

3rd PORTO MEETING 2020
September, 18th - 19th

ARRHYTHMO "GENETIC" CARDIOMYOPATHIES: PERSONALIZING SUDDEN DEATH RISK STRATIFICATION

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Advances in Heart Failure

SUMMARY

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- ✓ Sudden cardiac death in non-ischemic dilated cardiomyopathy
- ✓ New concepts – the arrhythmogenic cardiomyopathies
- ✓ Personalized arrhythmic risk assessment
- ✓ Clinical, electrocardiographic, electrophysiological and imaging parameters
- ✓ Importance of genetic profiling in sudden death risk stratification
- ✓ Conclusions

SUDDEN CARDIAC DEATH IN (NON-ISCHAEMIC) DILATED CARDIOMYOPATHY: A CHALLENGING ISSUE

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- Three-year mortality remains high (12-20%) resulting from heart failure or arrhythmic sudden cardiac death (SCD).
- Incidence of SCD in dilated cardiomyopathy (DCM) patients is ~1.5/100 PA.
- DCM accounts for a substantial proportion of SCD specially among people of working age (younger, less comorbidity, lower mortality risk from other causes).



SUDDEN CARDIAC DEATH IN (NON-ISCHAEMIC) DILATED CARDIOMYOPATHY: A CHALLENGING ISSUE

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| Study | n | Inclusion Criteria | Intervention | Follow-Up (Median), mo | All-Cause Mortality | Sudden Cardiac Death |
|------------------------------------|------|---|-----------------------|------------------------|--|---|
| CAT ³ | 104 | LVEF <30% NYHA II–III | ICD vs OMT | 23 | Terminated early | |
| AMIOVIRT ⁴ | 103 | LVEF ≤35% NYHA I–III NSVT | ICD vs amio | 24 | Terminated early | |
| SCD-HeFT (DCM cohort) ⁵ | 1211 | LVEF <35% NYHA II–III | ICD vs OMT vs amio | 46 | I, 21.4%; C, 27.9% (5 y) HR, 0.73; 95% CI, 0.50–1.07; P=0.06 | |
| DEFINITE ⁶ | 458 | LVEF <36% NYHA I–III NSVT or PVCs | ICD vs OMT | 29 | I, 12.2%; C, 17.4% HR, 0.65; 95% CI, 0.40–1.06; P=0.08 | I, 1.3%; C, 6.1% HR, 0.20; 95% CI, 0.06–0.71; P=0.006 |
| DANISH ⁷ | 1116 | LVEF <35% NYHA II–III (IV if CRT) NT-proBNP >200pg/mL | ICD vs OMT | 68 | I, 21.6%; C, 23.4%; HR, 0.87; 95% CI, 0.68–1.12; P=0.28 | I, 4.3%; C, 8.2% HR, 0.50; 95% CI, 0.31–0.82; P=0.005 |

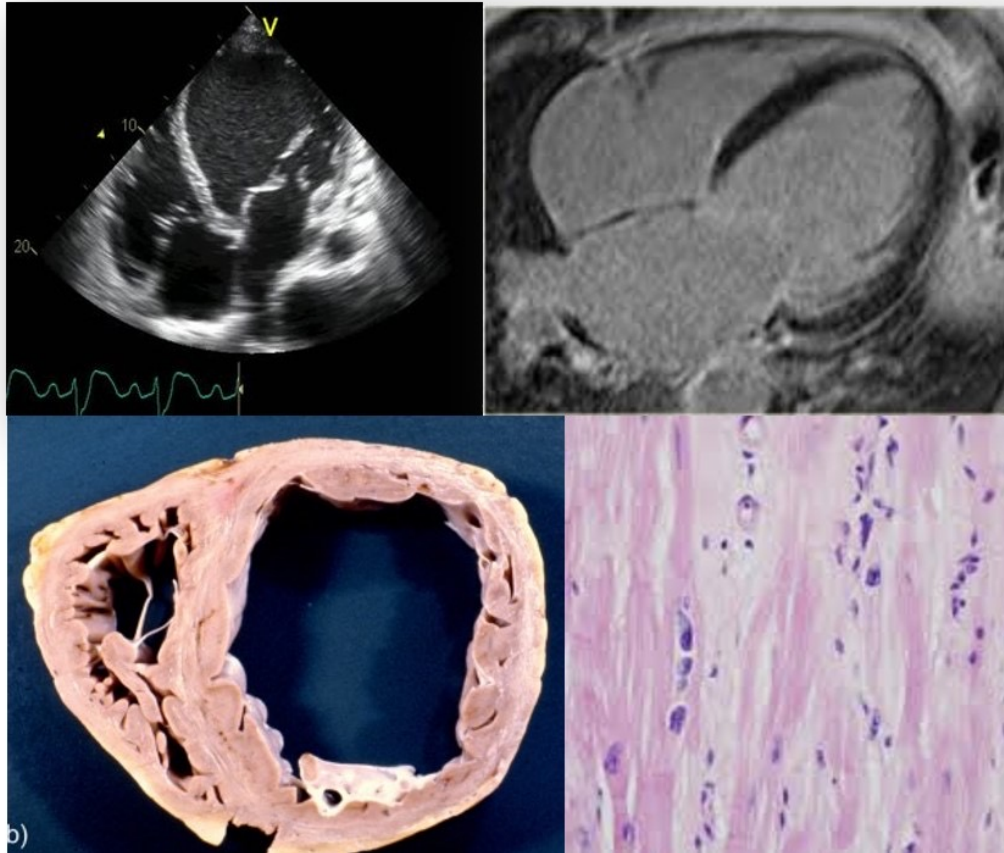
SUDDEN CARDIAC DEATH IN (NON-ISCHAEMIC) DILATED CARDIOMYOPATHY: A CHALLENGING ISSUE

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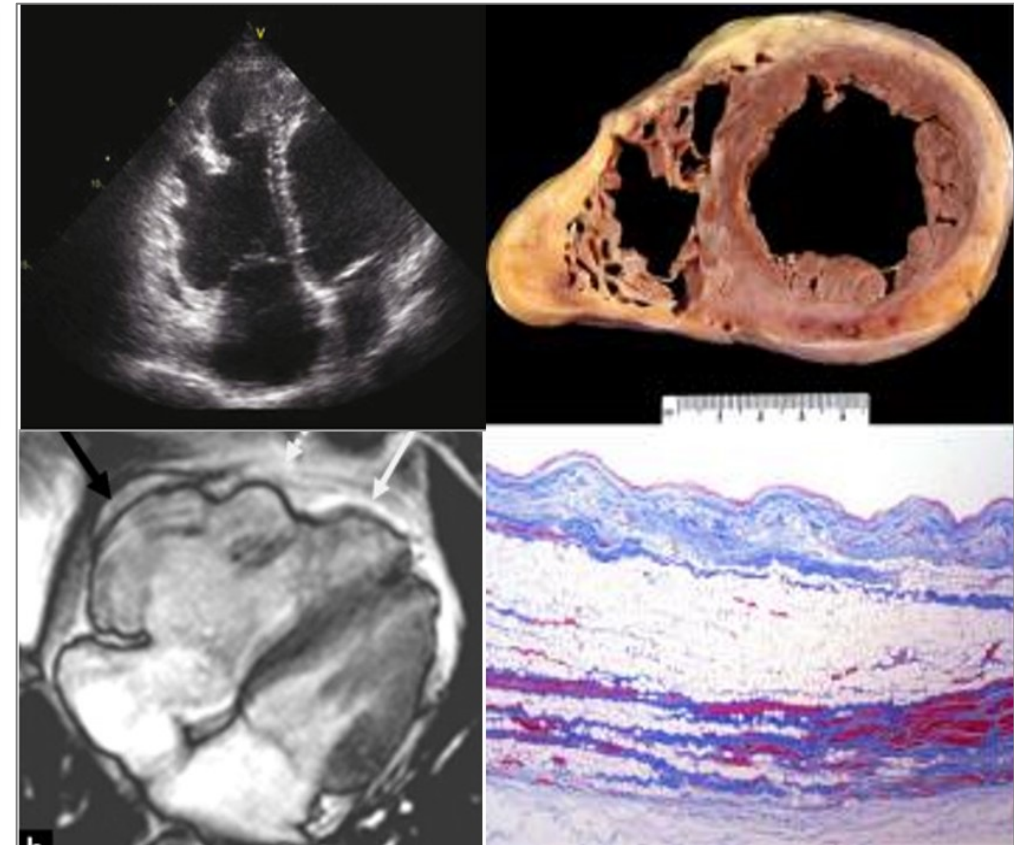
- Poor sensitivity and specificity of left ventricular ejection fraction (LVEF) for predicting SCD:
 - High proportion of cardiac arrest in patients with LVEF > 35%
 - Low incidence of appropriate ICD therapies
- Complications of ICD, inappropriate shocks and costs
- Inherent subjectivity in assessing LVEF and NYHA
- Up to 40% experience reverse remodelling under treatment

CLASSIC DEFINITIONS

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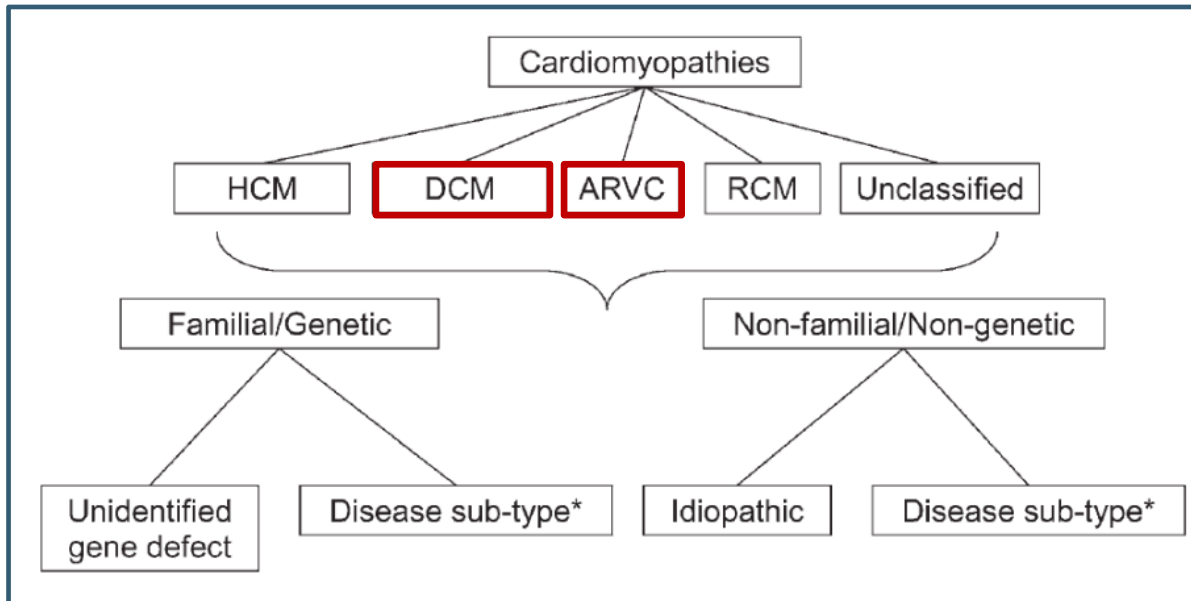


