

XII Meeting. State of the Art in

# HEART FAILURE

CLINICAL PRACTICE AND ORGANIZATIONAL MODELS

Venue: Hotel Meliá María Pita, A Coruña

A Coruña 26-27 September 2025



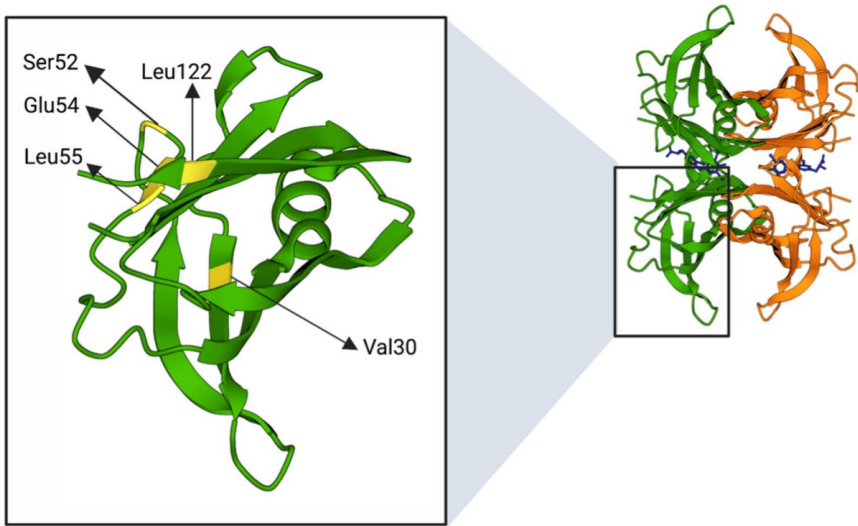
#ACORUÑAHF2025



## Genetic testing: what to look for, when to do it and what it brings to clinical management.

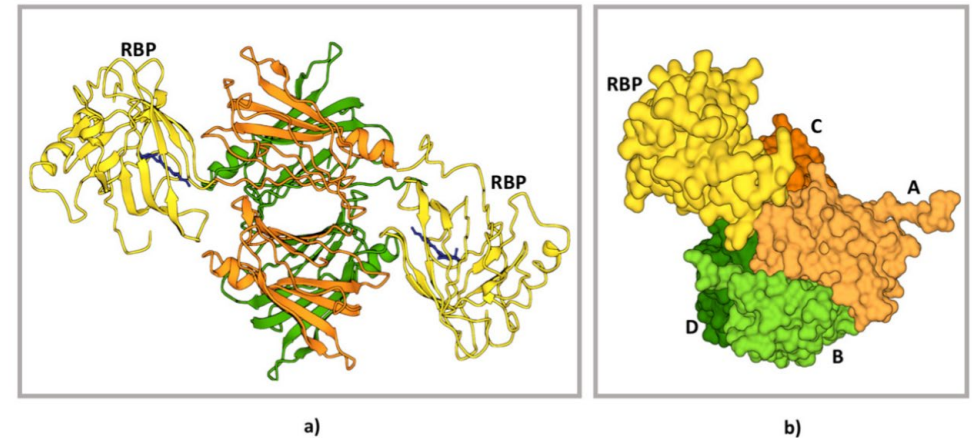
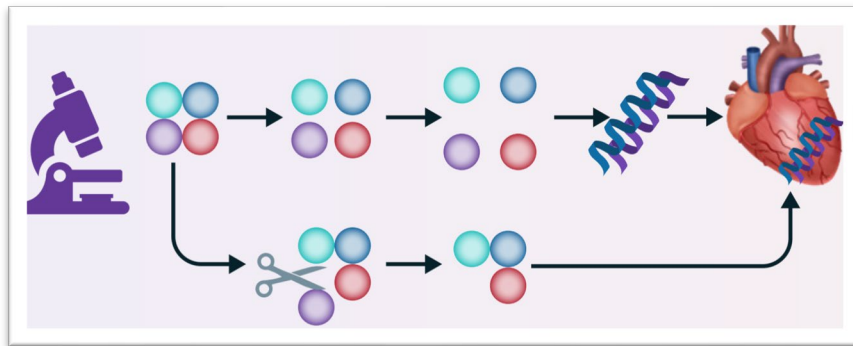
Esteban Martín Álvarez || *Complexo Hospitalario Universitario A Coruña*

# WHAT TO LOOK FOR?

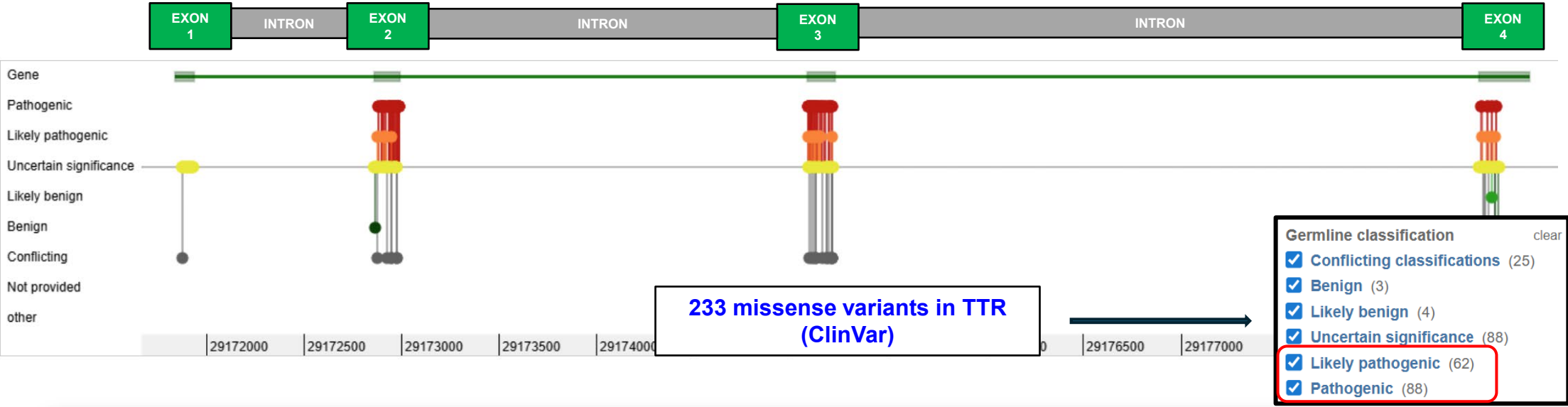


- **Human TTR** is a protein encoded by a single gene located on chromosome 18q.
- **Mature protein has 127 amino acids.** Pro-TTR monomers have 147 amino acids (including a 20-amino acid signal peptide). The current nomenclature, according to the Human Gene Organization, includes this peptide.
- **ATTRv** is caused by **pathogenic variants** in the **TTR gene** that **reduce stability** of the TTR tetramer, leading to **easier dissociation** in pro-amyloidogenic monomers.

**Figure 4.** Positions of amino acids (highlighted in yellow) subjected to mutations described in Table 1. The figure was created through “[www.rcsb.org](http://www.rcsb.org)” and “[biorender.com](http://biorender.com)” web sites (license agreement number KJ247URGZS), access date 22 June 2022.



