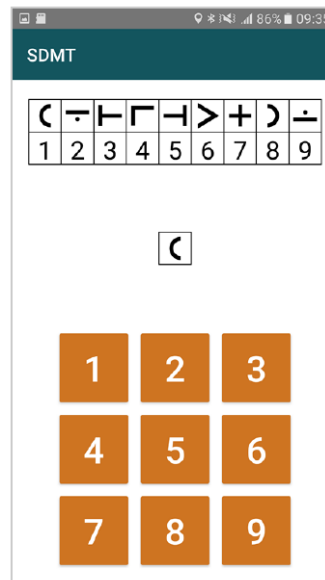


Permette registrazione continuativa dei movimenti del paziente durante la giornata

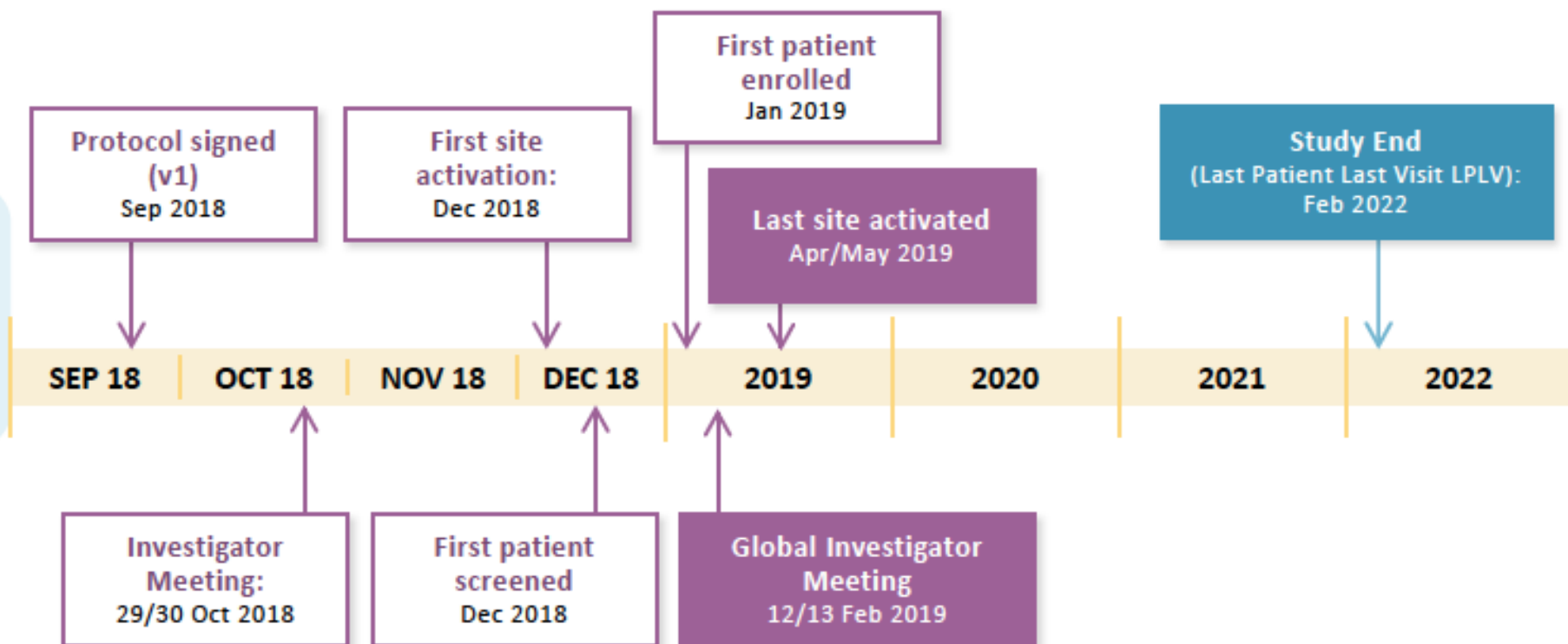
## Smart-phone



Da utilizzare al domicilio per eseguire test cognitivi e motori



