

Precision Medicine in cardiology: our classification of cardiomyopathies



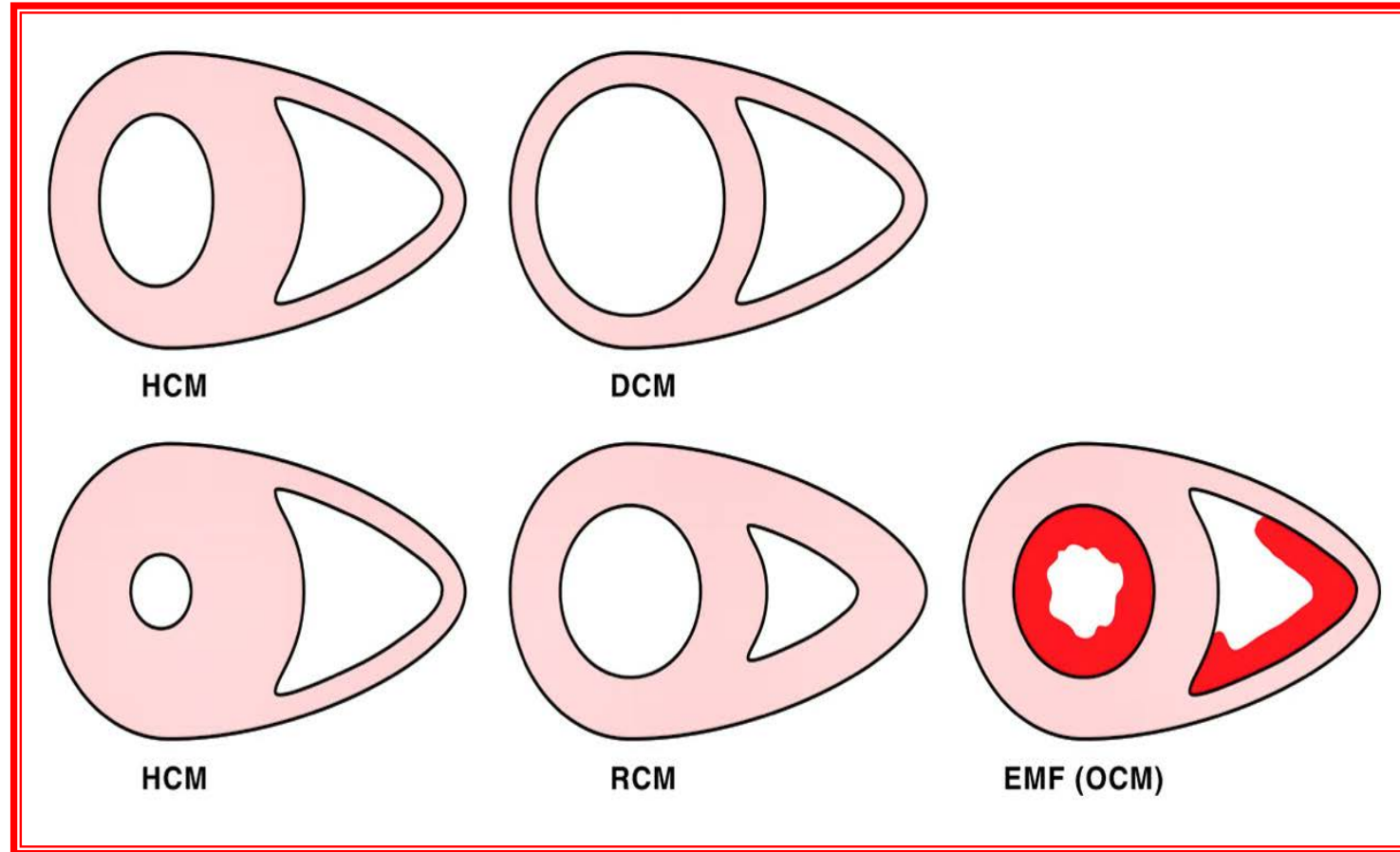
THE common cause of isolated myocardial disease is coronary disease ; but there are many other causes, mostly obscure and relatively rare. When patients develop apparently causeless heart-failure, diagnosis of myocarditis is sometimes made, but usually the condition is attributed either to a previous hypertension or painless coronary disease. Often such patients die unexpectedly and pathological investigation may not reveal the cause and nature of the disease. The present study is concerned with these myocardial diseases which are either isolated or so nearly isolated that disease in other systems is of a minor degree and significant only as an aid to diagnosis. The term cardiomyopathy is used here to indicate isolated non-coronary myocardial disease.

- The St.Cyres lecture, delivered in London on June 13, 1956.

Cardiomyopathies: The beginning.

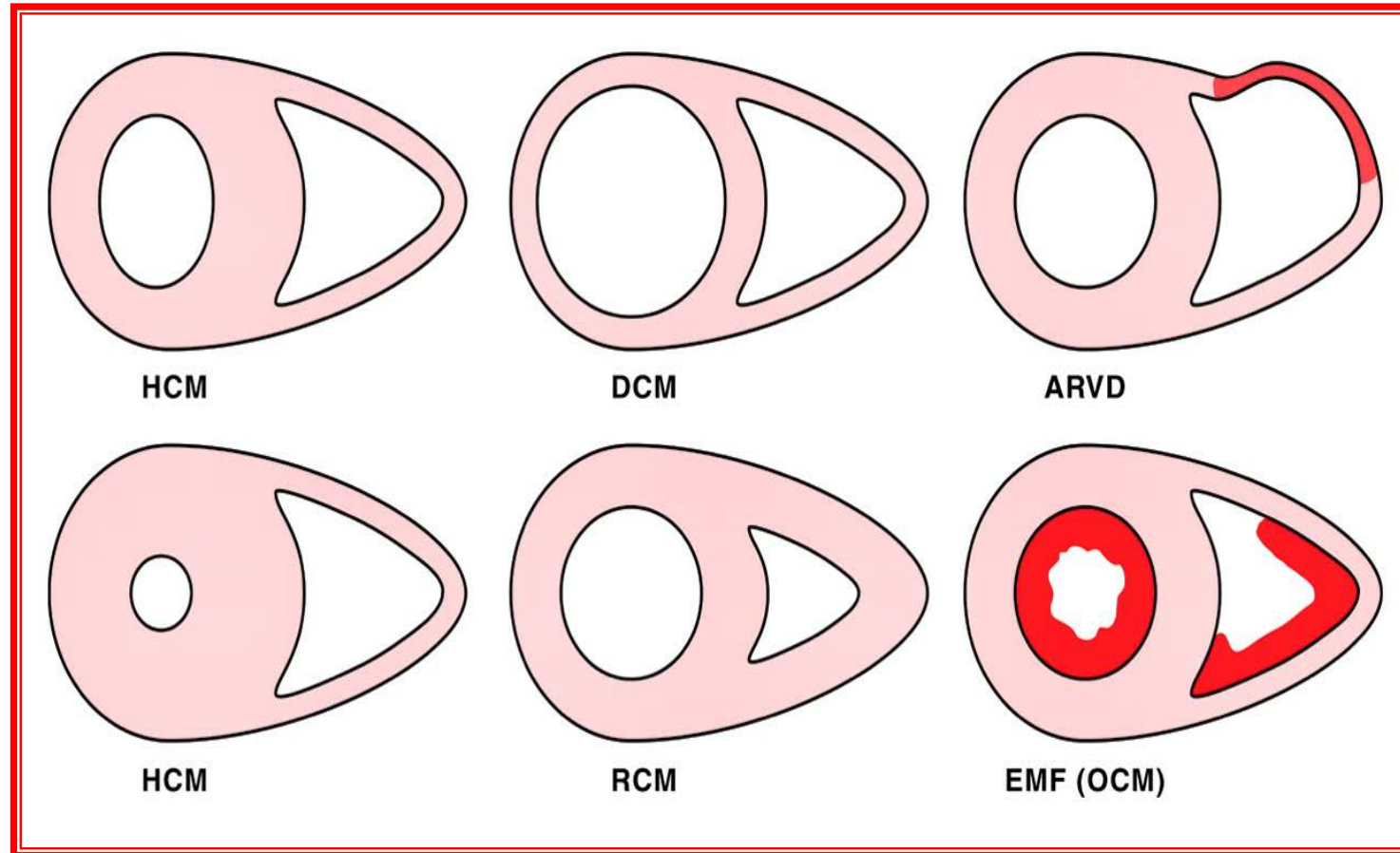
- “primary myocardial disorders not caused by clearly identifiable structural abnormalities, such as valve disease and coronary artery stenoses, or systemic disorders.”
 - J. Goodwin
- “A disorder of heart muscle of unknown cause.”
 - C. Oakley

Classification of Cardiomyopathies



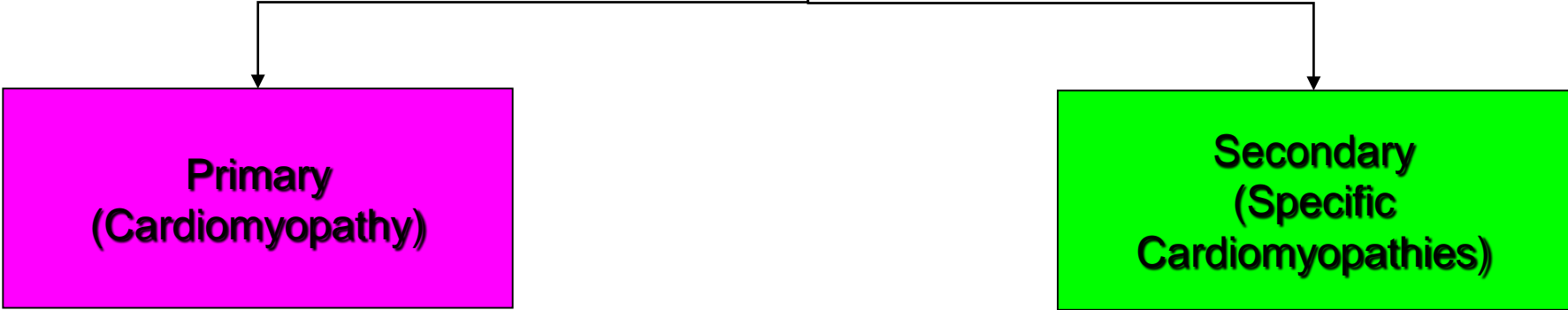
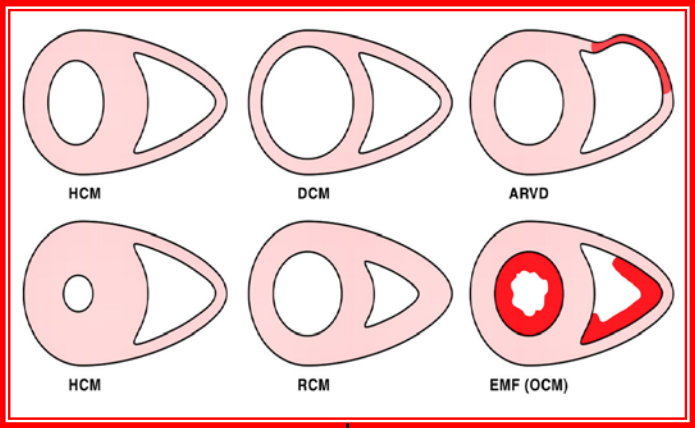
Adapted from Davies M. Heart 2000;83:469-474

Classification of Cardiomyopathies: 1995



Davies M. Heart 2000;83:469-474

Classification of Cardiomyopathies: 1995





Specific Heart Muscle Disease

Ischaemic

Valvular

Hypertensive


Inflammatory

Metabolic

Neuromuscular disorders


Sensitivity and toxic reactions

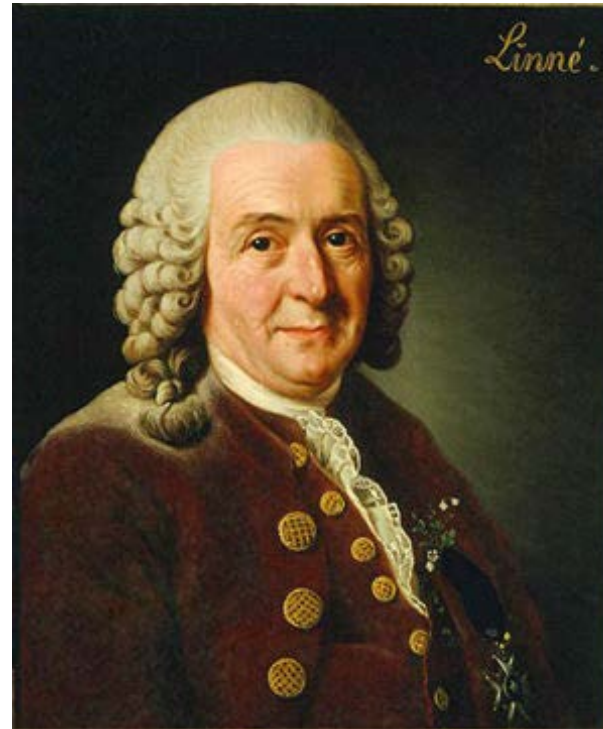
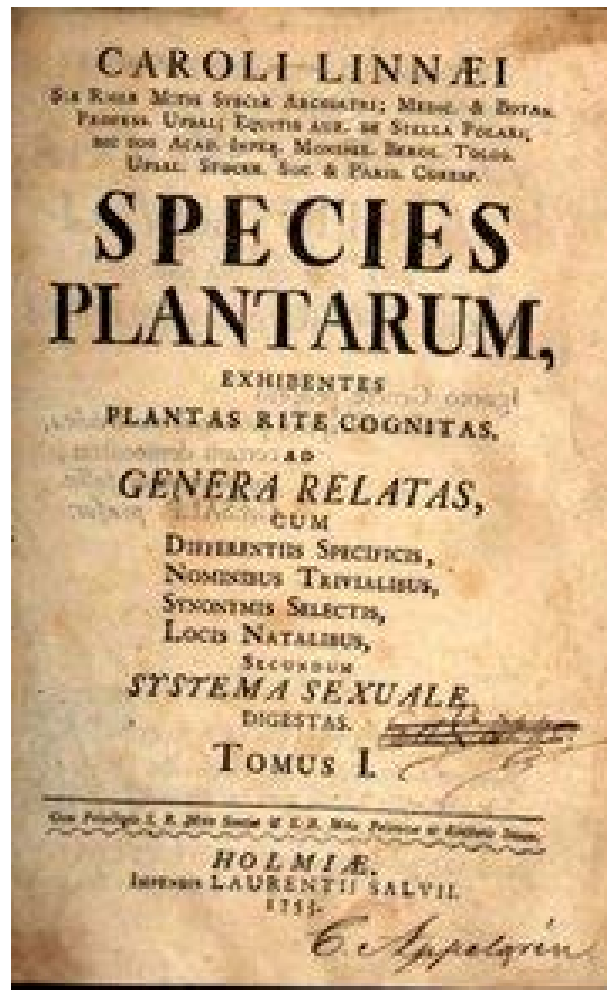
General system disease



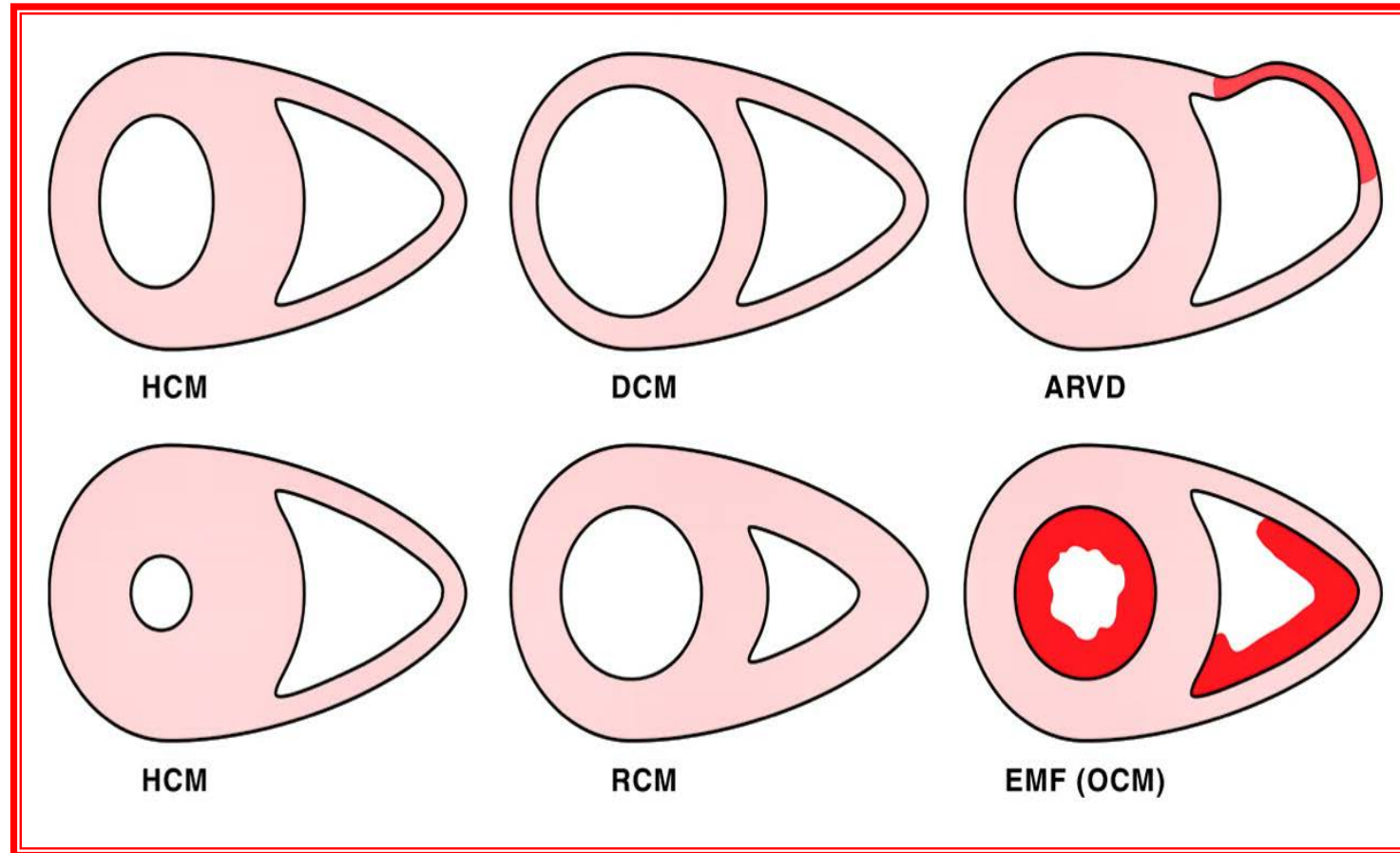


Limitations

- Historical legacy
 - Contradictory, confusing terminology, many situations (rare and common).
 - Evolving or early phenotypes
 - Emerging complexity (aetiology and phenotypes, overlapping diseases)
 - Aetiology specific treatments
- 



Classification of Cardiomyopathies: 1995



Davies M. Heart 2000;83:469-474

AHA Scientific Statement

Contemporary Definitions and Classification of the Cardiomyopathies

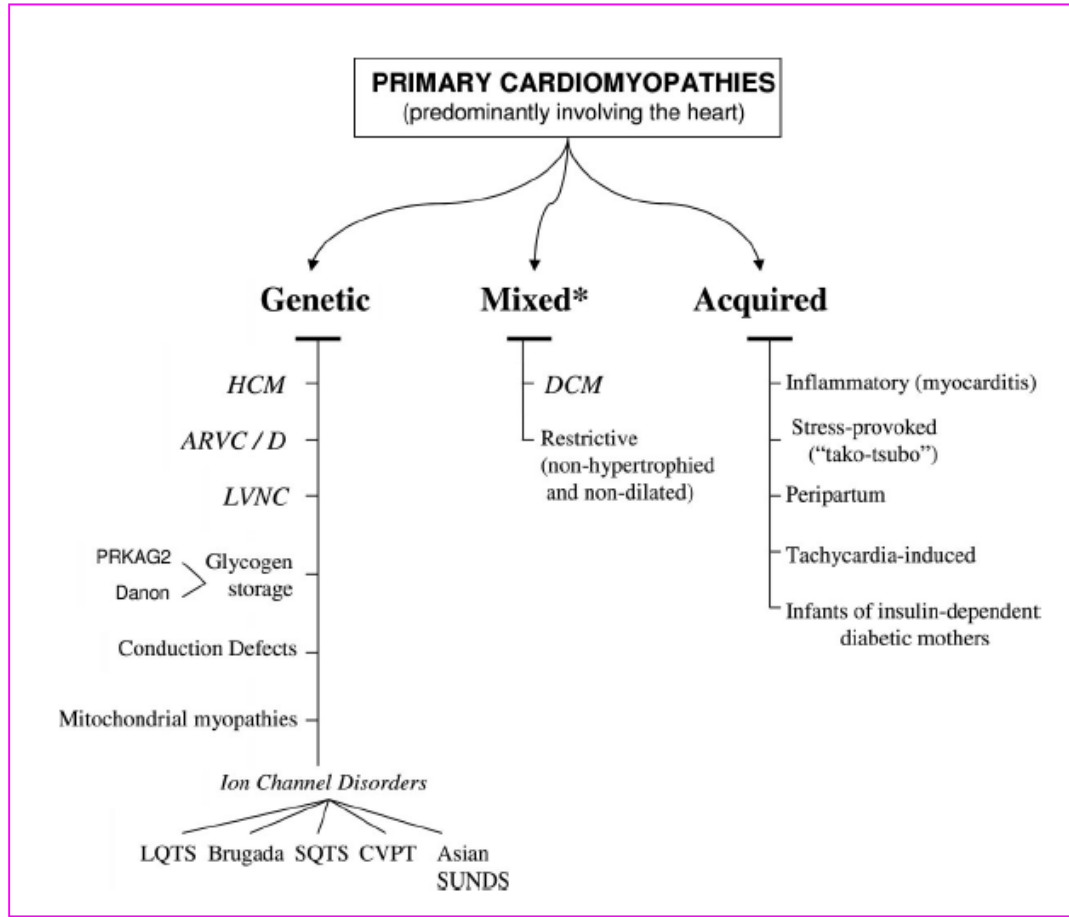
**An American Heart Association Scientific Statement From the Council on
Clinical Cardiology, Heart Failure and Transplantation Committee;
Quality of Care and Outcomes Research and Functional Genomics and
Translational Biology Interdisciplinary Working Groups; and Council on
Epidemiology and Prevention**

Barry J. Maron, MD, Chair; Jeffrey A. Towbin, MD, FAHA; Gaetano Thiene, MD;
Charles Antzelevitch, PhD, FAHA; Domenico Corrado, MD, PhD; Donna Arnett, PhD, FAHA;
Arthur J. Moss, MD, FAHA; Christine E. Seidman, MD, FAHA; James B. Young, MD, FAHA

Maron B et al. Circulation 2006;113:1807-1816

AHA Scientific Statement

Contemporary Definitions and Classification of the Cardiomyopathies



Secondary Cardiomyopathies

Infiltrative*

- Amyloidosis (primary, familial autosomal dominant†, senile, secondary forms)
- Gaucher disease‡
- Hurler's disease‡
- Hunter's disease‡
- Storage‡
- Hemochromatosis
- Fabry's disease‡
- Glycogen storage disease‡ (type II, Pompe)
- Niemann-Pick disease‡

Toxicity

- Drugs, heavy metals, chemical agents

Endomyocardial

- Endomyocardial fibrosis

- Hyper eosinophilic syndrome (Löffler's endocarditis)

Endomyocardial

Endomyocardial fibrosis

Hyper eosinophilic syndrome (Löffler's endocarditis)

Hypertrophy†

- Hyperparathyroidism
- Pheochromocytoma
- Acromegaly

Cardiofacial

- Noonan syndrome‡
- Lentiginosis‡

Neuromuscular/neurological

- Friedreich's ataxia†
- Duchenne-Becker muscular dystrophy†
- Emery-Dreifuss muscular dystrophy†
- Myotonic dystrophy†

Neurofibromatosis†

- Tuberous sclerosis†

Nutritional deficiencies

- Beriberi (thiamine), pellagra, scurvy, selenium, carnitine, kwashiorkor

Autoimmune/collagen

- Systemic lupus erythematosus
- Dermatomyositis
- Rheumatoid arthritis
- Scleroderma
- Polyarteritis nodosa

Electrolyte imbalance

Consequence of cancer therapy

- Anthracyclines: doxorubicin (adriamycin), daunorubicin
- Cyclophosphamide

Radiation

*Accumulation of abnormal substances between myocytes (ie, extracellular).