# WHAT TO LOOK FOR?

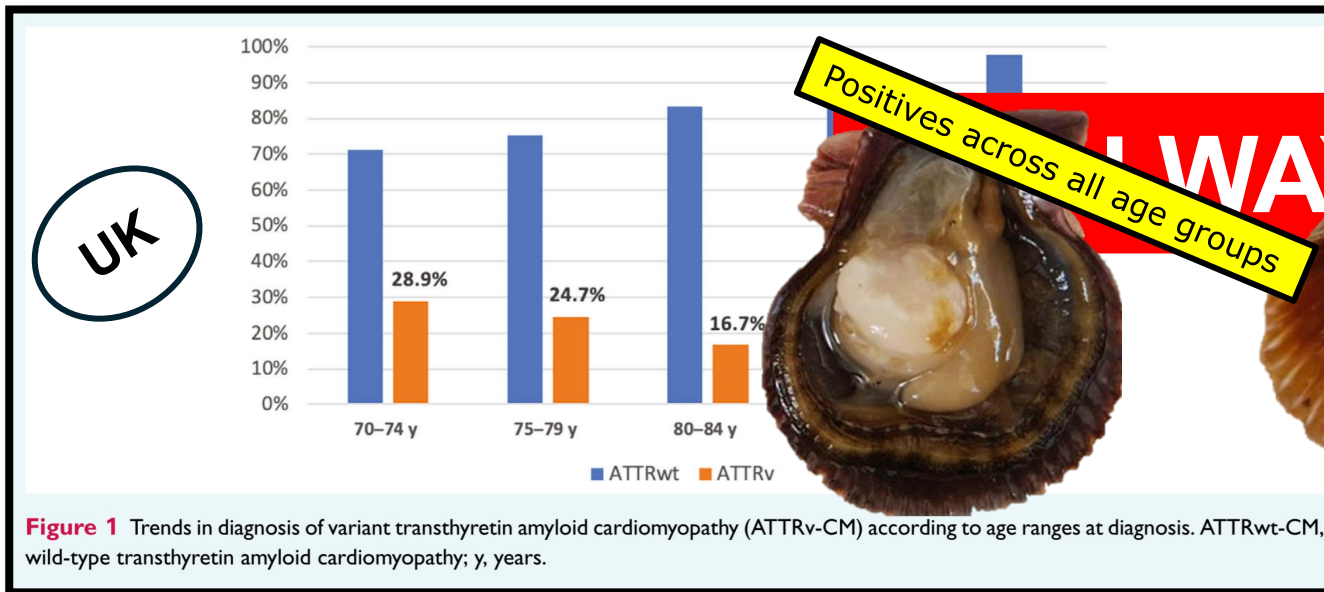


<input type="checkbox"/>	<a href="#">NM_000371.4(TTR):c.133G&gt;T (p.Ala45Ser)</a>	TTR (A45S)	Single nucleotide variant (missense variant)	Cardiovascular phenotype	<b>Likely pathogenic</b> ★
<input type="checkbox"/>	<a href="#">NM_000371.4(TTR):c.148G&gt;T (p.Val50Leu)</a>	TTR (V50L)	Single nucleotide variant (missense variant)	Familial amyloid neuropathy	<b>Pathogenic</b> ★
<input type="checkbox"/>	<a href="#">NM_000371.4(TTR):c.148G&gt;C (p.Val50Leu)</a>	TTR (V50L)	Single nucleotide variant (missense variant)	Cardiovascular phenotype +2 more	<b>Pathogenic</b> ★★
<input type="checkbox"/>	<a href="#">NM_000371.4(TTR):c.148G&gt;A (p.Val50Met)</a>	TTR (V50M)	Single nucleotide variant (missense variant)	Cardiovascular phenotype +7 more	<b>Pathogenic</b> ★★
<input type="checkbox"/>	<a href="#">NM_000371.4(TTR):c.149T&gt;G (p.Val50Gly)</a>	TTR (V50G)	Single nucleotide variant (missense variant)	Familial amyloid neuropathy	<b>Pathogenic</b> ★
<input type="checkbox"/>	<a href="#">NM_000371.4(TTR):c.149T&gt;C (p.Val50Ala)</a>	TTR (V50A)	Single nucleotide variant (missense variant)	Familial amyloid neuropathy	<b>Pathogenic</b> ★★
<input type="checkbox"/>	<a href="#">NM_000371.4(TTR):c.155T&gt;C (p.Val52Ala)</a>	TTR (V52A)	Single nucleotide variant (missense variant)	Cardiovascular phenotype +1 more	<b>Likely pathogenic</b> ★★

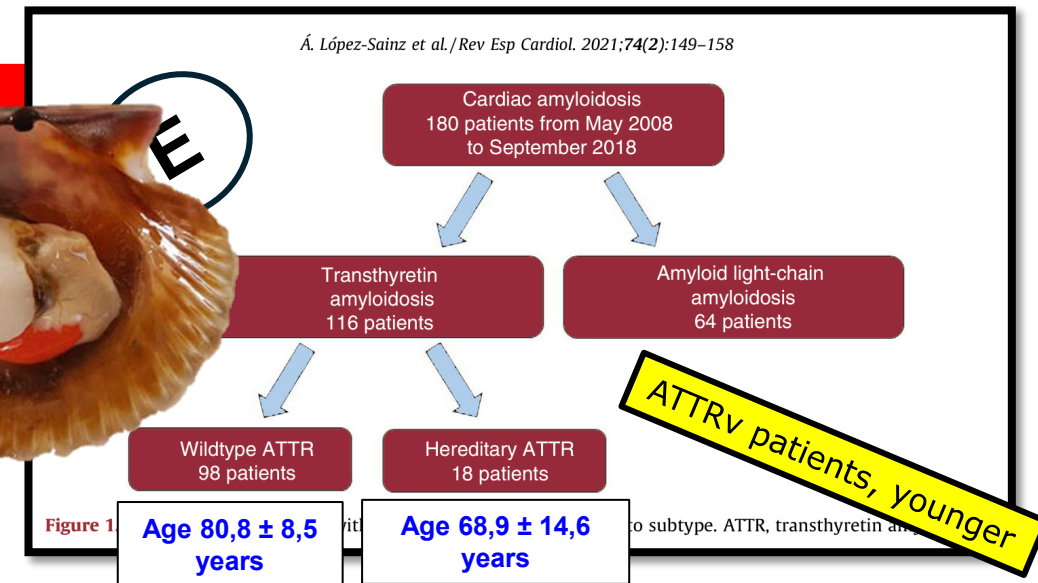
# WHEN TO DO A GENETIC TEST?

- As soon as **ATTR-CM** has been confirmed, the presence of **pathogenic variants** in the TTR gene should be evaluated and **genetic counseling** should be given.
- ATTRv** and **ATTRwt** are two different diseases. **100% of ATTRwt affect the heart**, but **not all ATTRv do so** (30-100%).
- Genetic testing should be done **regardless of age**.