Dilated cardiomyopathy

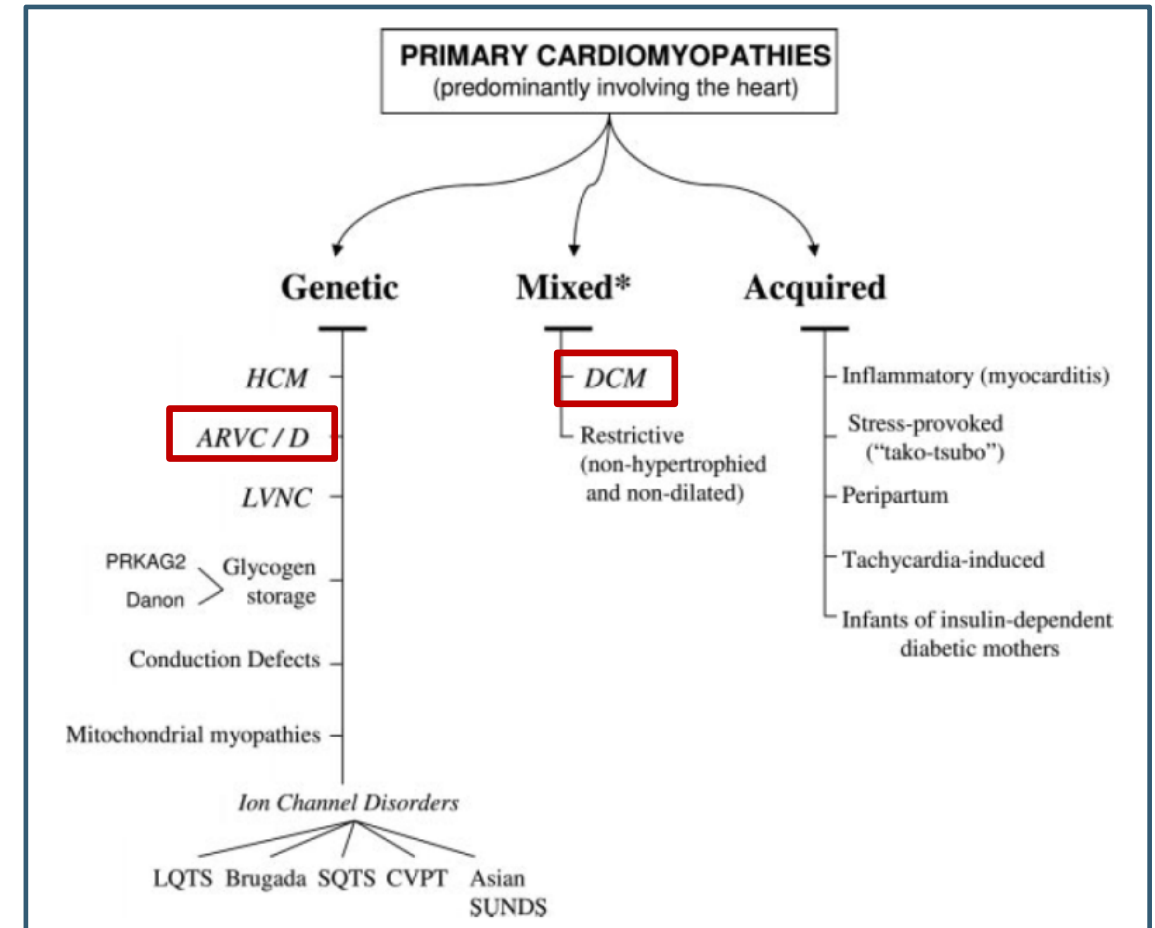


Arrhythmogenic (right ventricular) cardiomyopathy

CLASSIFICATION OF CARDIOMYOPATHIES



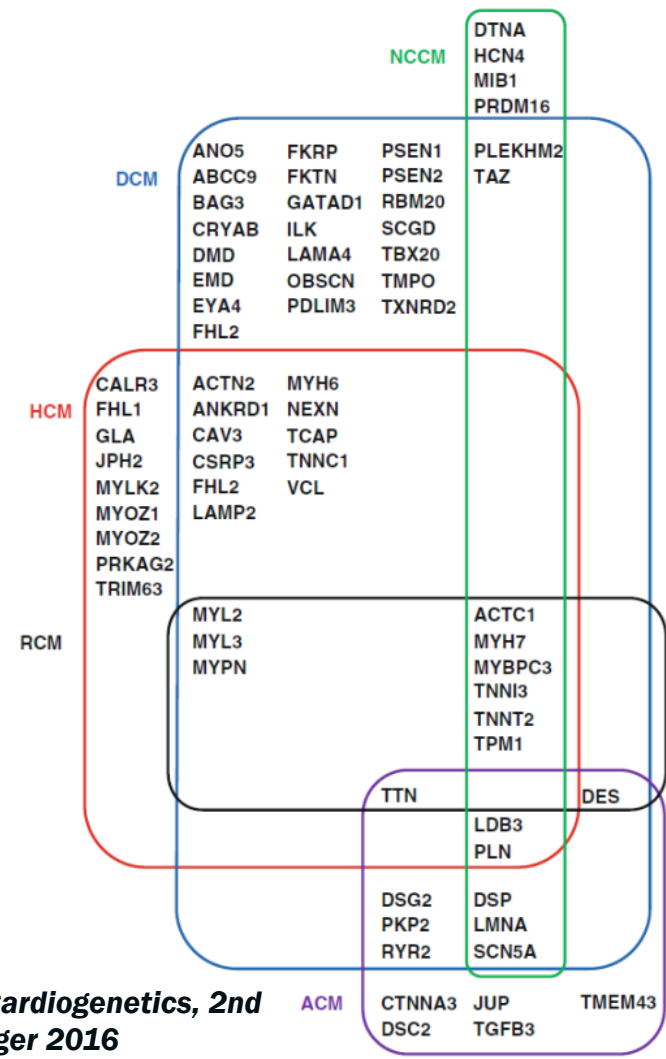
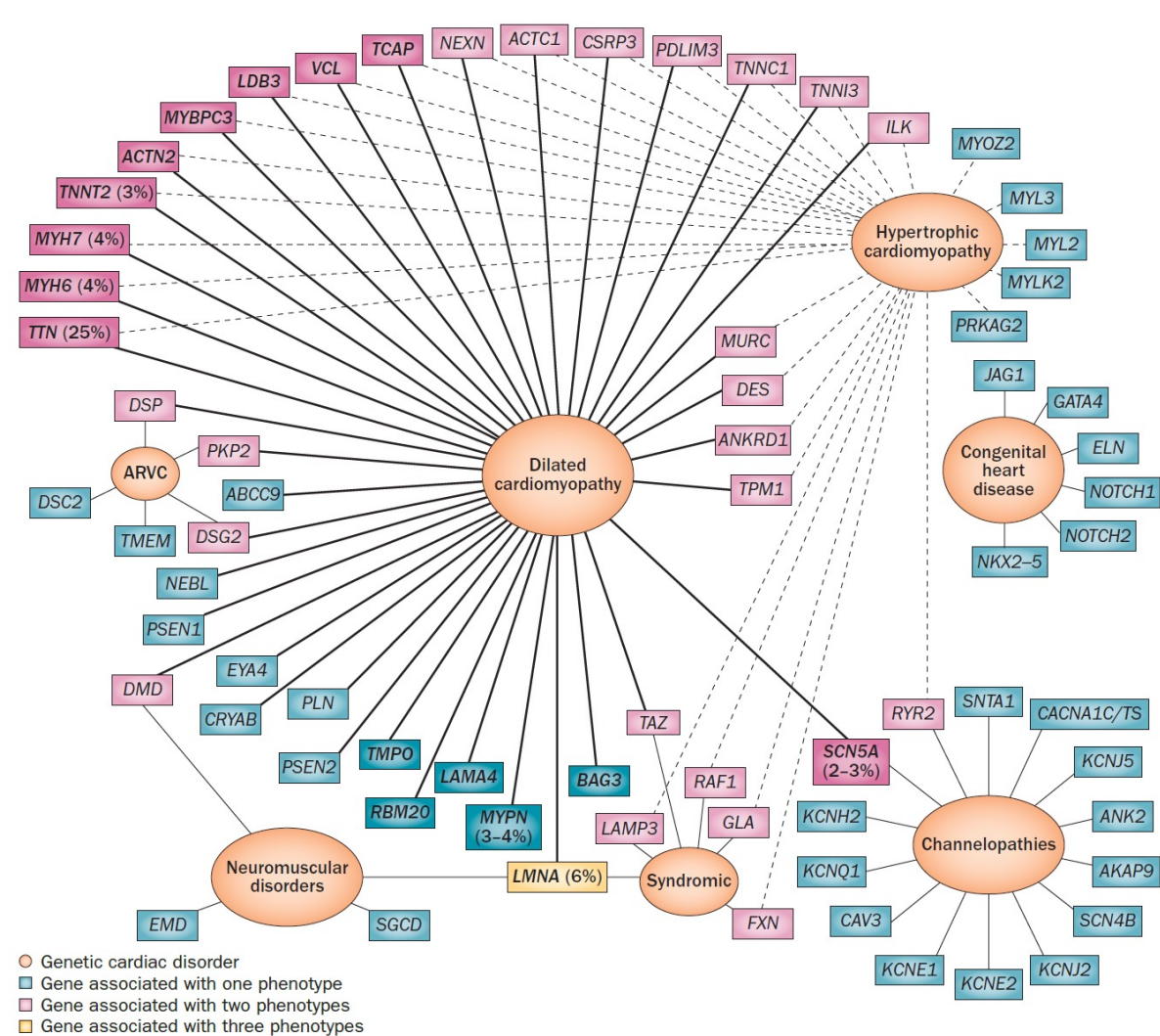
Elliot P, EHJ 2008



Maron BJ, Circ 2006

Genetic heterogeneity – numerous genetic variants, in multiple genes, with great overlap between “different” cardiomyopathies

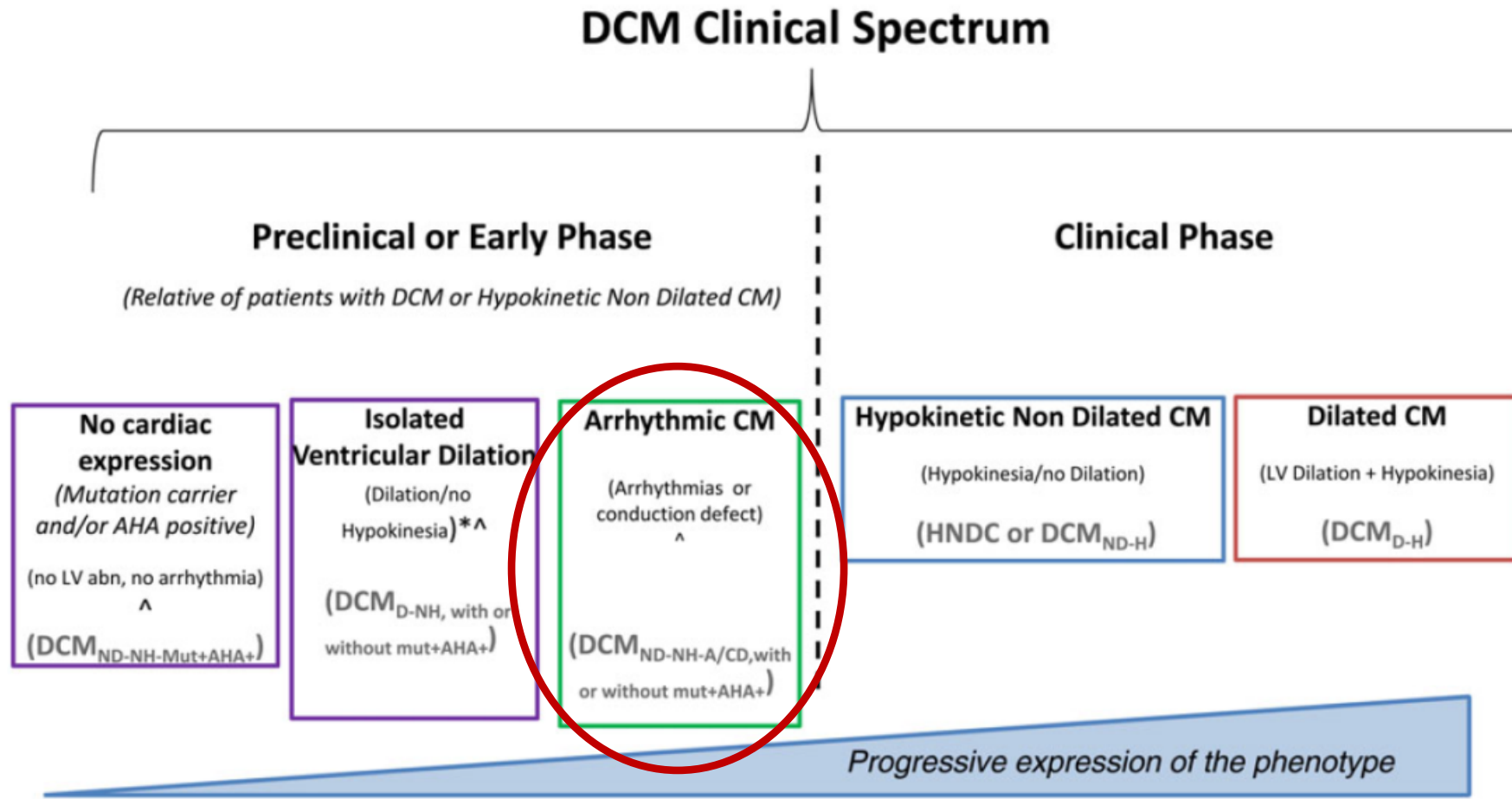
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Clinical Cardiogenetics, 2nd
Ed. Springer 2016