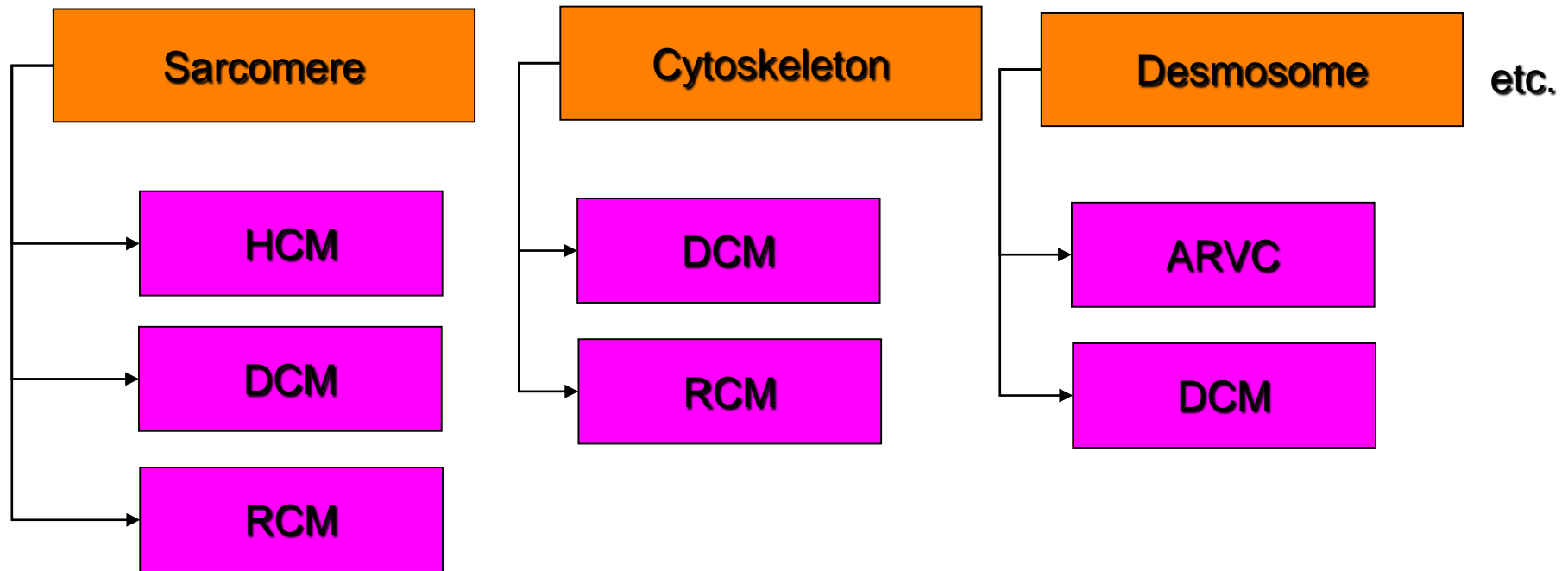
†Genetic (familial) origin.

‡Accumulation of abnormal substances within myocytes (ie, intracellular).

Classification of the cardiomyopathies: a position statement from the european society of cardiology working group on myocardial and pericardial diseases

Perry Elliott, Bert Andersson, Eloisa Arbustini, Zofia Bilinska, Franco Cecchi, Philippe Charron, Olivier Dubourg, Uwe Kühl, Bernhard Maisch, William J. McKenna, Lorenzo Monserrat, Sabine Pankuweit, Claudio Rapezzi, Petar Seferovic, Luigi Tavazzi, and Andre Keren*

“Molecular” classification



The clinical pathway

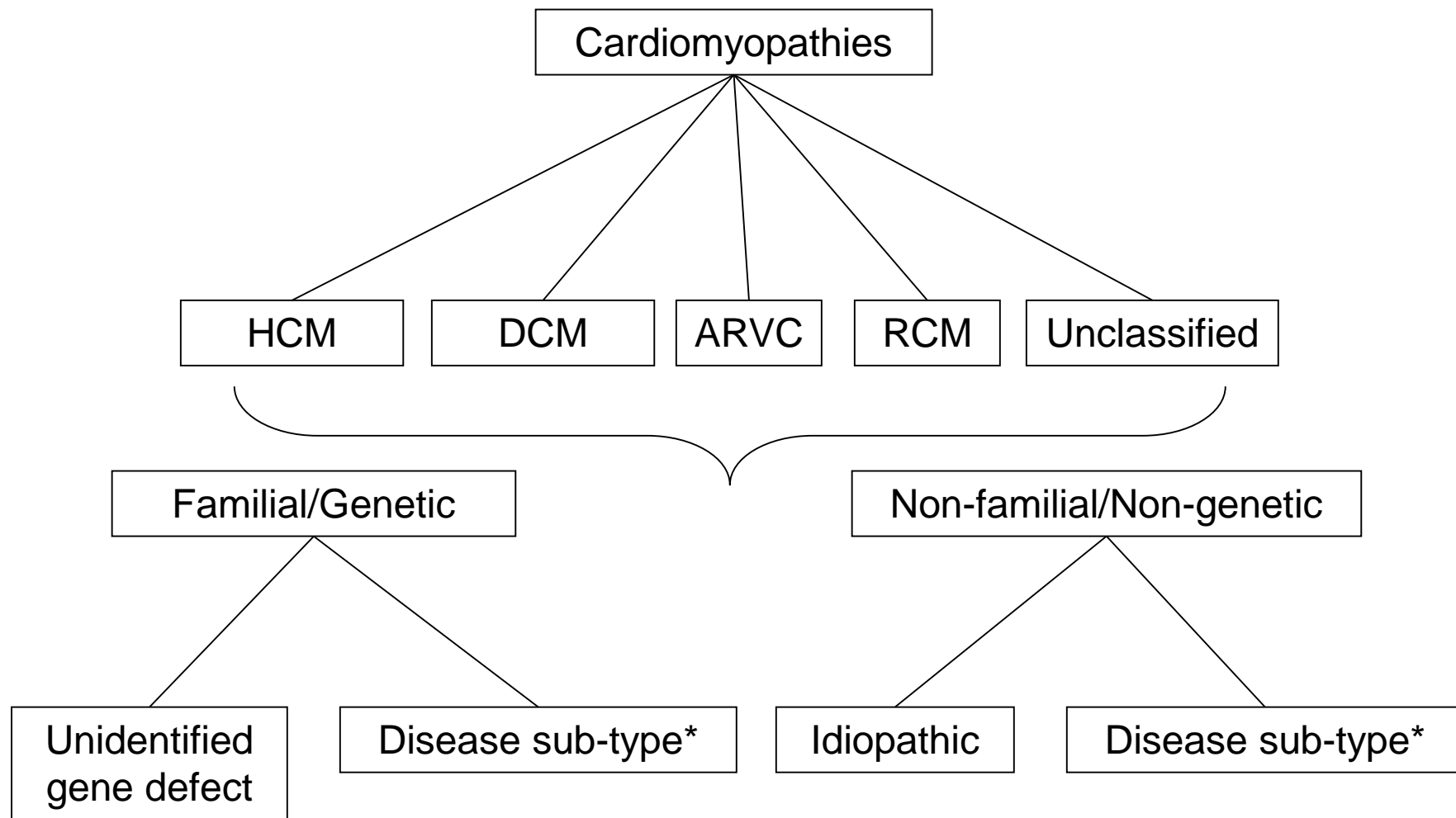


The patient is complaining of sarcomere protein disease.

**Symptoms/
Signs**

**Phenotype
(HCM/DCM...)**

Diagnosis

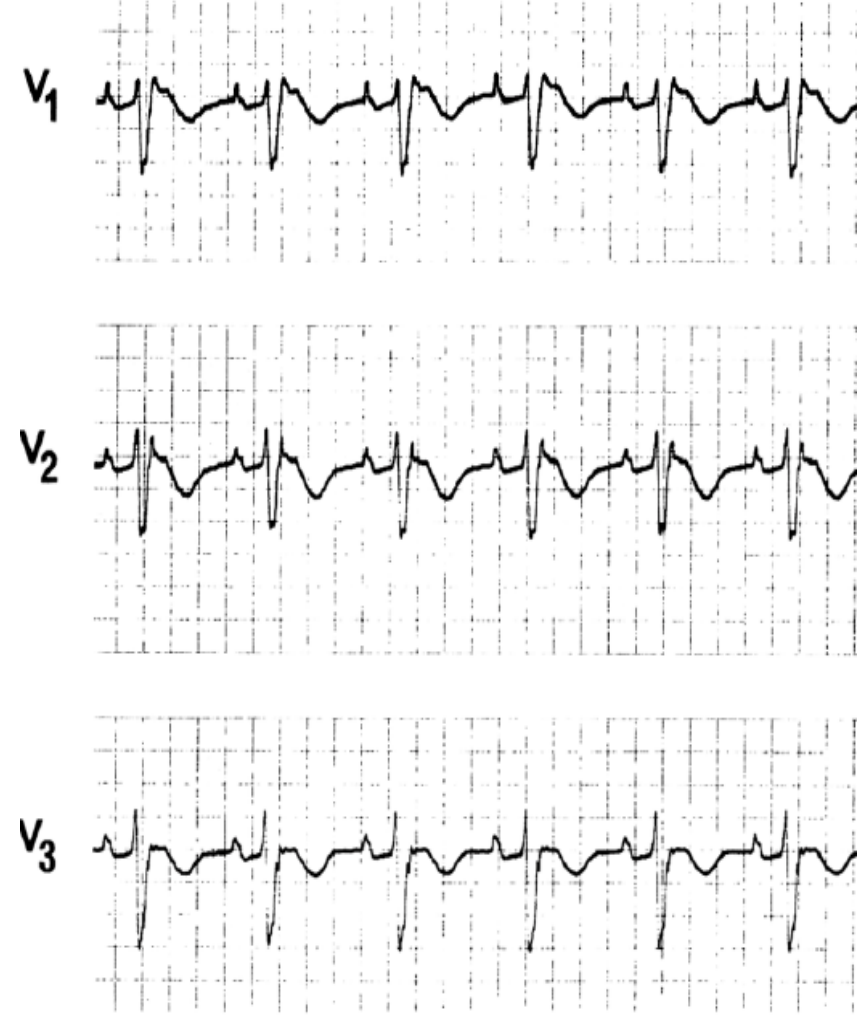
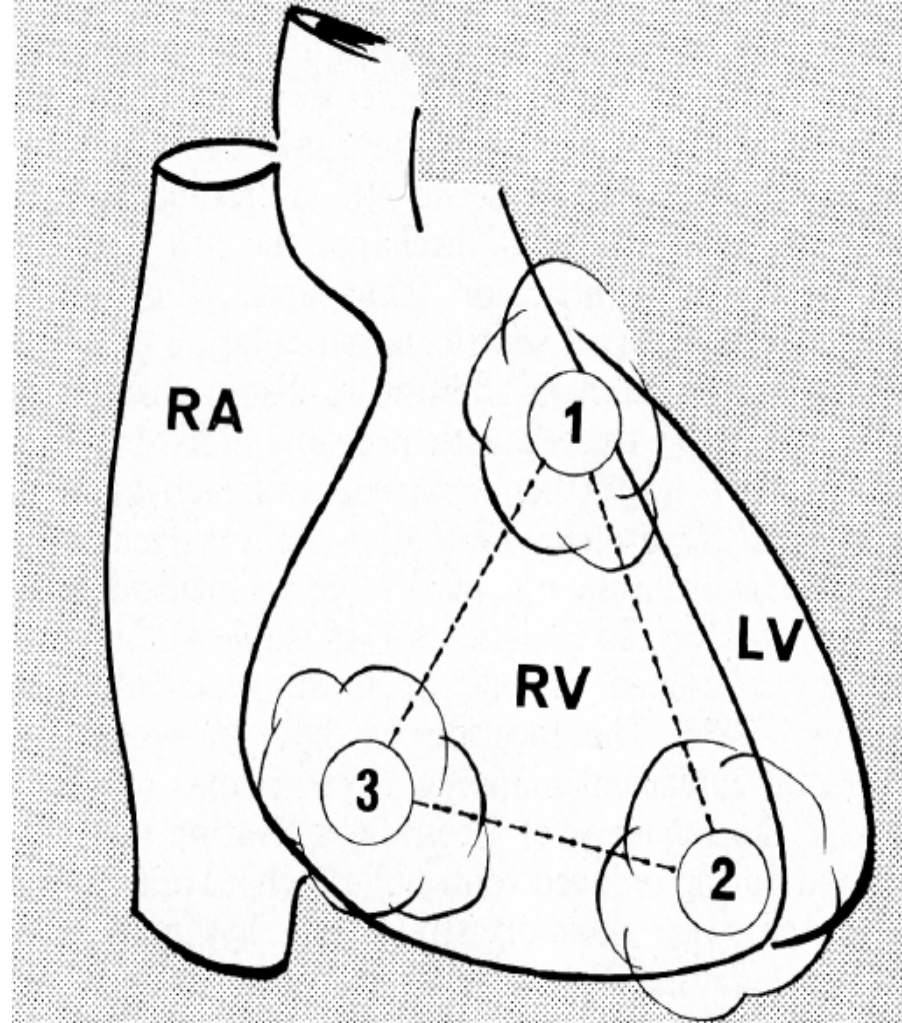


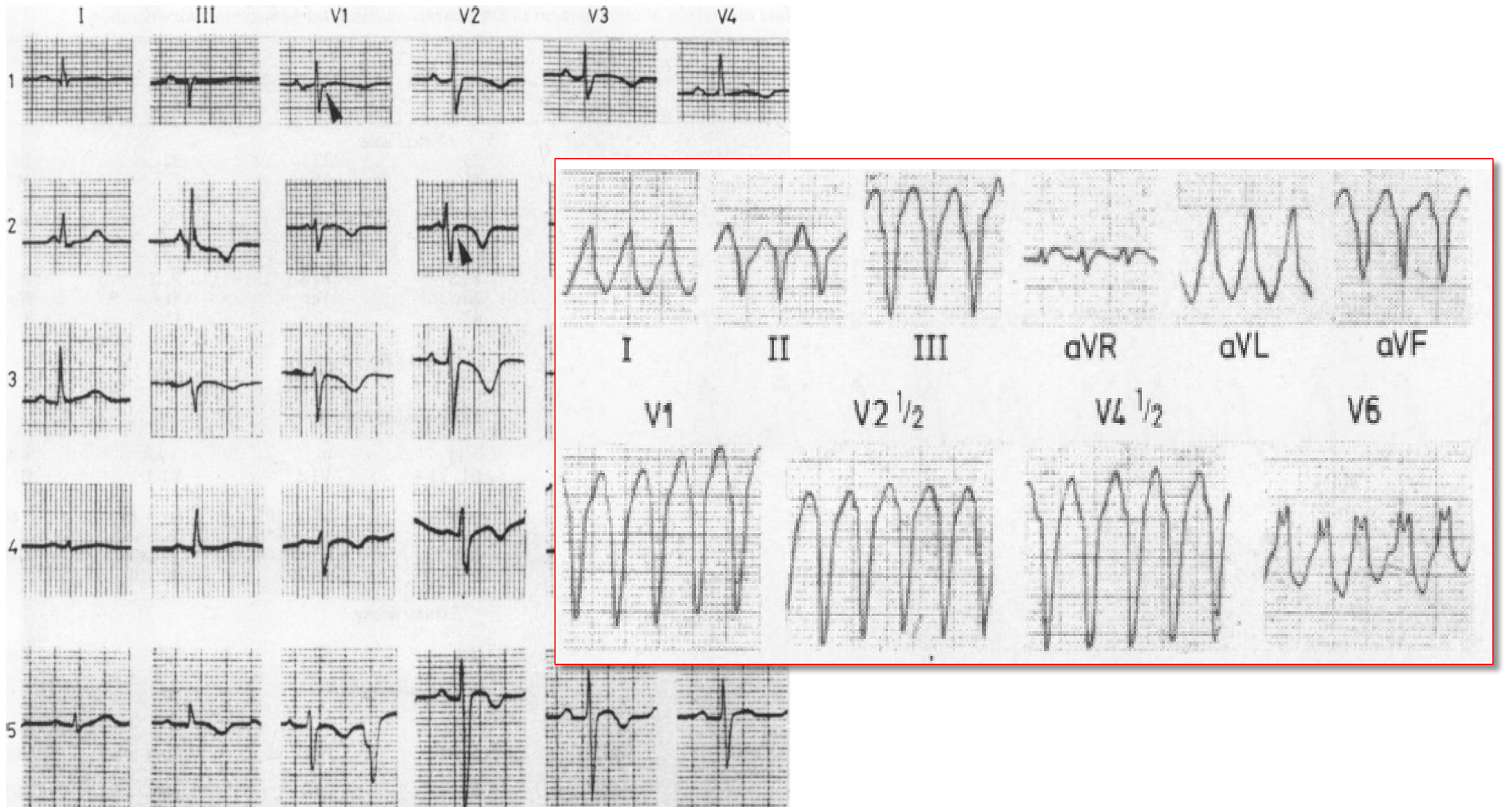
European WG on Myocardial and Pericardial Diseases (EHJ 2007)

Limitations: 2021

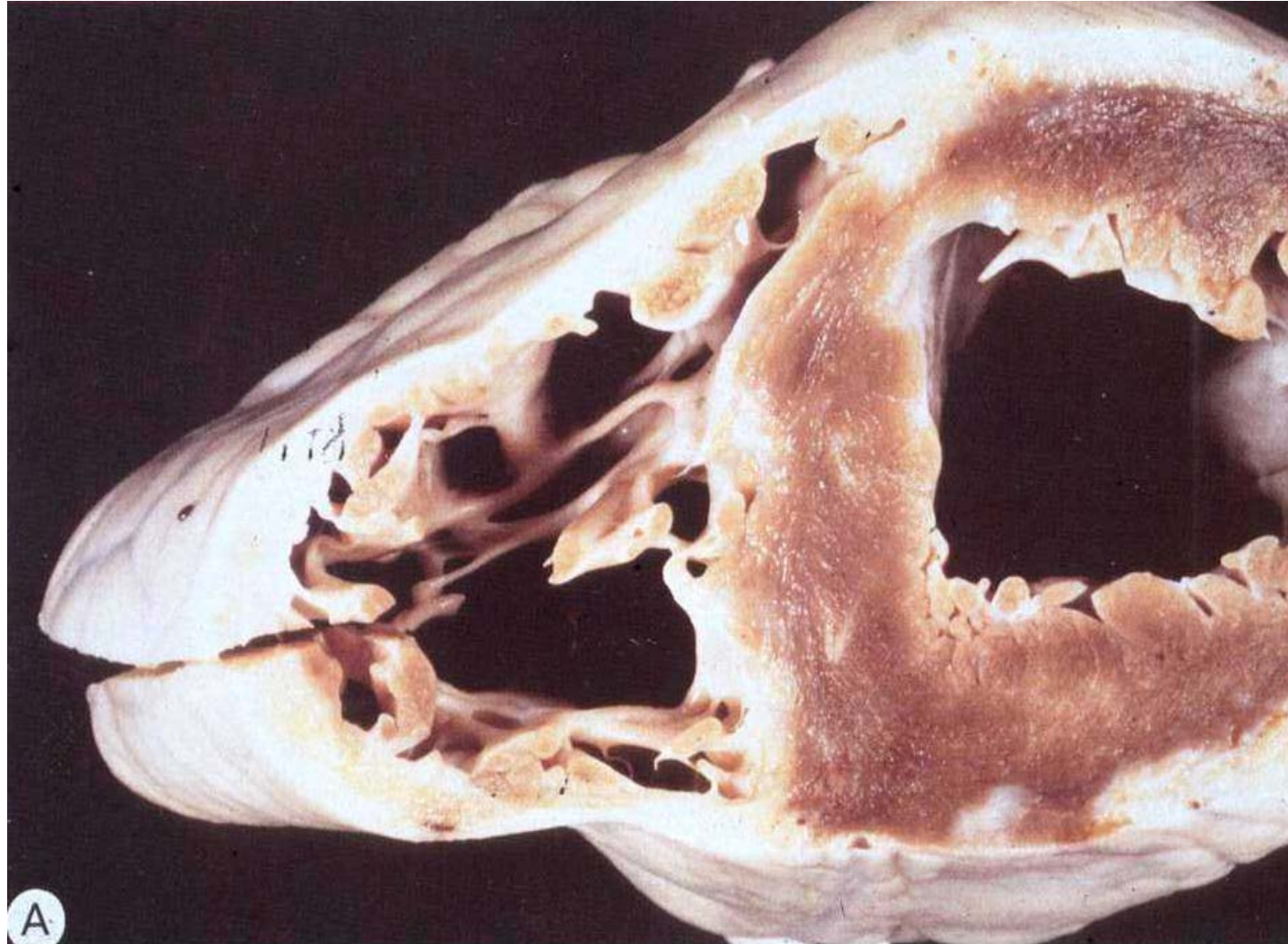
- Cardiomyopathies no longer “idiopathic”
- Overlapping phenotypes (“one gene many diseases, one disease many genes”)
- New phenotypes (LVNC, TTS)
- No consideration of advances in molecular biology