## ATTRv ≠ ATTRwt

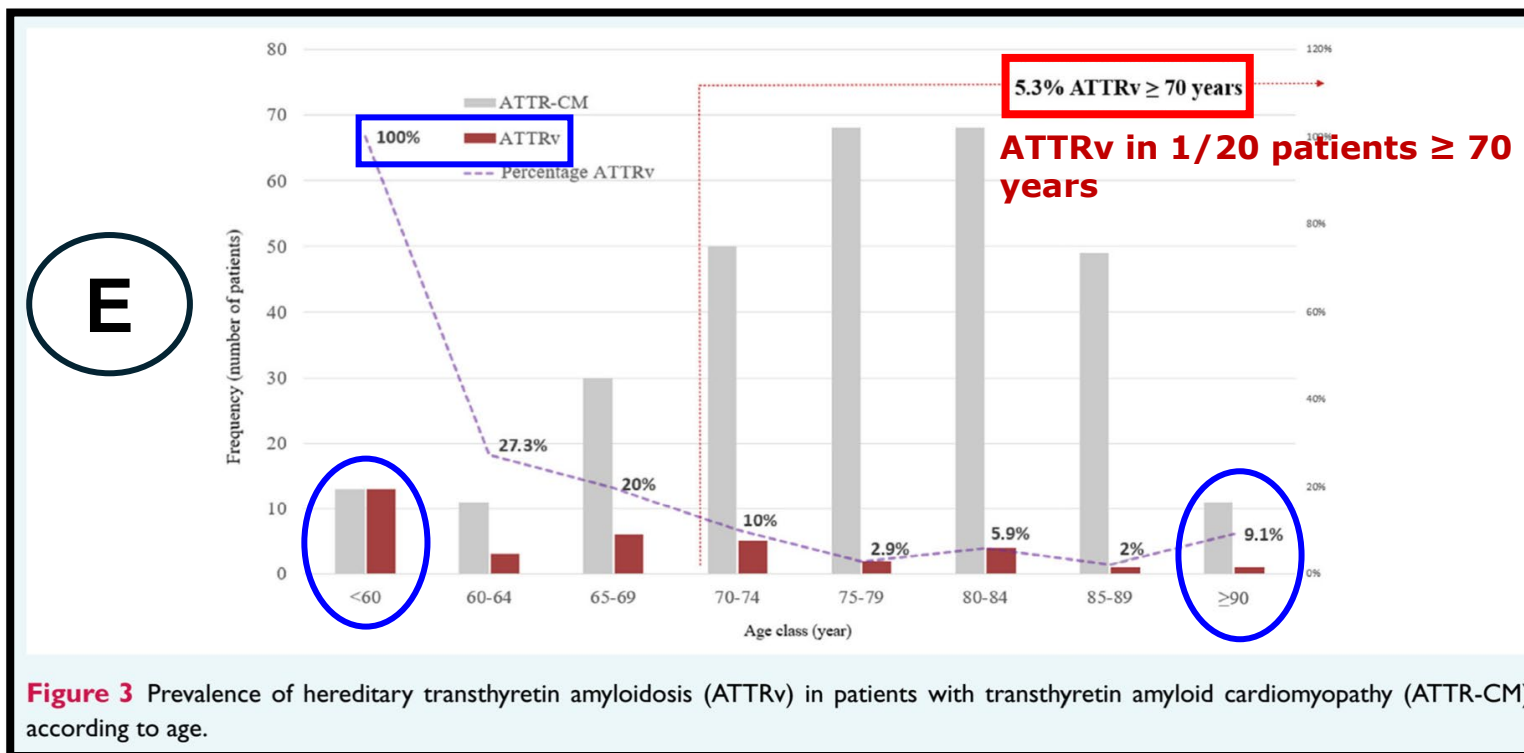


Porcari, Aldostefano et al. "Prevalence, characteristics and outcomes of older patients with hereditary versus wild-type transthyretin amyloid cardiomyopathy." *European journal of heart failure* vol. 25,4 (2023): 515-524.

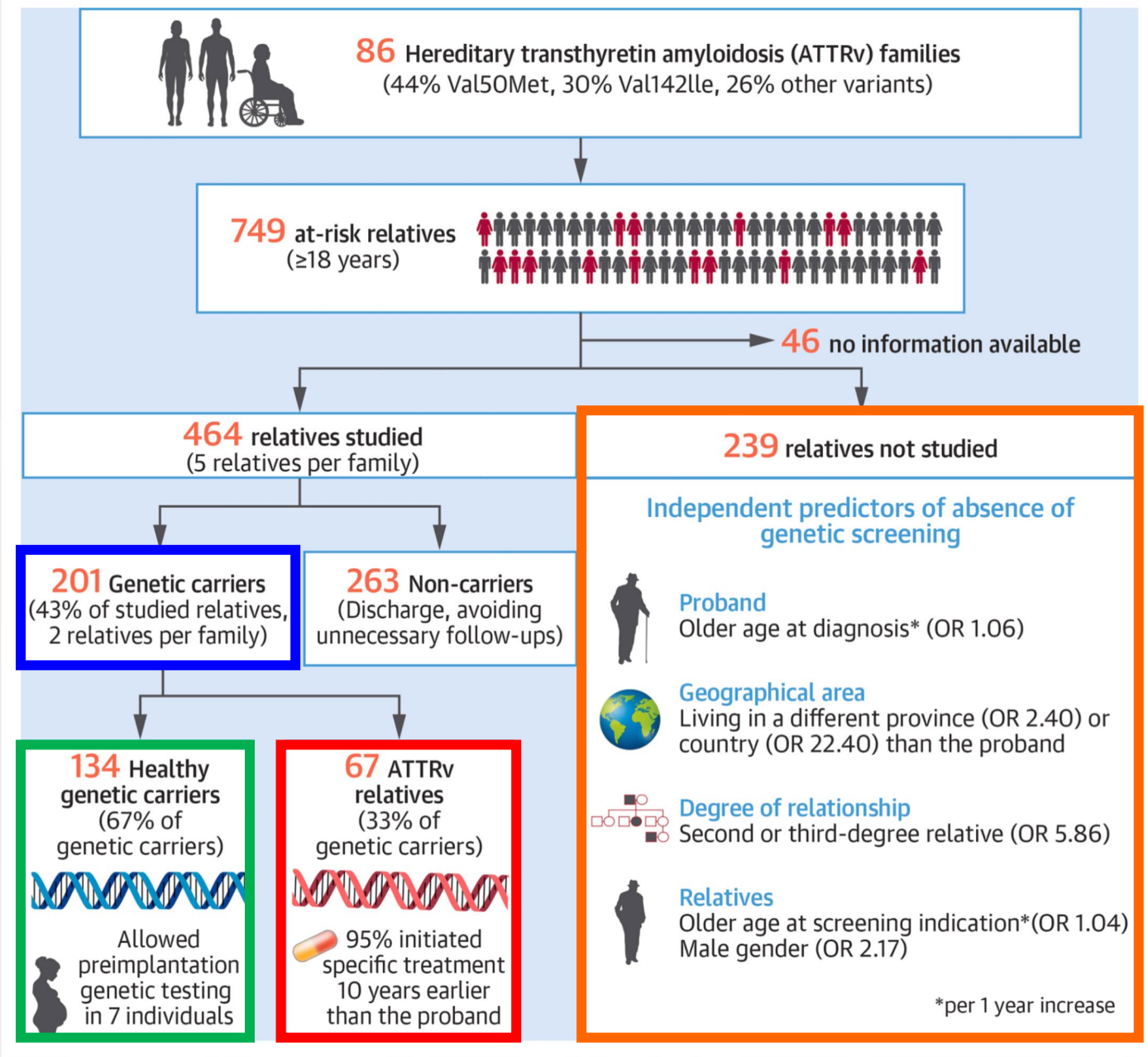


López-Sainz, Ángela et al. "Clinical profile and outcome of cardiac amyloidosis in a Spanish referral center." *Revista española de cardiología (English ed.)* vol. 74,2 (2021): 149-158.

- Genetic testing should be done **regardless of age**.



- Overall**, the prevalence of **ATTRv** among 300 patients with ATTR-CM was **12%**.
- In the cohort of ATTR-CM patients **≥ 70 years**, 13/246 had **ATTRv (5.3%)**.
- Prevalence of **ATTRv** among **elderly female** patients with ATTR-CM was **13%**.
- Oldest ATTRv patient was 93-year-old Caucasian female with the p.Val142Ile variant
- Prevalence in <60 years was 100%.
- Implications of ATTRv diagnosis: **transthyretin-specific** drug treatment, genetic screening in **relatives**, identification of **asymptomatic carriers**.



- Almost **half of studied relatives** were genetic carriers.
- Genetic carriers:
  - **2/3 healthy carriers.**
  - **1/3 ATTRv.**
- **1/3** of **at-risk** relatives **not studied.**