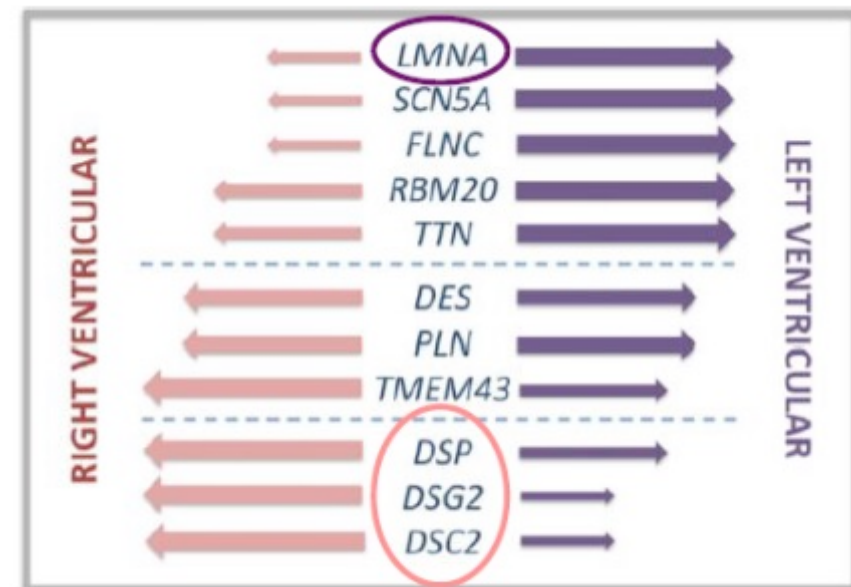
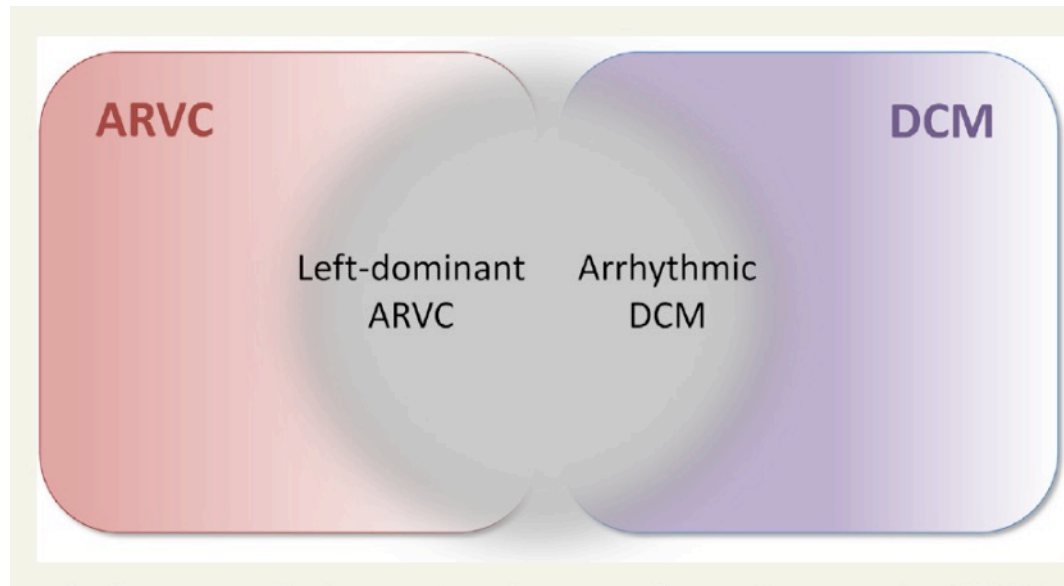
ESC NEW DEFINITION OF DILATED CARDIOMYOPATHY

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*Shown by two independent imaging modalities, ^mutation carrier or not; anti-heart autoantibody (AHA) positive or negative

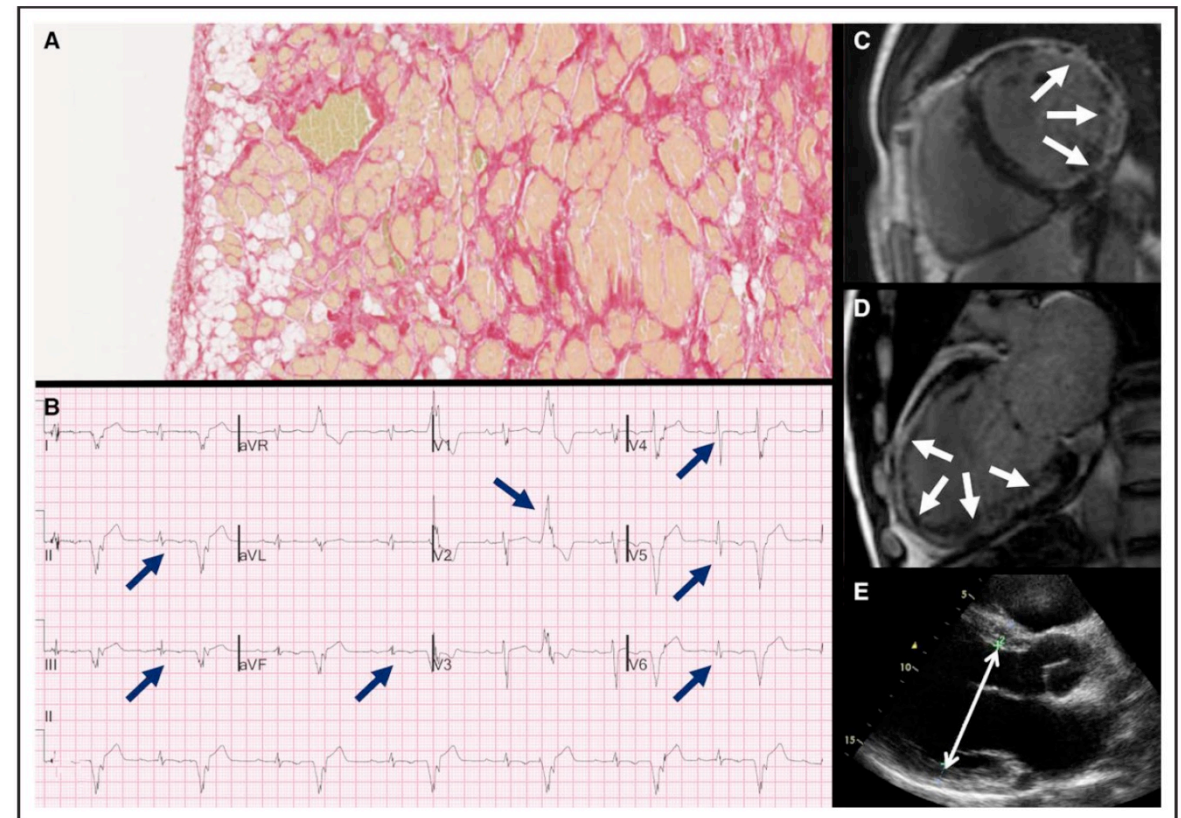
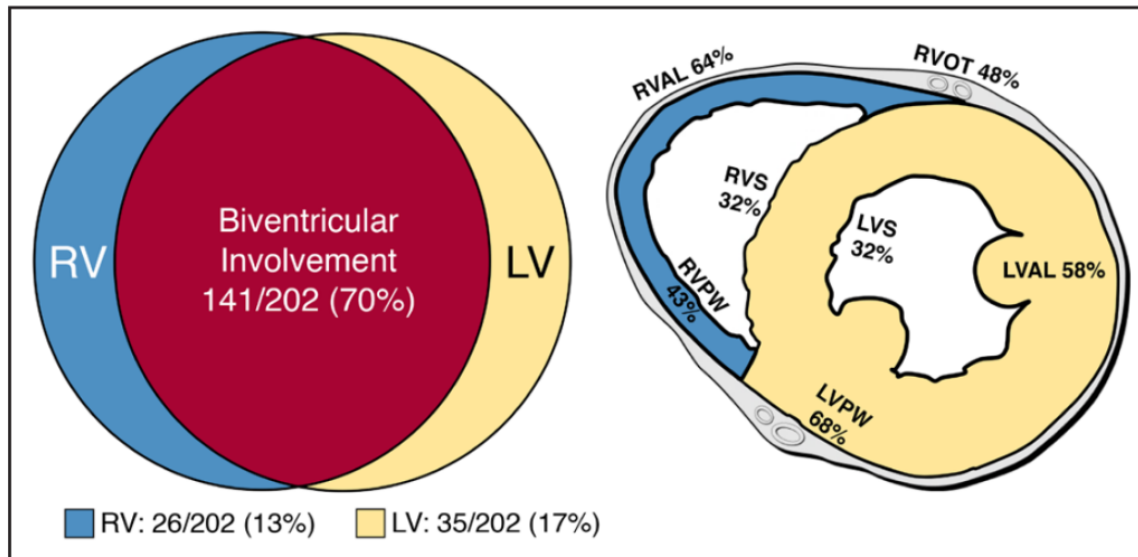
Arrhythmic Genotypes in Familial Dilated Cardiomyopathy: Implications for Genetic Testing and Clinical Management



LEFT VENTRICULAR INVOLVEMENT IN ARRHYTHMOGENIC CARDIOMYOPATHY

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Sudden Death and Left Ventricular Involvement in Arrhythmogenic Cardiomyopathy



NEW DEFINITIONS OF ARRHYTHMOGENIC CARDIOMYOPATHY

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European Journal of Heart Failure (2019)
doi:10.1002/ejhf.1534

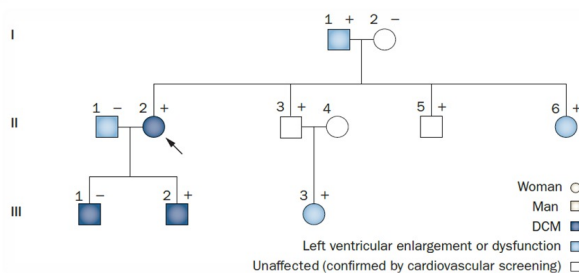
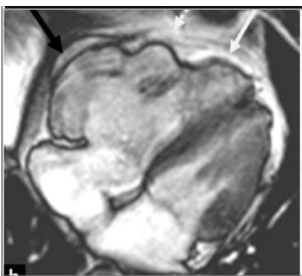
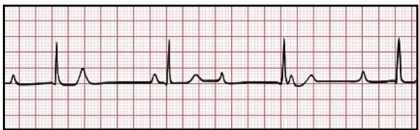
CONSENSUS DOCUMENT

Definition and treatment of arrhythmogenic cardiomyopathy: an updated expert panel report

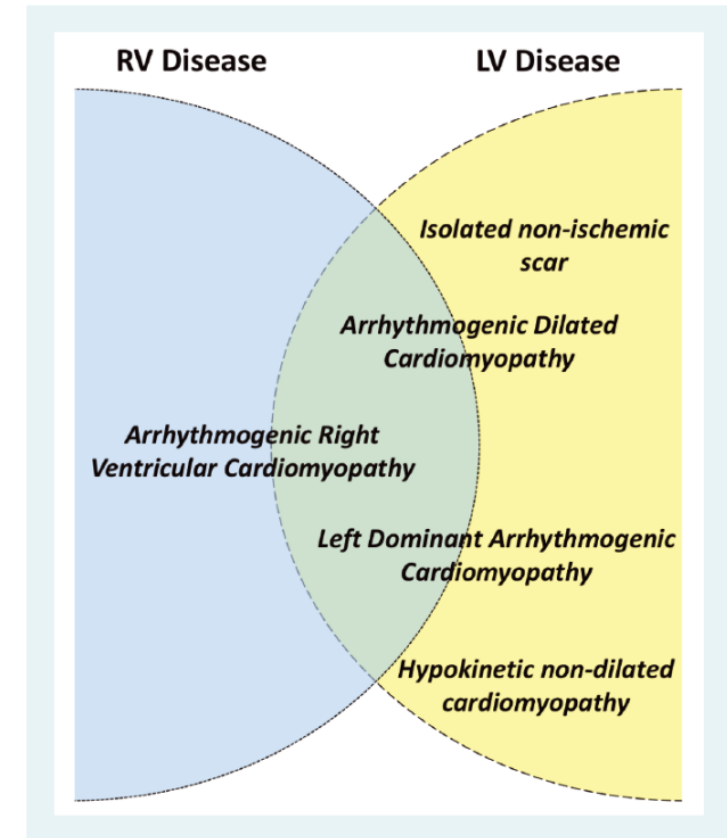
Perry M. Elliott^{1*}, Aris Anastasakis², Angeliki Asimaki³, Cristina Basso⁴, Barbara Bauce⁴, Matthew A. Brooke⁵, Hugh Calkins⁶, Domenico Corrado⁴, Firat Duru⁷, Kathleen J. Green⁸, Daniel P. Judge⁹, David Kelsell⁵, Pier D. Lambiase¹, William J. McKenna¹⁰, Kalliopi Pilichou⁴, Alexandros Protonotarios¹, Jeffrey E. Saffitz¹¹, Petros Syrris¹⁰, Hari Tandri⁶, Anneline Te Riele¹², Gaetano Thiene⁴, Adalena Tsatsopoulou¹³, and J. Peter van Tintelen^{14,15}

NEW DEFINITIONS OF ARRHYTHMOGENIC CARDIOMYOPATHY

The term 'arrhythmogenic cardiomyopathy' is used to describe a family of diseases that feature structural myocardial abnormalities (identified by macro- and microscopic pathological examination besides cardiac imaging) and ventricular arrhythmia. In Figure 1,



- 1) Arrhythmia
- 2) Electrical abnormalities
- 3) Structural abnormalities
- 4) Heritability
- 5) Phenocopy exclusion