Arrhythmogenic Right Ventricular Cardiomyopathy

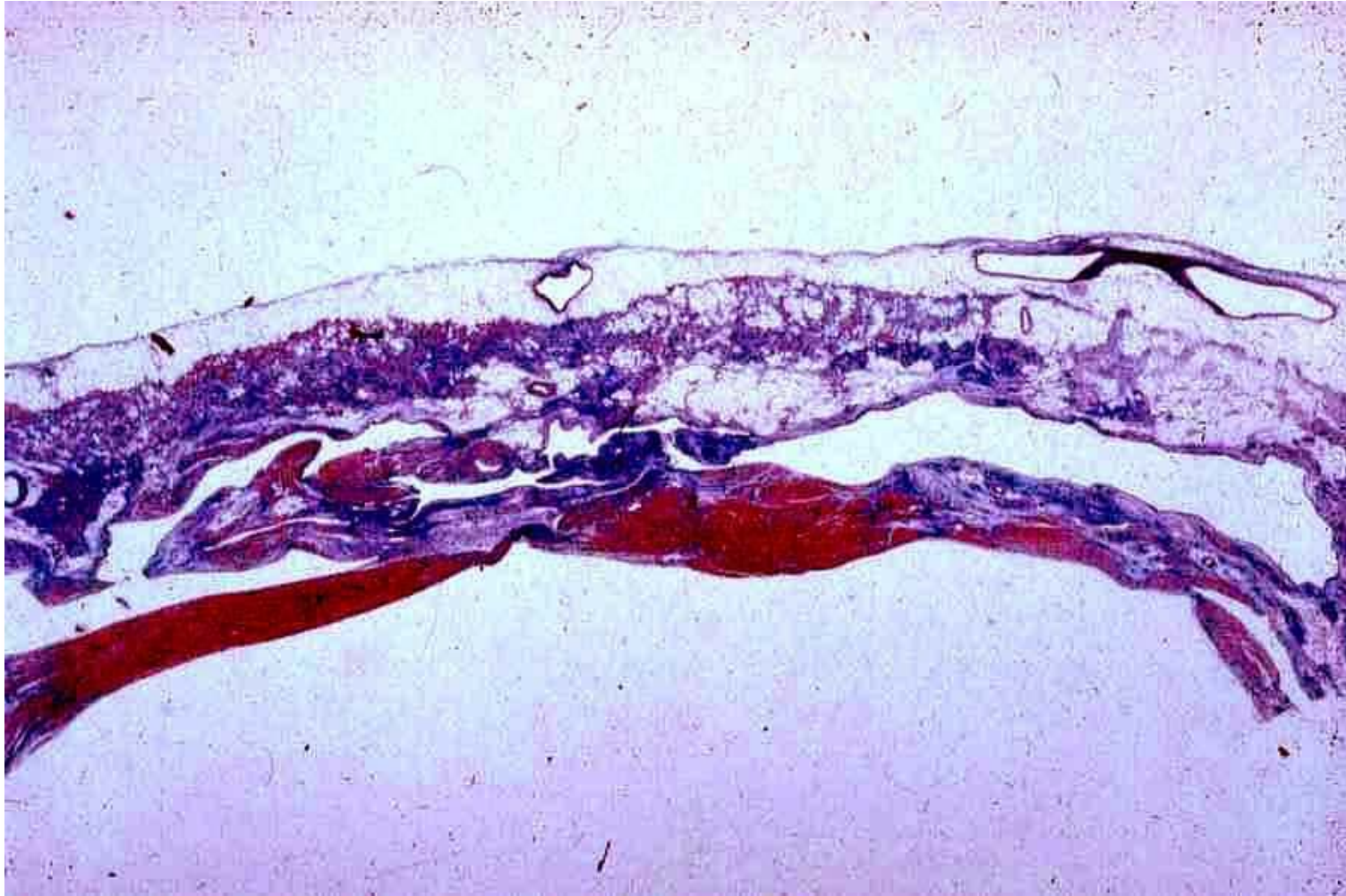




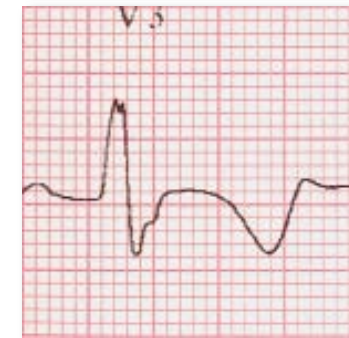
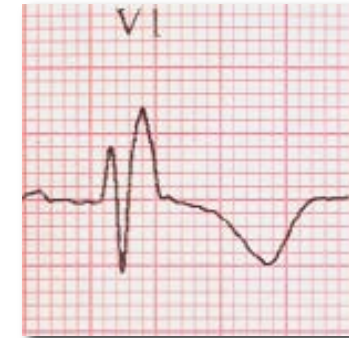
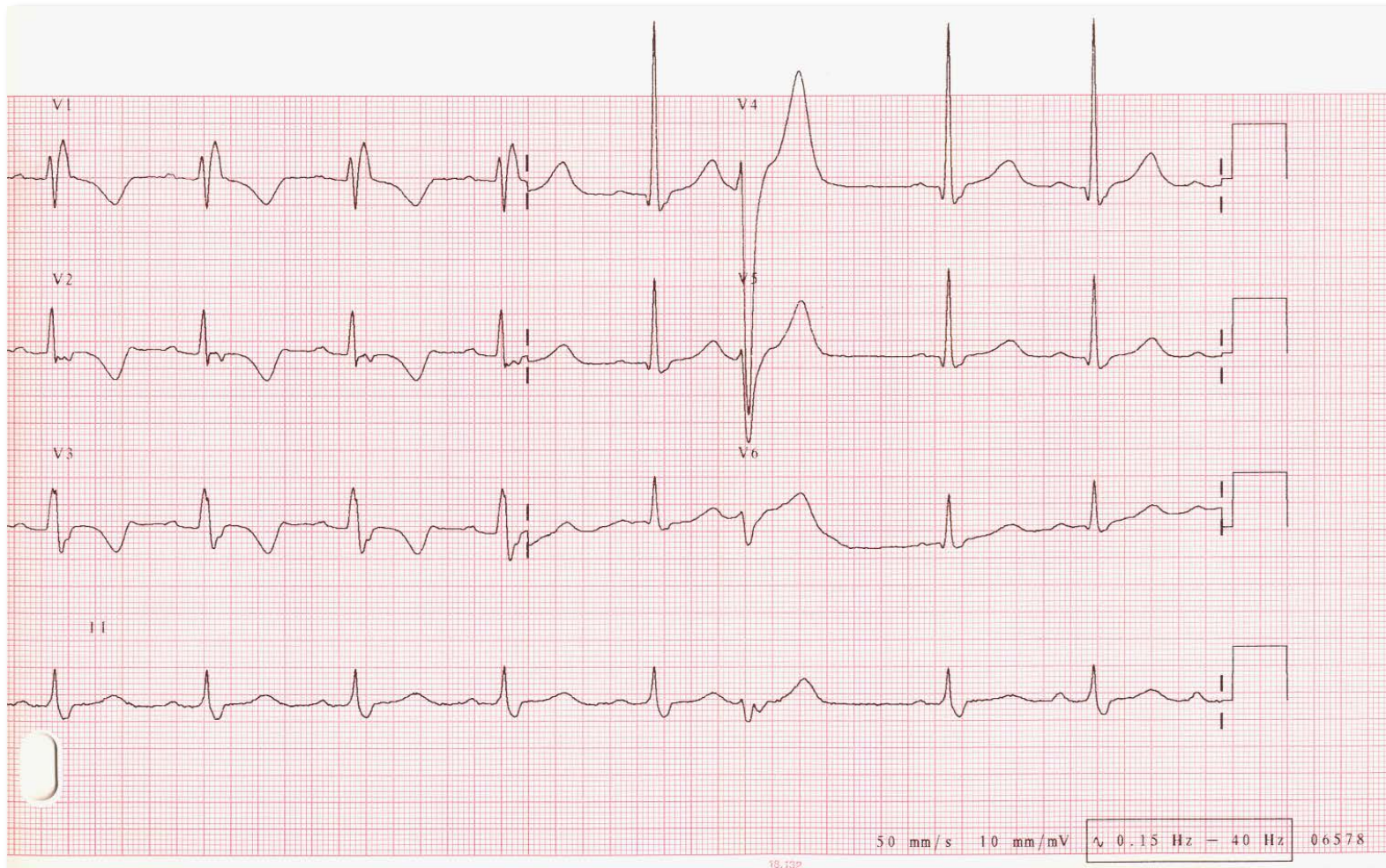
Arrhythmogenic Right Ventricular Cardiomyopathy

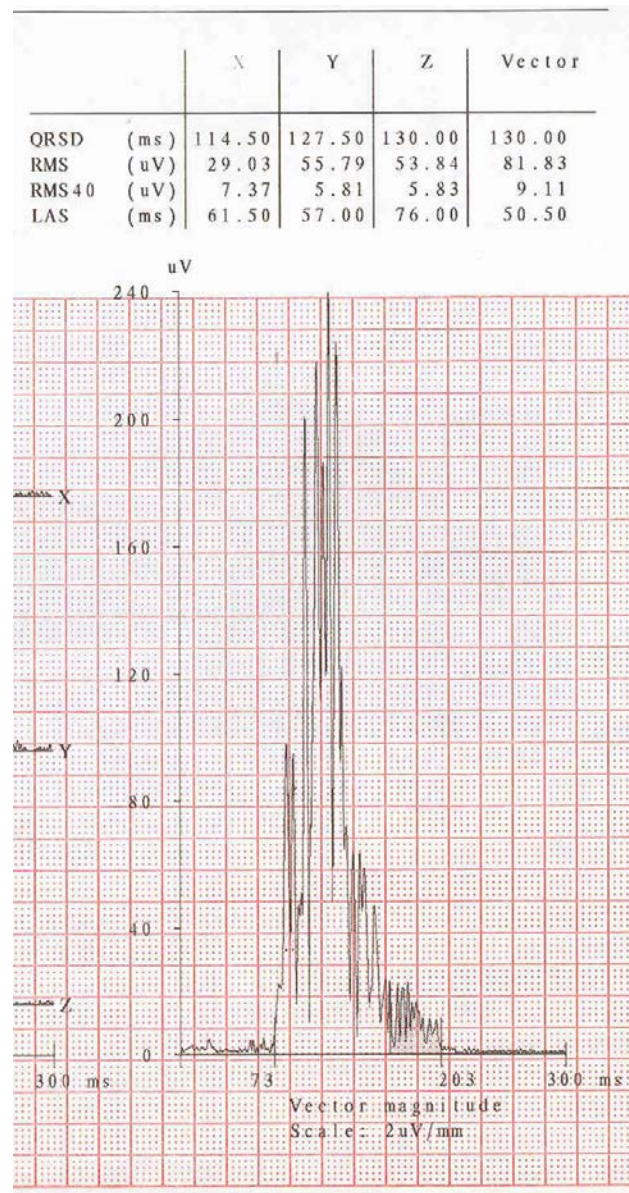
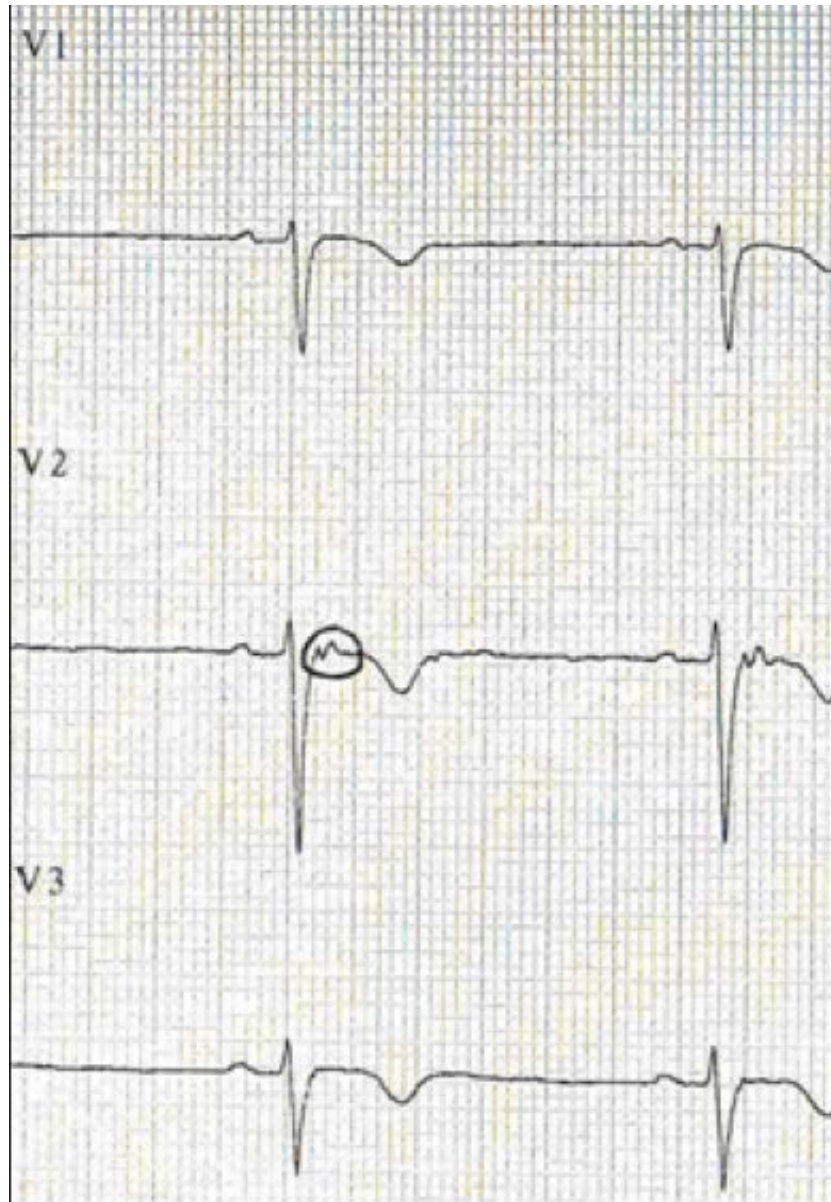


Arrhythmogenic Right Ventricular Cardiomyopathy



ARVC: Electrocardiography

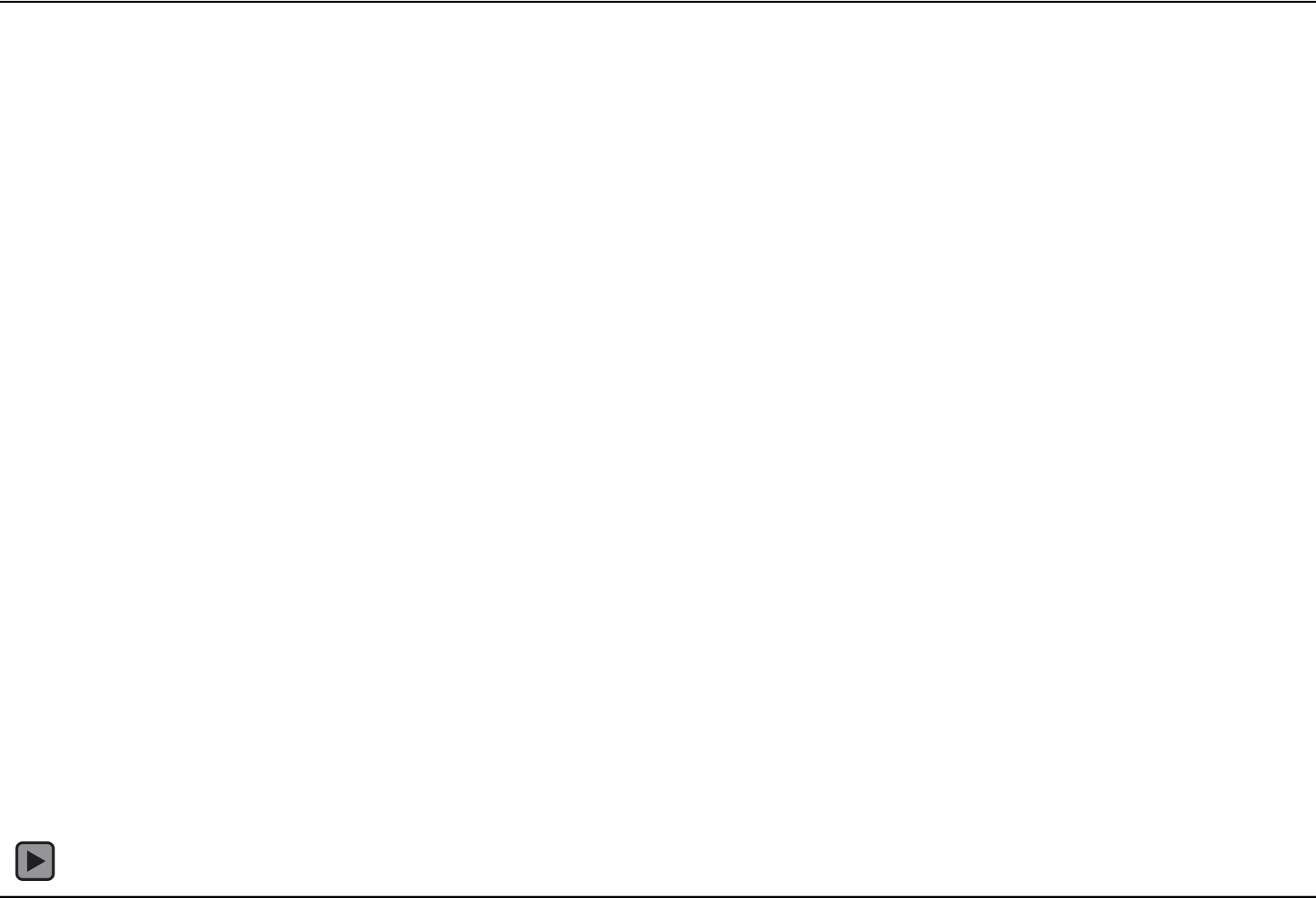


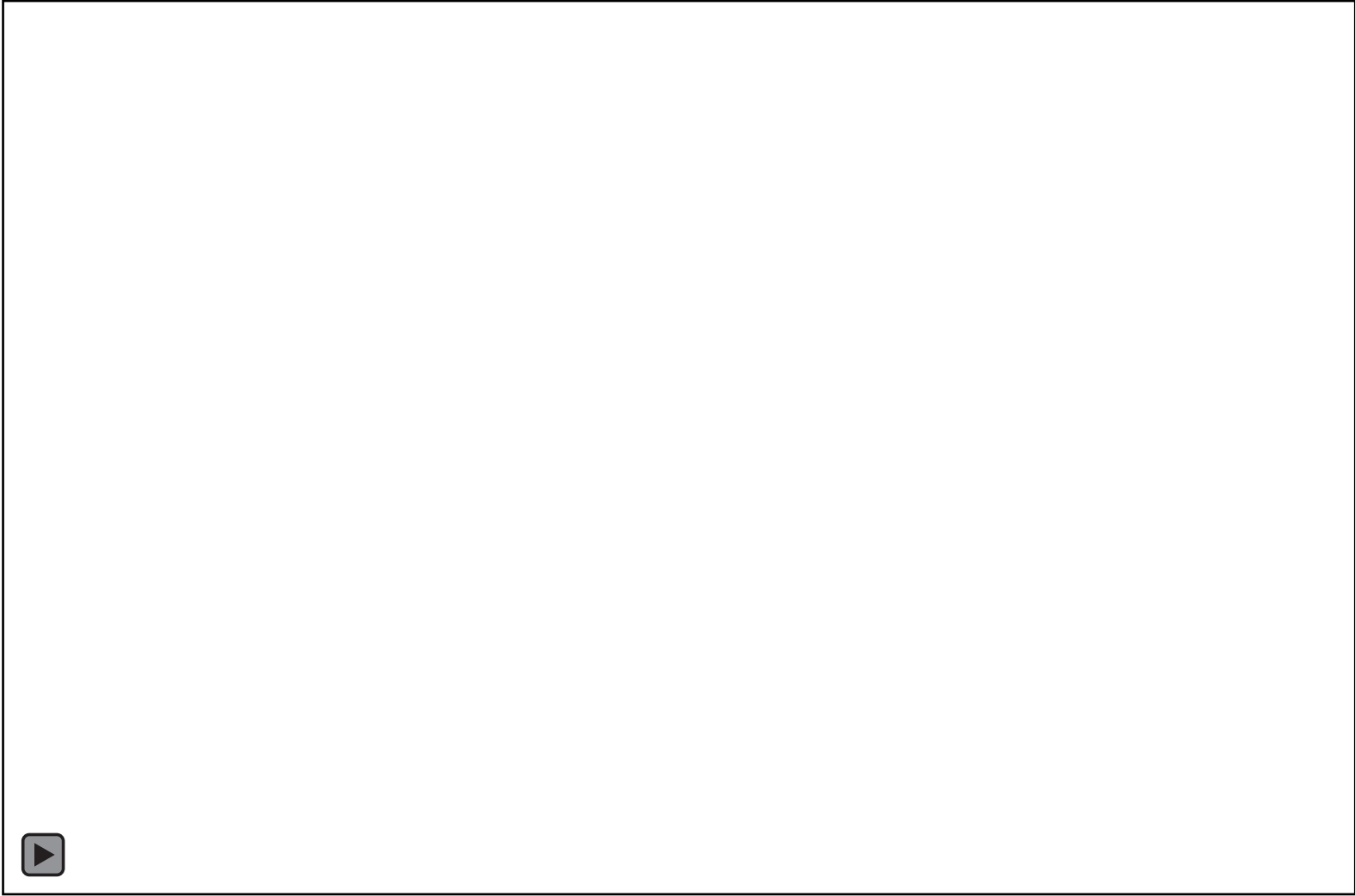


>114 ms

< 20 uV

>38 msec



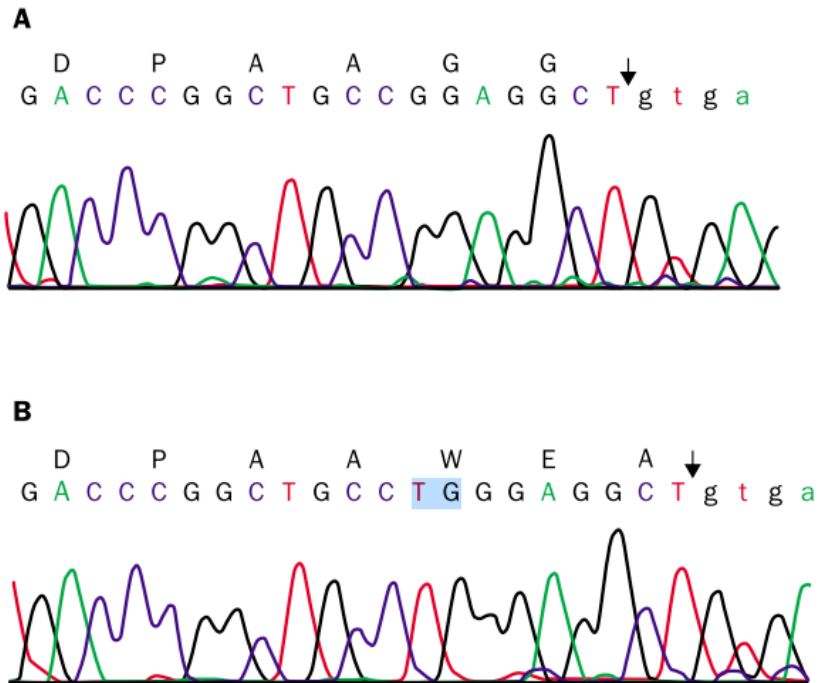


Articles

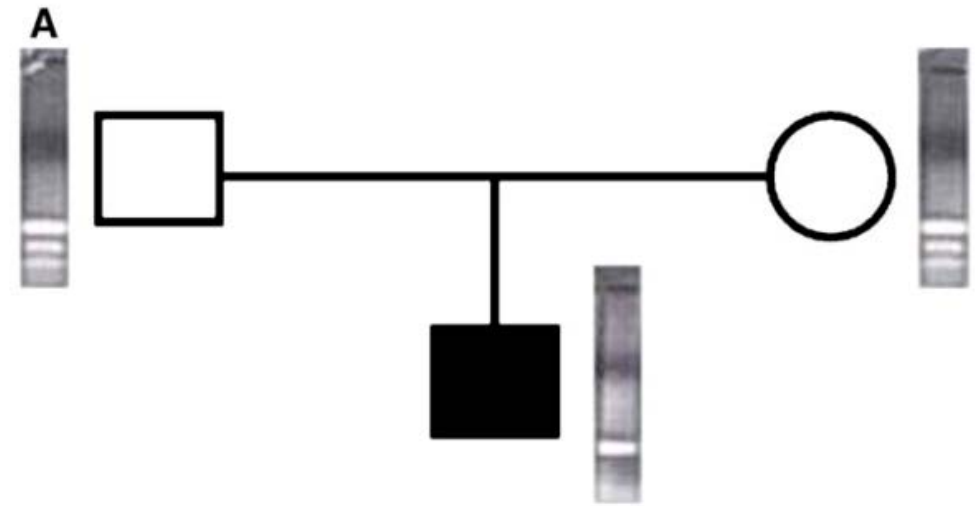
Identification of a deletion in plakoglobin in arrhythmogenic right ventricular cardiomyopathy with palmoplantar keratoderma and woolly hair (Naxos disease)

Godfrina McKoy, PhD, Nikos Protonotarios, MD, Andrew Crosby, PhD, Adalena Tsatsopoulou, MD, Aris Anastasakis, MD, Aman Coonar, MB, Mark Norman, MD, Christina Baboonian, PhD, Steve Jeffery, PhD, Prof William J McKenna, MD

Published: 17 June 2000



frameshift and premature termination of translation



+ ARVC

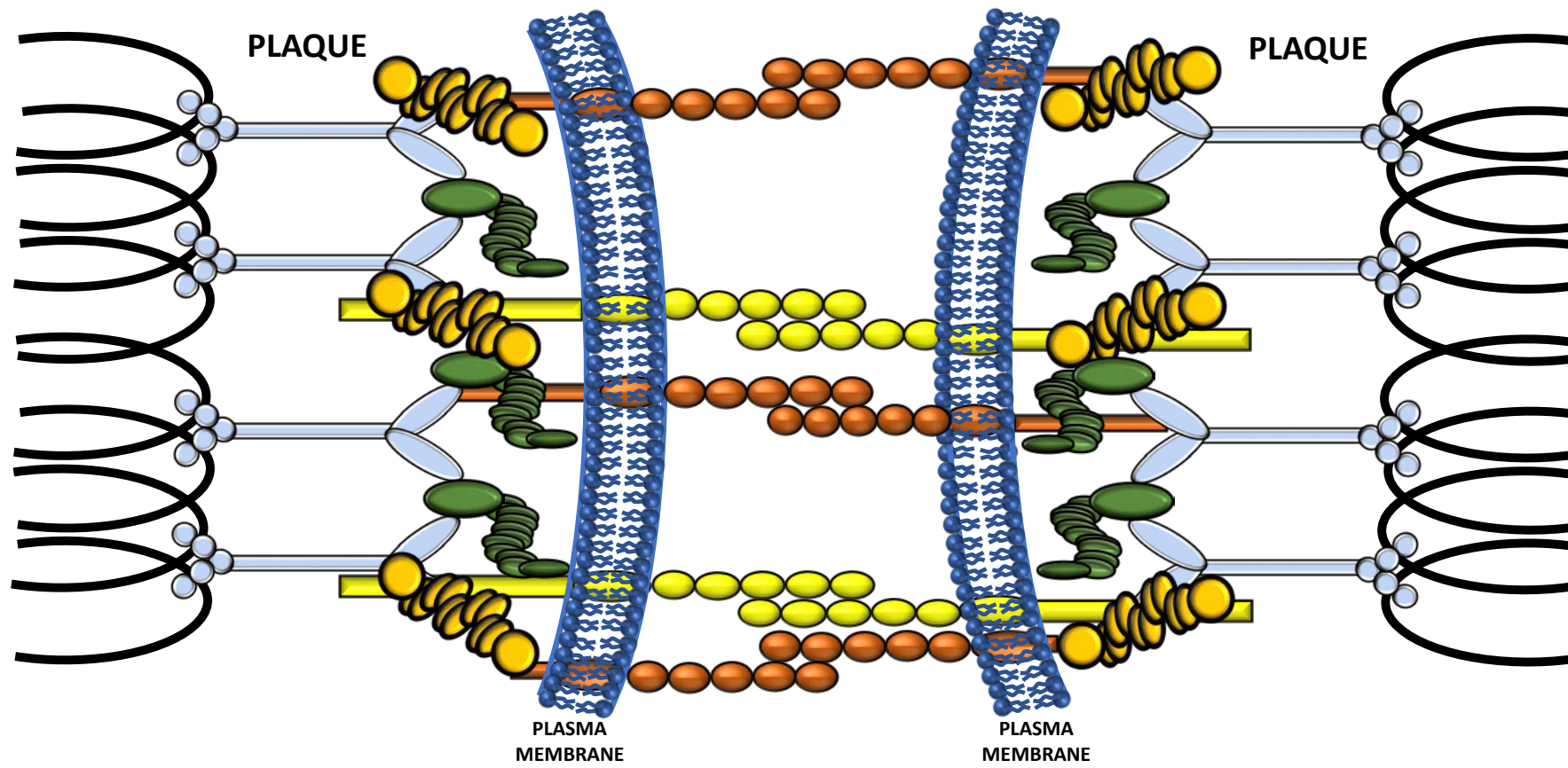






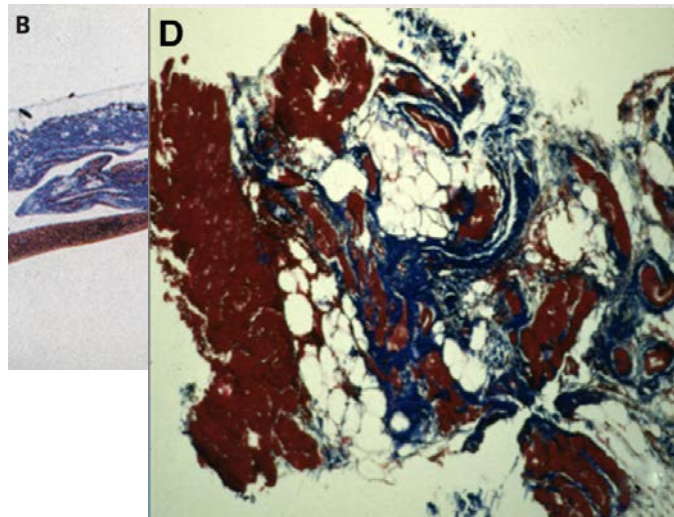
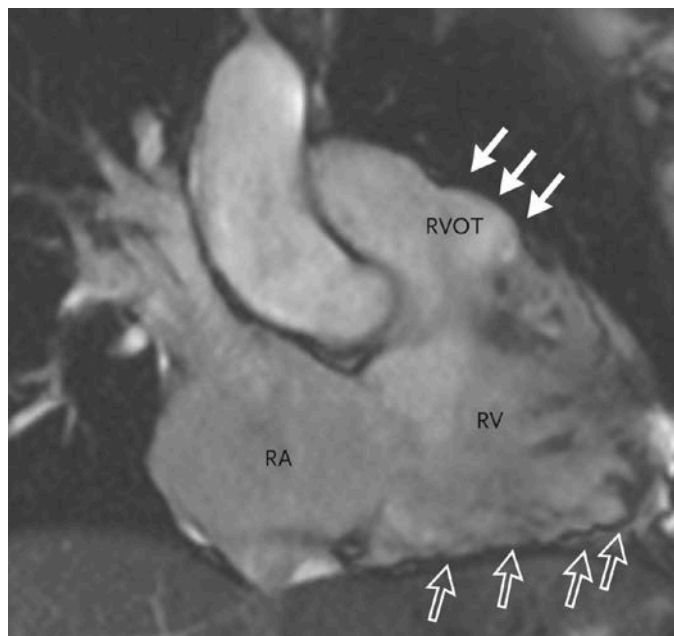