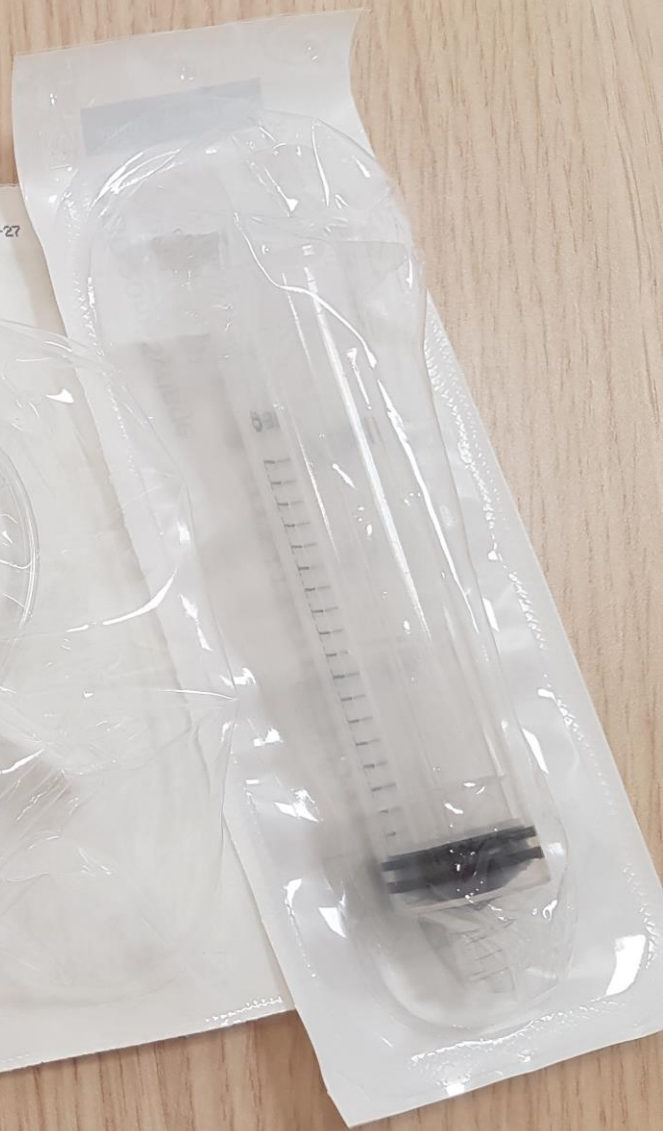
## GENERATION HD1 Study Start-up Timelines







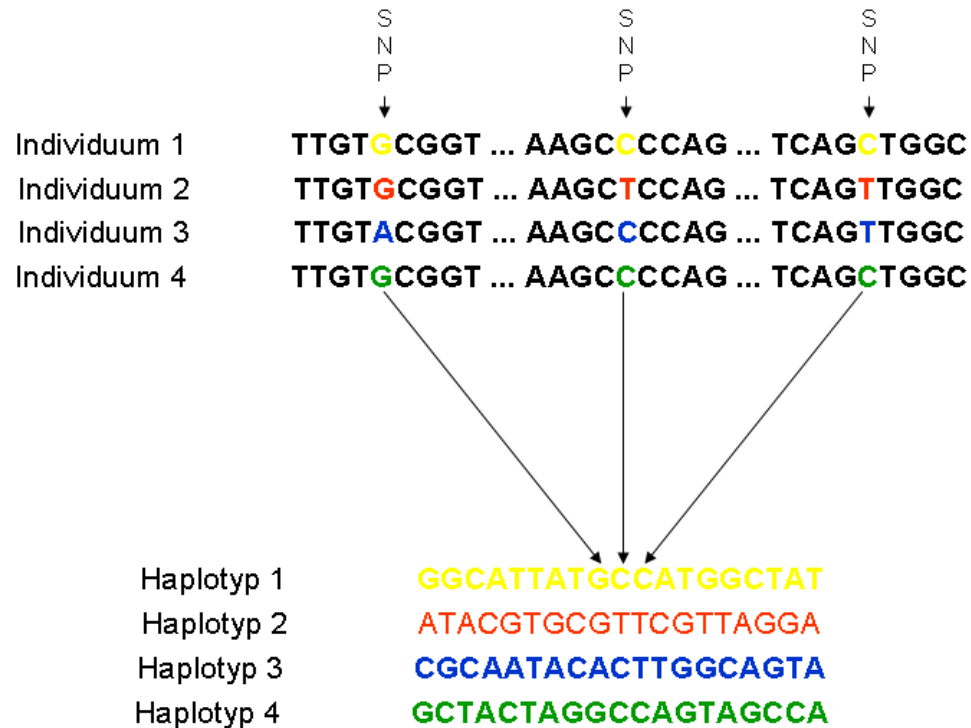
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DATE OF MANUFACTURE 2018-03-27