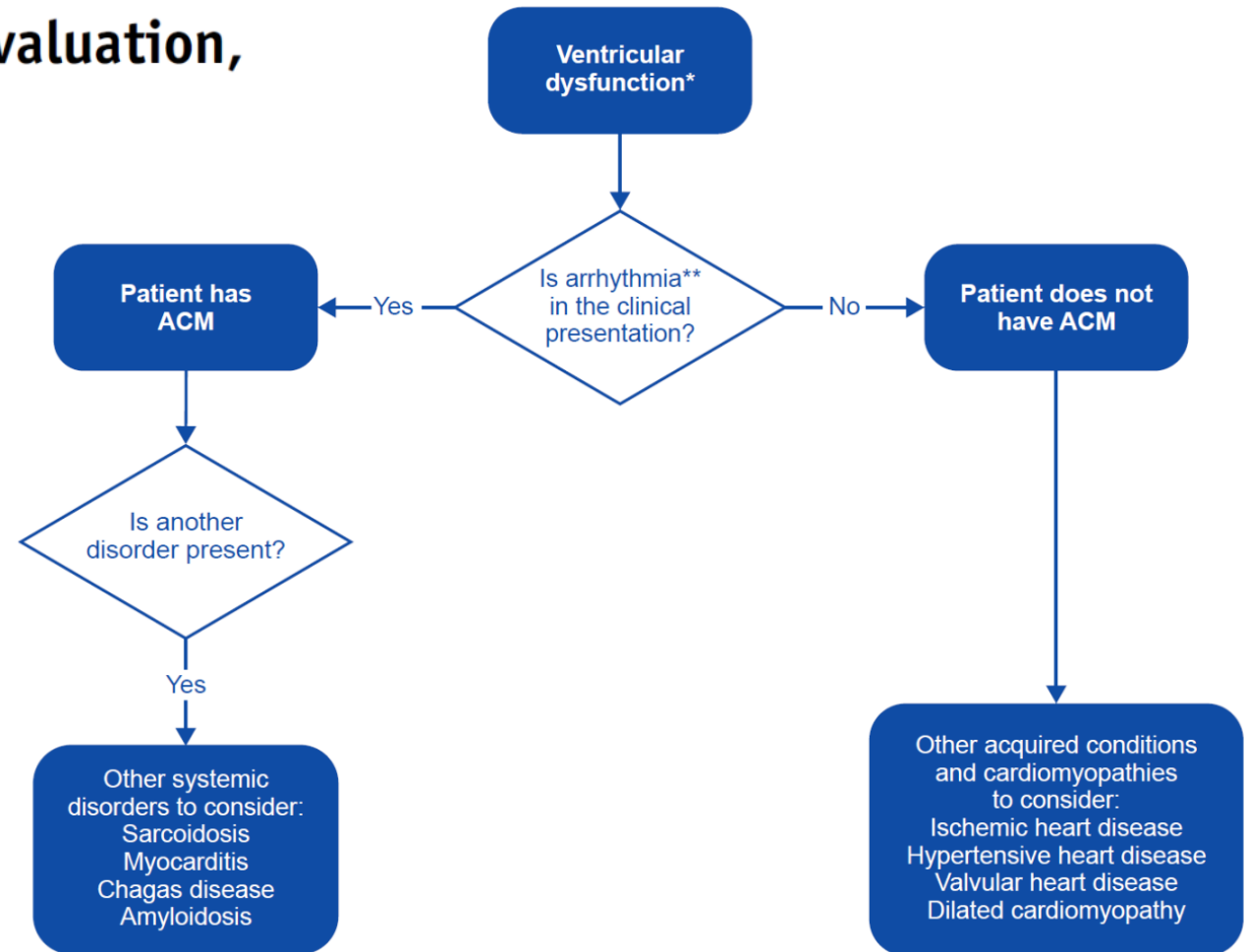


NEW DEFINITIONS OF ARRHYTHMOGENIC CARDIOMYOPATHY

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

- Arrhythmogenic heart muscle disease **not explained by ischemia, hypertension or valvopathies**
- **Arrhythmia is the presenting feature** (AF, CD, RV or LV arrhythmias)
- **Overlapping syndromes**
- **Includes:** ARVC/ALVC, DCM, Chagas disease, Myocarditis, Sarcoidosis, Amyloidosis, LV non-compaction



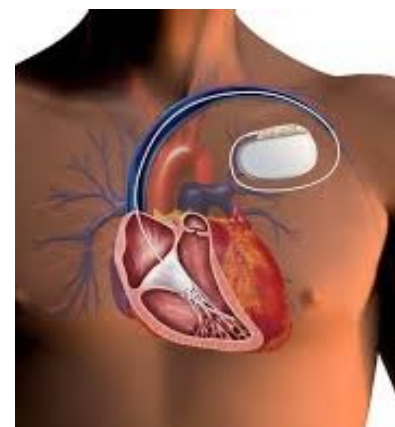
*Not explained by ischemic, hypertensive, or valvular heart disease

**Arrhythmia includes conduction disease, atrial arrhythmias, ventricular arrhythmias

ARRHYTHMIC RISK ASSESSMENT

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LVEF
NYHA class



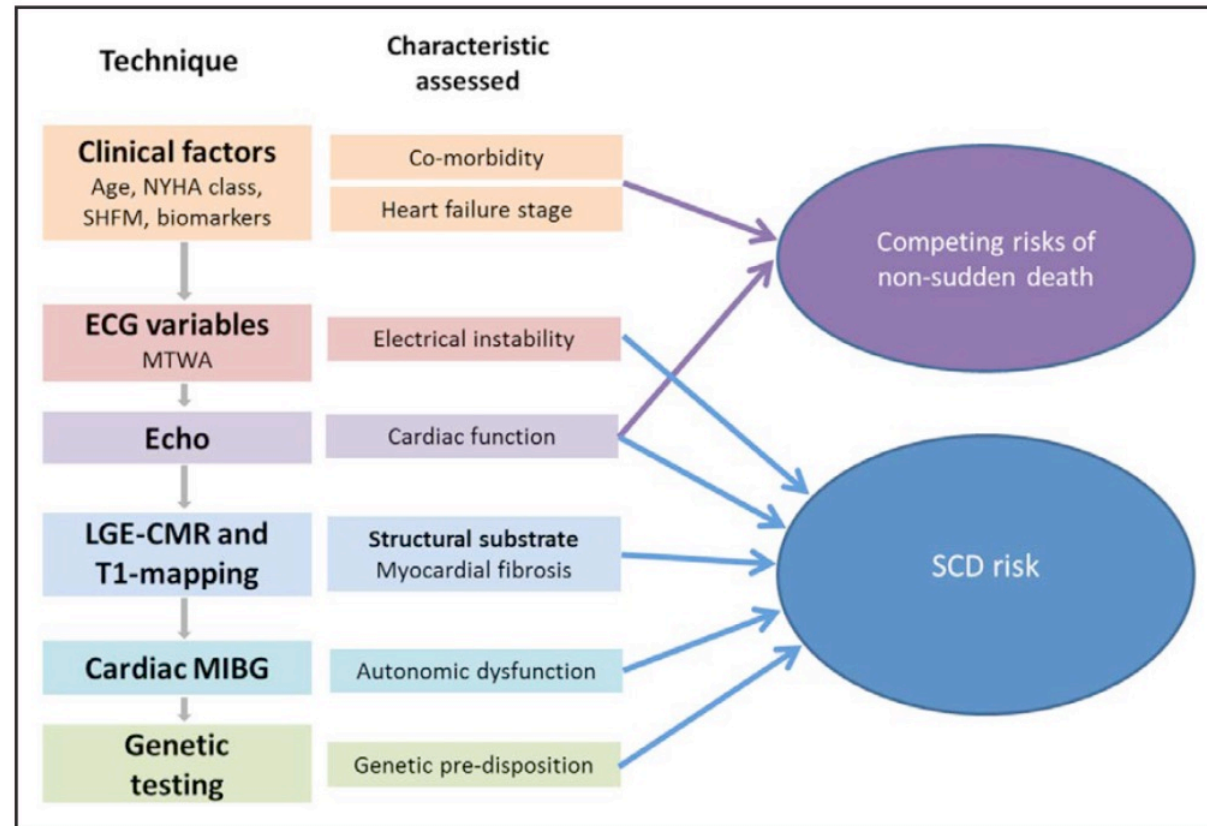
Is it enough

PERSONALIZED ARRHYTHMIC RISK ASSESSMENT

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Personalizing Risk Stratification for Sudden Death in Dilated Cardiomyopathy

The Past, Present, and Future



PERSONALIZED ARRHYTHMIC RISK ASSESSMENT

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| ARRHYTHMIC PREDICTORS IN DCM | |
|--|--|
| CLINICAL | <ul style="list-style-type: none"> Etiological characterization Personal and family history (syncope, familial history of SCD) |
| ECG and Holter ECG | <ul style="list-style-type: none"> Fragmented QRS Long QRS Low QRS amplitude Anterolateral T wave inversion NSVT (≥ 5 beats, ≥ 150 bpm) Frequent PVCs (1000/24h) and couplets ($\geq 50/24h$) |
| ECHOCARDIOGRAPHY | <ul style="list-style-type: none"> LVEF $\leq 35\%$ Reduced GLS |
| CMR | <ul style="list-style-type: none"> LGE presence LGE extent and location (i.e. septal and free-wall) T1-mapping (increased T1 relaxation time) |
| GENETIC TESTING | <ul style="list-style-type: none"> LMNA FLNC, PLN, RBM20 SCN5A TTN * |
| ELECTROPHYSIOLOGICAL PARAMETERS | <ul style="list-style-type: none"> Abnormal mTWA test |

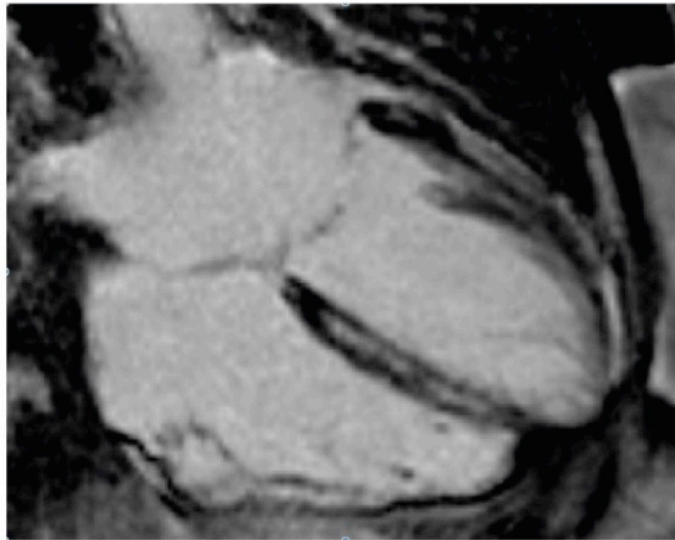
Weak evidence



Strong evidence

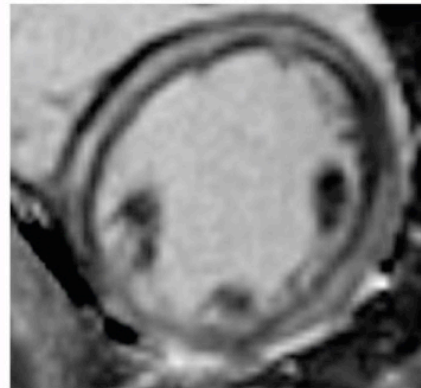
CMR – LATE GADOLINIUM ENHANCEMENT

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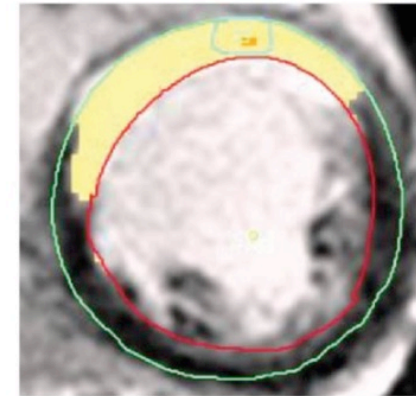
PRESENCE / ABSENCE

Strong association with worse outcomes



LOCALIZATION

Concomitant presence of septal and free wall LGE associated with worse outcome. In the future LGE localization might provide further prognostic impact