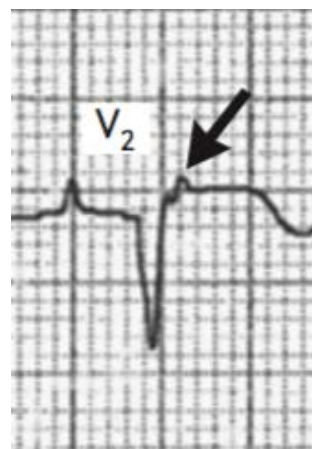
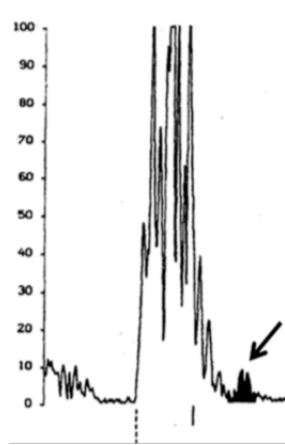
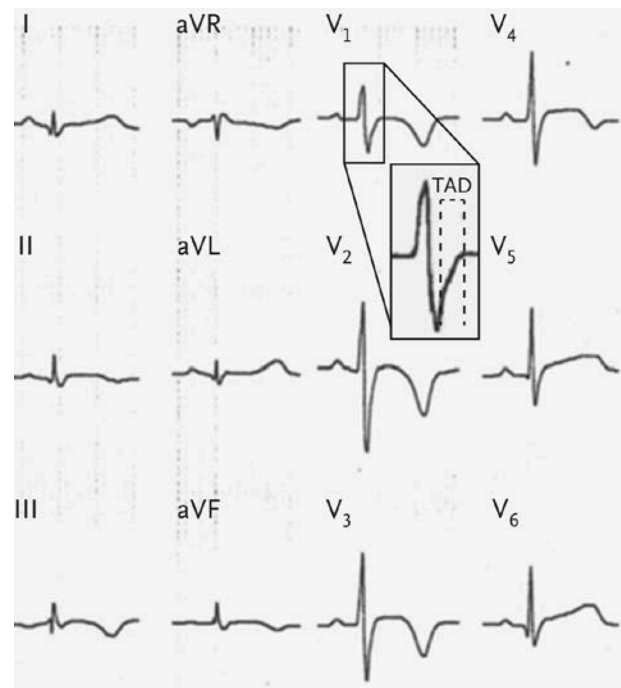
Figure 1. Schematic model of a desmosome.

	Desmoplakin		Desmoglein		Desmocollin		Plakoglobin		Plakophilin
									Intermediate filaments

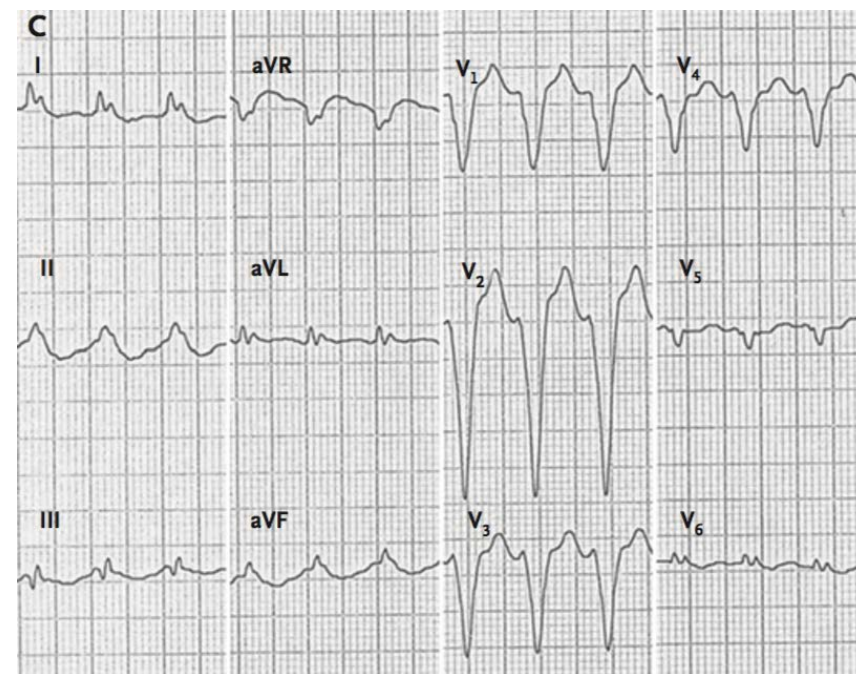
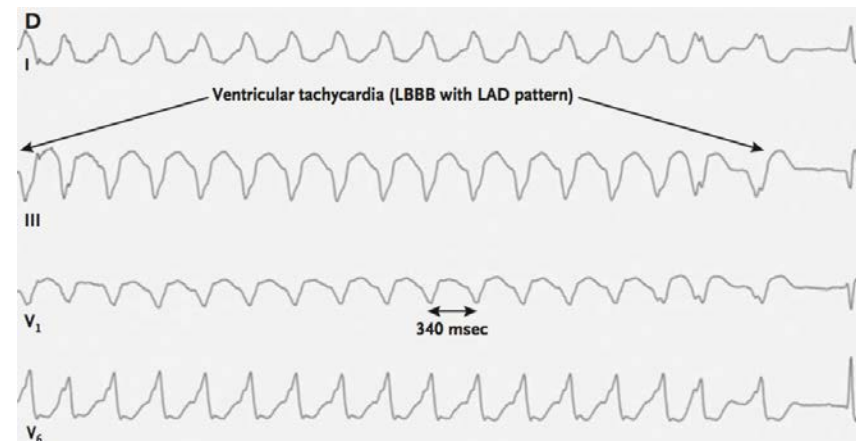
Structural



Electrical



Arrhythmic



Diagnosis of arrhythmogenic right ventricular cardiomyopathy/dysplasia

Proposed Modification of the Task Force Criteria

**Frank I. Marcus^{1*} Chair, William J. McKenna² Co-Chair, Duane Sherrill¹,
Cristina Basso³, Barbara Bauce³, David A. Bluemke⁴, Hugh Calkins⁵,
Domenico Corrado³, Moniek G.P.J. Cox⁶, James P. Daubert⁷, Guy Fontaine¹⁰,
Kathleen Gear¹, Richard Hauer⁶, Andrea Nava³, Michael H. Picard¹¹,
Nikos Protonotarios¹³, Jeffrey E. Saffitz¹², Danita M. Yoerger Sanborn¹¹,
Jonathan S. Steinberg⁹, Harikrishna Tandri⁵, Gaetano Thiene³, Jeffrey A. Towbin¹⁴,
Adalena Tsatsopoulou¹³, Thomas Wichter¹⁵, and Wojciech Zareba⁸**

Table 1 Comparison of original and revised task force criteria

Original task force criteria	Revised task force criteria
I. Global or regional dysfunction and structural alterations*	
Major	
<ul style="list-style-type: none"> Severe dilatation and reduction of RV ejection fraction with no (or only mild) LV impairment Localized RV aneurysms (akinetic or dyskinetic areas with diastolic bulging) Severe segmental dilatation of the RV 	<p>By 2D echo:</p> <ul style="list-style-type: none"> Regional RV akinesia, dyskinesia, or aneurysm and 1 of the following (end diastole): <ul style="list-style-type: none"> PLAX RVOT ≥ 32 mm (corrected for body size [PLAX/BSA] ≥ 19 mm/m²) PSAX RVOT ≥ 36 mm (corrected for body size [PSAX/BSA] ≥ 21 mm/m²) or fractional area change $\leq 33\%$ <p>By MRI:</p> <ul style="list-style-type: none"> Regional RV akinesia or dyskinesia or dyssynchronous RV contraction and 1 of the following: <ul style="list-style-type: none"> Ratio of RV end-diastolic volume to BSA ≥ 110 mL/m² (male) or ≥ 100 mL/m² (female) or RV ejection fraction $\leq 40\%$ <p>By RV angiography:</p> <ul style="list-style-type: none"> Regional RV akinesia, dyskinesia, or aneurysm
Minor	
<ul style="list-style-type: none"> Mild global RV dilatation and/or ejection fraction reduction with normal LV Mild segmental dilatation of the RV Regional RV hypokinesia 	<p>By 2D echo:</p> <ul style="list-style-type: none"> Regional RV akinesia or dyskinesia and 1 of the following (end diastole): <ul style="list-style-type: none"> PLAX RVOT ≥ 29 to < 32 mm (corrected for body size [PLAX/BSA] ≥ 16 to < 19 mm/m²) PSAX RVOT ≥ 32 to < 36 mm (corrected for body size [PSAX/BSA] ≥ 18 to < 21 mm/m²) or fractional area change $> 33\%$ to $\leq 40\%$ <p>By MRI:</p> <ul style="list-style-type: none"> Regional RV akinesia or dyskinesia or dyssynchronous RV contraction and 1 of the following: <ul style="list-style-type: none"> Ratio of RV end-diastolic volume to BSA ≥ 100 to < 110 mL/m² (male) or ≥ 90 to < 100 mL/m² (female) or RV ejection fraction $> 40\%$ to $\leq 45\%$
II. Tissue characterization of wall	
Major	
<ul style="list-style-type: none"> Fibrofatty replacement of myocardium on endomyocardial biopsy 	<ul style="list-style-type: none"> Residual myocytes $< 60\%$ by morphometric analysis (or $< 50\%$ if estimated), with fibrous replacement of the RV free wall myocardium in ≥ 1 sample, with or without fatty replacement of tissue on endomyocardial biopsy
Minor	
	<ul style="list-style-type: none"> Residual myocytes 60% to 75% by morphometric analysis (or 50% to 65% if estimated), with fibrous replacement of the RV free wall myocardium in ≥ 1 sample, with or without fatty replacement of tissue on endomyocardial biopsy
III. Repolarization abnormalities	
Major	
	<ul style="list-style-type: none"> Inverted T waves in right precordial leads (V₁, V₂, and V₃) or beyond in individuals > 14 years of age (in the absence of complete right bundle-branch block QRS ≥ 120 ms)
Minor	
<ul style="list-style-type: none"> Inverted T waves in right precordial leads (V₂ and V₃) (people age > 12 years, in absence of right bundle-branch block) 	<ul style="list-style-type: none"> Inverted T waves in leads V₁ and V₂ in individuals > 14 years of age (in the absence of complete right bundle-branch block) or in V₄, V₅, or V₆ Inverted T waves in leads V₁, V₂, V₃, and V₄ in individuals > 14 years of age in the presence of complete right bundle-branch block

Continued

Table 1 Continued

Original task force criteria	Revised task force criteria
IV. Depolarization/conduction abnormalities	
Major	
<ul style="list-style-type: none"> Epsilon waves or localized prolongation (> 110 ms) of the QRS complex in right precordial leads (V₁ to V₃) 	<ul style="list-style-type: none"> Epsilon wave (reproducible low-amplitude signals between end of QRS complex to onset of the T wave) in the right precordial leads (V₁ to V₃)
Minor	
<ul style="list-style-type: none"> Late potentials (SAECG) 	<ul style="list-style-type: none"> Late potentials by SAECG in ≥ 1 of 3 parameters in the absence of a QRS duration of ≥ 110 ms on the standard ECG Filtered QRS duration (fQRS) ≥ 114 ms Duration of terminal QRS < 40 μV (low-amplitude signal duration) ≥ 38 ms Root-mean-square voltage of terminal 40 ms ≤ 20 μV Terminal activation duration of QRS ≥ 55 ms measured from the nadir of the S wave to the end of the QRS, including R', in V₁, V₂, or V₃, in the absence of complete right bundle-branch block
V. Arrhythmias	
Major	
	<ul style="list-style-type: none"> Nonsustained or sustained ventricular tachycardia of left bundle-branch morphology with superior axis (negative or indeterminate QRS in leads II, III, and aVF and positive in lead aVL)
Minor	
<ul style="list-style-type: none"> Left bundle-branch block-type ventricular tachycardia (sustained and nonsustained) (ECG, Holter, exercise) Frequent ventricular extrasystoles (> 1000 per 24 hours) (Holter) 	<ul style="list-style-type: none"> Nonsustained or sustained ventricular tachycardia of RV outflow configuration, left bundle-branch block morphology with inferior axis (positive QRS in leads II, III, and aVF and negative in lead aVL) or of unknown axis > 500 ventricular extrasystoles per 24 hours (Holter)
VI. Family history	
Major	
<ul style="list-style-type: none"> Familial disease confirmed at necropsy or surgery 	<ul style="list-style-type: none"> ARVC/D confirmed in a first-degree relative who meets current Task Force criteria ARVC/D confirmed pathologically at autopsy or surgery in a first-degree relative Identification of a pathogenic mutation[†] categorized as associated or probably associated with ARVC/D in the patient under evaluation
Minor	
<ul style="list-style-type: none"> Family history of premature sudden death (< 35 years of age) due to suspected ARVC/D Familial history (clinical diagnosis based on present criteria) 	<ul style="list-style-type: none"> History of ARVC/D in a first-degree relative in whom it is not possible or practical to determine whether the family member meets current Task Force criteria Premature sudden death (< 35 years of age) due to suspected ARVC/D in a first-degree relative ARVC/D confirmed pathologically or by current Task Force Criteria in second-degree relative

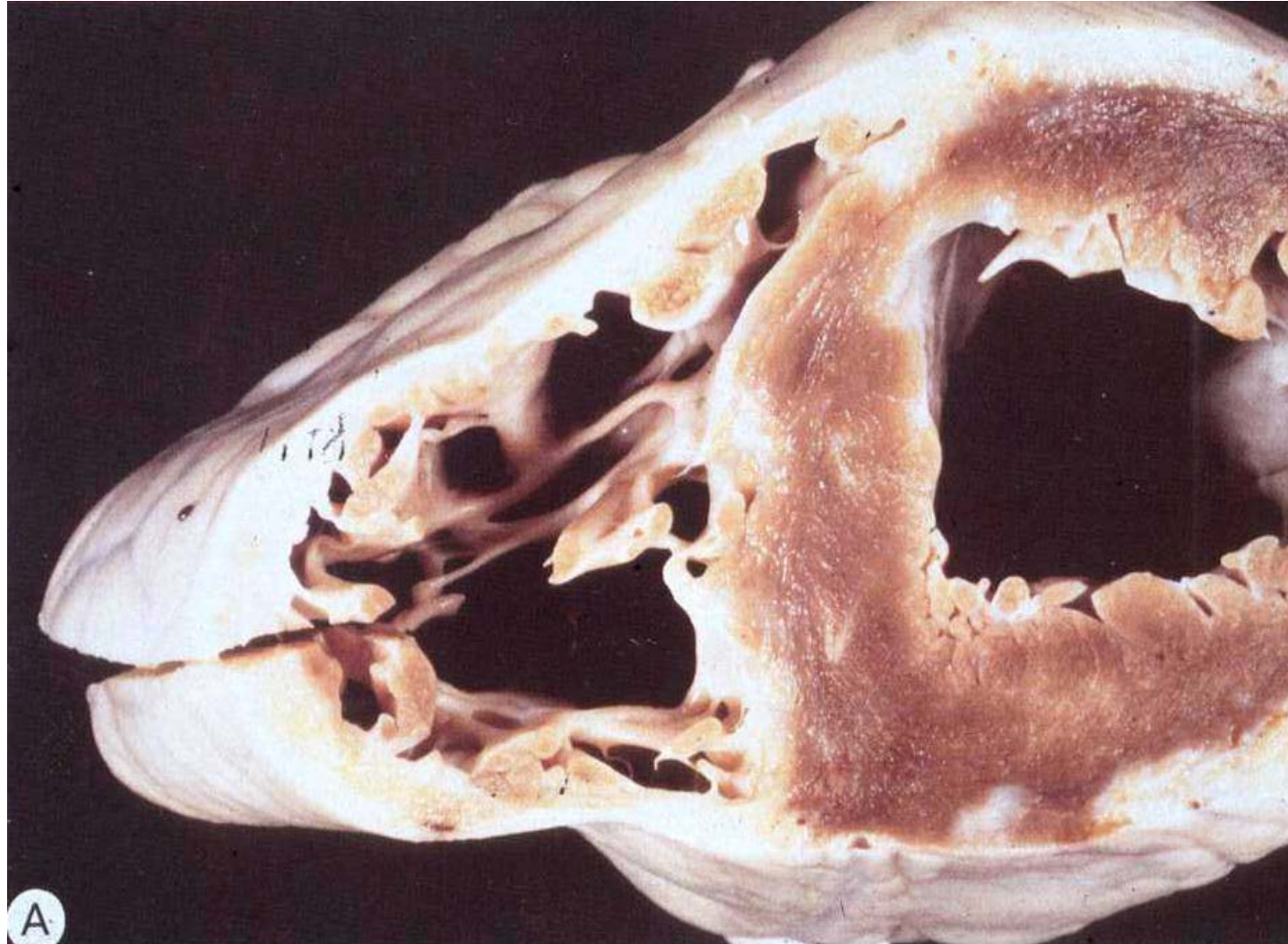
PLAX indicates parasternal long-axis view; RVOT, RV outflow tract; BSA, body surface area; PSAX, parasternal short-axis view; aVF, augmented voltage unipolar left foot lead; and aVL, augmented voltage unipolar left arm lead.

Diagnostic terminology for original criteria: This diagnosis is fulfilled by the presence of 2 major, or 1 major plus 2 minor criteria or 4 minor criteria from different groups. Diagnostic terminology for revised criteria: definite diagnosis: 2 major or 1 major and 2 minor criteria or 4 minor from different categories; borderline: 1 major and 1 minor or 3 minor criteria from different categories; possible: 1 major or 2 minor criteria from different categories.

[†]Hypokinesia is not included in this or subsequent definitions of RV regional wall motion abnormalities for the proposed modified criteria.

^{††}A pathogenic mutation is a DNA alteration associated with ARVC/D that alters or is expected to alter the encoded protein, is unobserved or rare in a large non-ARVC/D control population, and either alters or is predicted to alter the structure or function of the protein or has demonstrated linkage to the disease phenotype in a conclusive pedigree.

Arrhythmogenic Right Ventricular Cardiomyopathy



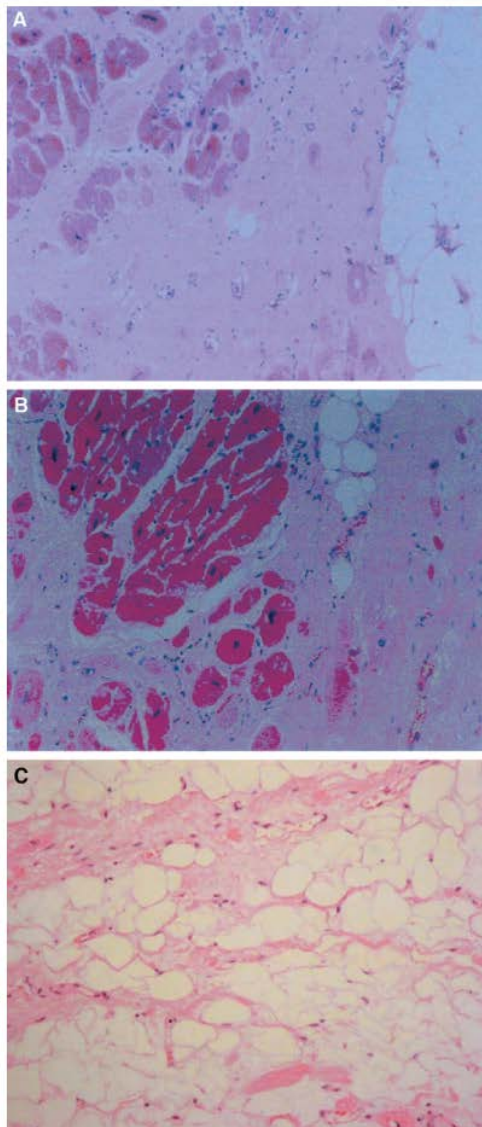


Figure 2. Histological example of LV myocardium from proband IV:1, showing myocyte loss with fibrofatty replacement (Hematoxylin and eosin staining). A, Low-power ($\times 4$) view; B, slightly higher low-power ($\times 10$) view; C, high-power view ($\times 20$).

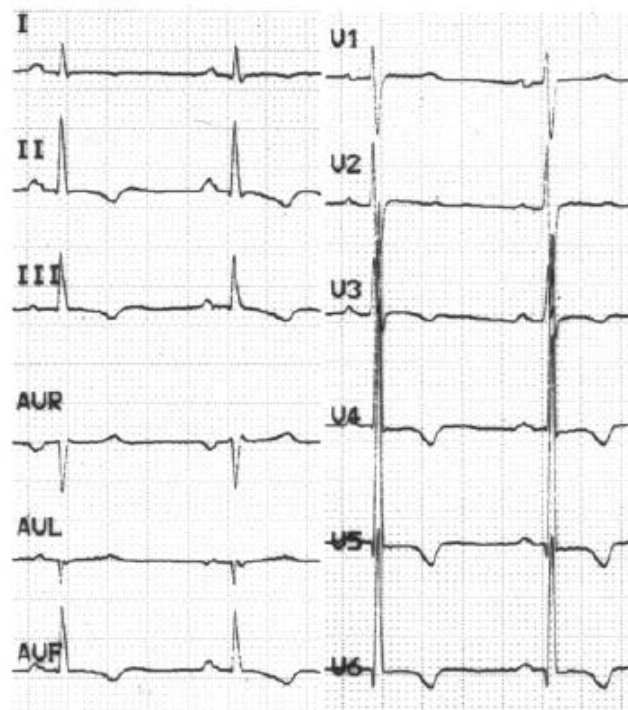


Figure 3. Typical resting 12-lead ECG findings with inferior and lateral T-wave inversion.

