European Reference Networks



# IMPROVING DIAGNOSIS FOR RARE DISEASES IN EUROPE – IMPACT OF EUROPEAN REFERENCE NETWORKS

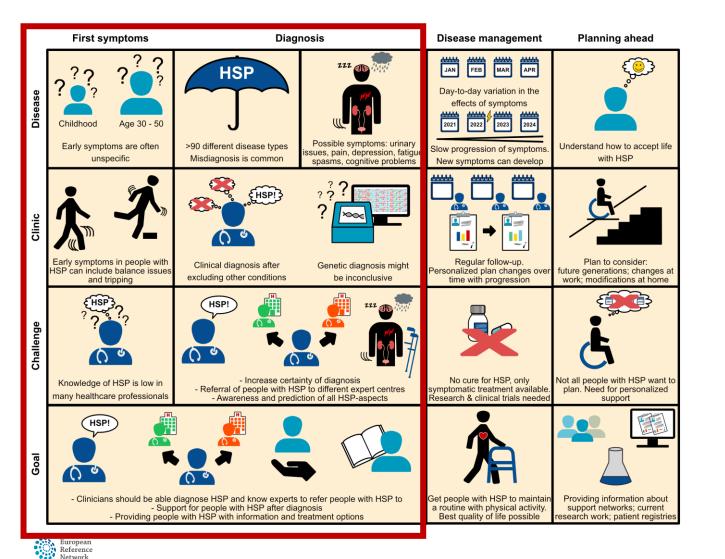
Holm Graessner University Hospital Tübingen, Germany ERN-RND Coordinator





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# **DIAGNOSIS** - MAIN HEALTHCARE PROBLEM FOR RARE DISEASES



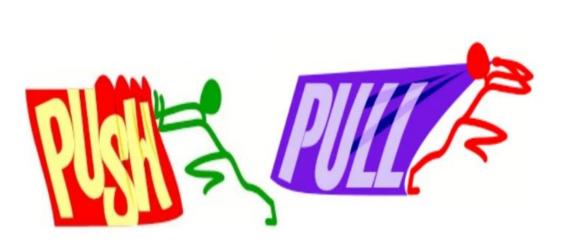
# Main issues:

- Diagnostic journey takes years
- A large fraction of RD patients are misdiagnosed (>50% of all patients)
- Major impact of misdiagnosis
- For about 50% of genetically tested patients the molecular disease cause can not be confirmed

#### (Ref. EURORDIS: Global Rare barometer survey On the journey to diagnosis for people living with a rare disease.)

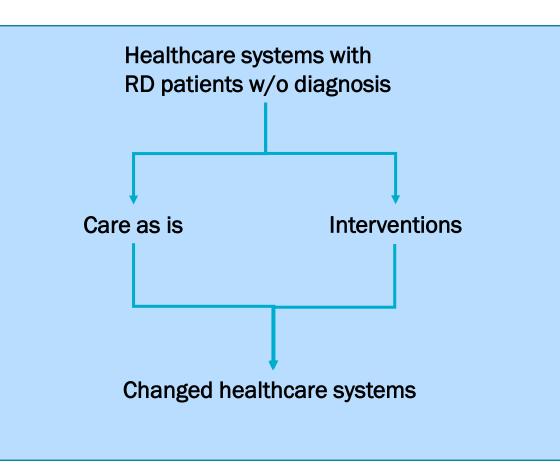
# **RD** DIAGNOSIS – PUSH AND PULL FACTORS

- European Reference Networks
- Diagnostic technology development
- Precision therapies for RD patients



- Medical need of RD patients
- Costs for healthcare systems



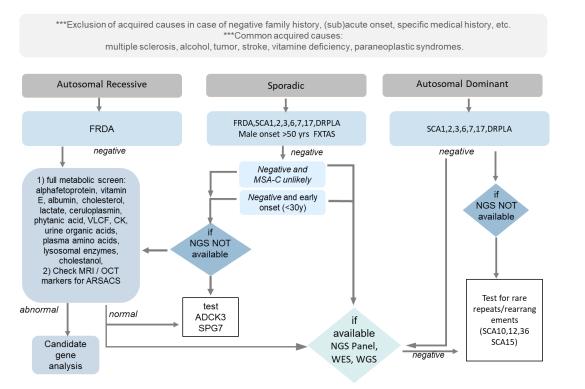


## Interventions

- European Reference Networks
- Data sharing
- Novel Diagnostic Technology
- New Born Screening
- Undiagnosed Disease Programs
- Etc.