	Allele selectivity	Delivery	Vector	Sponsor	Key advantages	Key disadvantages	References
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Pre-mRNA degradation	None	Intrathecal	None	Ionis Pharmaceuticals (Carlsbad, CA, USA)	Single drug for all carriers of the Huntington's disease mutation	Theoretical risk from reducing wild-type HTT	Bennett, 2010; <sup>25</sup> Kordasiewicz, 2012; <sup>26</sup> Leavitt, 2016; <sup>27</sup> NCT02519036
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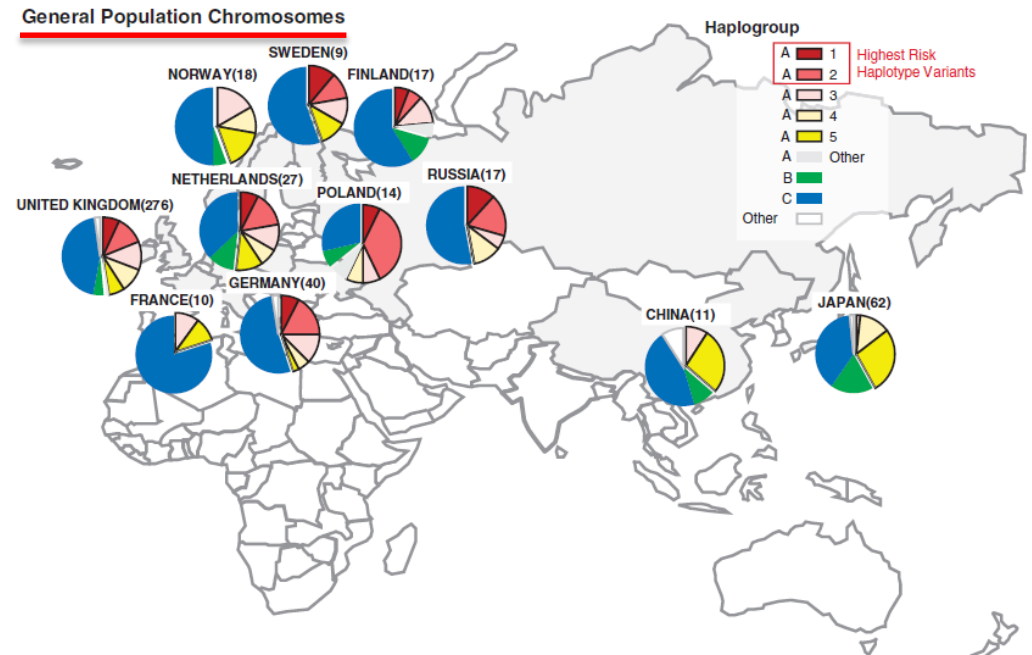
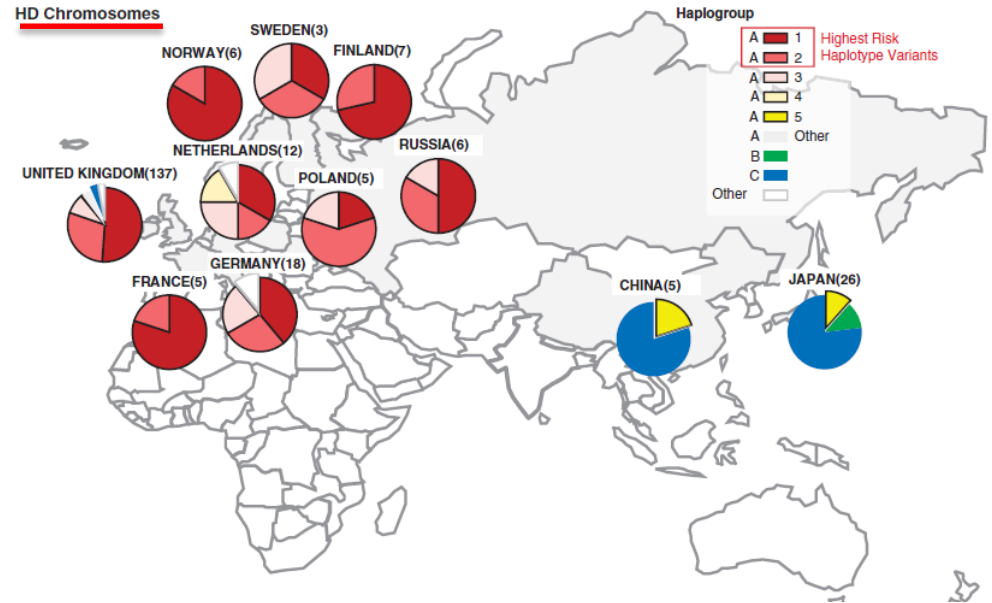
**Table 1: Huntington-lowering programmes targeting mRNA, by class and mechanism**