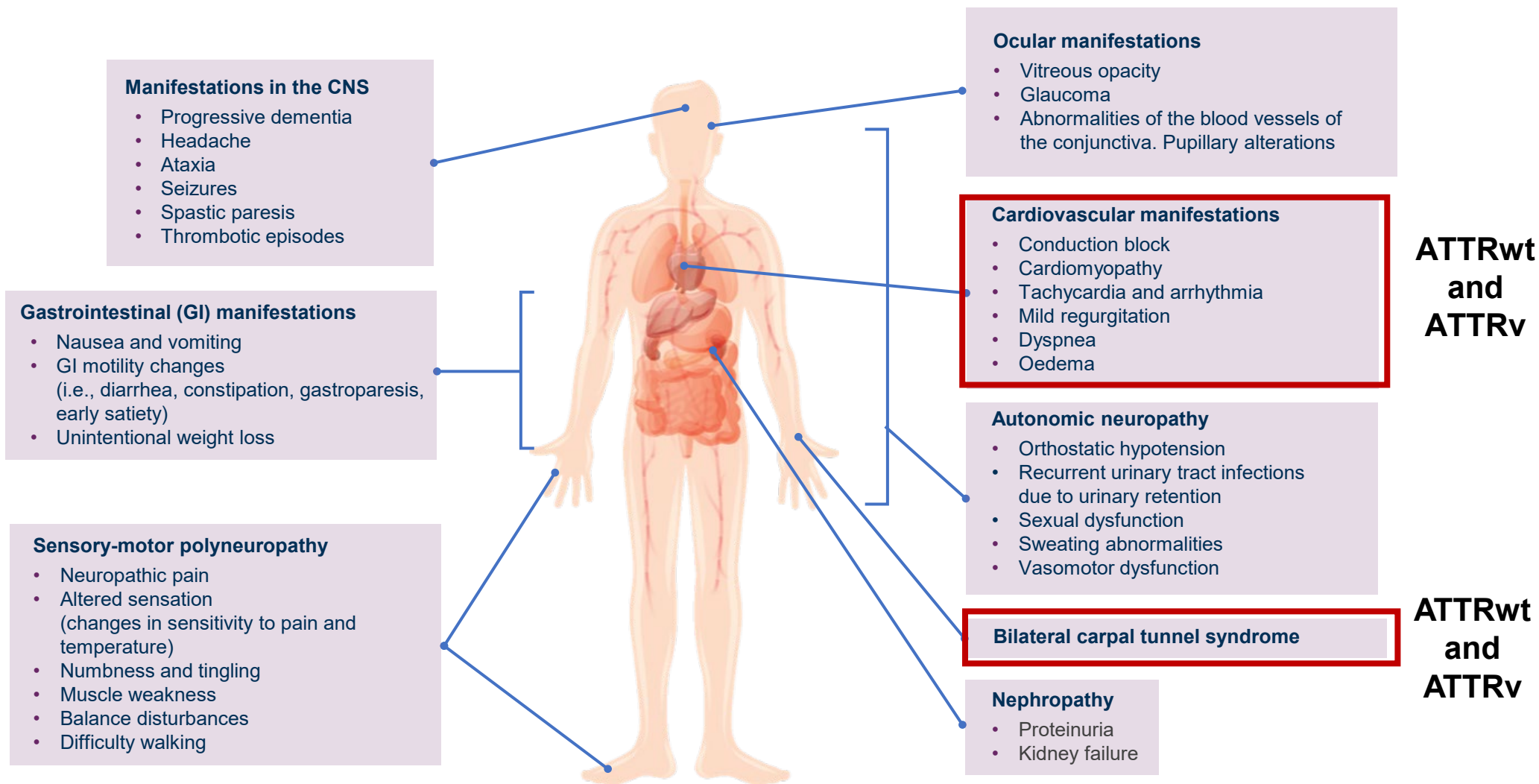
**Early diagnosis**

**Reproductive planning**

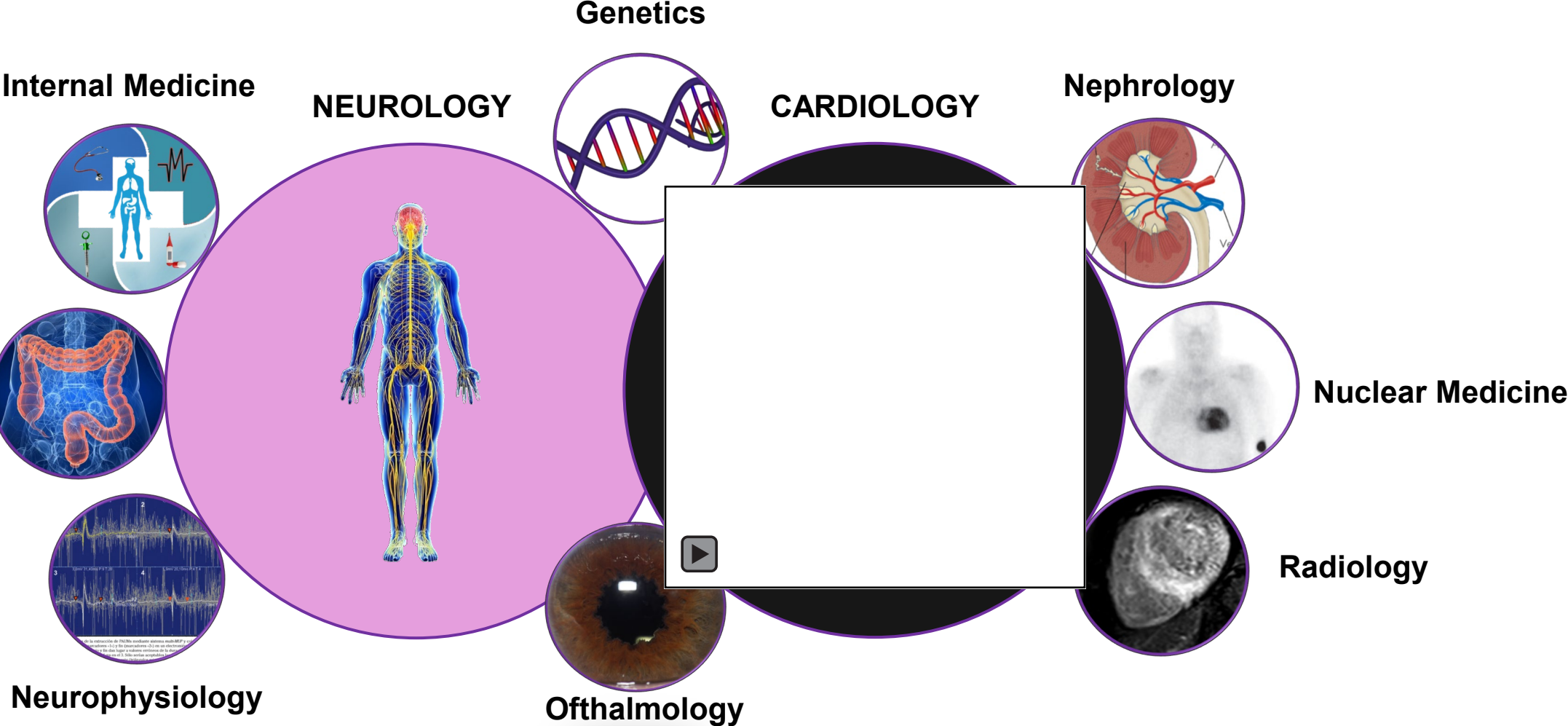
**Early treatment**

# RELEVANCE IN CLINICAL MANAGEMENT

## ATTRv ≠ ATTRwt

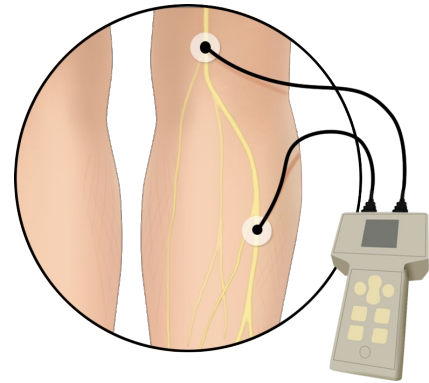


# ATTRv: Multidisciplinary management



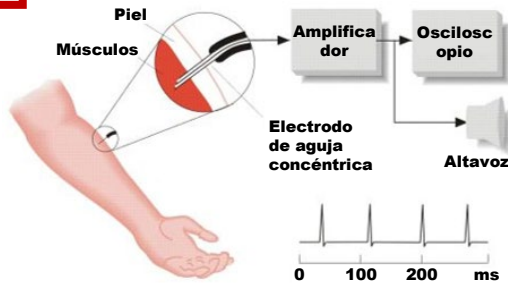
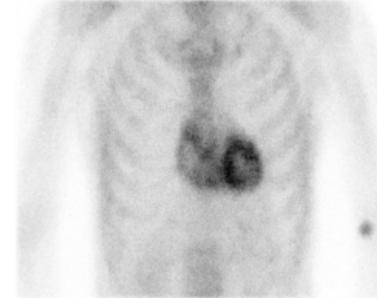
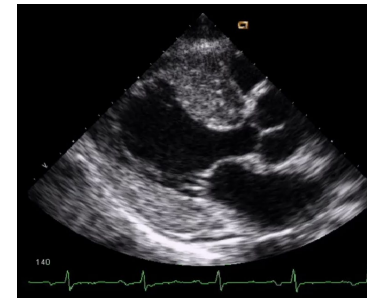
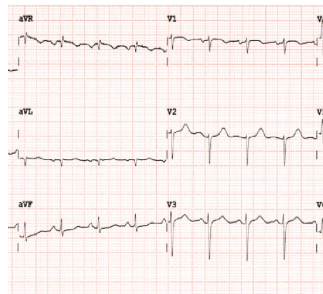
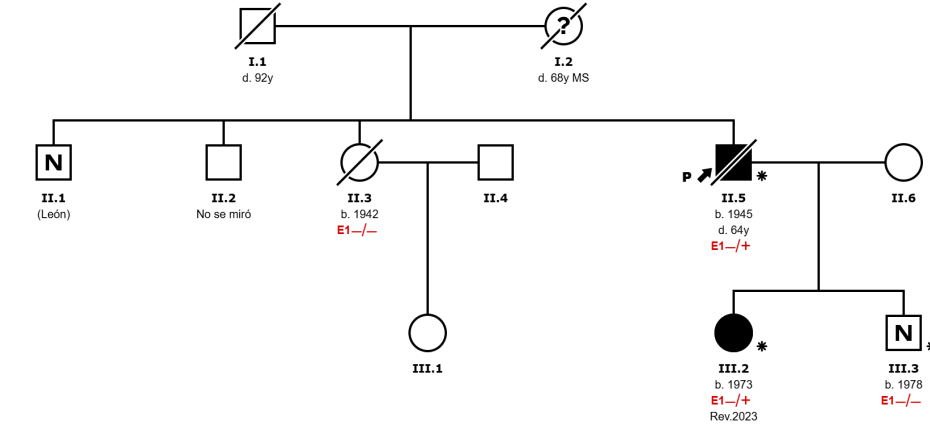
# ATTRv: Multidisciplinary management