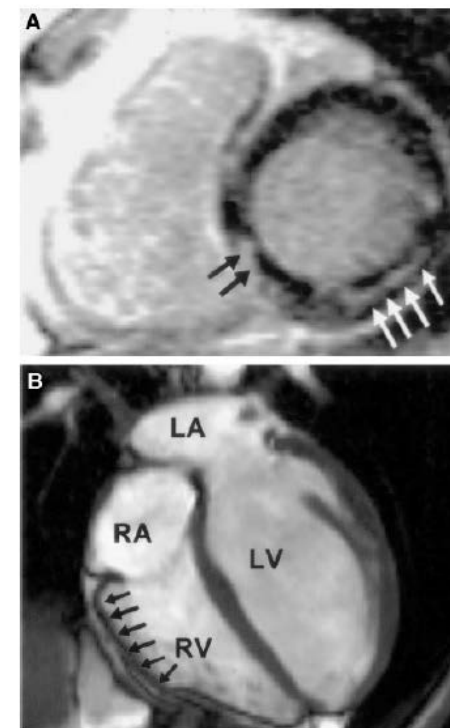
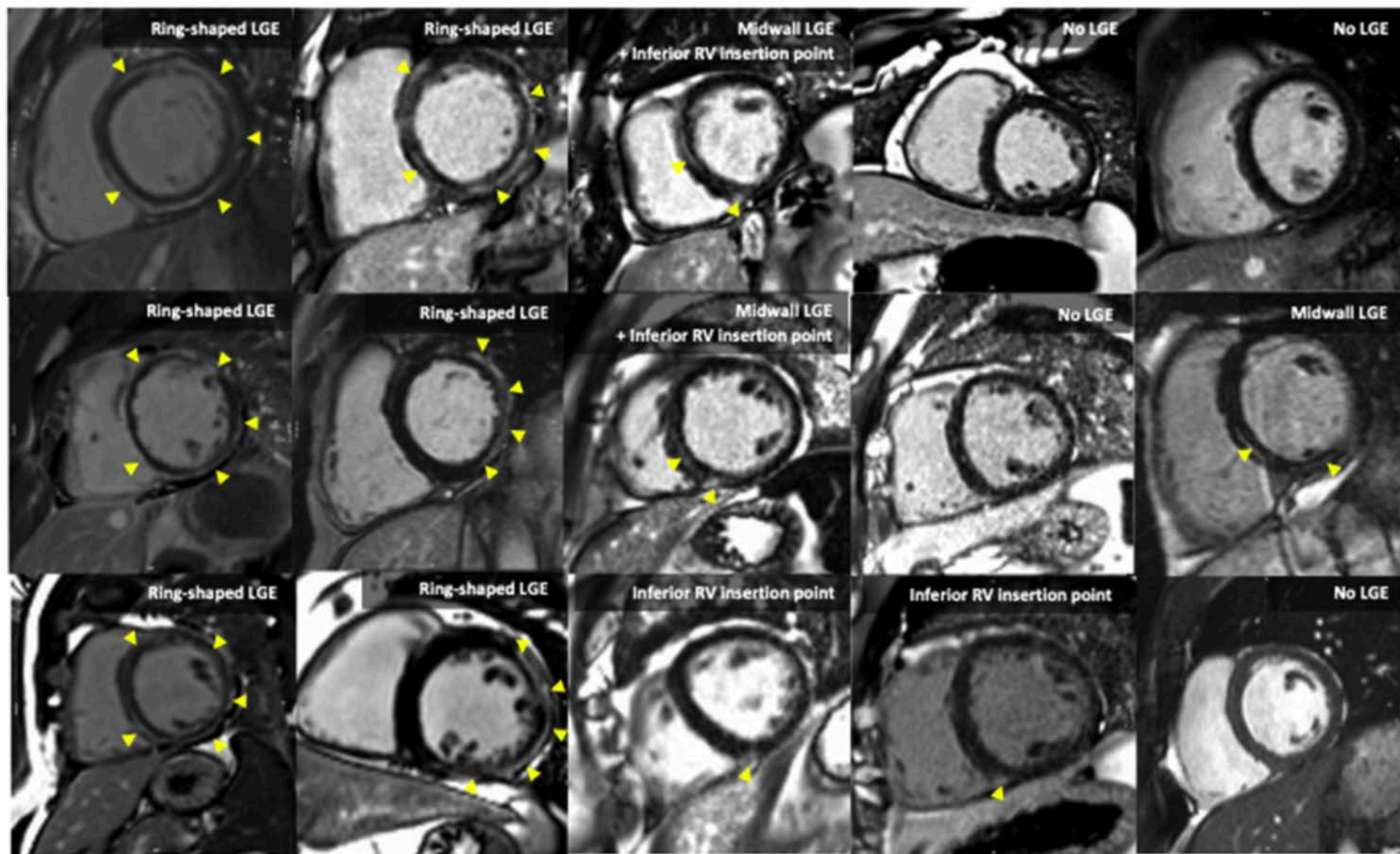


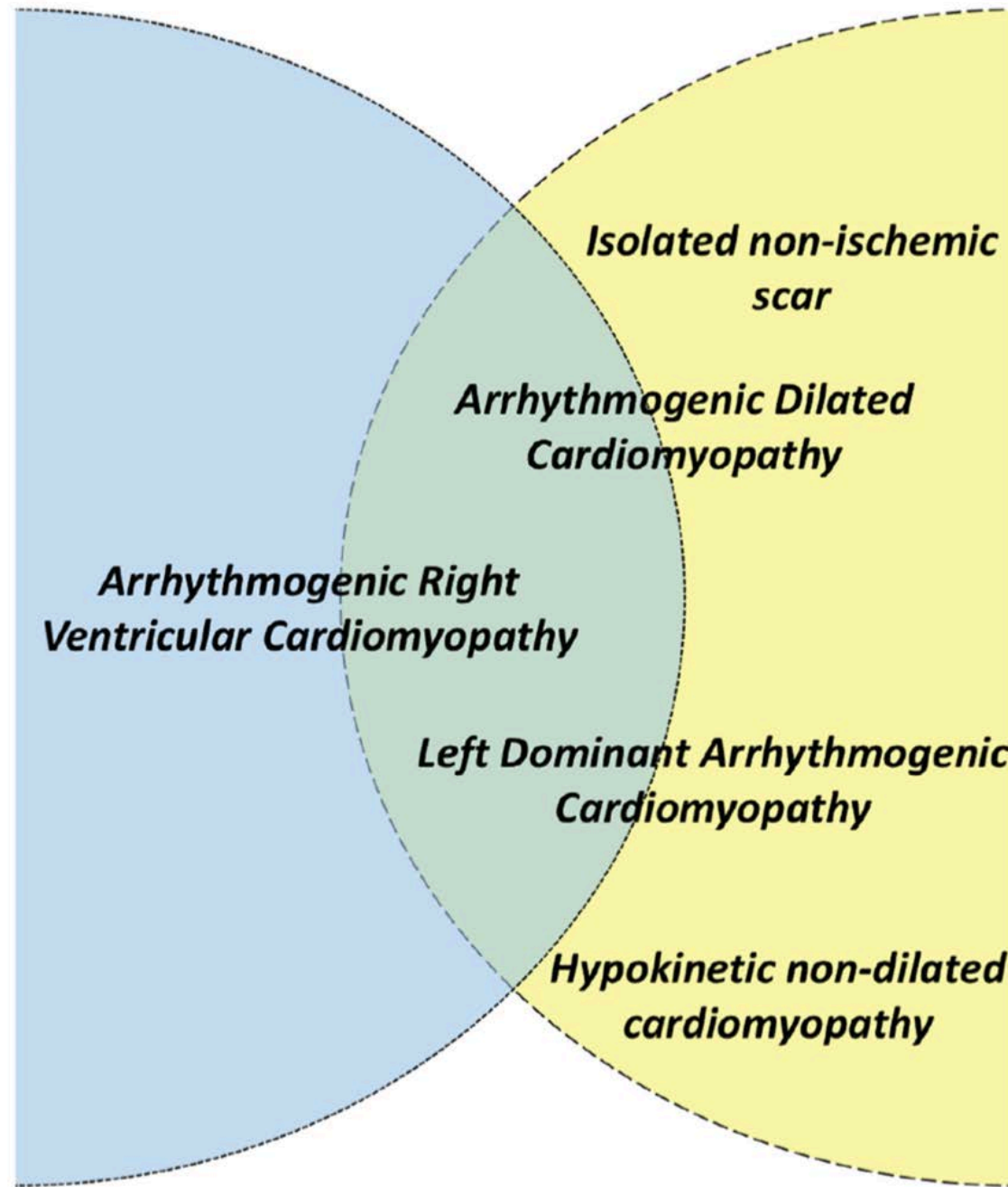
Figure 4. CMR images from 2 patients (III:7 and IV:9). A, Late gadolinium enhancement at inferolateral LV wall (white arrows) and septum (black arrows). B, Localized dilation in subtricuspid region (black arrows) of RV. This segment was markedly hypokinetic during systole. RA indicates right atrium; LA, left atrium.

Norman M et al. Circulation. 2005;112:636-642



RV Disease

LV Disease



Arrhythmogenic Right Ventricular Cardiomyopathy

Arrhythmogenic Dilated Cardiomyopathy

Left Dominant Arrhythmogenic Cardiomyopathy

Hypokinetic non-dilated cardiomyopathy

Isolated non-ischemic scar

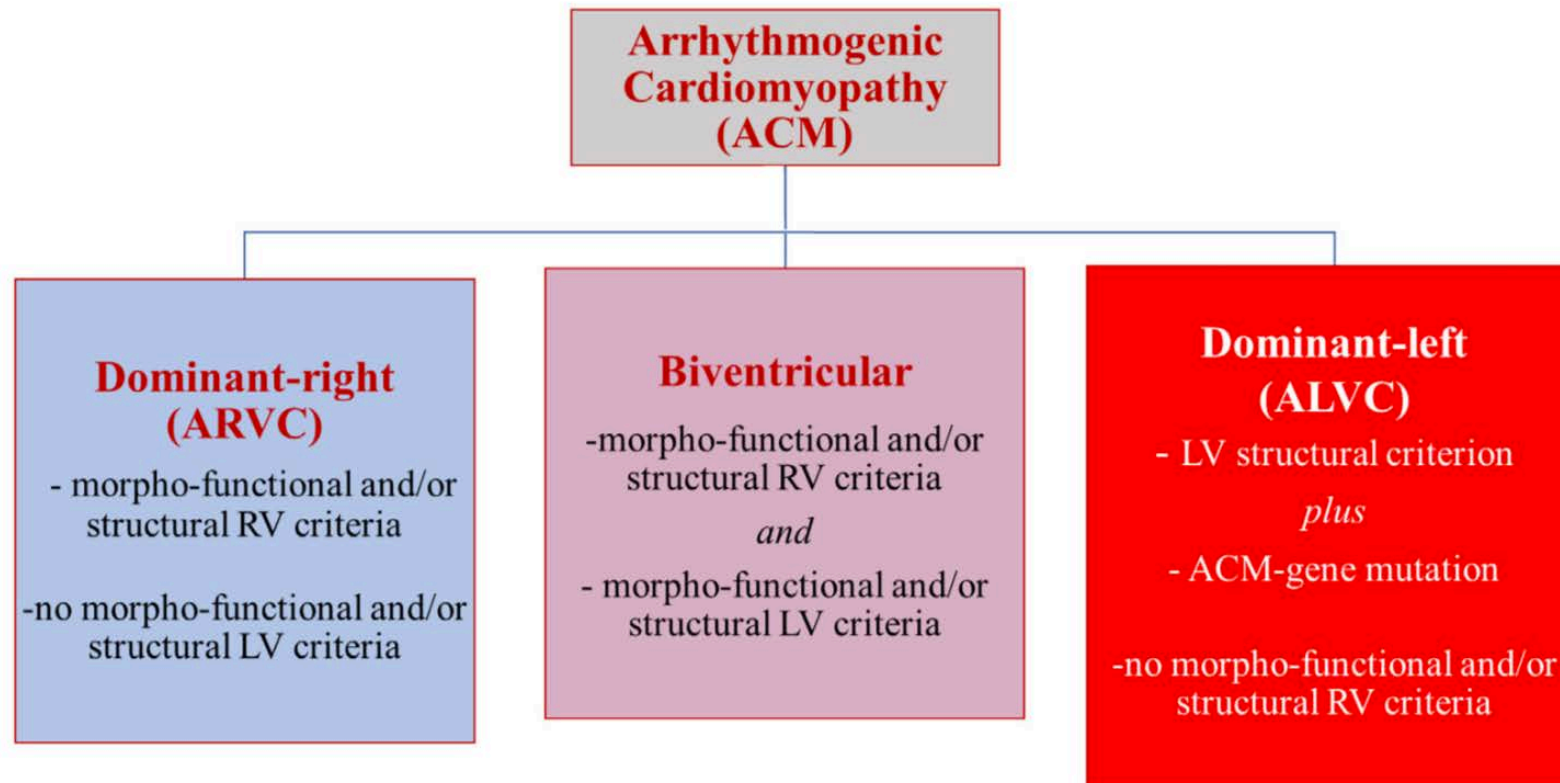


Arrhythmogenic Cardiomyopathy?

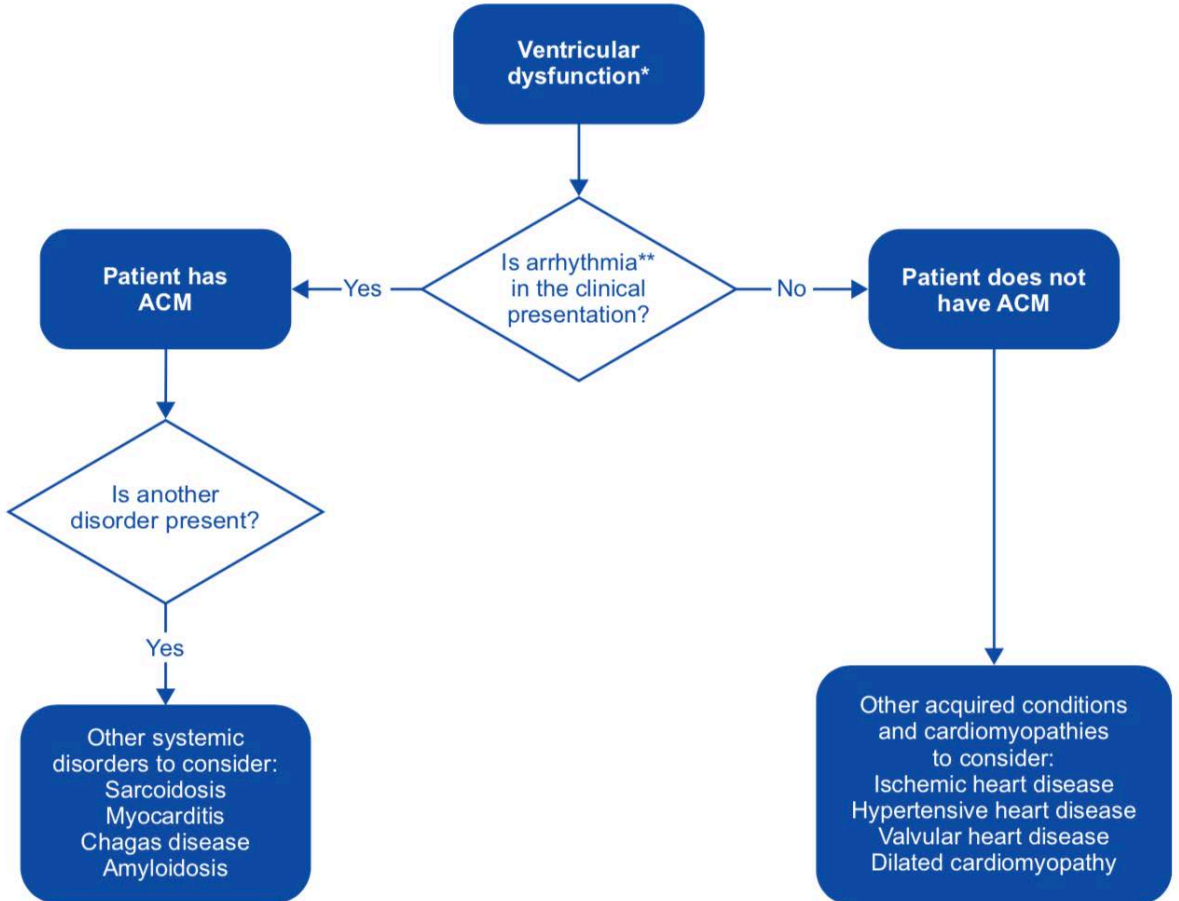
Diagnosis of arrhythmogenic cardiomyopathy: The Padua criteria



Domenico Corrado ^{a,*}, Martina Perazzolo Marra ^a, Alessandro Zorzi ^a, Giorgia Beffagna ^a, Alberto Cipriani ^a, Manuel De Lazzari ^a, Federico Migliore ^a, Kalliopi Pilichou ^a, Alessandra Rampazzo ^b, Ilaria Rigato ^a, Stefania Rizzo ^a, Gaetano Thiene ^a, Aris Anastasakis ^c, Angeliki Asimaki ^d, Chiara Bucciarelli-Ducci ^e, Kristine H. Haugaa ^f, Francis E. Marchlinski ^g, Andrea Mazzanti ^h, William J. McKenna ⁱ, Antonis Pantazis ^j, Antonio Pelliccia ^k, Christian Schmied ^l, Sanjay Sharma ^m, Thomas Wichter ⁿ, Barbara Bauce ^a, Cristina Basso ^a



2019 HRS expert consensus statement on evaluation, risk stratification, and management of arrhythmogenic cardiomyopathy



*Not explained by ischemic, hypertensive, or valvular heart disease
**Arrhythmia includes conduction disease, atrial arrhythmias, ventricular arrhythmias



If it doesn't fit, then do not force it



TALES FROM THE CLINIC (1)

Clinical History

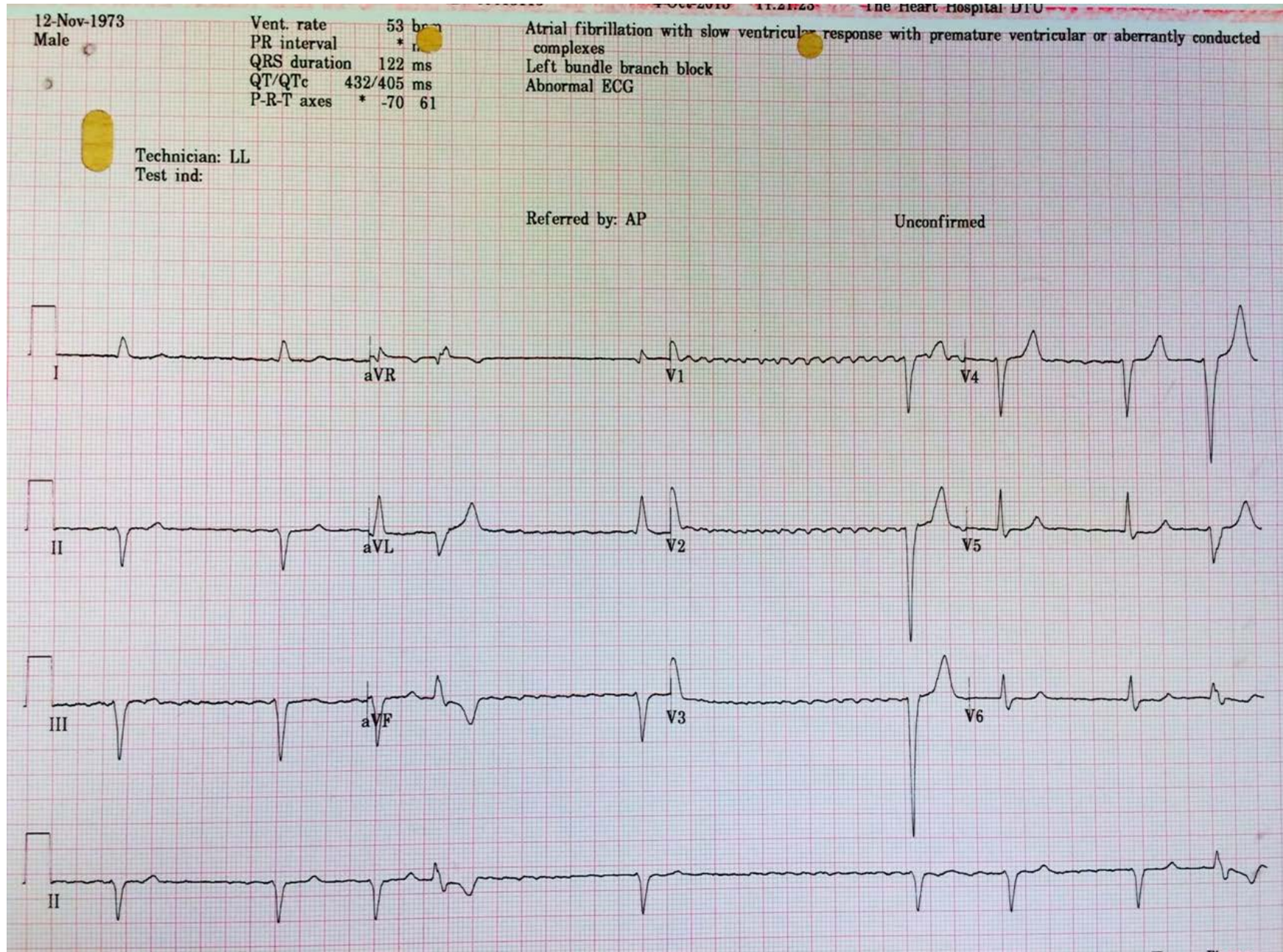
42y, male

- AF 2007
 - Incidental finding
 - DCCV 2008
 - Medication: Aspirin 75mg
 - Holter: NSVT x 5 beats
- CMR: Mild impairment of LV function. Biatrial dilatation with LA diameter of 43mm.