EXTENSION

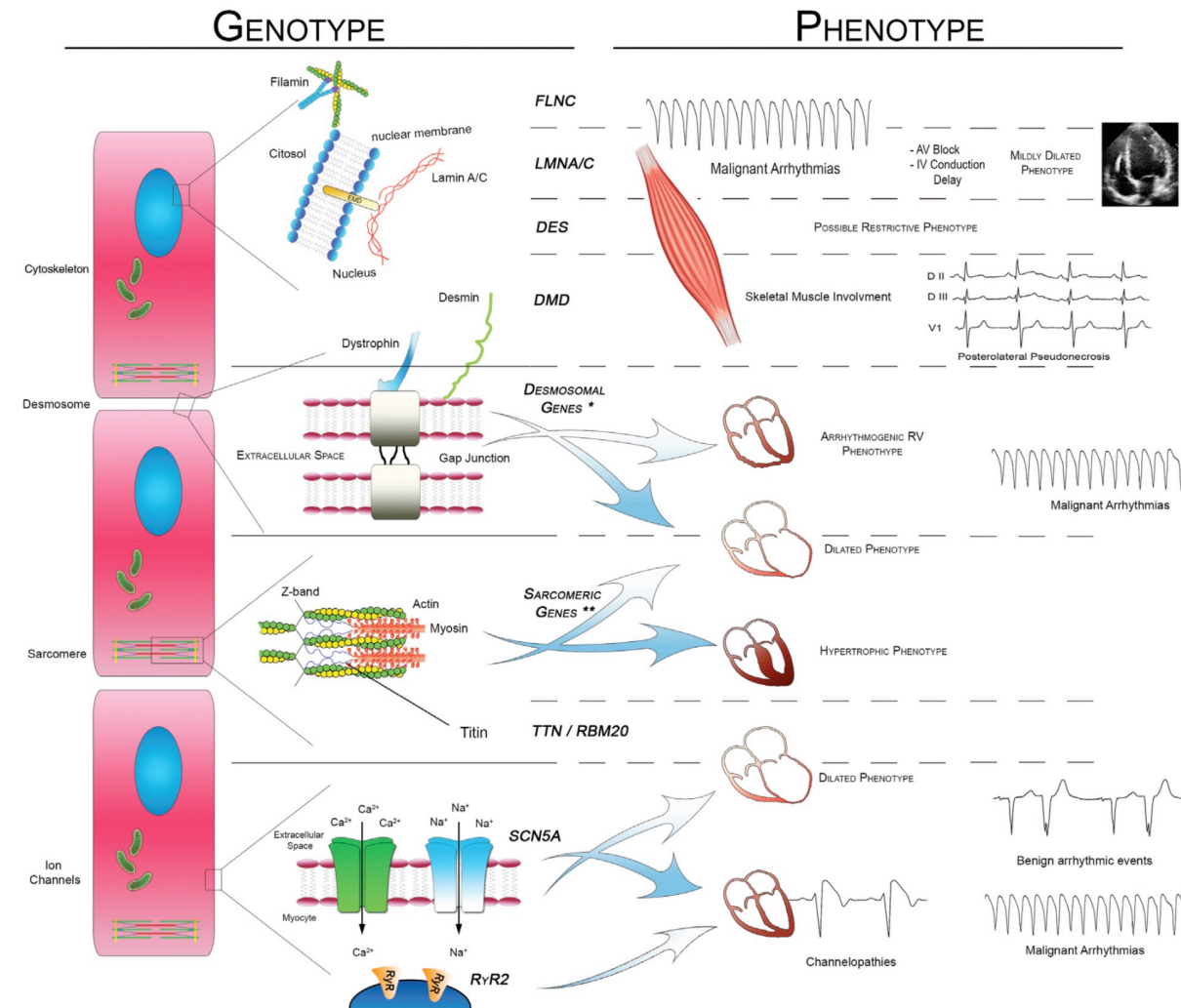
Quantitative assessment might provide further prognostic impact in the future

Strong evidences in large studies and metanalyses

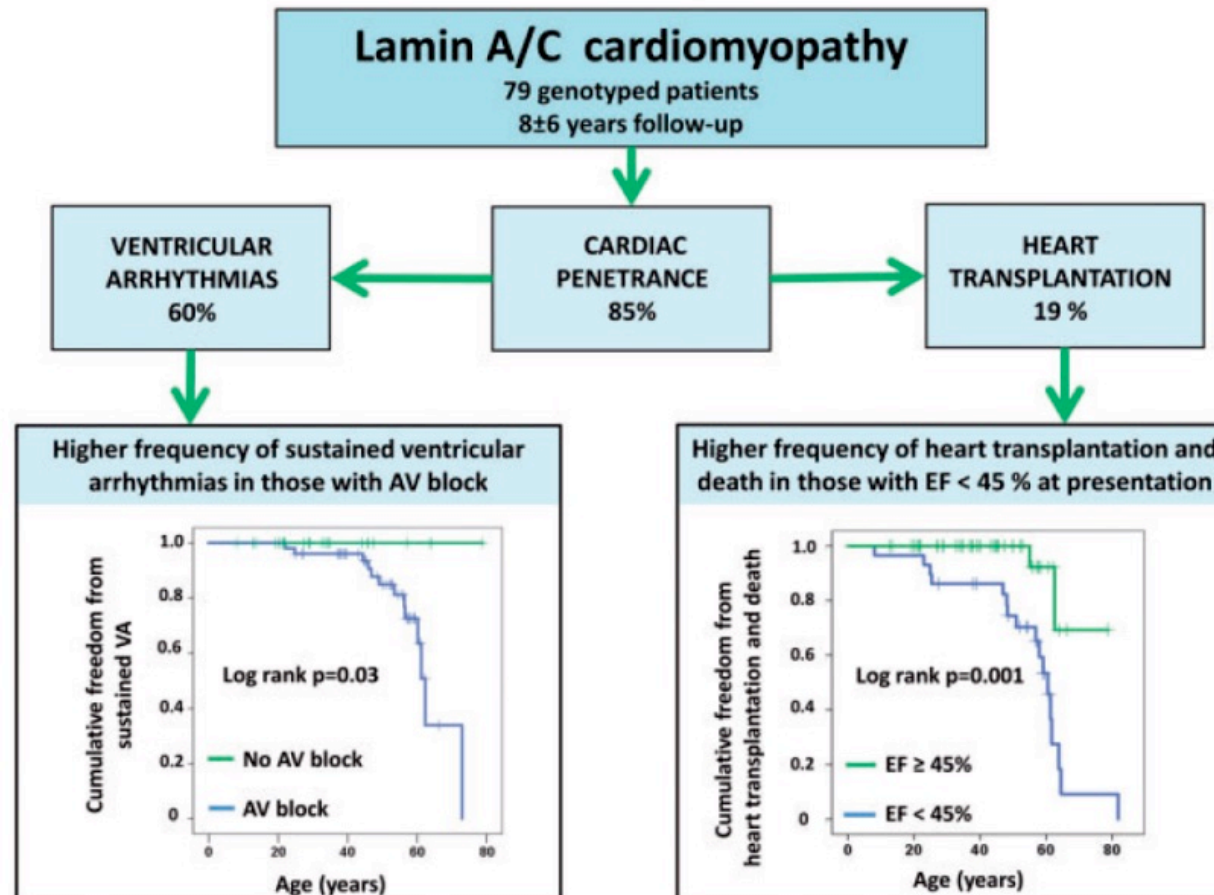
Conflicting evidences from smaller studies

IMPORTANCE OF GENETIC PROFILING IN SUDDEN DEATH RISK STRATIFICATION

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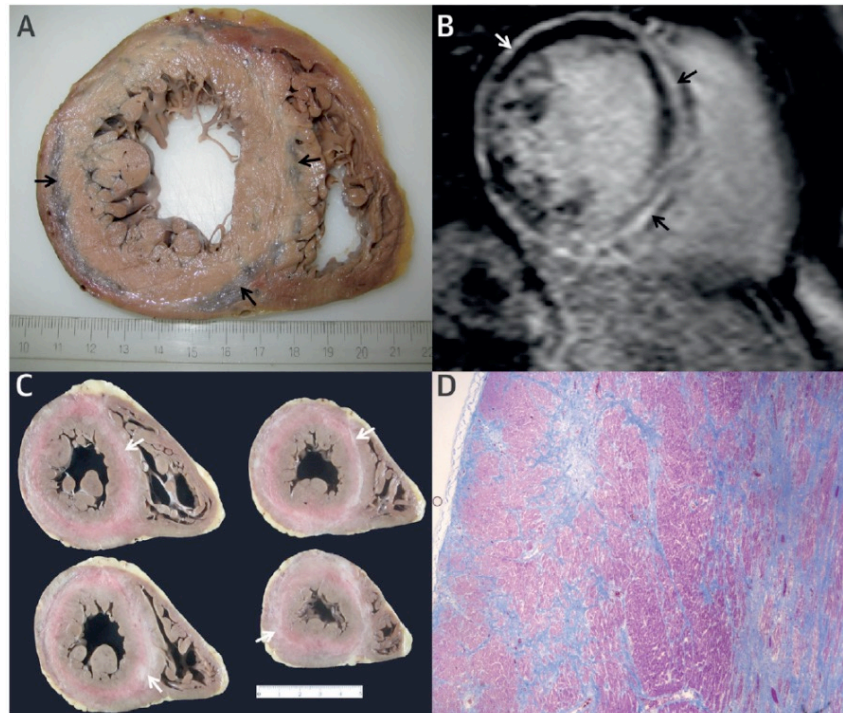
Lamin A/C cardiomyopathy: young onset, high penetrance, and frequent need for heart transplantation



FLNC – filamin C

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

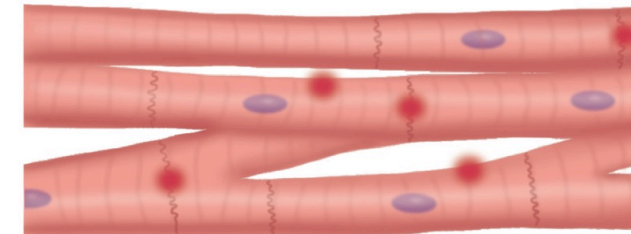


Ortiz-Genga MF, JACC 2016

Truncating FLNC Mutation Produces an Abnormal Protein

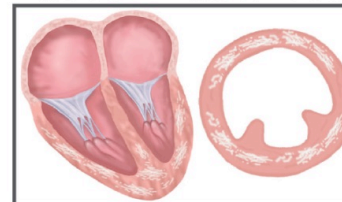


Alteration of Intercalated Disks and Costameres Weakens Myocytes' Adhesion

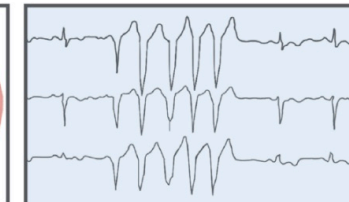


Dilated/Arrhythmogenic Cardiomyopathies

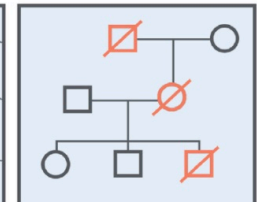
Left Ventricular Dilation and Systolic Dysfunction with Myocardial Fibrosis



Ventricular Arrhythmias



Familial Sudden Cardiac Death



Alexandra Sousa

PLN - phospholamban

Phospholamban R14del mutation in patients diagnosed with dilated cardiomyopathy or arrhythmogenic right ventricular cardiomyopathy: evidence supporting the concept of arrhythmogenic cardiomyopathy

**R14del mutation
(15% DCM, 12% ACM Netherlands)**

We screened a cohort of 97 ARVC and 257 DCM unrelated index patients for *PLN* mutations and evaluated their clinical characteristics. *PLN* mutation R14del was identified in 12 (12%) ARVC patients and in 39 (15%) DCM patients. Haplotype analysis revealed a common founder, estimated to be between 575 and 825 years old. A low voltage elec-

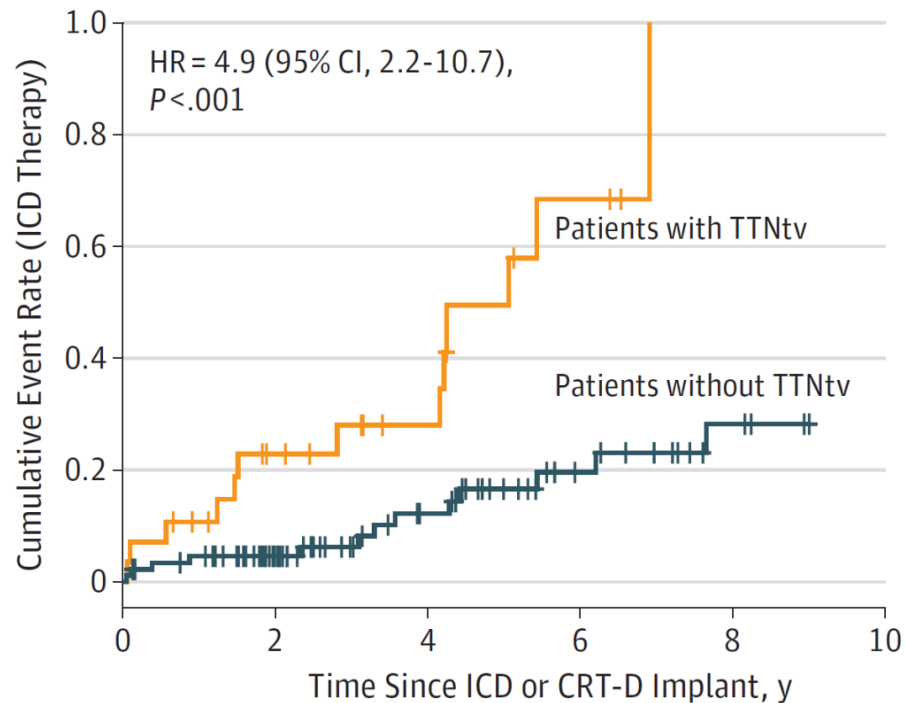
The *PLN* R14del founder mutation is present in a substantial number of patients clinically diagnosed with DCM or ARVC. R14del+ patients diagnosed with DCM showed an arrhythmogenic phenotype, and SCD at young age can be the presenting symptom. These findings support the concept of 'arrhythmogenic cardiomyopathy'.