## Sudoscan

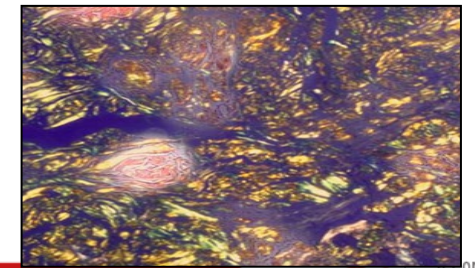
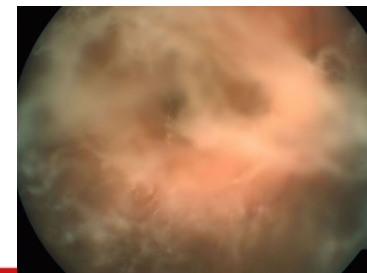
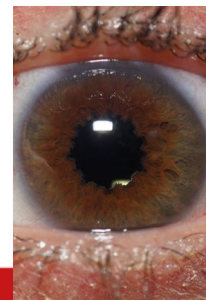


nerve  
conduction  
study(NCS)

**BIOMARKERS**  
(TnI, NT-  
proBNP)  
**RENAL**  
**FUNCTION,**  
**uACR**



**Electromyography**  
**(EMG)**



**COMPASS 31**

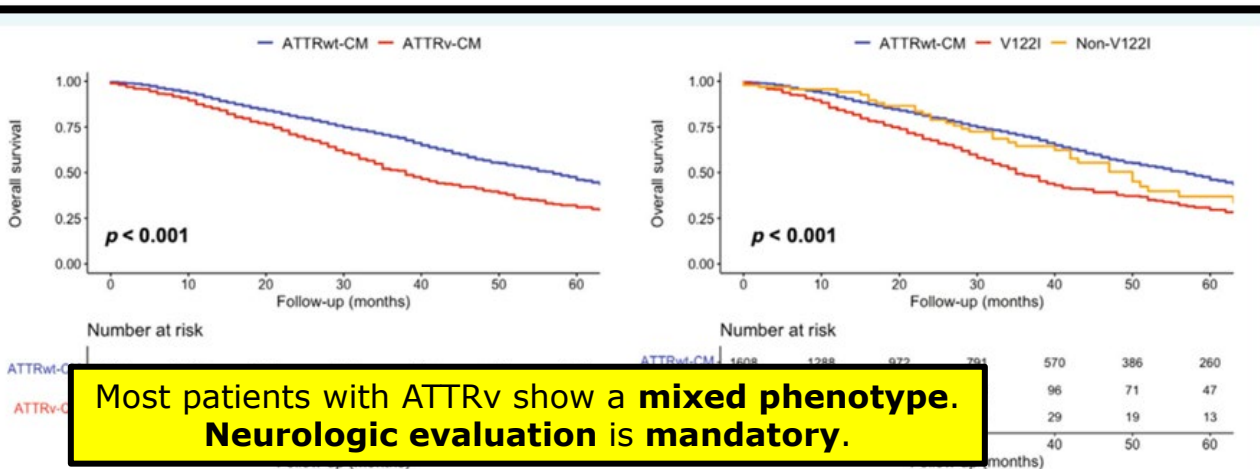
# RELEVANCE IN CLINICAL MANAGEMENT

ATTRv is a **multisystem autosomal dominant disease**. **Phenotypic expression** differs between different **variants**, and even for the **same variant** (early-onset p.Val50Met vs. late-onset p.Val50Met)

**"Cardiac variants":**  
**Val142Ile, Leu131Met, Thr80Ala, Ile88Leu**

**TABLE 1** Demographic, Genetic

Average age, y (range)
Male, %
Race/ethnicity/country of origin
Genetics
Phenotype



**Most patients with ATTRv show a mixed phenotype. Neurologic evaluation is mandatory.**

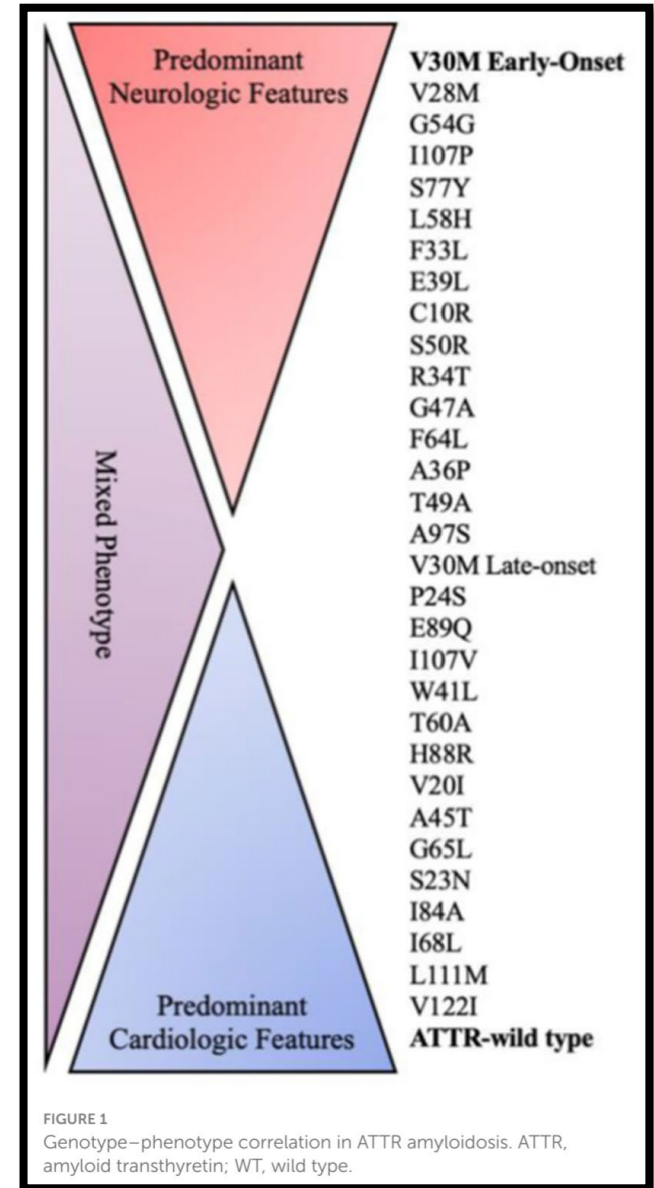
**Figure 2** Prognostic impact of variant transthyretin amyloid cardiomyopathy (ATTRv-CM) and specific transthyretin variant on overall survival compared to wild-type transthyretin amyloid cardiomyopathy (ATTRwt-CM). V122I, valine-to-isoleucine substitution at position 122.

Porcari, Aldostefano et al. "Prevalence, characteristics and outcomes of older patients with hereditary versus wild-type transthyretin amyloid cardiomyopathy." *European journal of heart failure* vol. 25,4 (2023): 515-524.

<sup>a</sup>Depends on endemic vs nonendemic areas, early vs late age onset. <sup>b</sup>Age of disease penetrance is later in women than in men.