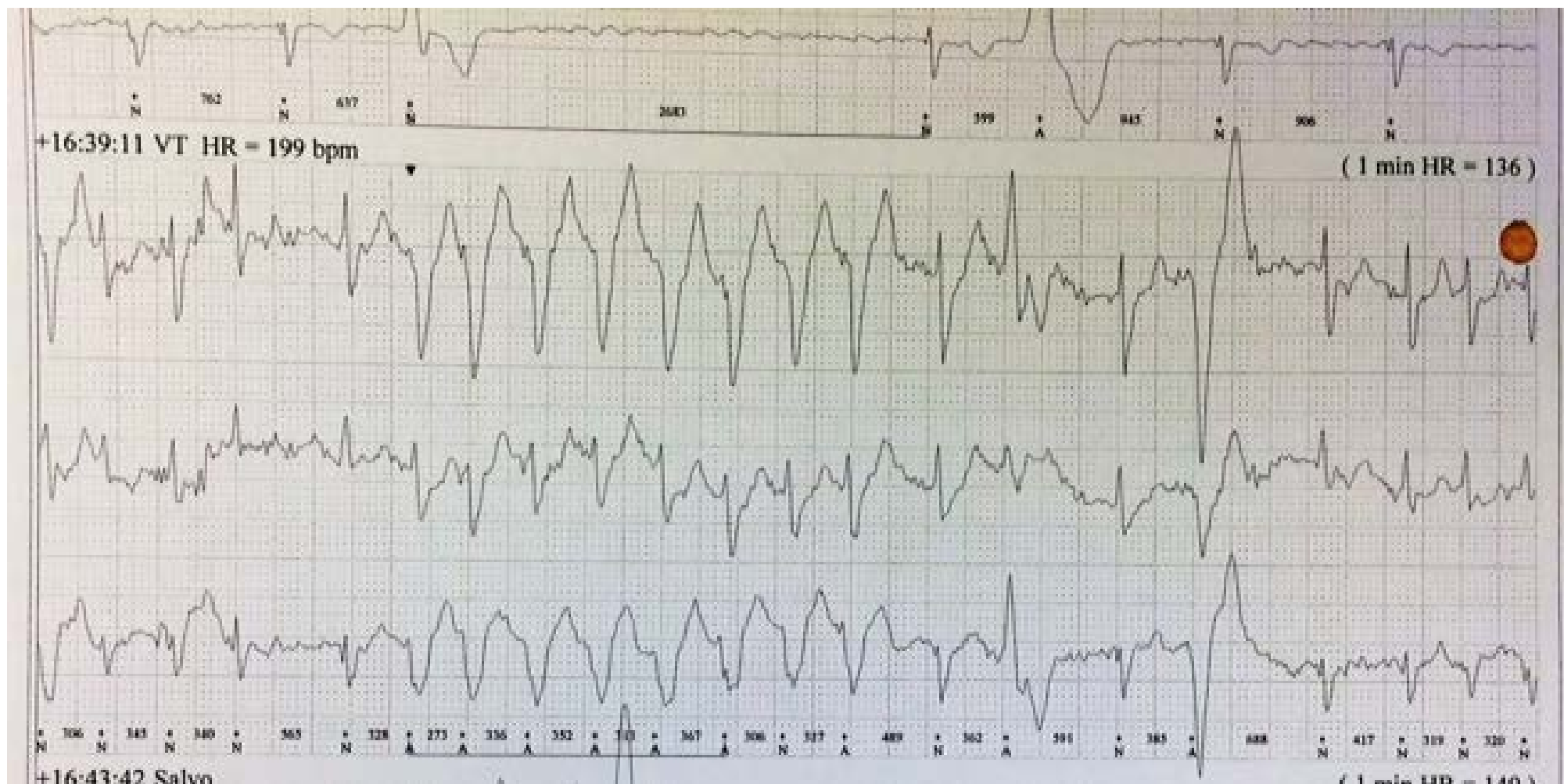
Clinical History

- Family History
 - Father:
 - AF
 - PPM – CHB
 - CRT-P 2011
 - RIP aged 63
 - Paternal grandfather:
 - AF and PPM
 - RIP aged 64 ?cause

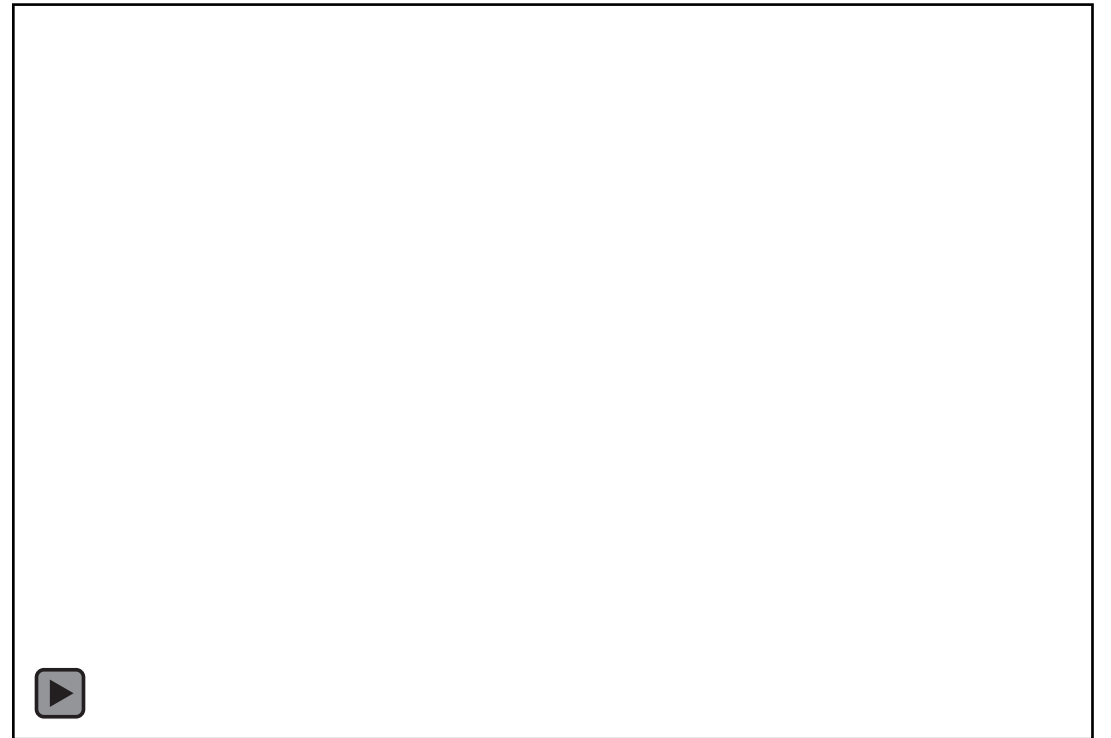
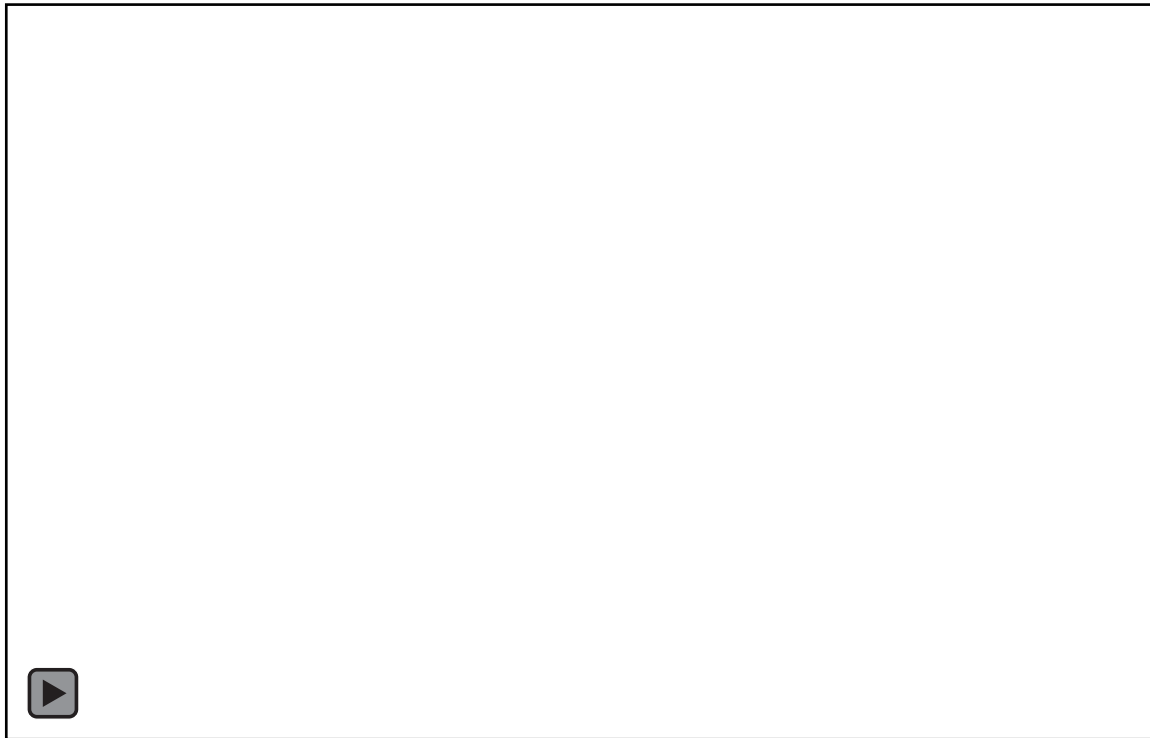
ECG



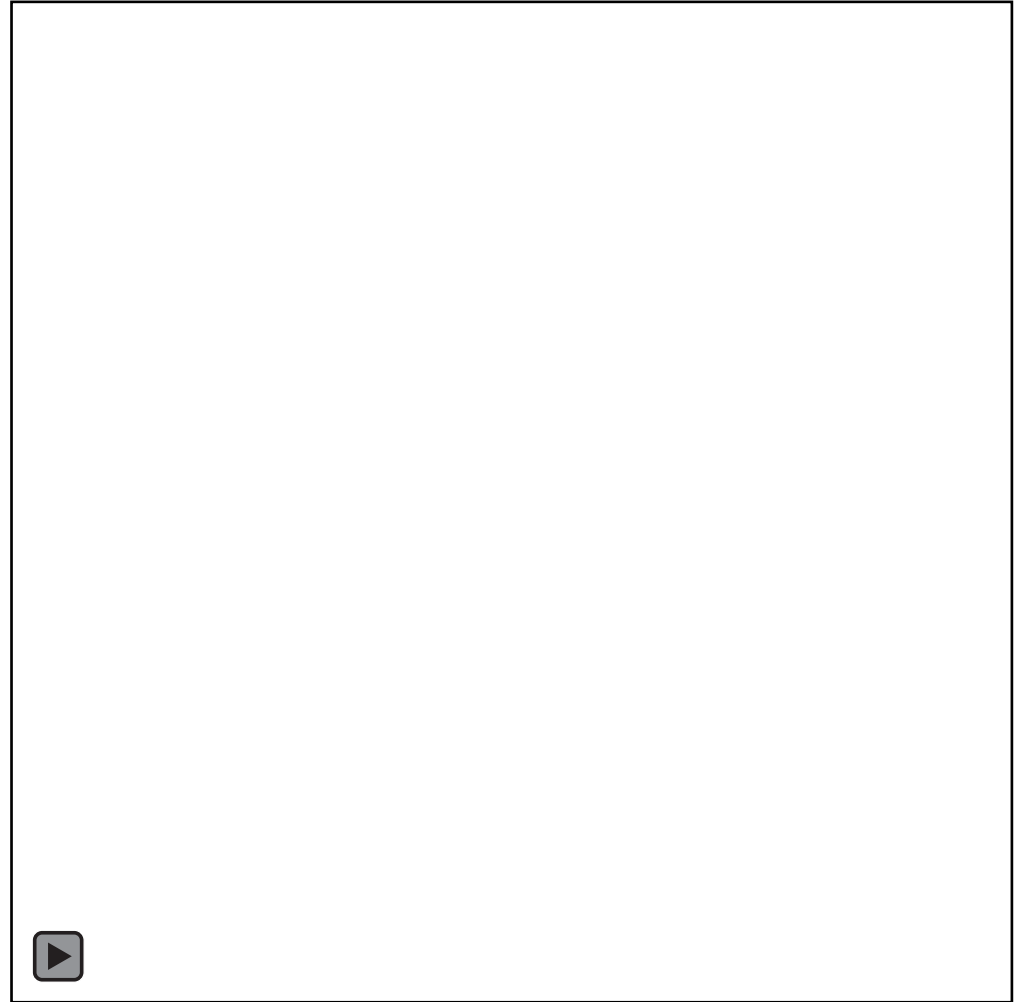
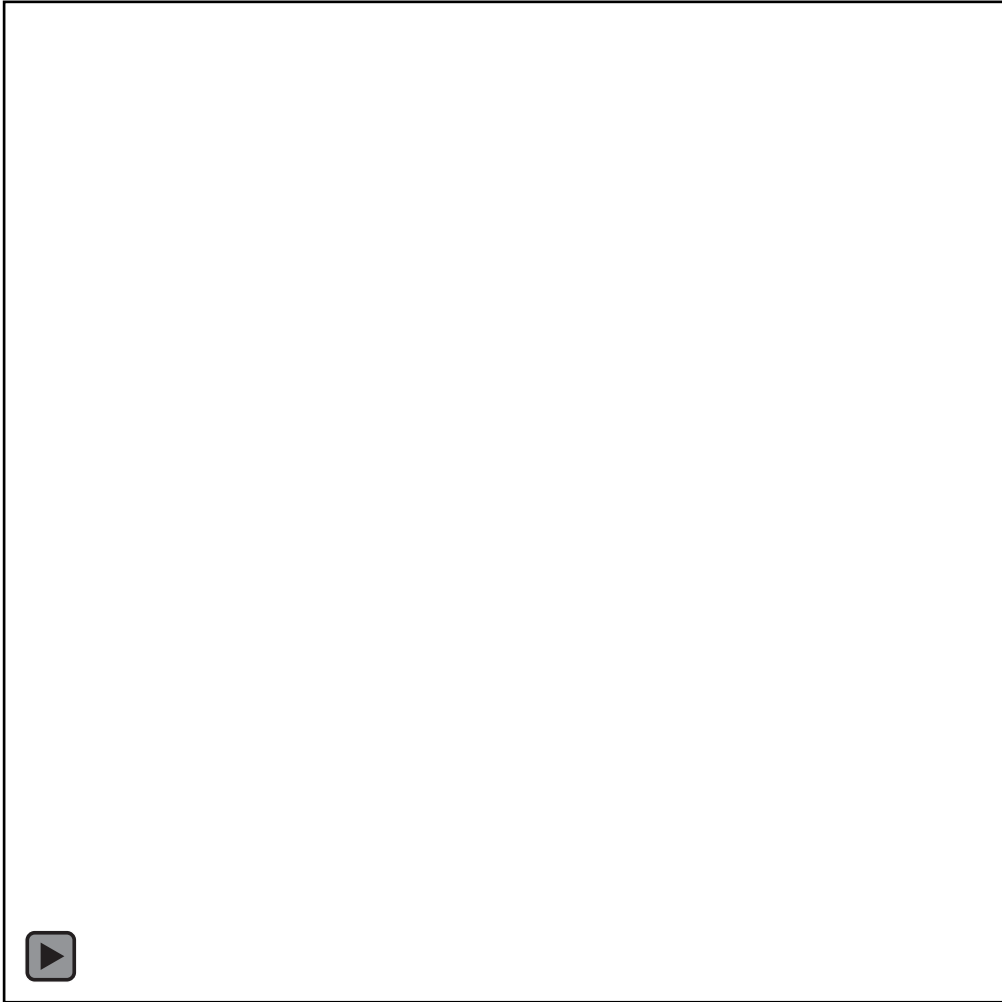
24 hour Holter



Echocardiogram



CMR



The Phenotype

- Young
- Family History of AF, PM
- NSVT
- Mild LV impairment

• WHY?

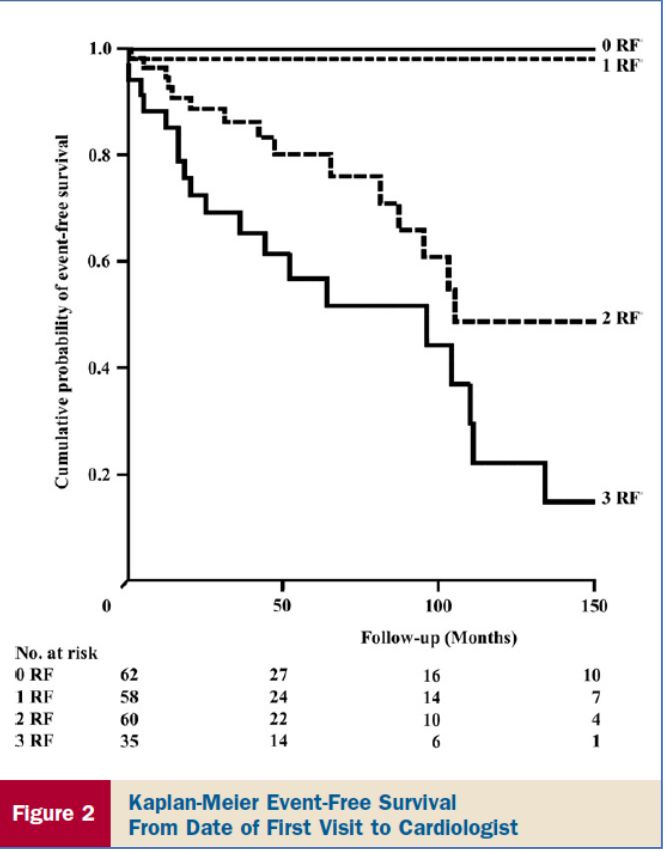
Diagnosis

- **Genetic Testing**

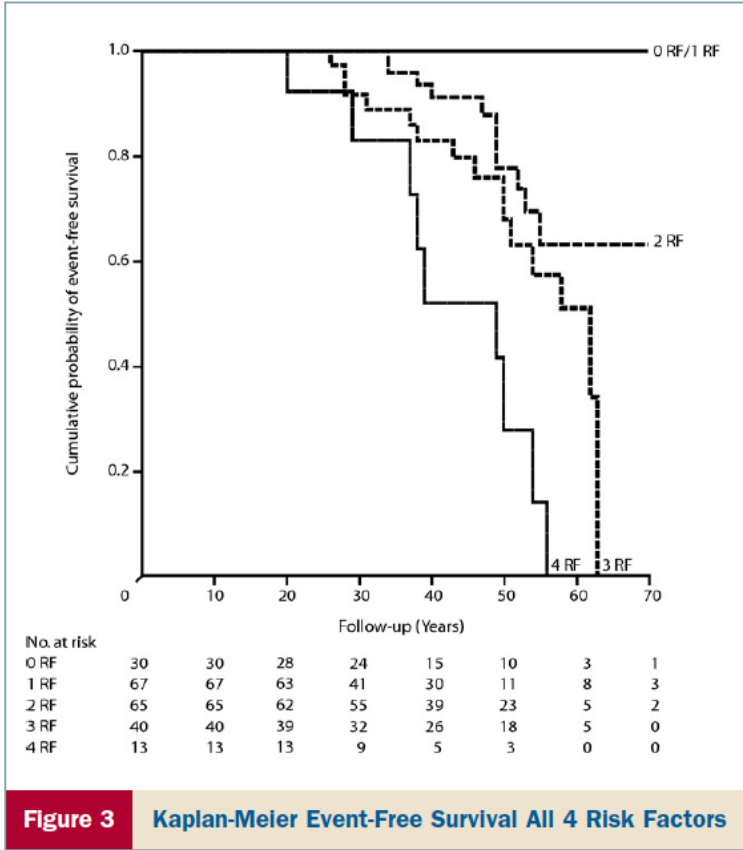
- **Lamin A/C : c.1489-1 G>A**

Risk Factors for Malignant Ventricular Arrhythmias in Lamin A/C Mutation Carriers

A European Cohort Study



NSVT, LVEF 45%, male



+ non-missense mutations (ins-del/truncating or mutations affecting splicing)

(J Am Coll Cardiol 2012;59:493–500)

TALES FROM THE CLINIC (2)

Background - 71y Male

Asymptomatic: tennis 3/week

Family history of DCM

- Nephew RIP 27 SCD – PM DCM, enlarged LV with uniform fine fibrosis
- Brother RIP 71 – Chemo for hairy cell leukaemia
- Niece – DCM and ICD

Holter 4-5% VE, no NSVT

Echo 2011

- EF 45%

Medications

- Ramipril 2.5mg





Test(s) requested: Dilated cardiomyopathy - Familial "screening"

RESULT: POSITIVE

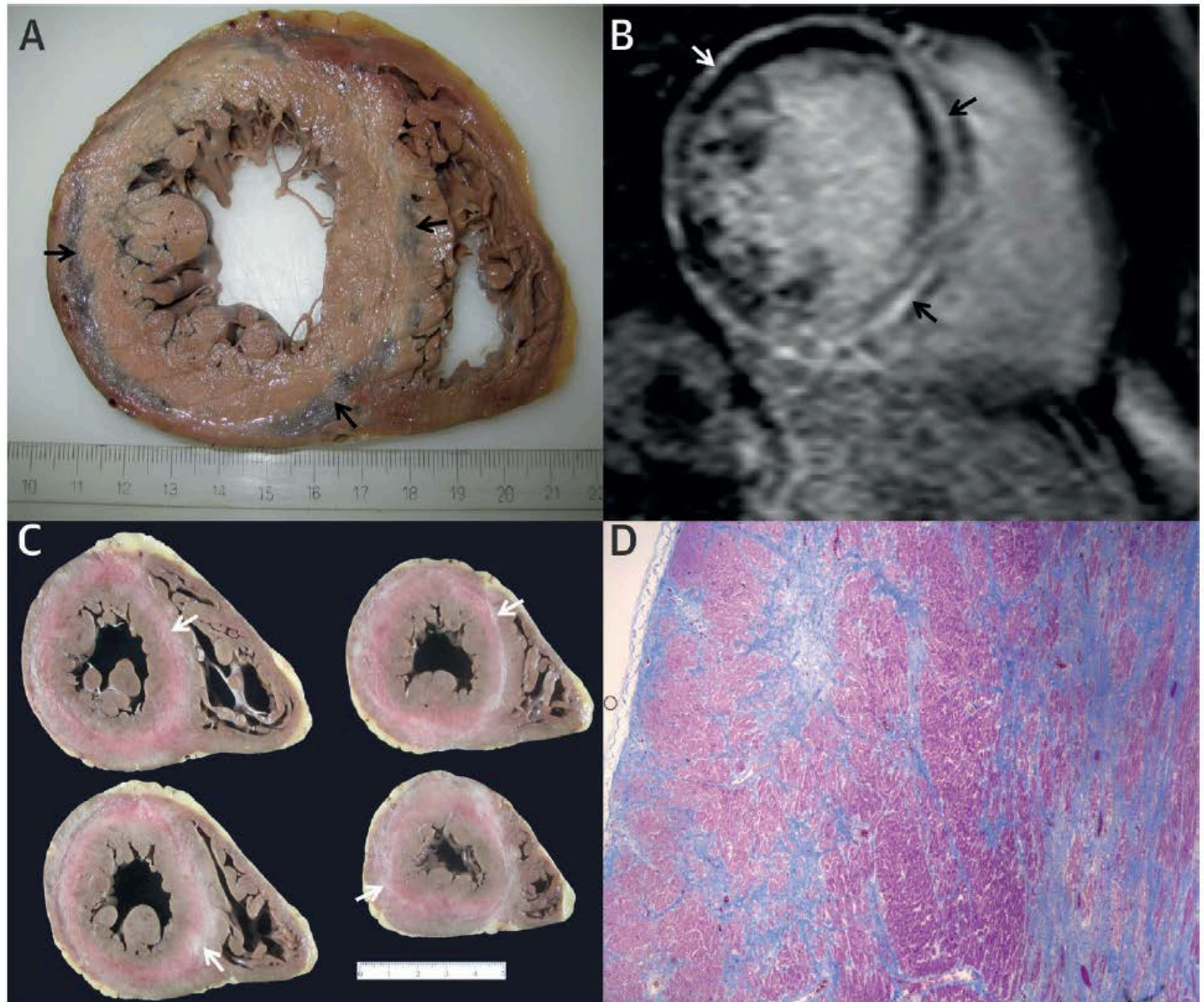
In the sample of this individual, we have confirmed the presence of the variant in *FLNC* that was previously identified in the index case.

Gene	Variant	Result	Pathogenicity	Population frequency	Number of references
<i>FLNC</i>	NP_001449.3:p.Arg482* NM_001458.4:c.1444C>T NC_000007.13:g.128480109C>T	Heterozygosis	Very likely to be pathogenic or disease-causing (++)	Mutation (not found in controls)	0

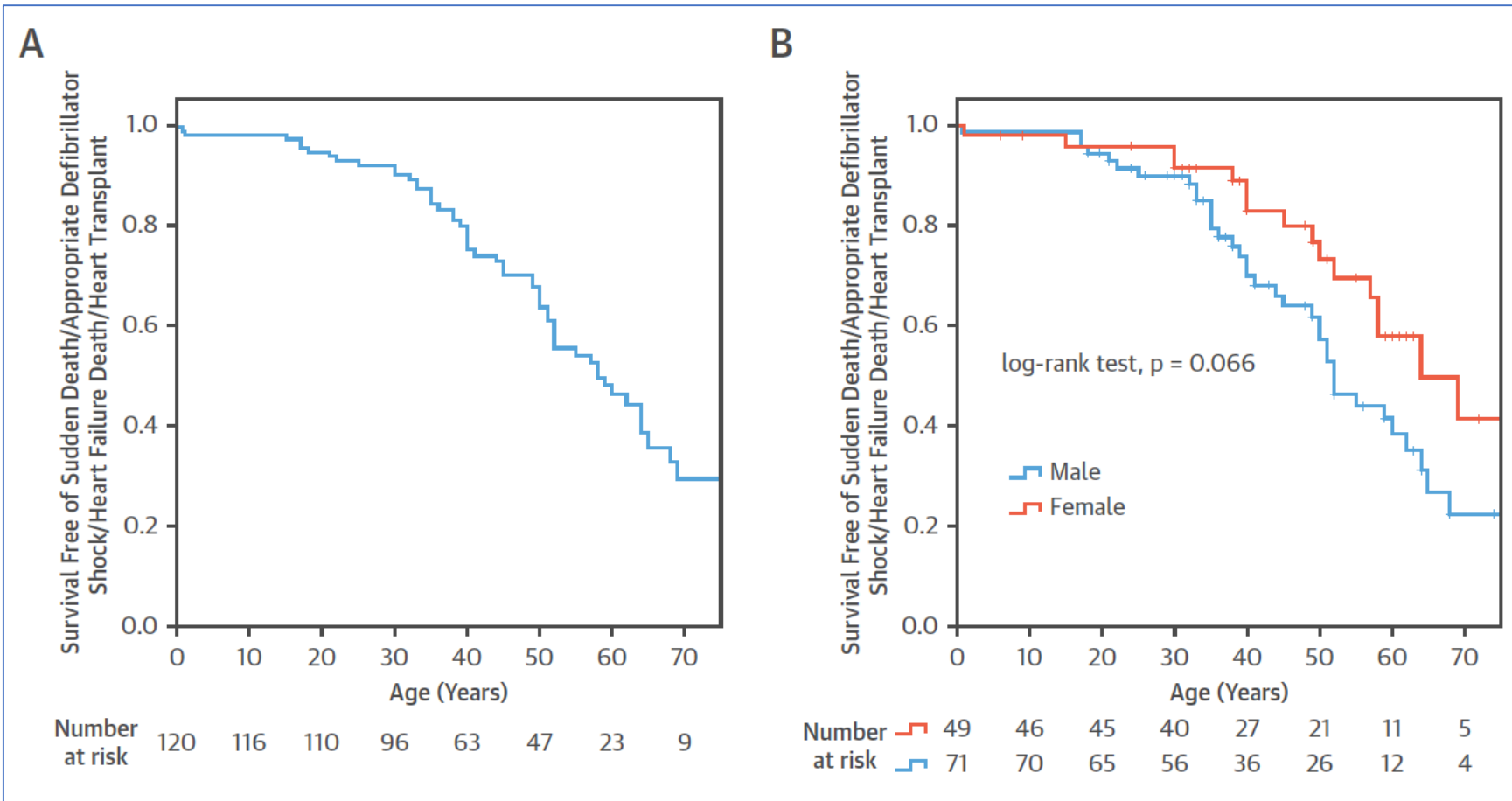
Truncating *FLNC* Mutations Are Associated With High-Risk Dilated and Arrhythmogenic Cardiomyopathies