PLN-R14Del -

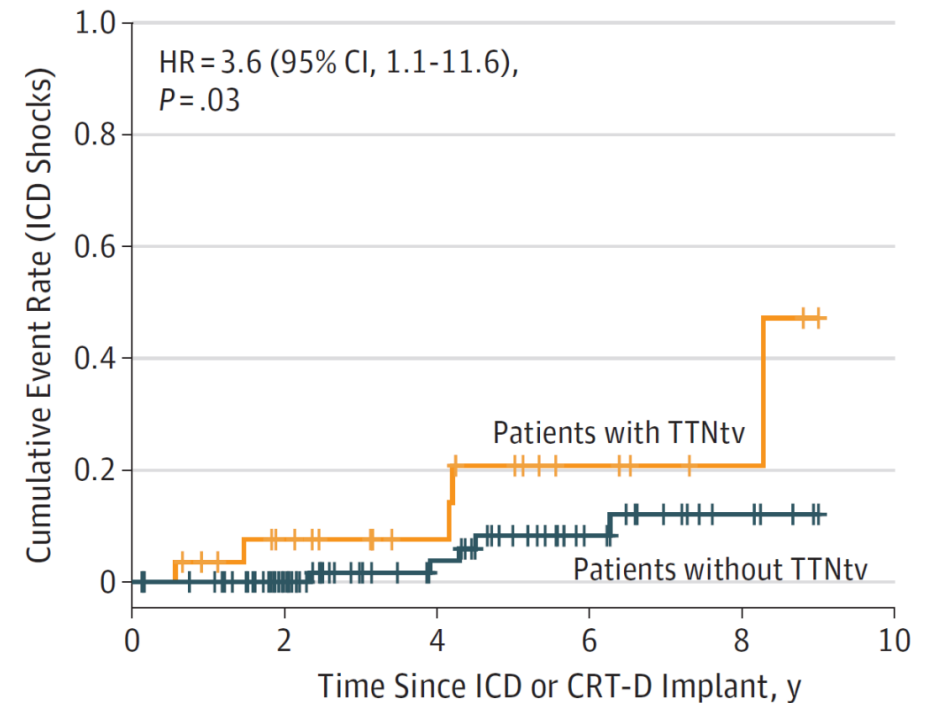


Association of Titin-Truncating Genetic Variants With Life-threatening Cardiac Arrhythmias in Patients With Dilated Cardiomyopathy and Implanted Defibrillators

A Antitachycardia pacing or shock



B Shock only

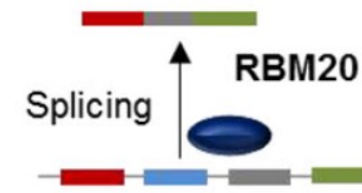


The association of TTNtv with ventricular arrhythmias was independent of the presence of midwall LGE or other covariates.

RBM20 – RNA binding motif protein 20

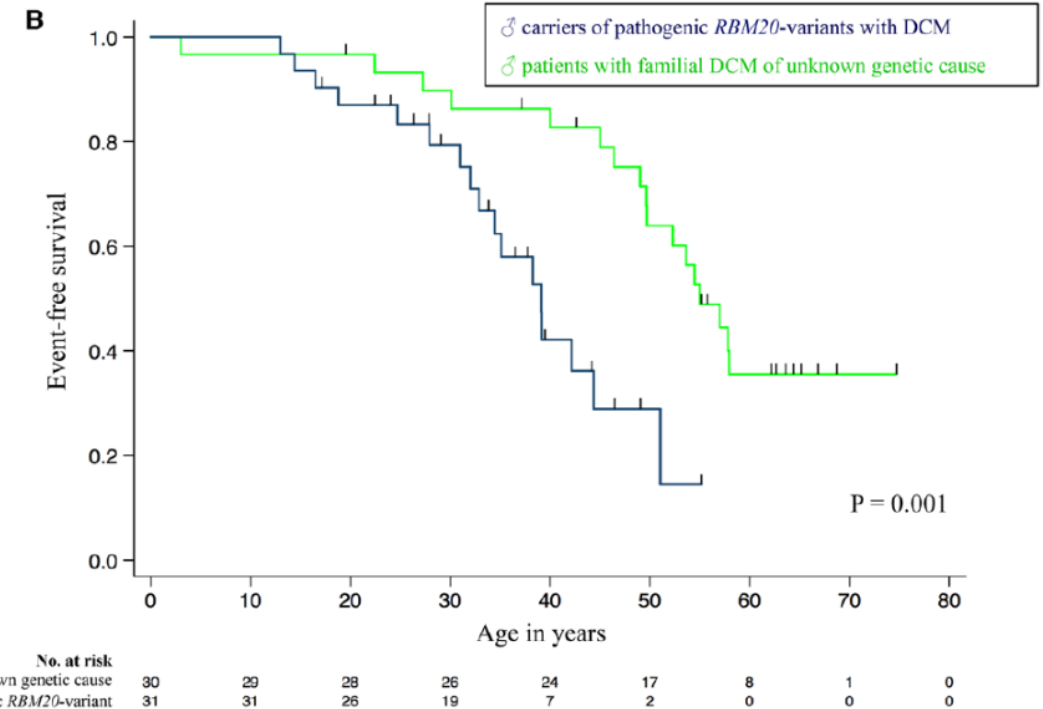
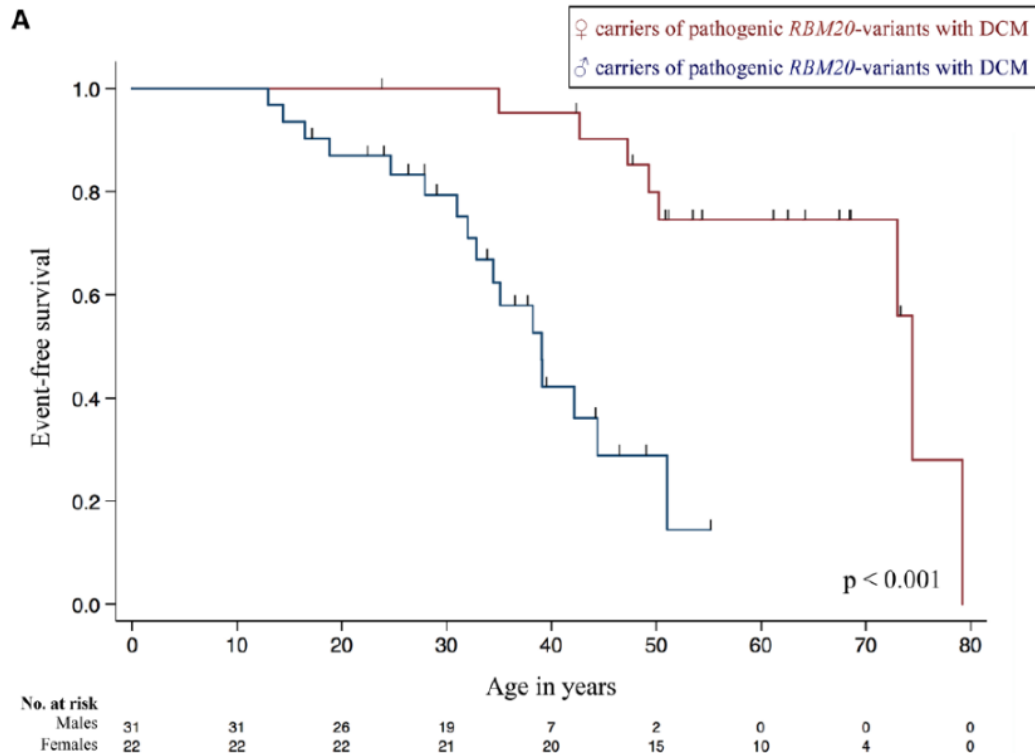
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Pathogenic *RBM20*-Variants Are Associated With a Severe Disease Expression in Male Patients With Dilated Cardiomyopathy



- Frequency: 2%
- Male predominance: 59%
- Conduction disease: 30%
- HTx at younger age

Kayvanpour E, Clin Res Cardiol 2017



TMEM43 – transmembrane protein 43

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Arrhythmogenic Right Ventricular Cardiomyopathy Type 5 Is a Fully Penetrant, Lethal Arrhythmic Disorder Caused by a Missense Mutation in the *TMEM43* Gene

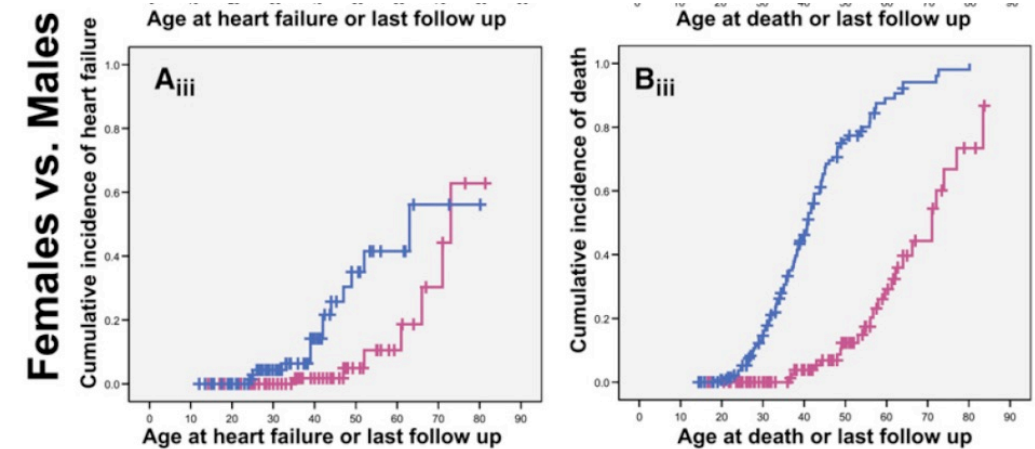
Merner ND, *Am J Hum Genet* 2008

TMEM43 mutations associated with arrhythmogenic right ventricular cardiomyopathy in non-Newfoundland populations