**In general, variants with a predominantly cardiac phenotype have a worse prognosis**

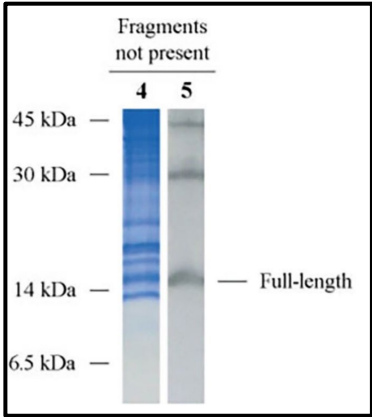
Griffin, Jan M et al. *JACC. CardioOncology* vol. 5,4 (2023): 475-484.



**FIGURE 1** Genotype-phenotype correlation in ATTR amyloidosis. ATTR, amyloid transthyretin; WT, wild type.

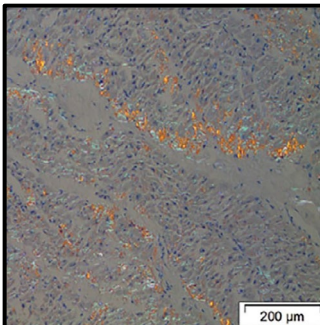
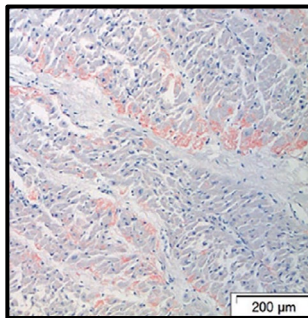
# RELEVANCE IN CLINICAL MANAGEMENT

ATTRv is a **multisystem autosomal dominant disease**. **Phenotypic expression** differs between different **variants**, and even for the **same variant (early-onset p.Val50Met vs. late-onset p.Val50Met)**



**Type B: Long and thick**

**Small non myelinated fibres**



Some variants are associated with **false negatives on the 99mTc-DPD scintigraphy scan**, even in cases with cardiac involvement (**early-onset p.Val50Met**, p.Phe84Leu, p.Ser97Tyr).

**Type A: Short and thin**

**Large myelinated fibres**

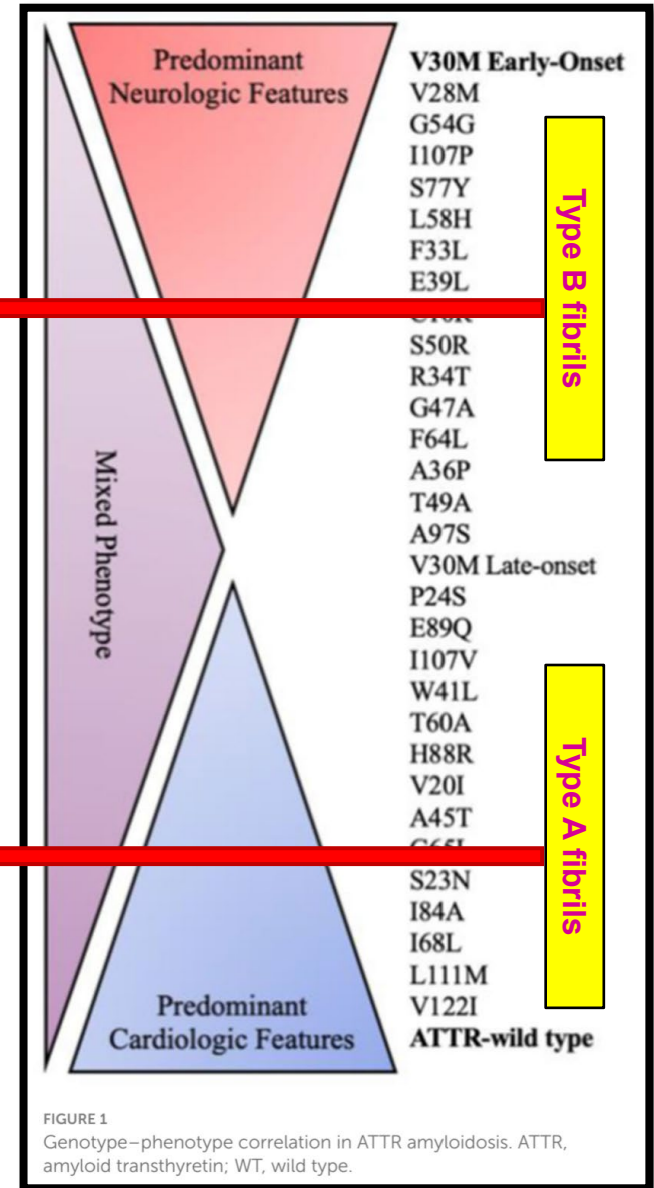
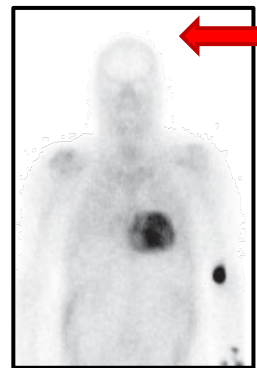
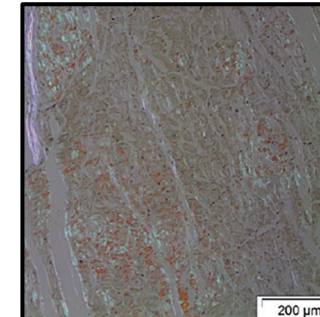
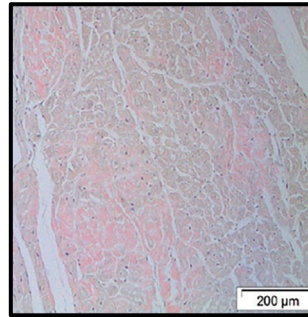
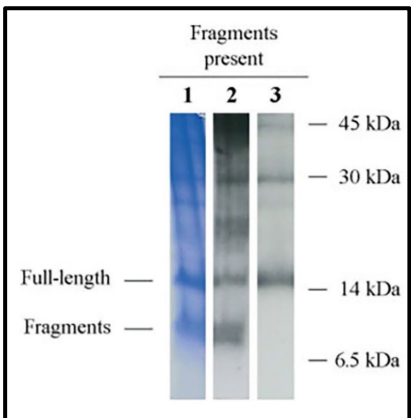
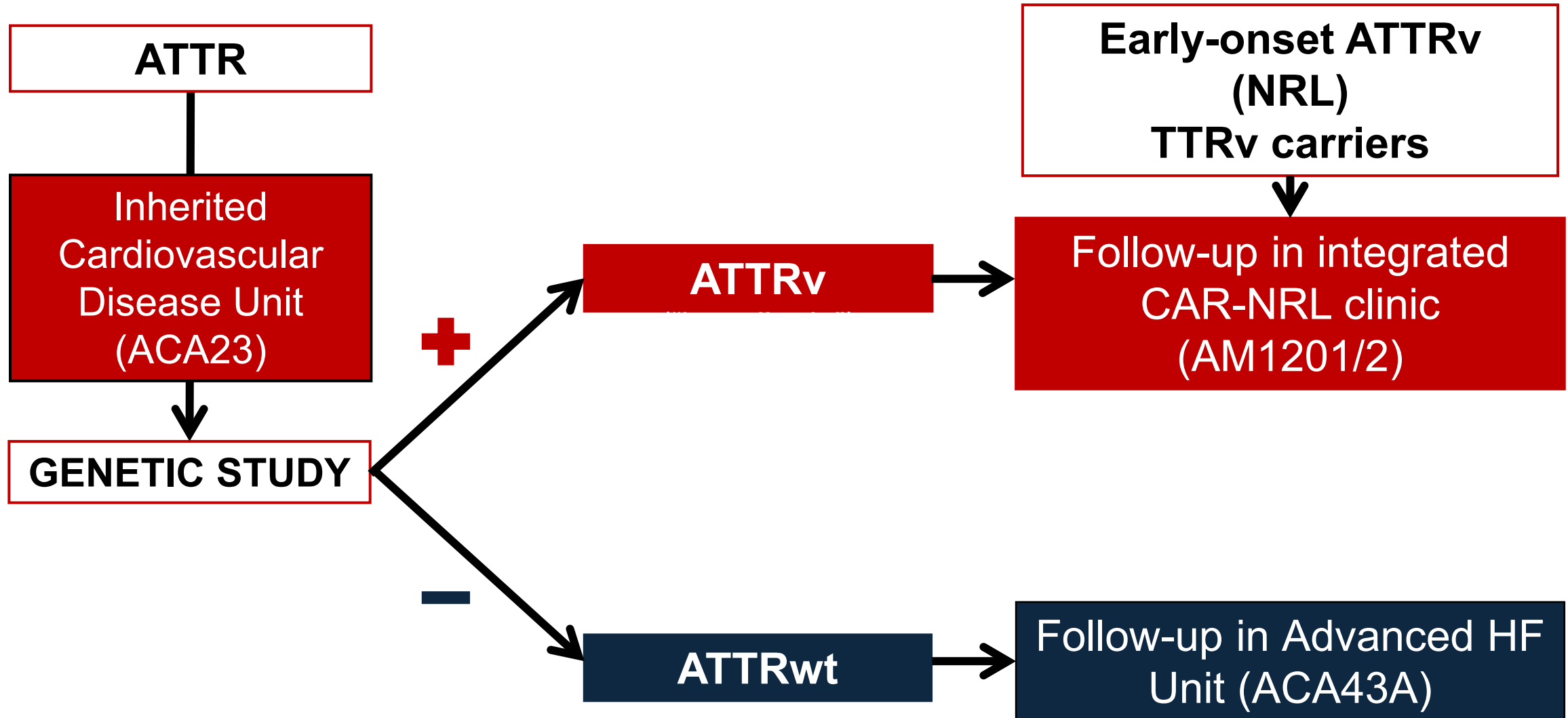