Martín F. Ortiz-Genga, MD,^{a,b} Sofía Cuenca, MD, PhD,^c Matteo Dal Ferro, MD,^d Esther Zorio, MD, PhD,^e Ricardo Salgado-Aranda, MD,^f Vicente Climent, MD,^g Laura Padrón-Barthe, PhD,^h Iria Duro-Aguado, MD,ⁱ Juan Jiménez-Jáimez, MD, PhD,^j Víctor M. Hidalgo-Olivares, MD,^k Enrique García-Campo, MD,^l Chiara Lanzillo, MD, PhD,^m M. Paz Suárez-Mier, MD, PhD,ⁿ Hagith Yonath, MD,^o Sonia Marcos-Alonso, MD, PhD,^p Juan P. Ochoa, MD,^b José L. Santomé, BSc,^b Diego García-Giustiniani, MD,^b Jorge L. Rodríguez-Garrido, MD,^{b,p} Fernando Domínguez, MD,^c Marco Merlo, MD,^d Julián Palomino, MD, PhD,^l María L. Peña, MD,^q Juan P. Trujillo, MD, PhD,^p Alicia Martín-Vila, PharmD,^l Davide Stolfo, MD,^d Pilar Molina, MD, PhD,^r Enrique Lara-Pezzi, PhD,^{h,s} Francisco E. Calvo-Iglesias, MD, PhD,^l Eyal Nof, MD,^o Leonardo Calò, MD,^m Roberto Barriales-Villa, MD, PhD,^{a,p} Juan R. Gimeno-Blanes, MD, PhD,^t Michael Arad, MD, PhD,^o Pablo García-Pavía, MD, PhD,^{c,u} Lorenzo Monserrat, MD, PhD^{a,b}

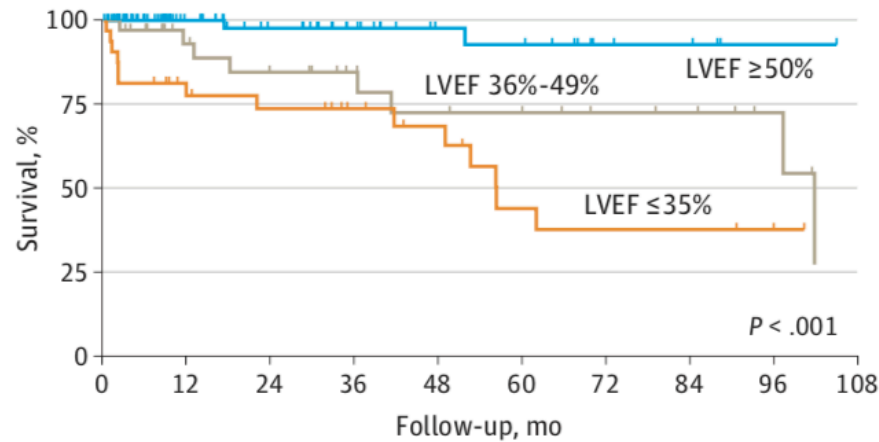


J Am Coll Cardiol. 2016 Dec 6;68(22):2440-2451



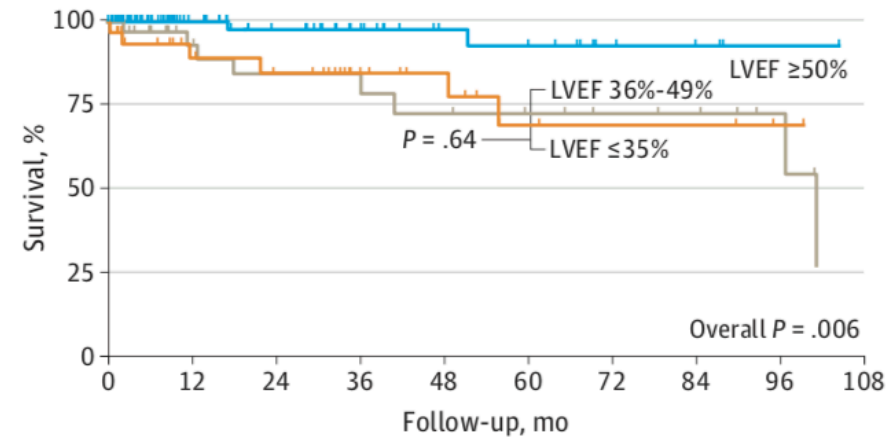
Association of Left Ventricular Systolic Dysfunction Among Carriers of Truncating Variants in Filamin C With Frequent Ventricular Arrhythmia and End-stage Heart Failure

A Survival free from primary composite end point

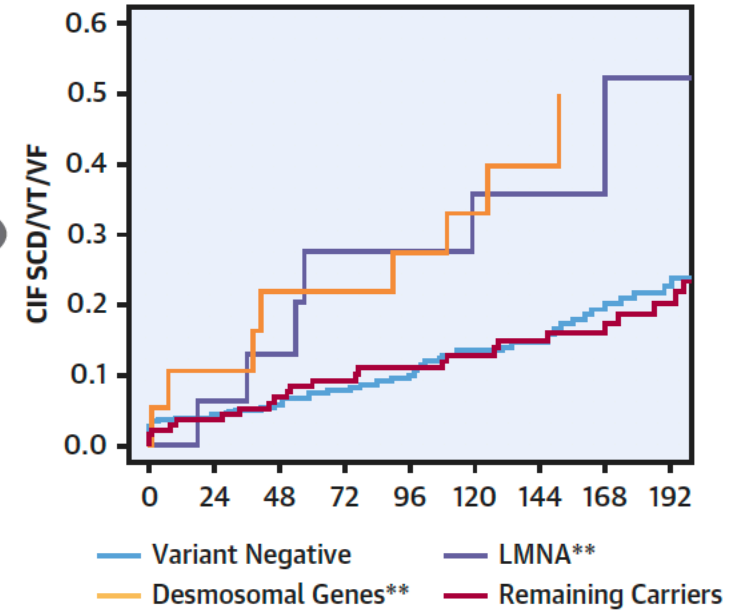
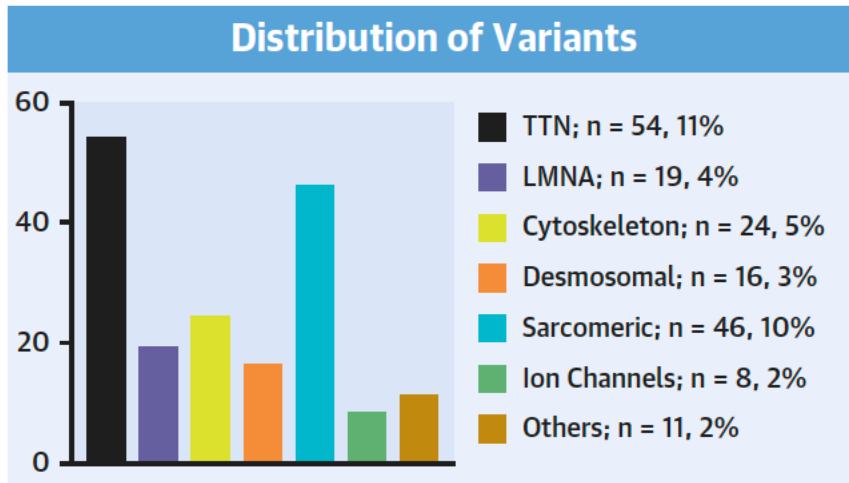
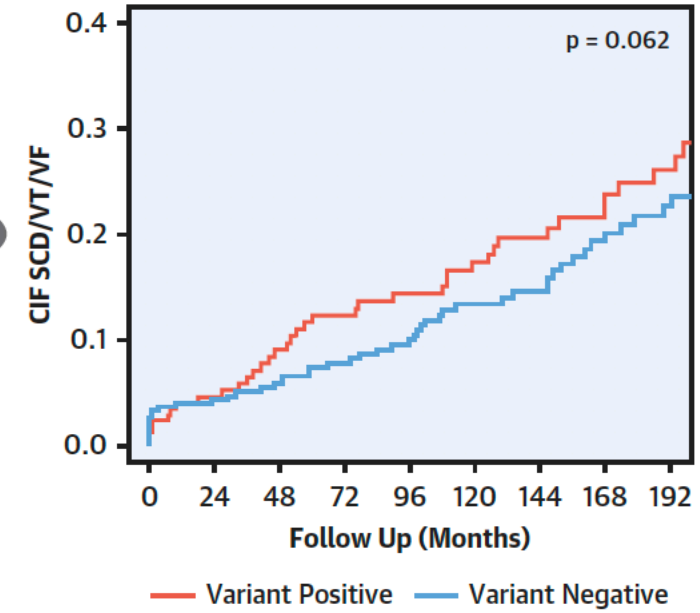
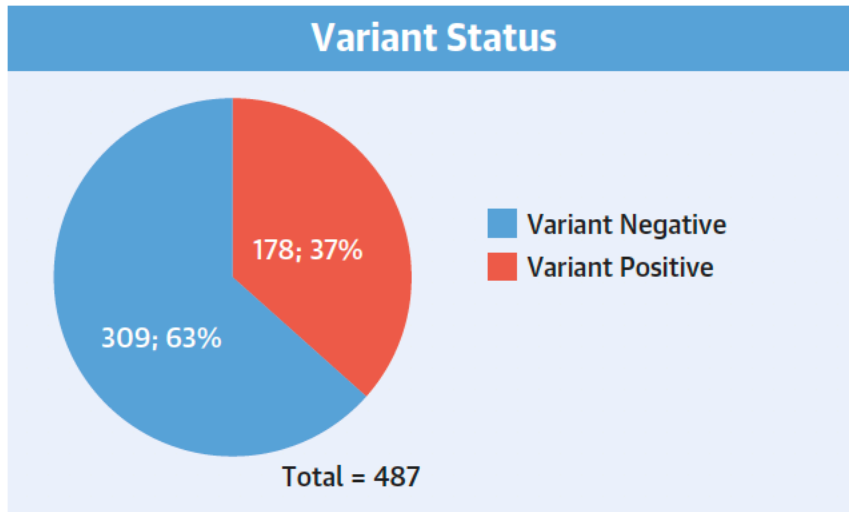


No. at risk	0	12	24	36	48	60	72	84	96	108
LVEF $\geq 50\%$	95	50	38	28	20	19	13	12	9	8
LVEF 36%-49%	36	22	19	15	12	10	8	7	4	1
LVEF $\leq 35\%$	33	21	19	15	12	7	6	6	4	3

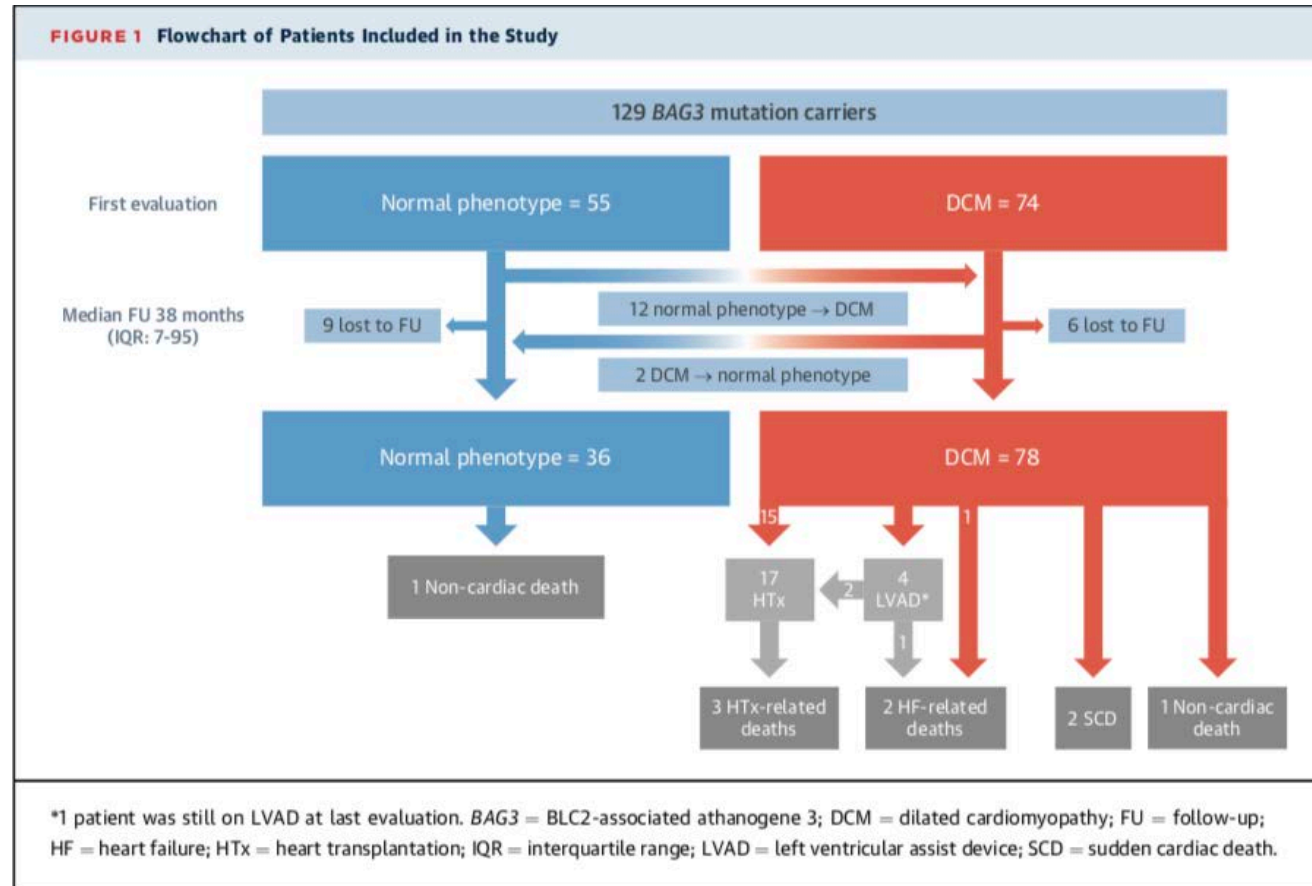
B Survival free from secondary composite arrhythmic end point



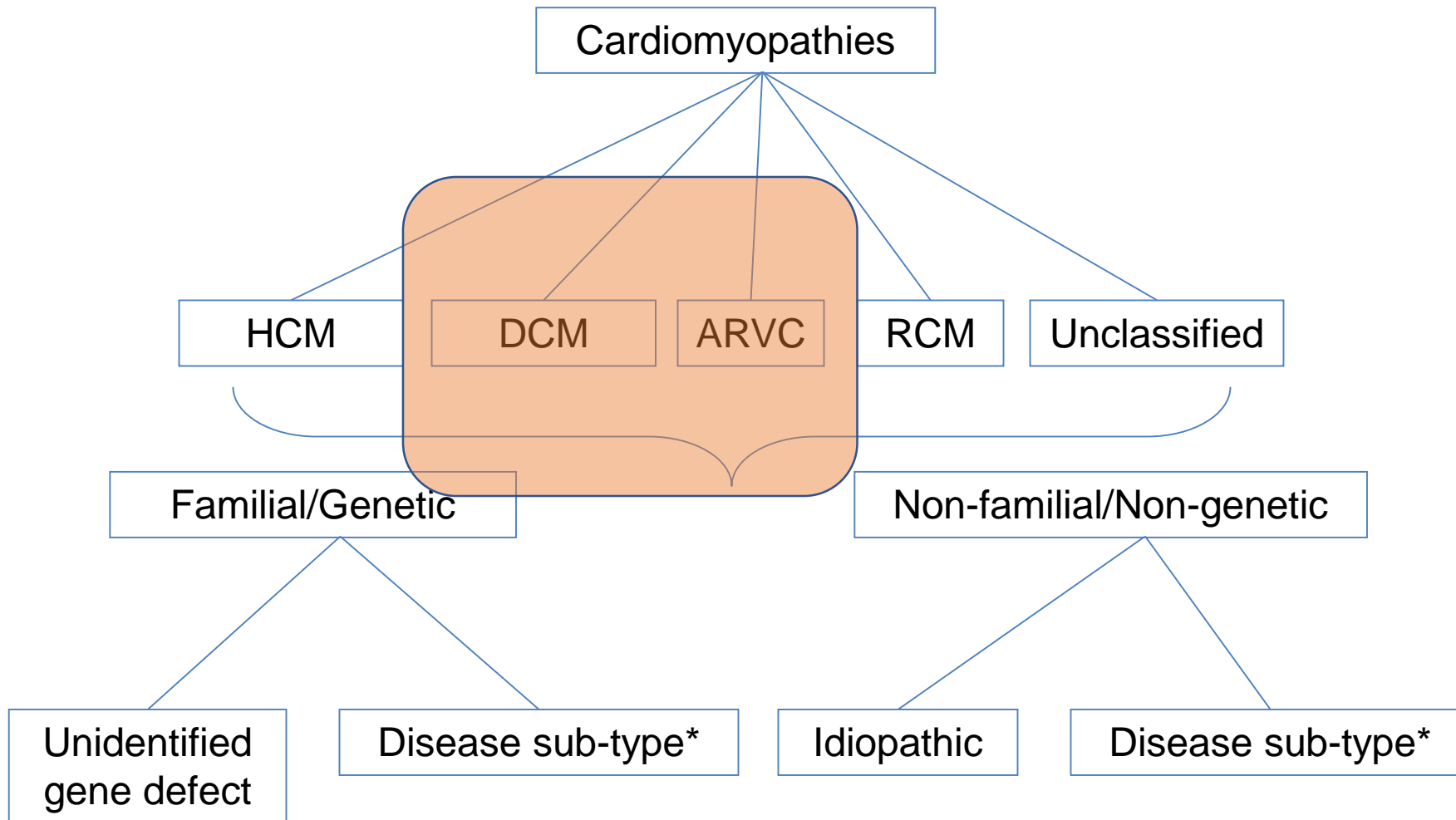
No. at risk	0	12	24	36	48	60	72	84	96	108
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LVEF $\leq 35\%$	33	21	19	15	12	7	6	6	4	3



Dilated Cardiomyopathy Due to BLC2-Associated Athanogene 3 (*BAG3*) Mutations

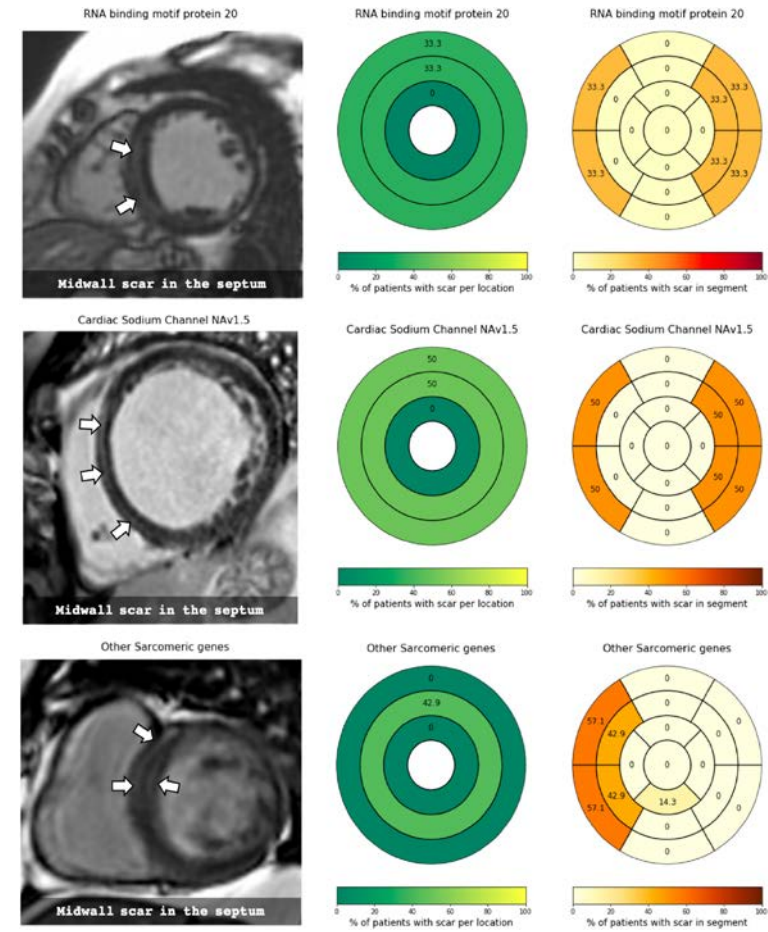
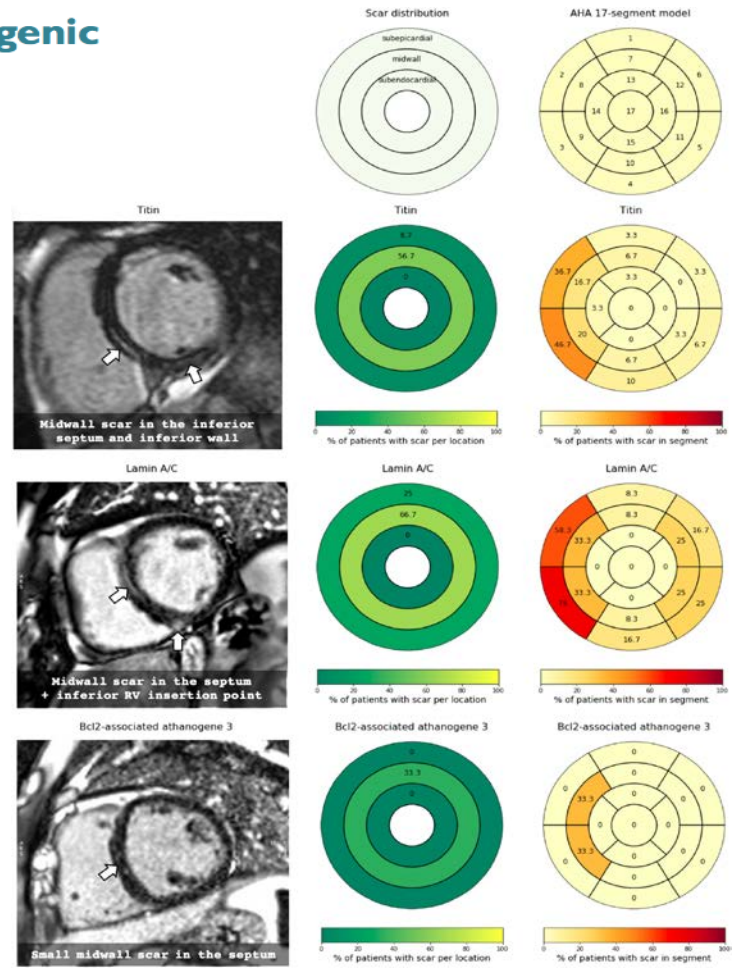
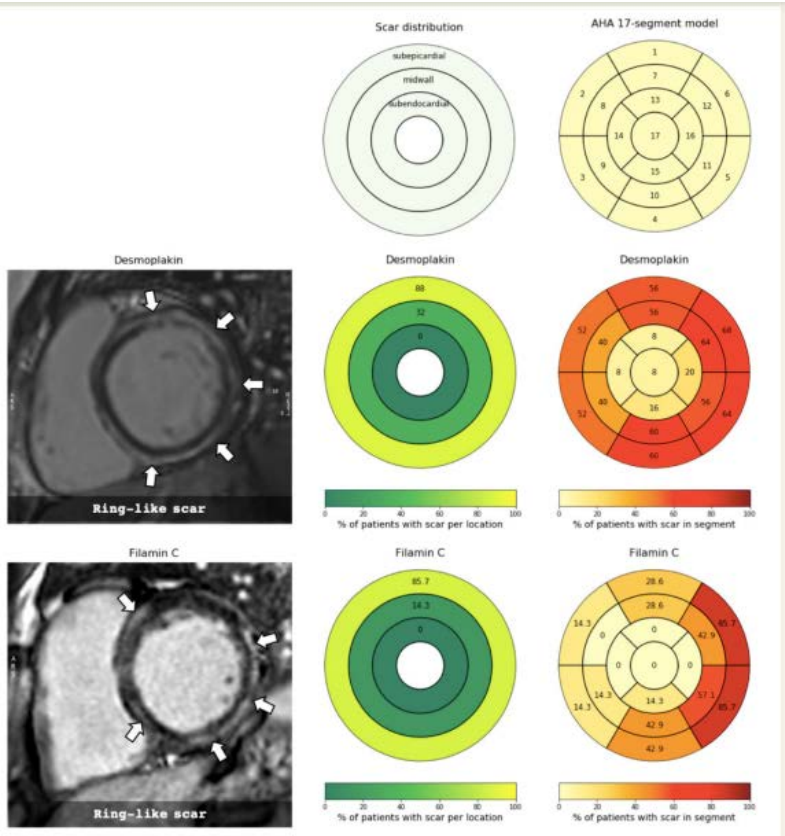


What do we learn from all this?