**Polimorfismo a singolo nucleotide** (single-nucleotide polymorphism o SNP): variazione del materiale genico a carico di un unico nucleotide presente nella popolazione in una proporzione superiore all'1%

**Aplotipo**: combinazione di varianti alleliche (SNP o mutazioni genetiche) lungo un segmento cromosomico contenente loci in «linkage disequilibrium» (strettamente associati e trasmessi insieme alla progenie)

Espansione di triplette CAG nel gene *huntingtina* è in linkage disequilibrium (strettamente associato e trasmesso insieme alla progenie) con alcuni SNP. Tali SNP variano a seconda della popolazione di riferimento.





	TARGET	MECHANISM	DISCOVERY	CANDIDATE	CLINICAL	TRIAL PHASE	WAVE'S COMMERCIAL RIGHTS	PARTNER
<b>CNS</b>								
Huntington's disease	mHTT SNP1	(A)	●	●	●	Phase 1b/2a	50% Global	Takeda
Huntington's disease	mHTT SNP2	(A)	●	●	●	Phase 1b/2a	50% Global	Takeda
Amyotrophic lateral sclerosis	C9orf72	(A)	●	●	○		50% Global	Takeda
Frontotemporal dementia	C9orf72	(A)	●	●	○		50% Global	Takeda
Spinocerebellar ataxia 3	ATXN3	(S)	●	○	○		50% Global	Takeda
CNS diseases	Multiple*	○	●	○	○		Milestones & Royalties	Takeda
<b>MUSCLE</b>								
Duchenne muscular dystrophy	Exon 51	(E)	●	●	●	Phase 1/OLE	100% Global	—
Duchenne muscular dystrophy	Exon 53	(E)	●	●	○		100% Global	—
Duchenne muscular dystrophy	Exons 44, 45, 52, 54, 55	(E)	●	○	○		100% Global	—
Neuromuscular diseases	Multiple	○	●	○	○		100% Global	—
<b>OPHTHALMOLOGY</b>								
Retinal diseases	RHO, USH2A, ABCA4, CEP290	○	●	○	○		100% Global	—
<b>HEPATIC</b>								
Metabolic liver diseases	APOC3 and Multiple (4) <sup>‡</sup>	(S)	●	○	○		Milestones & Royalties	Pfizer

(S) = silencing. (A) = allele-specific silencing. (E) = exon skipping. OLE = Open label extension.