FIGURE 1 Genotype-phenotype correlation in ATTR amyloidosis. ATTR, amyloid transthyretin; WT, wild type.



1. Poli, Loris et al. "Hereditary transthyretin amyloidosis: a comprehensive review with a focus on peripheral neuropathy." *Frontiers in neurology* vol. 14 1242815. 5 Oct. 2023. //2. Garcia-Pavia, Pablo et al. "Diagnosis and treatment of cardiac amyloidosis: a position statement of the ESC Working Group on Myocardial and Pericardial Diseases." *European heart journal* vol. 42,16 (2021): 1554-1568. //3. Adams, David et al. "Hereditary transthyretin amyloidosis: a model of medical progress for a fatal disease." *Nature reviews. Neurology* vol. 15,7 (2019): 387-404. //4. Morfino, Paolo et al. "Amyloid seeding as a disease mechanism and treatment target in transthyretin cardiac amyloidosis." *Heart failure reviews* vol. 27,6 (2022): 2187-2200.

# How do we do it?



# Family Study and Asymptomatic Carriers of Pathogenic Variants

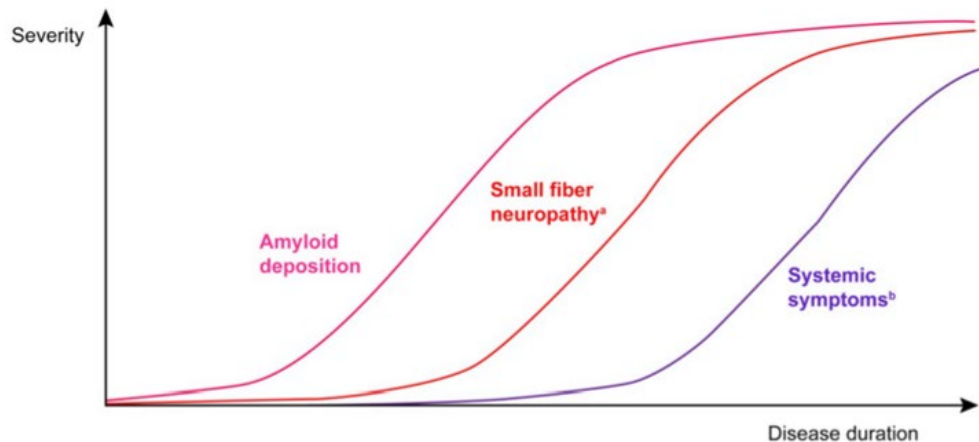
- In hereditary TTR amyloidosis, we do not see isolated patients, but **families**.
- The **age of presentation, phenotype, penetrance, and progression** depend on the specific variant.
- To decide when to initiate penetrance assessment in carriers, we are guided by the **specific variant** and the **age of onset in affected relatives (PADO)**.

**Table 2.** Estimating the predicted age of disease onset in patients with ATTR amyloidosis.

Phenotype group	Genotype	Penetrance	Typical age of onset	Rate of progression
Neurologic	V30M early onset	>90%	<40 years	++++++
Neurologic/mixed	V30M late onset	>60%	>50 years	++++++
Cardiac	V122I	Unknown	55 years	++
	L111M	>90%	35–40 years	++
	T60A	>90%	55 years	++
	I68L	>90%	55 years	++
Mixed	S77Y	>90%	55 years	++++
	E89Q	>90%	50 years	++++
	G47E	>90%	30 years	++++++

The number of “+” provides an indication of how fast the disease progresses, with “+” representing slow progression of ATTR amyloidosis.

Amyloid deposition begins **before** the onset of symptoms. **Monitoring of carriers** is essential for early diagnosis.



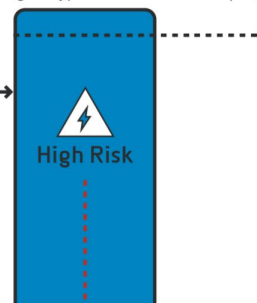
Ueda, Mitsuharu et al. *Journal of the neurological sciences* vol. 414 (2020): 116813.

## Establishing the follow-up and frequency

**10 years before** the age of debut in family members. **Sooner**, if you have suspicious symptoms or signs.

Educate the carrier to understand the early clinical signs associated with the specific mutation and increase frequency of surveillance if signs suspected

Increased frequency of follow-up, particularly in those genotypes associated with rapidly progressing disease



Predicted age of disease onset } Typical age of onset for the specific genotype  
 · Family history  
 · Proband age of onset

Baseline assesment 10 years prior to predicted age of onset

Figure 1. Establishing the start and frequency of follow-up of carriers of a TTR mutation.

# Follow-up of gene variant carriers

Evaluation	Frequency
<b>Medical interview</b> (sensation, movement, autonomic nerve function [orthostatic hypotension, gastrointestinal symptoms, and dysuria], weight loss, heart failure, arrhythmia, and eye symptoms)	<b>ANNUAL</b>
<b>Physical examination</b> (neurological, autonomic, cardiac, orthostatic test, NIS scales, mBMI)	<b>ANNUAL</b>
<b>Neurological Evaluations:</b> EMG/ENG and <b>Sudoscan</b>	<b>ANNUAL</b>
<b>Asymptomatic ATTRv patients vs asymptomatic TTRv carriers</b>	
<b>Ophthalmological examination</b>	<b>ANNUAL</b>
<b>Laboratory:</b> NT-proBNP, Troponin, Creatinine, GFR, Albumin, Microalbuminuria.	<b>ANNUAL</b>
<b>ECG</b>	<b>ANNUAL</b>
<b>Echocardiogram</b>	<b>ANNUAL</b>
<b>Holter-ECG</b>	<b>Every 2 years</b>
<b>DPD ± Cardiac MRI Scan</b> <b>MIBG scan?</b>	<b>Every 3 years/if suspicion</b>
<b>Biopsy</b> (sural nerve, salivary gland, abdominal fat, skin, heart)	<b>If suspected and negative non-invasive diagnosis</b>
<b>LIGHT-CHAIN NEUROFILAMENTS?</b>	<b>ANNUAL?</b>