#### Diagnostic flowcharts -Ataxias

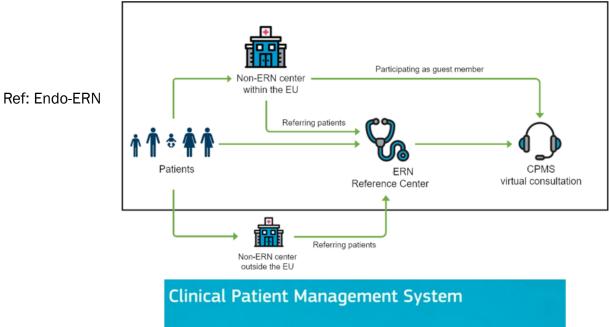


#### Interventions

- European Reference Networks
  - Standards of care including care pathways, referral pathways and guidelines
  - Cross-border diagnostic care pathways including multidisciplinary case discussions (CPMS)
  - Quality assurance for next generation sequencing

#### Collaborative research based on data sharing







# Interventions

- European Reference Networks
  - Standards of care including care pathways, referral pathways and guidelines

 Cross-border diagnostic care pathways including multidisciplinary case discussions (CPMS)

 Quality assurance for next generation sequencing

#### Collaborative research based on data sharing

Quality assurance for the next-generation sequencing diagnostics of rare neurological diseases in the European Reference Network

#### **Pilot scheme results**

Participation 25 laboratories from 17 countries



#### Approaches to diagnosis of RNDs The majority of participating labs employ (clinical) exome

 Image: Trajentry of participating labs energy (clinical) exercise

 sequencing (76%)

 14
 5
 6

 Whole exome
 Clinical exome
 Panels

The majority of participating analyse data using internally developed pipelines (72%)



#### Interpretation

Sufficient evidence

ACMG evidence codes listed

Reference

A wider variability was observed in adherence to variant interpretation standards

Several (32%) labs did not report using an accepted variant interpretation system



· 28% labs presented incomplete evidence to support variant's pathogenicity

18 7 Missing or partial evidence

· A minority of labs provided evidence codes supporting pathogenicity assertion

### Interventions

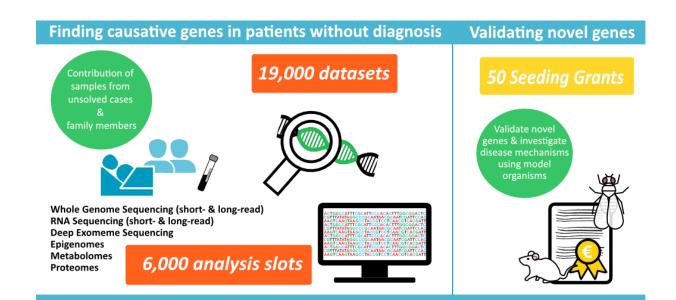
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  - Standards of care including care pathways, referral pathways and guidelines
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#### Quality assurance for Next Generation Sequencing

#### Collaborative research based on data sharing

ACMG evidence codes not listed





## Interventions

- European Reference Networks
  - Standards of care including care pathways, referral pathways and guidelines
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  - Quality assurance for next generation sequencing

### Collaborative research based on data sharing









#### Neurological Diseases (ERN-RND)

EURO-NMD

Building bridges and breaking barriers in rare neuromuscular diseases



ERN-ITHACA focuses on rare congenital malformation syndromes and intellectual disability



European Reference Network for rare and complex epilepsies









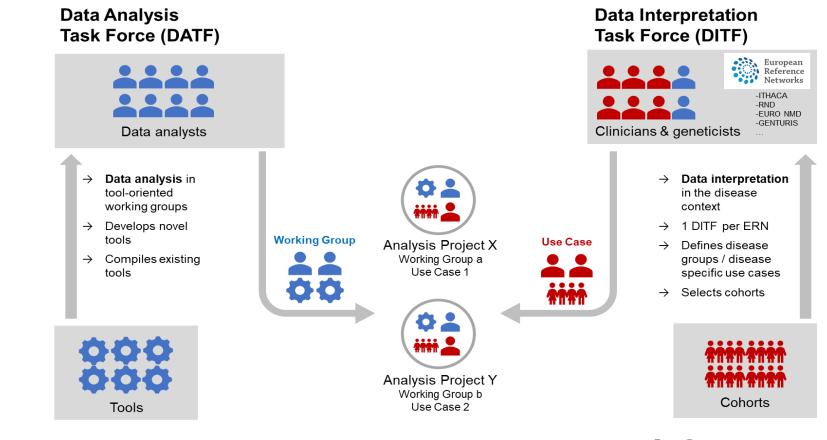


Only WES and WGS datasets

- Data~ 21.000 WES~ 2.350 WGSSharing+ phenotypic data + pedigrees+ metadata
  - ~ 620.000.000.000 byte ~ 620 Terabyte