Baskin B, *Hum Genet* 2013

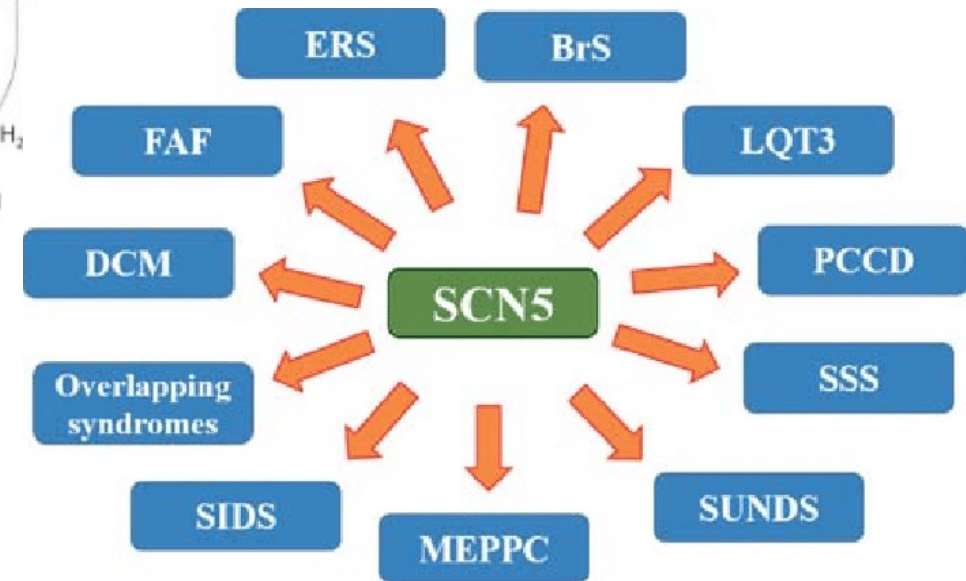
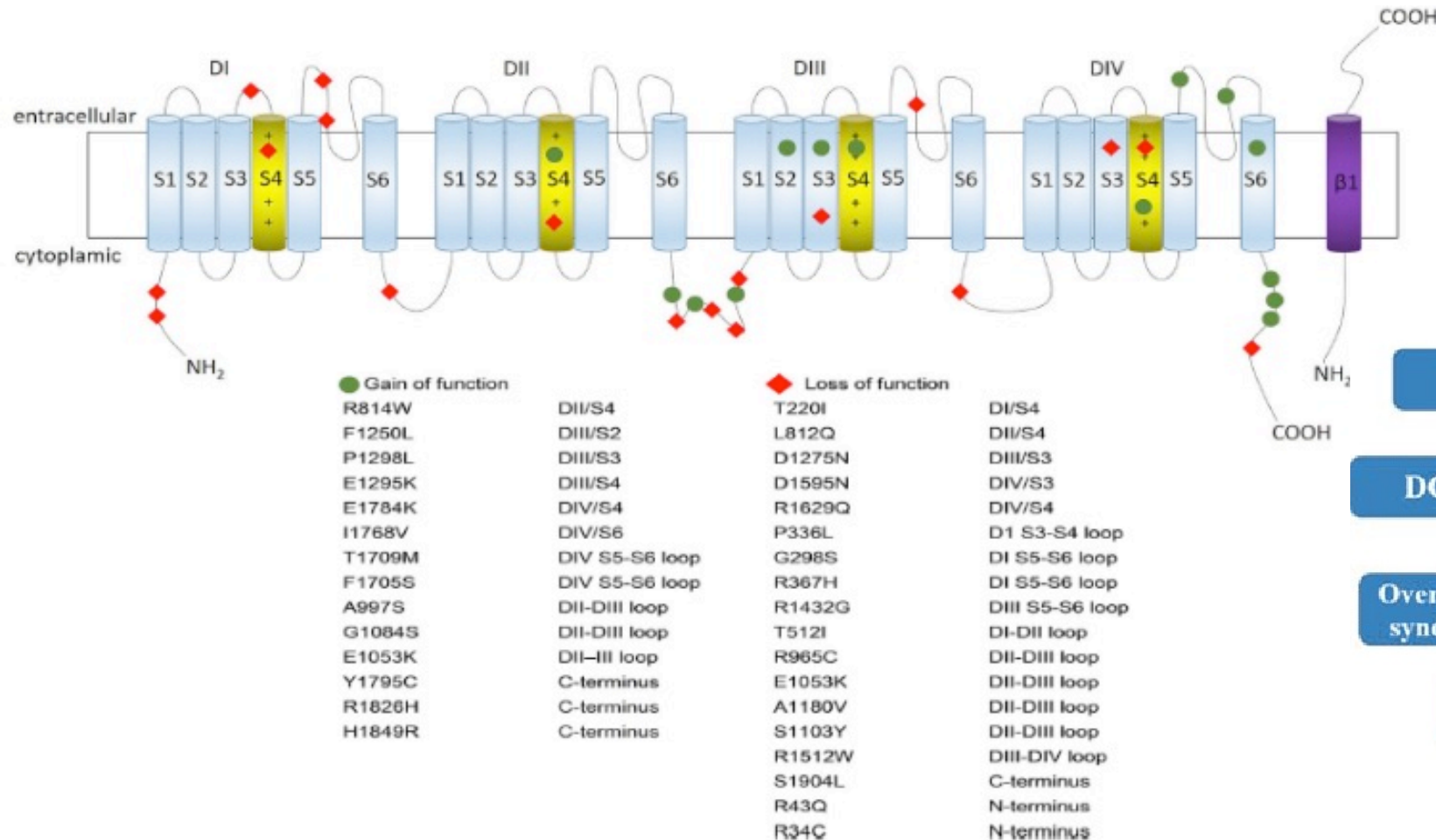
The natural history of a genetic subtype of arrhythmogenic right
ventricular cardiomyopathy caused by a p.S358L mutation in
TMEM43

Hodgkinson KA, *Clin Genet* 2013



11% fulfilled DCM criteria

SCN5A – sodium channel

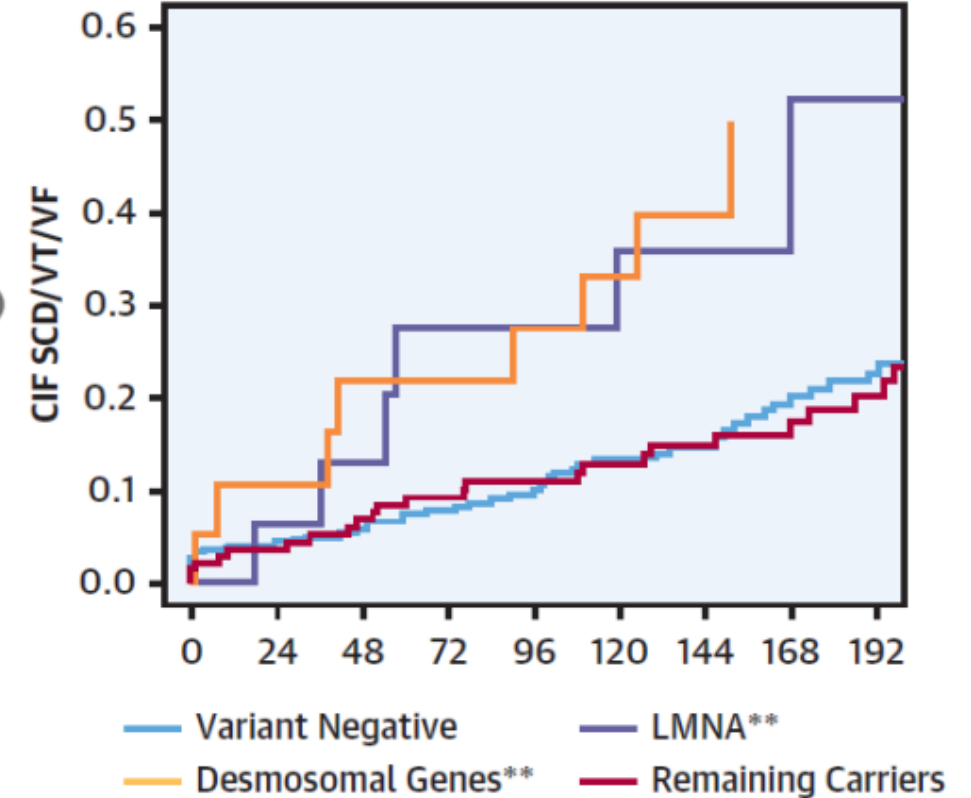
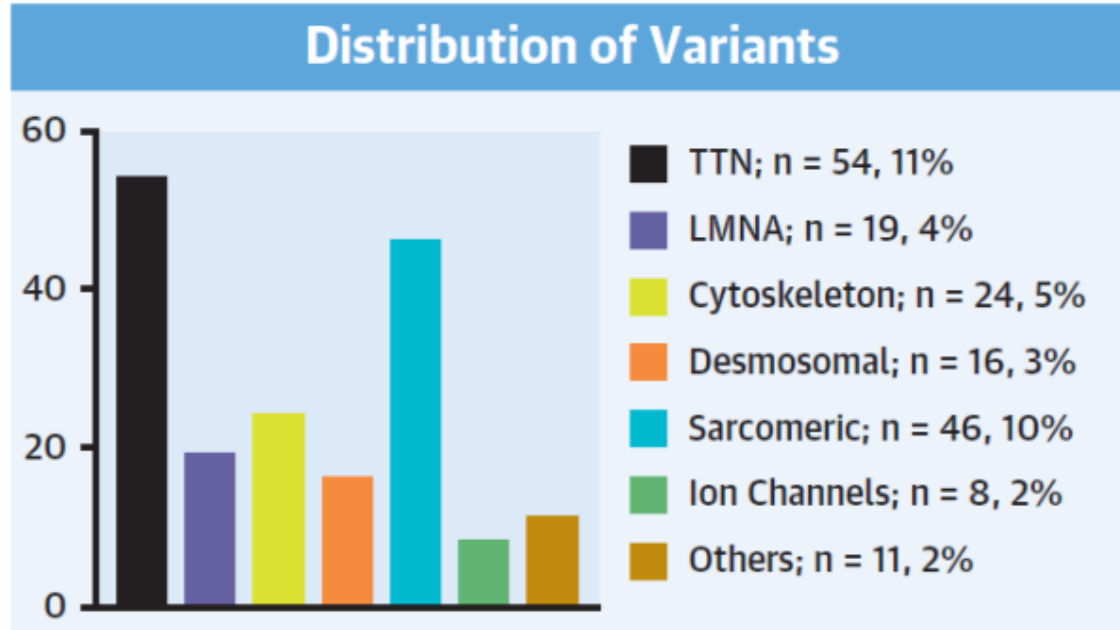


Pérez-Riera AR, J Hum Growth Dev 2016

Desmosomal genes – *DSP*, *DSC2*, *DSG2*, *PKP2*

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Genetic Risk of Arrhythmic Phenotypes in Patients With Dilated Cardiomyopathy



GENETICALLY-GUIDED SUDDEN DEATH RISK STRATIFICATION

Towards precision Medicine

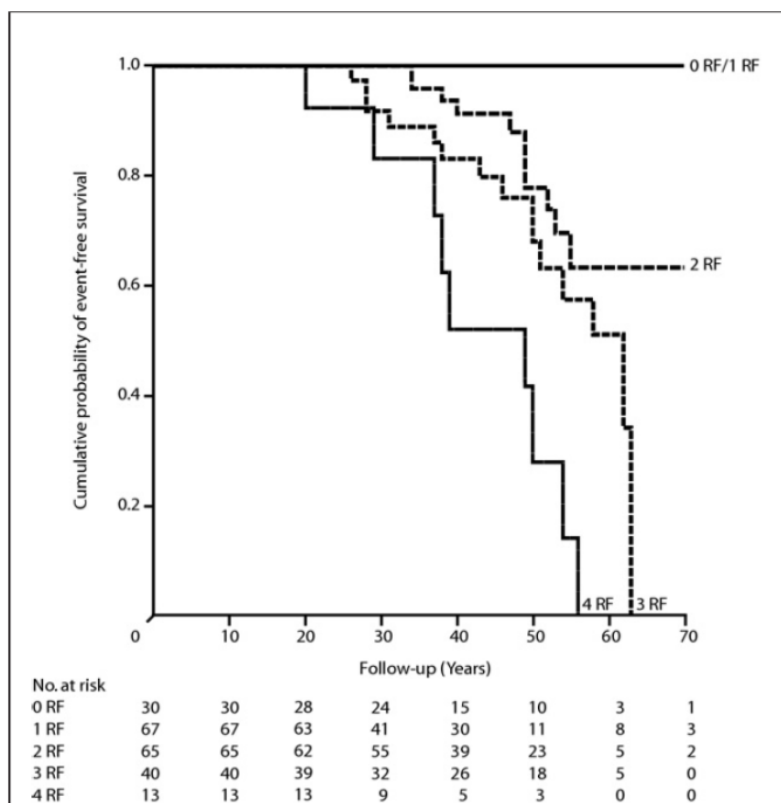


Sudden death risk stratification – *LMNA*

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

A European Cohort Study



An ICD should be considered in patients with DCM and a confirmed disease-causing *LMNA* mutation and clinical risk factors.^d

IIa

B

^dRisk factors in patients with a confirmed *LMNA* mutation: NSVT during ambulatory electrocardiogram monitoring, LVEF <45% at first evaluation, male sex and non-missense mutations (insertion, deletion, truncations or mutations affecting splicing).

Priori S, Europace 2015

In individuals with lamin A/C ACM and two or more of the following: LVEF <45%, NSVT, male sex, an ICD is reasonable.

IIa

B-NR

Towbin JA, Heart Rhythm 2019

Sudden death risk stratification – LMNA

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LMNA-risk VTA calculator

Risk Prediction Score for Life-Threatening Ventricular Tachyarrhythmias in Laminopathies

Sex

☐ Male ☐ Female

Non-missense LMNA mutation

☐ Yes ☐ No

Non-missense mutations include insertions, deletions, truncating mutations or mutations affecting splicing

Atrio-ventricular block

☐ Absent ☐ 1st degree ☐ High degree

Please select the highest degree. 1st degree AV block corresponds to ≥ 0.20 sec PR interval and high degree AV block to type II 2nd degree or 3rd degree (and not type I 2nd degree)

Non-sustained ventricular tachycardia

☐ Yes ☐ No

NSVT corresponds to ≥ 3 consecutive ventricular complexes at a rate ≥ 120 bpm on 24-h ambulatory electrocardiographic monitoring

Left ventricular ejection fraction

%

Left ventricular ejection fraction measurement derived from echocardiogram

Risk of Life-Threatening Ventricular Tachyarrhythmias at 5 years

___ %

[Reset](#)