# New tools in ATTRv

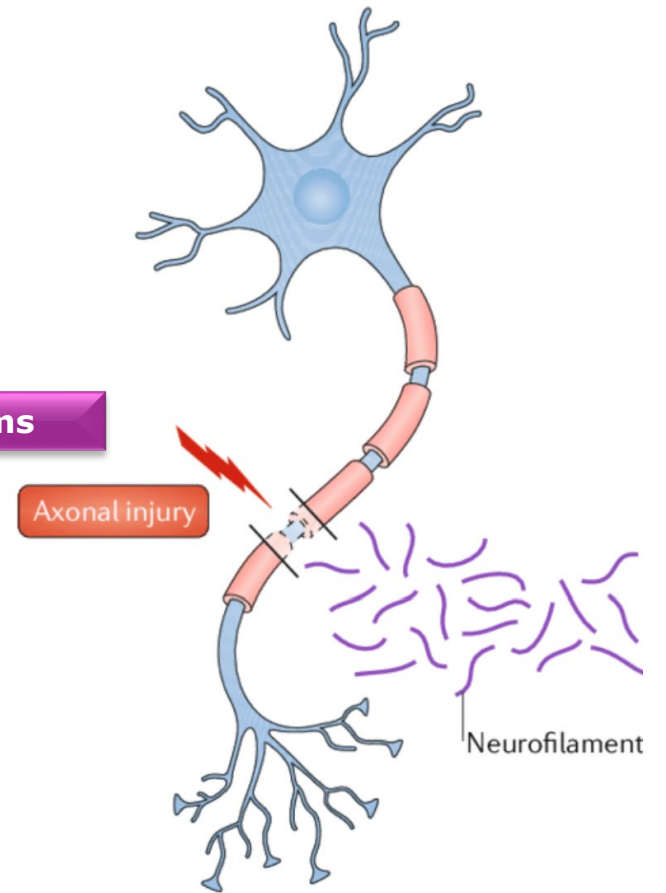
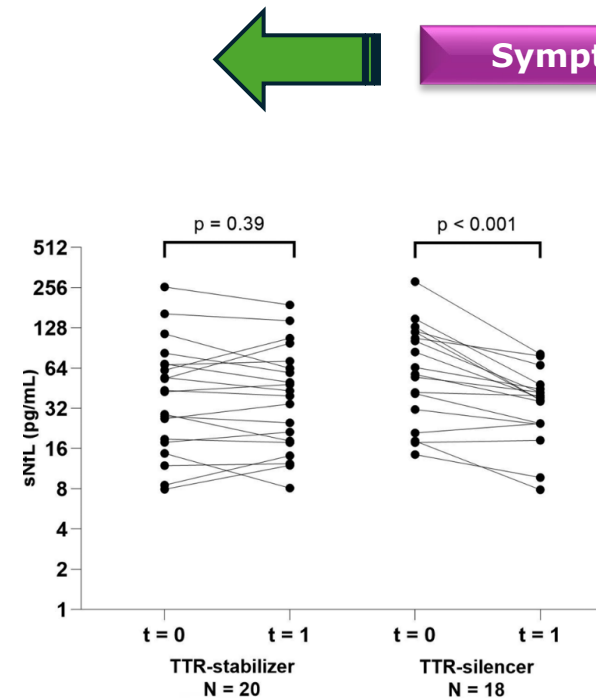
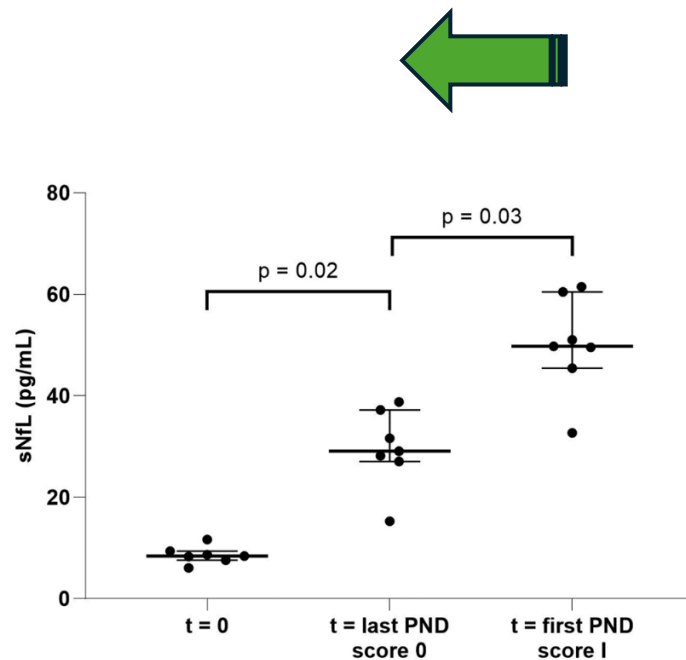
RESEARCH ARTICLE

OPEN ACCESS

Check for updates

## Longitudinal analysis of serum neurofilament light chain levels as marker for neuronal damage in hereditary transthyretin amyloidosis

- Elevated sNfL levels identify **asymptomatic ATTRv patients**.
- **Clinical progression** correlates with sNfL progression.
- **Stabilizers stabilize sNfL; silencers reduce sNfL**.



# Genetics are a must in ATTR!!

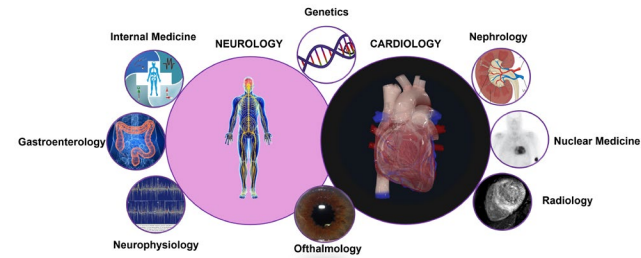
## DIAGNOSIS (ATTRv vs ATTRwt)



## CARRIERS

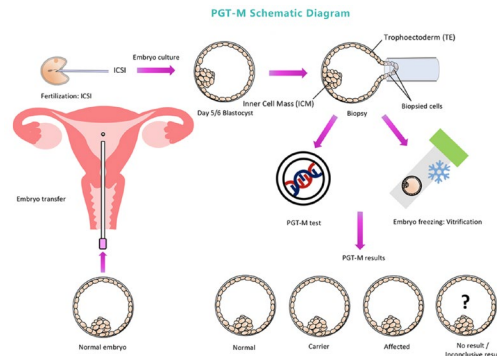


## FOLLOW-UP

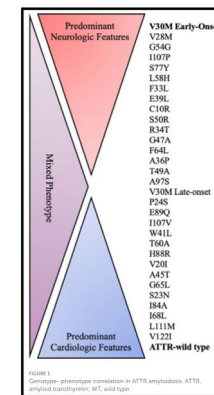


## TREATMENT

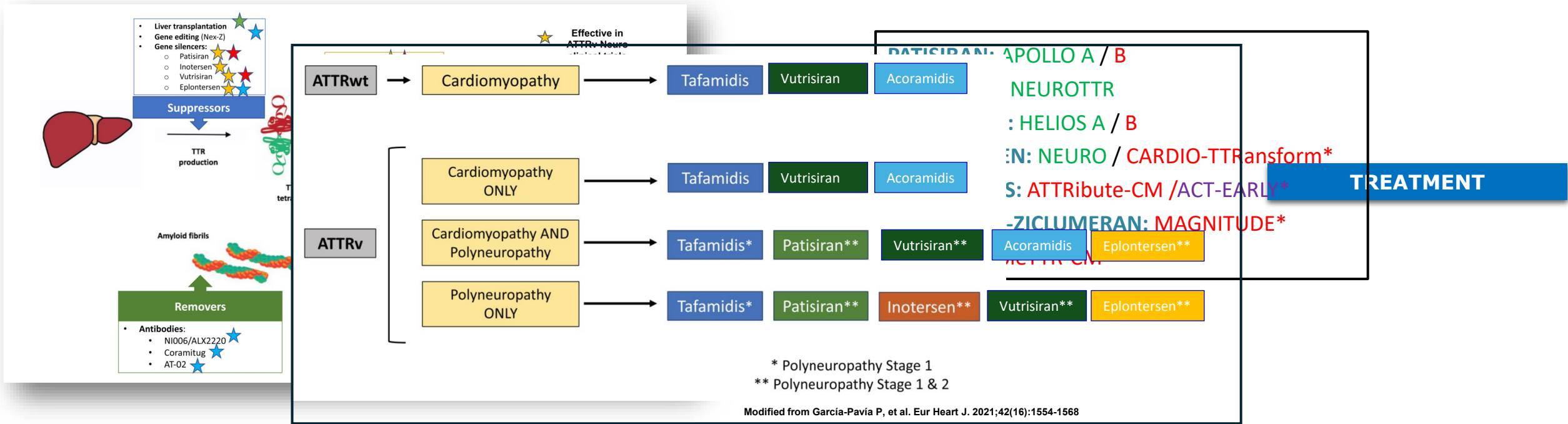
## GENETIC COUNSELING/REPRODUCTIVE PLANNING



## PROGNOSIS



# Genetics are a must in ATTR!!





**¡MUCHAS GRACIAS POR LA ATENCIÓN!**



**Complejo Hospitalario Universitario A Coruña**