\* During a four-year term, Wave and Takeda may collaborate on up to six preclinical targets at any one time.  
<sup>‡</sup> Pfizer has nominated four undisclosed targets in addition to APOC3.

## PRECISION-HD1 and HD2

- Trattamento con ASO allele-specifico con somministrazione intratecale (**non viene silenziato l'allele sano**)
- ASO hanno come «target» 2 diversi SNP presenti nel 70% dei pazienti con malattia di Huntington
- 2 studi randomizzati in doppio cieco fase I-II (50 pazienti per studio)
- Risultati attesi per la fine del 2019



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<b>RNA Interference compounds</b>							
mRNA degradation	None	Intracranial	AAV2	Spark (Philadelphia, PA, USA)	Single treatment provides sustained HTT reduction	Invasive delivery; limited treatment volume; cannot be reversed if adverse events occur	Harper, 2005; <sup>15</sup> Franich, 2008; <sup>16</sup> McBride, 2011 <sup>17</sup>
mRNA degradation	None	Intracranial	AAV1	Voyager (Cambridge MA, USA)	As above	As above	Stanek, 2015 <sup>31</sup>
mRNA degradation	None	Intracranial	AAV5	UniQure NV (Amsterdam, Netherlands)	As above	As above	Miniarikova, 2016; <sup>32</sup> Samaranch, 2017 <sup>33</sup>
<b>Small molecules</b>							
Screening programme*	Unknown	Potentially oral	None	CHDI Foundation (New York, NY, USA)	Potentially highly accessible route of delivery; potentially readily reversible	More difficult to achieve selectivity for HTT than with nucleotide approaches	Doherty, 2017 <sup>34</sup>

SNP= single nucleotide polymorphism. AAV1, 2, and 5= adeno-associated virus 1, 2, and 5. HTT= huntingtin protein. \*The mechanisms of action, route of delivery, and advantages and disadvantages of the small molecules being investigated by CHDI Foundation remain to be determined because the programme is currently at the phenotypic screen stage.

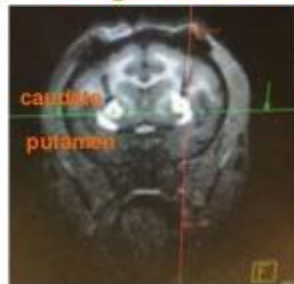
**Table 1: Huntington-lowering programmes targeting mRNA, by class and mechanism**



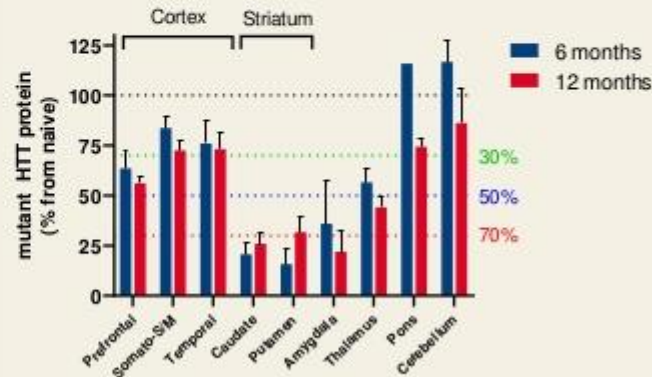
## Libechov transgenic (tgHD) minipigs:

- Life-span: 12-20 years
- Body weight: 50-140 kg
- Brain weight: 90-100 g
- Extra copy of human mutant HTT gene

### MRI-guided CED



## Comparable mutant huntingtin protein knockdown at 6 and 12 months post-injection



Bars represent average ± SEM of n=3-4 animals/group

## First clinical study in HD patients with AMT-130

- Phase 1/2a clinical study:
  - Objective is to test **safety** of AMT-130
  - This will be tested in **adult HD patients**
  - The first study will take place in the **US**
- *More information over details of the study coming soon in ClinicalTrials.gov and through patient organisations*  
<http://uniqure.com/patients/overview-contact-us.php>