SOLVE-RD



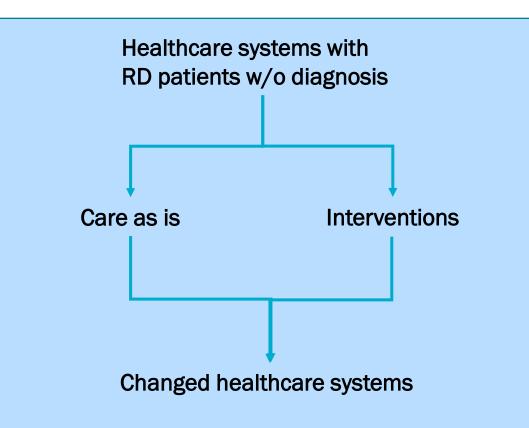
Research based Data Re-analysis

# ~ 10% additionally solved cases



SOLVE-RD

# **RD** DIAGNOSIS – TECHNOLOGY PUSH



# Main issue

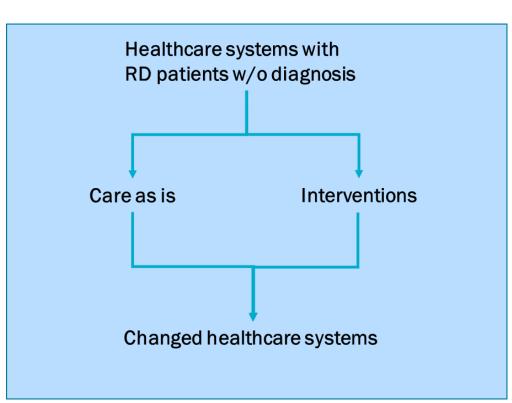
 For about 50% of genetically tested patients the molecular disease cause can not be confirmed

#### Interventions

Novel Diagnostic Technology
 → LONG-READ SEQUENCING



Goal 1: All patients coming to medical attention with a suspected rare disease will be diagnosed within one year if their disorder is known in the medical literature; all currently undiagnosable individuals will enter a globally coordinated diagnostic and RESEARCH pipeline



# How to measure the change in healthcare systems

- European Reference Network Registries

   Interoperable for ERDRI common dataset
  - (at least) All patients seen in ERNs (1.3 million)

#### $_{\odot}$ Data point for undiagnosed cases

6 Diagnosis	6.1.	Diagnosis of the rare disease	Diagnosis retained by the specialised centre	Orpha code (strongly recommended – see link) / Alpha code/ ICD-9 code/ ICD-9- CM code / ICD-10 code
	6.2.	Genetic diagnosis	Genetic diagnosis retained by the specialised centre	International classification of mutations (HGVS) (strongly recommended – see link) /
				HGNC / OMIM code
	6.3	Undiagnosed case	How the undiagnosed case is defined	<ul><li>Phenotype (HPO)</li><li>Genotype (HGVS)</li></ul>
	7.1.	Agreement to be contacted for	Patient's permission exists for being contacted for research	<ul><li>YES</li><li>NO</li></ul>



# How to measure the change in healthcare systems

### - European Reference Network Registries

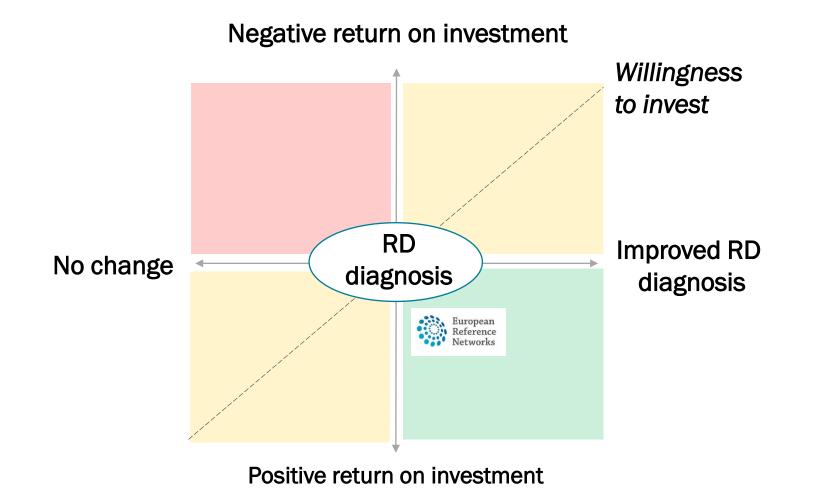
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# → European cohort of RD patients w/o (confirmed) diagnosis

# → Member state cohorts of RD patients w/o (confirmed) diagnosis



# IMPACT OF EUROPEAN REFERENCE NETWORKS ON RD DIAGNOSIS







European Reference Networks