Clinical translation of omics platforms in inherited heart muscle disease

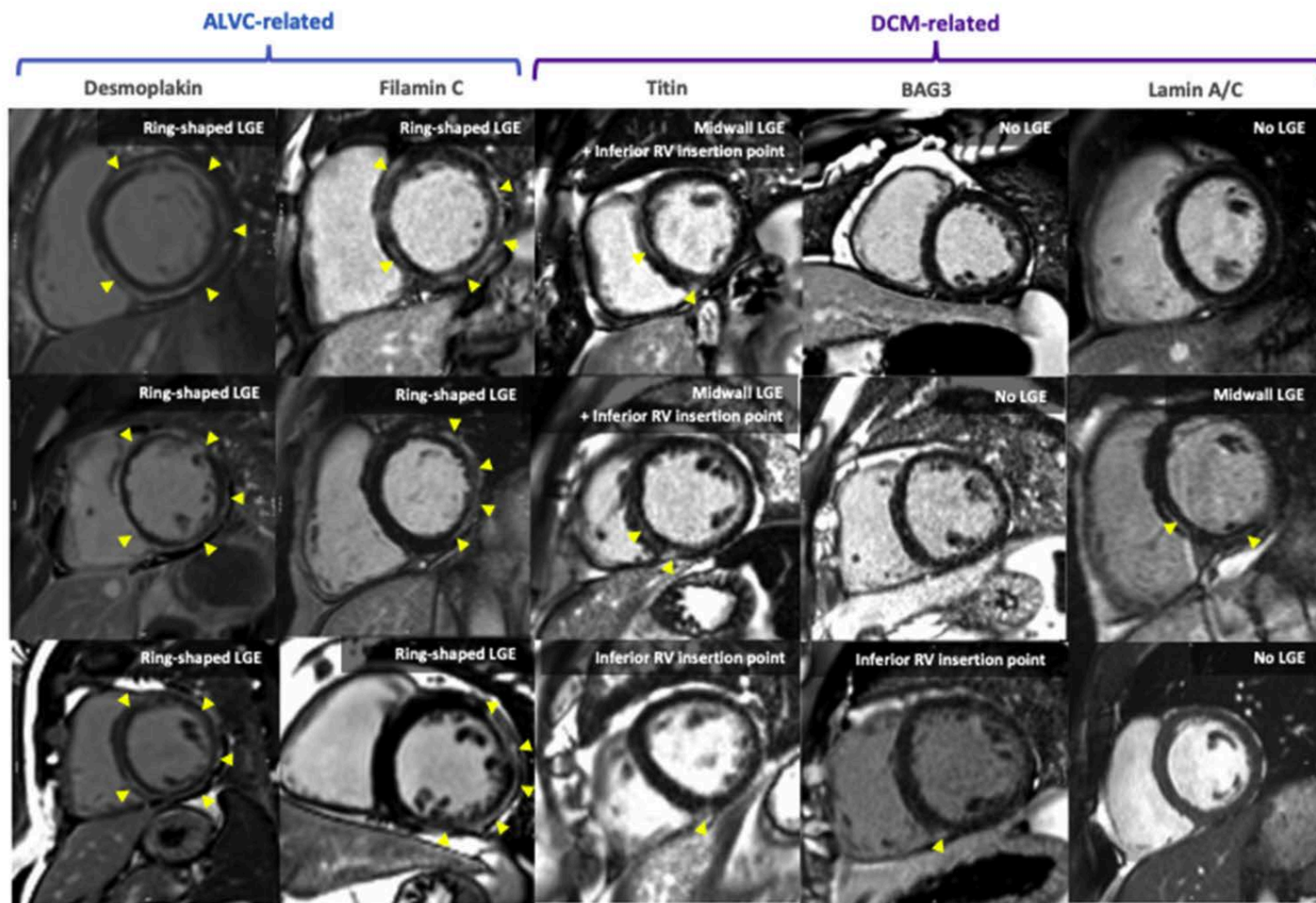
Dilated cardiomyopathy and arrhythmogenic left ventricular cardiomyopathy: a comprehensive genotype-imaging phenotype study



Distinctive imaging phenotypes define arrhythmogenic left ventricular cardiomyopathy (ALVC)



**YOU SEE,
BUT YOU
DO NOT
OBSERVE**



Some possible options?

1

Do nothing

2

Tinker with
existing
definitions

3

Redesign to
resolve current
confusion

4

Move to a new
aetiology-based
classification

Cardiomyopathy: Definition

- *A structural and or functional abnormality of left and or right ventricle (or atria) unexplained solely by abnormal loading conditions or coronary artery disease.*

- **STRUCTURAL**=hypertrophy, dilatation, systolic/diastolic impairment (global/regional), scar and fat.

- **NOT DEFINED BY PRESENCE OR ABSENCE OF ARRHYTHMIA**

- +
 - · Cardiomyopathy:
Diagnostic
Pathway



Diagnostic Pathway

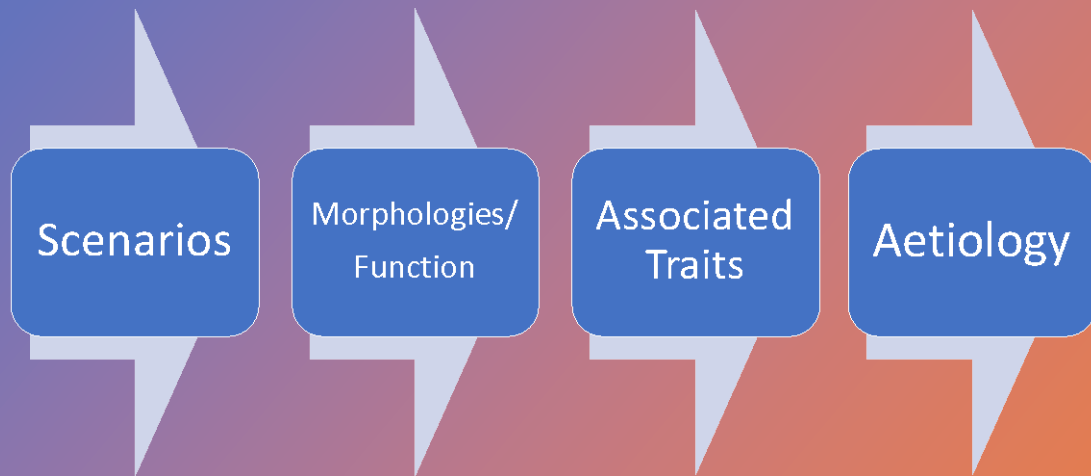


Scenarios

Morphologies/
Function

Associated
Traits

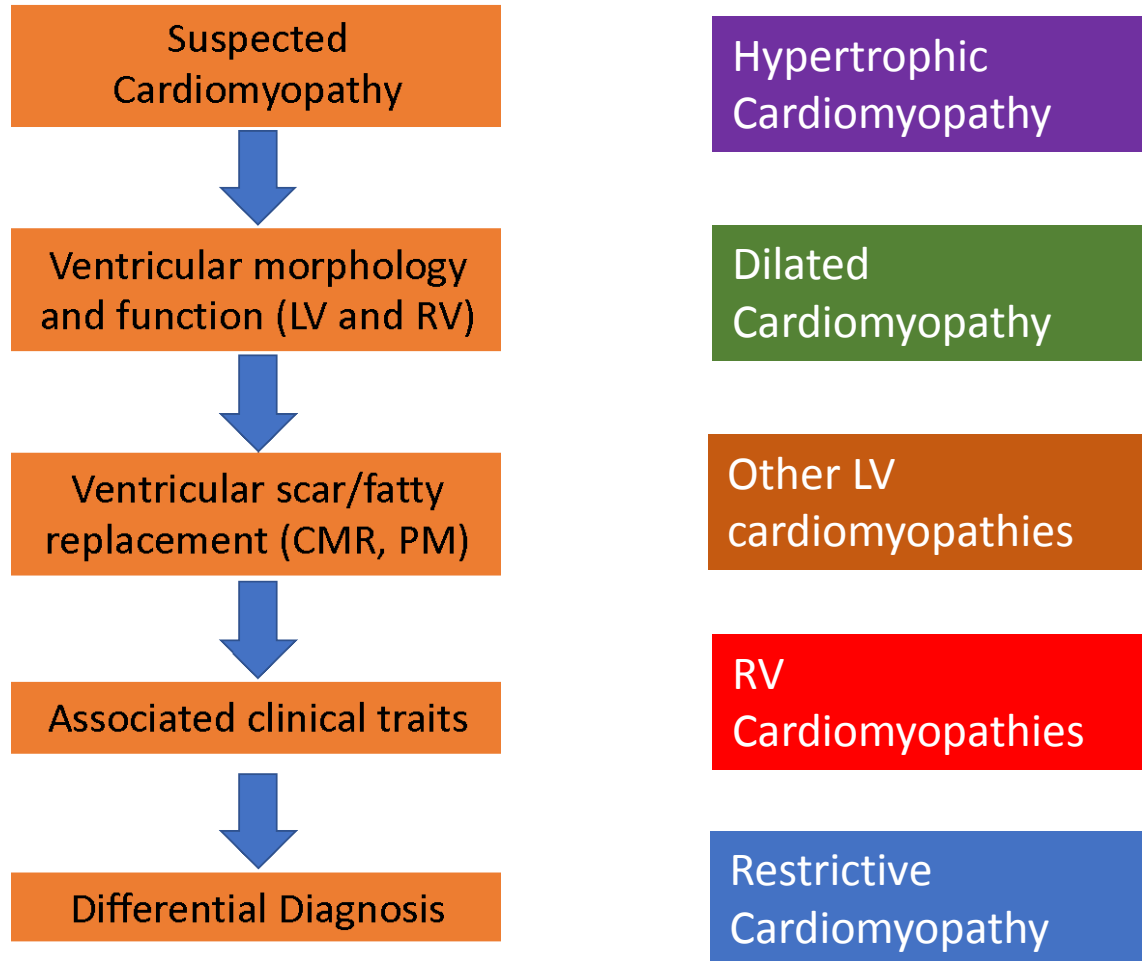
Aetiology

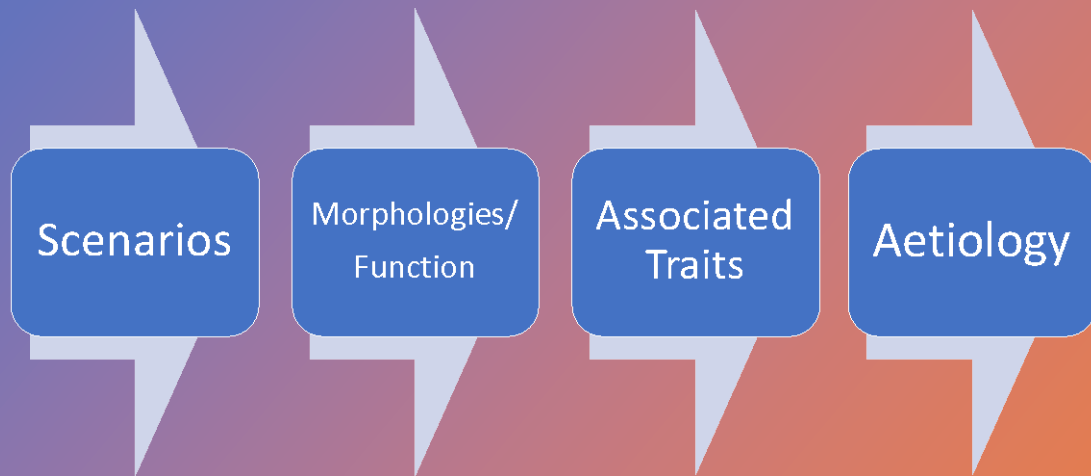


Scenarios

- Symptomatic
- Arrhythmia
- Family Screening
- Incidental

Morphological and functional subtypes of cardiomyopathy





Associated Traits*

Cardiac

Atrial Arrhythmia

Ventricular Arrhythmia

AV Block

Excessive Trabeculation

Extra-cardiac

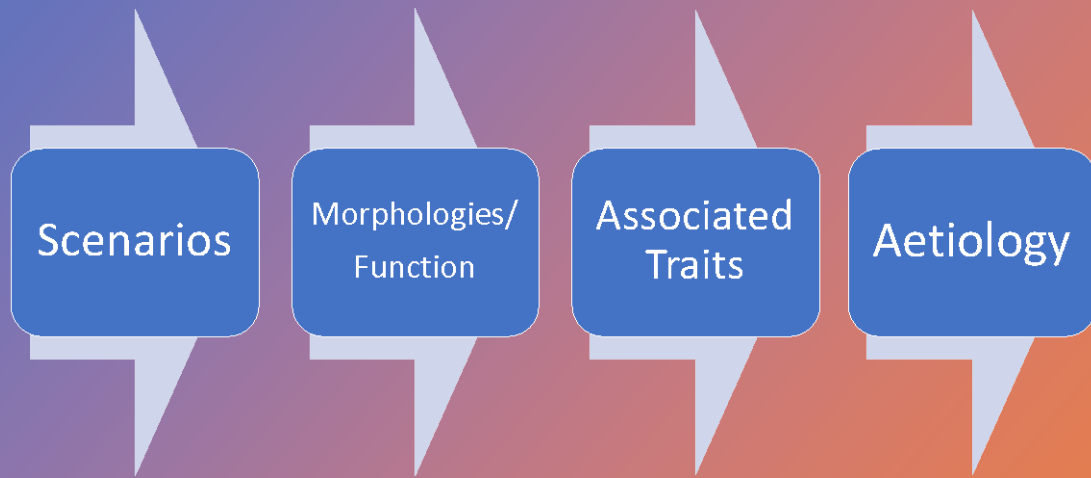
Somatic dysmorphism

Neuropathy

Renal dysfunction

Muscular dystrophy

(*including family history)



Aetiology



History

46 year old male
 Myocarditis presentation
 No family history
 Normal physical examination

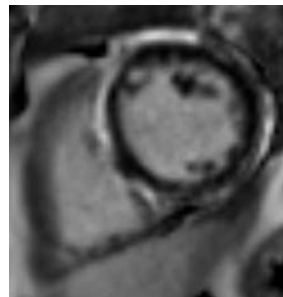
ECG/Arrhythmia



Sinus rhythm, low voltage

900 VEs/24 h

Cardiac Structure/function



Non-dilated LV
 cardiomyopathy LVEF
 subepicardial scar

Genotype

DSP p.Arg425*

Truncating variant in
 desmoplakin

APPROACHES TO 'CLASSIFICATION'

Modified Existing Morphological Classification

Arrhythmic cardiomyopathy

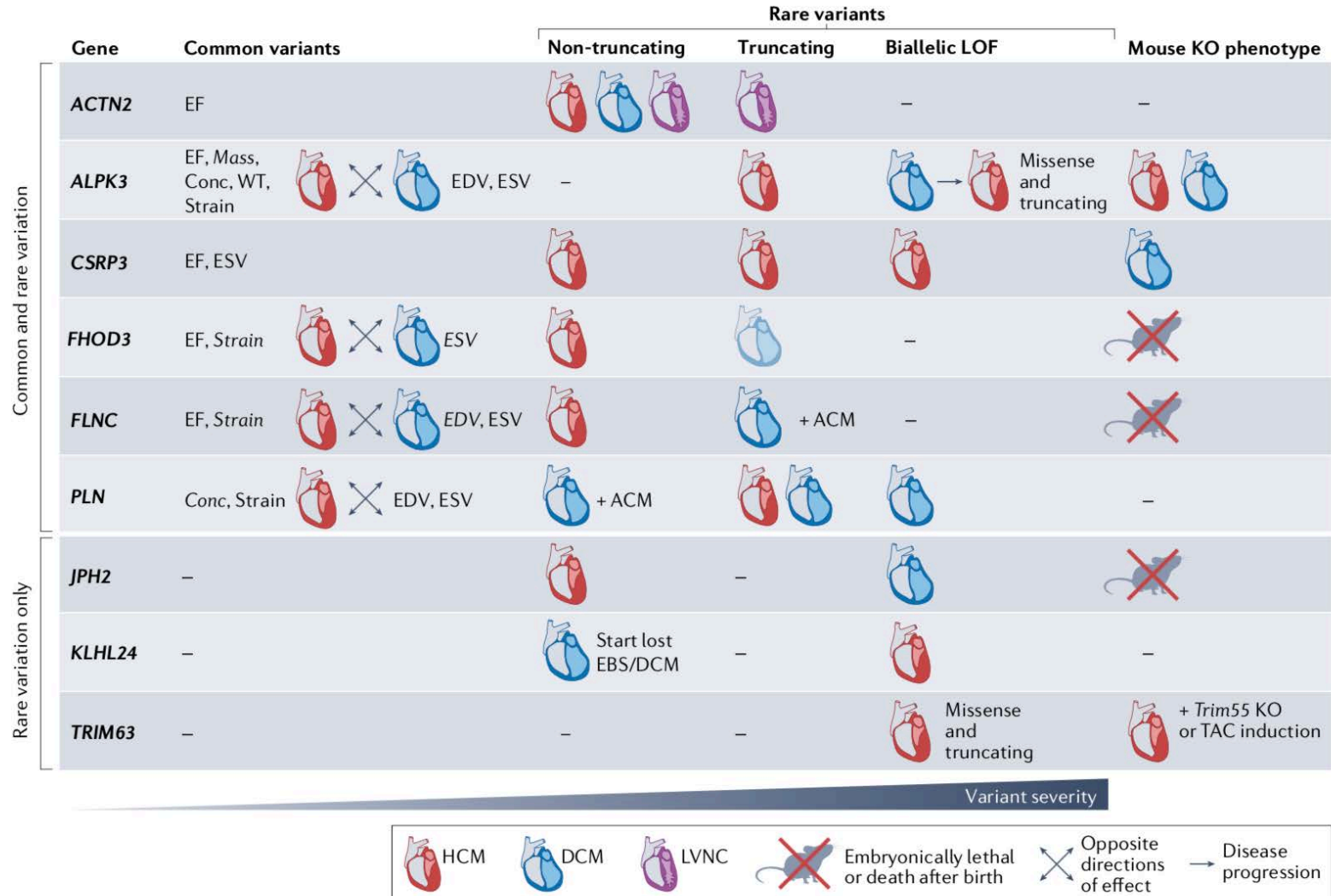
ALVC

Integrated aetiology/phenotype description

Autosomal dominant DSP-related non-dilated LV cardiomyopathy with mildly impaired LVEF and subepicardial scar

Minor hypertrophic cardiomyopathy genes, major insights into the genetics of cardiomyopathies

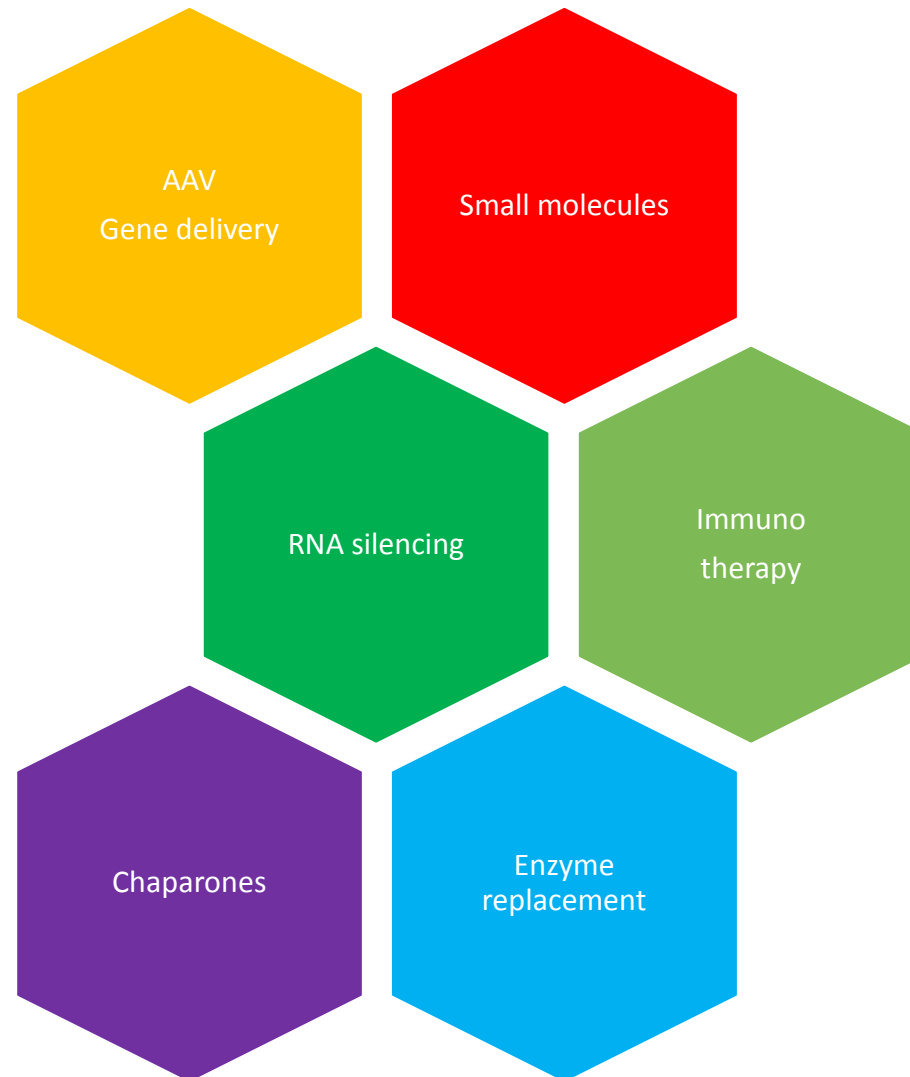
Roddy Walsh¹, Joost A. Offerhaus¹, Rafik Tadros² and Connie R. Bezzina¹



NOTATION	MORPHO-FUNCTIONAL PHENOTYPE	ORGAN/SYSTEM INVOLVEMENT	GENETIC INHERITANCE PATTERN		ETIOLOGY	STAGE
CHARACTERISTICS	Proband's cardiomyopathy (CM) diagnosis (DCM, HCM, RCM, ARVC/D, LVNC)	Clinical history and evaluation Organ involvement: Extracardiac organs/tissues Multidisciplinary evaluation according per clinical needs or diagnostic hypothesis	Genetic counseling with pedigree Familial Inheritance AD, AR XL (R or D) or Matrilineal	Clinical family screening Non-familial; Phenotypically sporadic Informative and non-informative families Consultant non-informed about family history Affected, asymptomatic relative unaware of the disease Relatives with ECG and/or Echo abnormalities Healthy family members with normal ECG and ECHO	Genetic testing in the proband Positive Cascade genetic testing in relatives Negative New tests novel genes Regular monitoring in relatives	Functional status ACC/AHA, NYHA
SUBSCRIPT	D Dilated H Hypertrophic R Restrictive R EMF Endomyocardial fibrosis LV=left ventricle RV=right ventricle	H Heart LV=left ventricle RV=right ventricle RLV=biventricular M Muscle (skeletal) N Nervous C Cutaneous	N Family history negative U Family history unknown AD Autosomal dominant AR Autosomal recessive XLD X-linked dominant XLR X-linked recessive		G Genetic cause OC Obligate carrier ONC Obligate non-carrier DN <i>De novo</i> Neg Genetic test negative for the known familial mutation N Genetic defect not identified	ACC-AHA stage represented as letter A, B, C, D NA not applicable NU

Frontiers: New Armoury

- β -blockers
- RAAS inhibitors
- Statins
- Vasodilators
- Antiplatelets
- Anticoagulants
- Devices



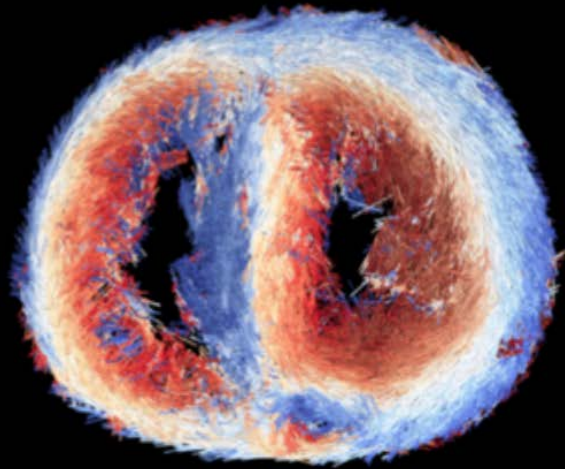


If it doesn't fit, then do not force it





CARDIOVASCULAR SCIENCE





THANK YOU

Summary: Is it time for a new classification?

- Update required to account for LV disease not described in current system
- No single scheme can meet every purpose
- For clinicians, focus should be on logical pursuit of a diagnosis guided by an iterative description of a phenotype

Ave Imperator, morituri te salutant