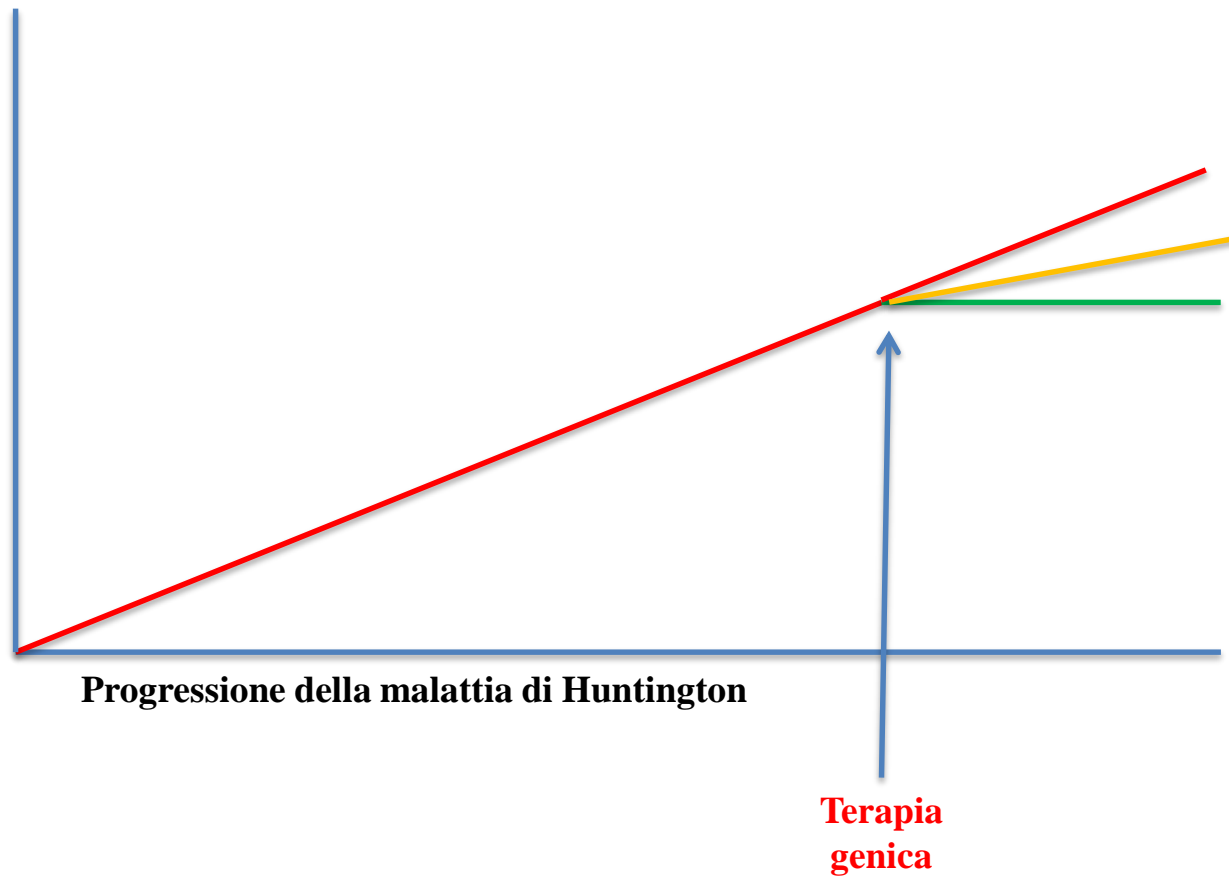
### For patients:

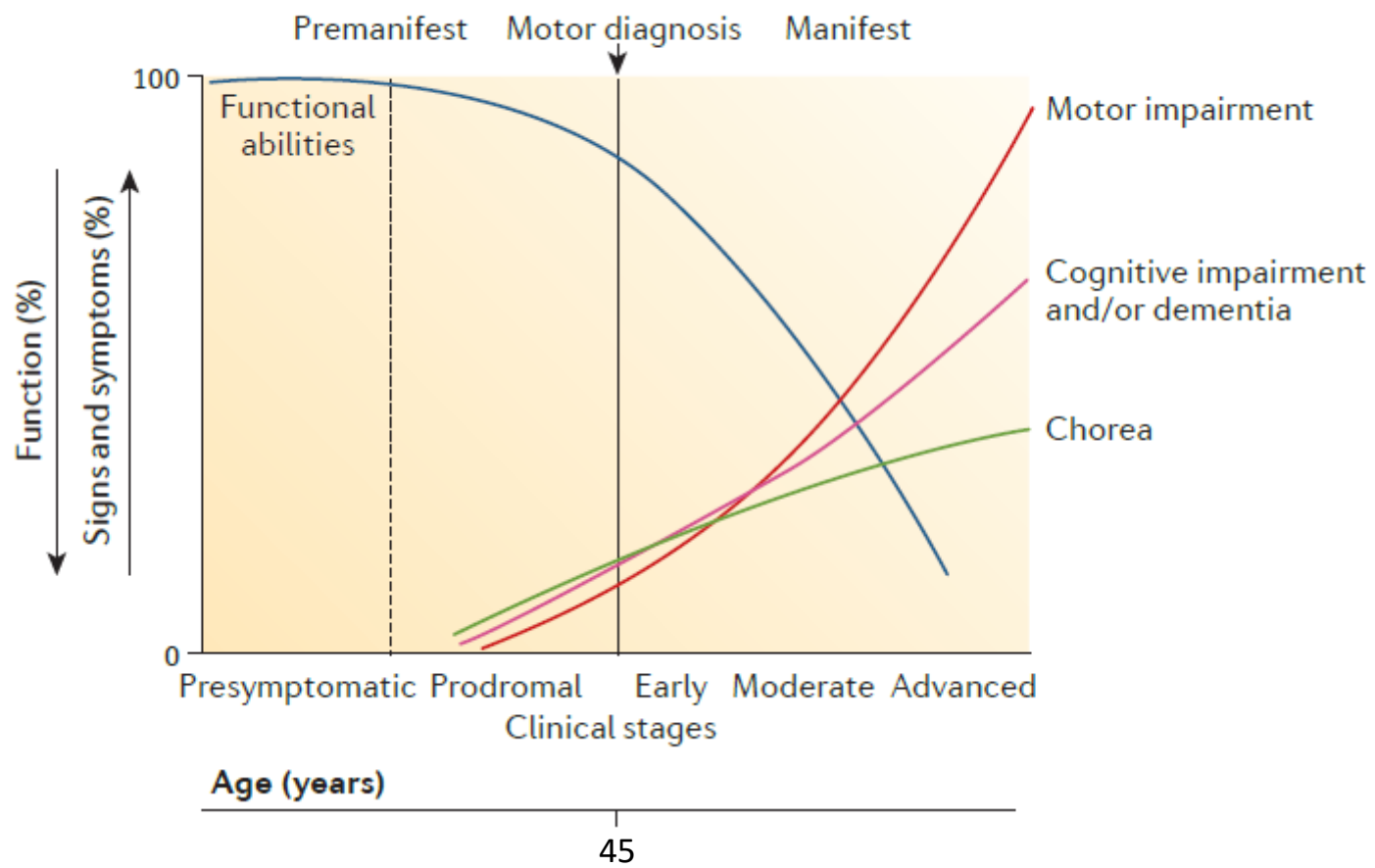
Daniel Leonard, *Director of Global Patient Advocacy*  
[Patients@uniQure.com](mailto:Patients@uniQure.com)



- Inizio sperimentazione ad ottobre 2019
- 26 pazienti seguiti per 5 anni (studio randomizzato in doppio cieco)
- Somministrazione intrastriatale

## Cosa ci aspettiamo come risultato delle sperimentazioni in corso?





Grazie dell'attenzione