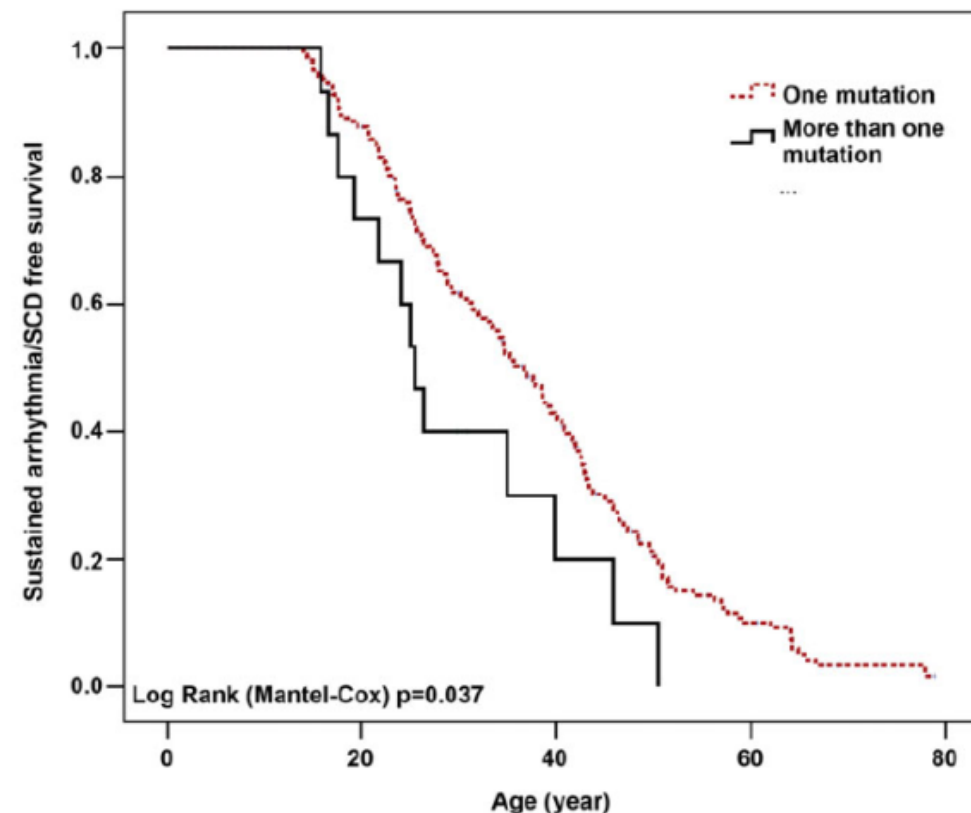
Life-Threatening Ventricular Tachyarrhythmias is defined as 1) sudden cardiac death, 2) appropriate ICD therapy, defined as a shock to terminate a VTA, or 3) other manifestations of hemodynamically unstable VTA

<https://lmna-risk-vta.fr/>

Sudden death risk stratification – desmosomal genes

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Impact of genotype on clinical course
in arrhythmogenic right ventricular
dysplasia/cardiomyopathy-associated
mutation carriers



| | | | | | |
|------------------------|-----|-----|----|----|----|
| One mutation | 215 | 186 | 80 | 14 | -- |
| More than one mutation | 15 | 11 | 2 | -- | -- |

Sudden death risk stratification – desmosomal genes

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COMPLEX GENETIC STATUS RELATED TO WORSE PROGNOSIS

- ❑ **Compound or digenic heterozygosity** up to 25% of cases.
- ❑ **Worse clinical course:**
 - greater penetrance
 - early ventricular arrhythmias
 - higher risk of SD
 - frequent LV involvement
 - increased risk of end-stage HF

ICD implantation may be considered in patients with one or more recognized risk factors for VA in adult patients with a life expectancy > 1 year following detailed clinical assessment that takes into account the lifelong risk of complications and the impact of an ICD on lifestyle, socioeconomic status and psychological health.



IIb

C

during EPS.^{113,114,387,389,395,404–406} **Compound or digenic heterozygosity occurs** in >10% of carriers of the ARVC-causing desmosomal gene mutation and **may be a risk factor for major arrhythmic events and SCD.**⁴⁰⁷ As the studies examining outcomes

Sudden death risk stratification – other genes

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

IIa

B-NR

In individuals with **phospholamban** cardiomyopathy and LVEF <45% or NSVT, an ICD is reasonable.

IIa

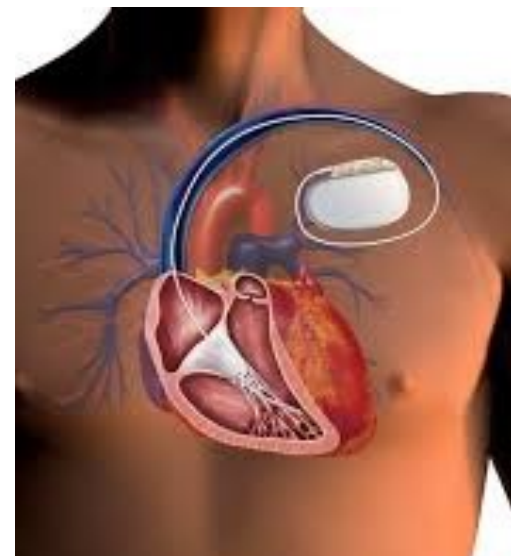
C-LD

In individuals with **FLNC** ACM and an LVEF <45%, an ICD is reasonable.

Sudden death risk stratification – other genes

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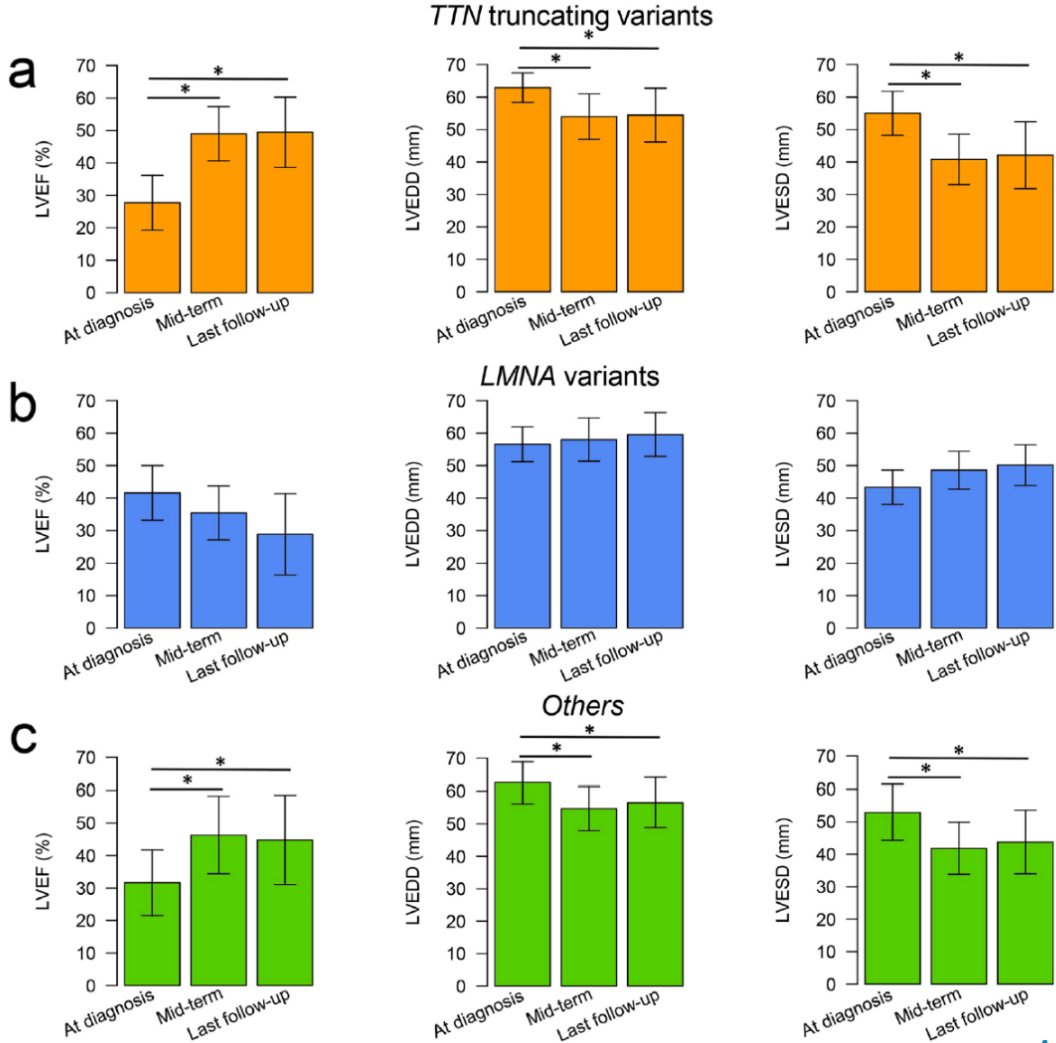
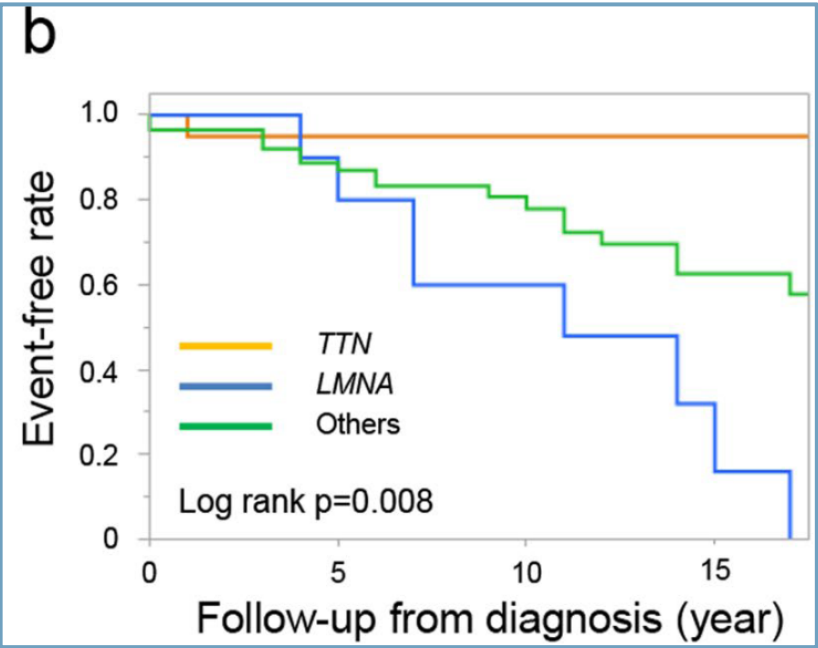
RBM20
TMEM43
SCN5A
DES
TTNtv



Hey TM, Circ Heart Fail 2019
Peters S, Heart Lung Circ 2019
Corden B, JAMA Network Open 2019
Bondue A, Cardiovasc Res 2018
Hodgkinson KA, Clin Genet 2013

INFLUENCE OF GENOTYPE IN REVERSE REMODELLING

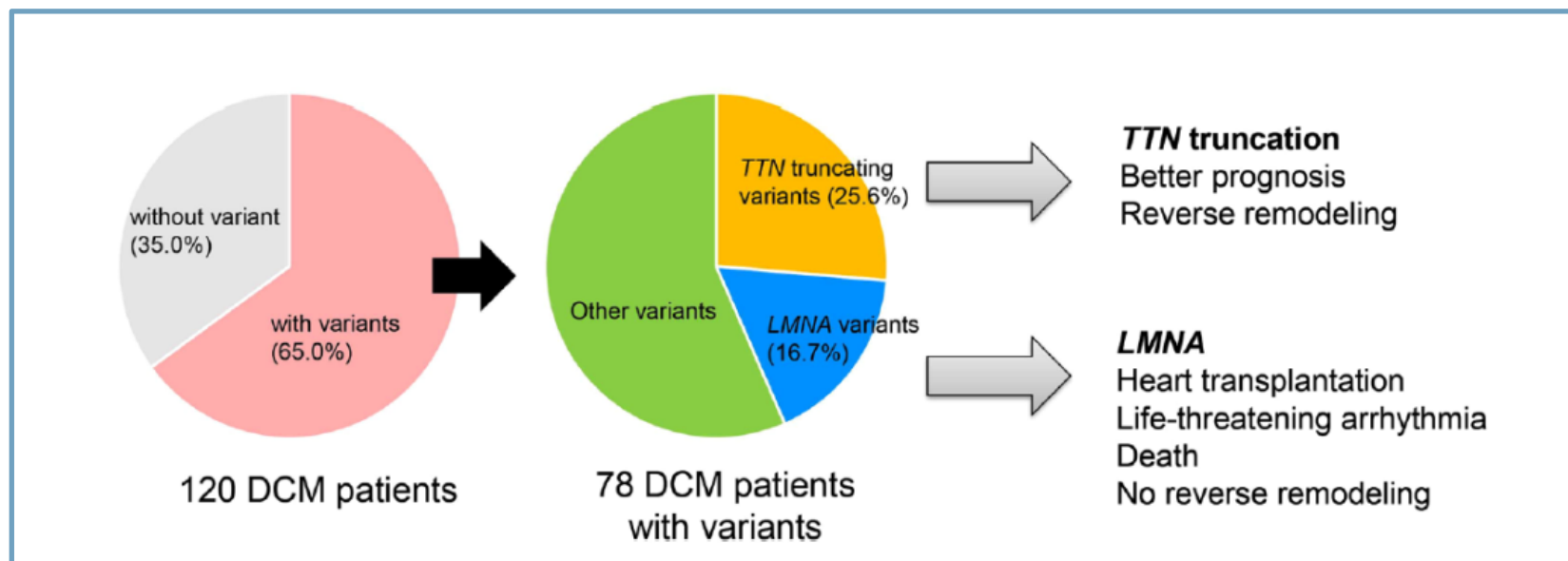
Genetic basis of cardiomyopathy and the genotypes involved in prognosis and left ventricular reverse remodeling



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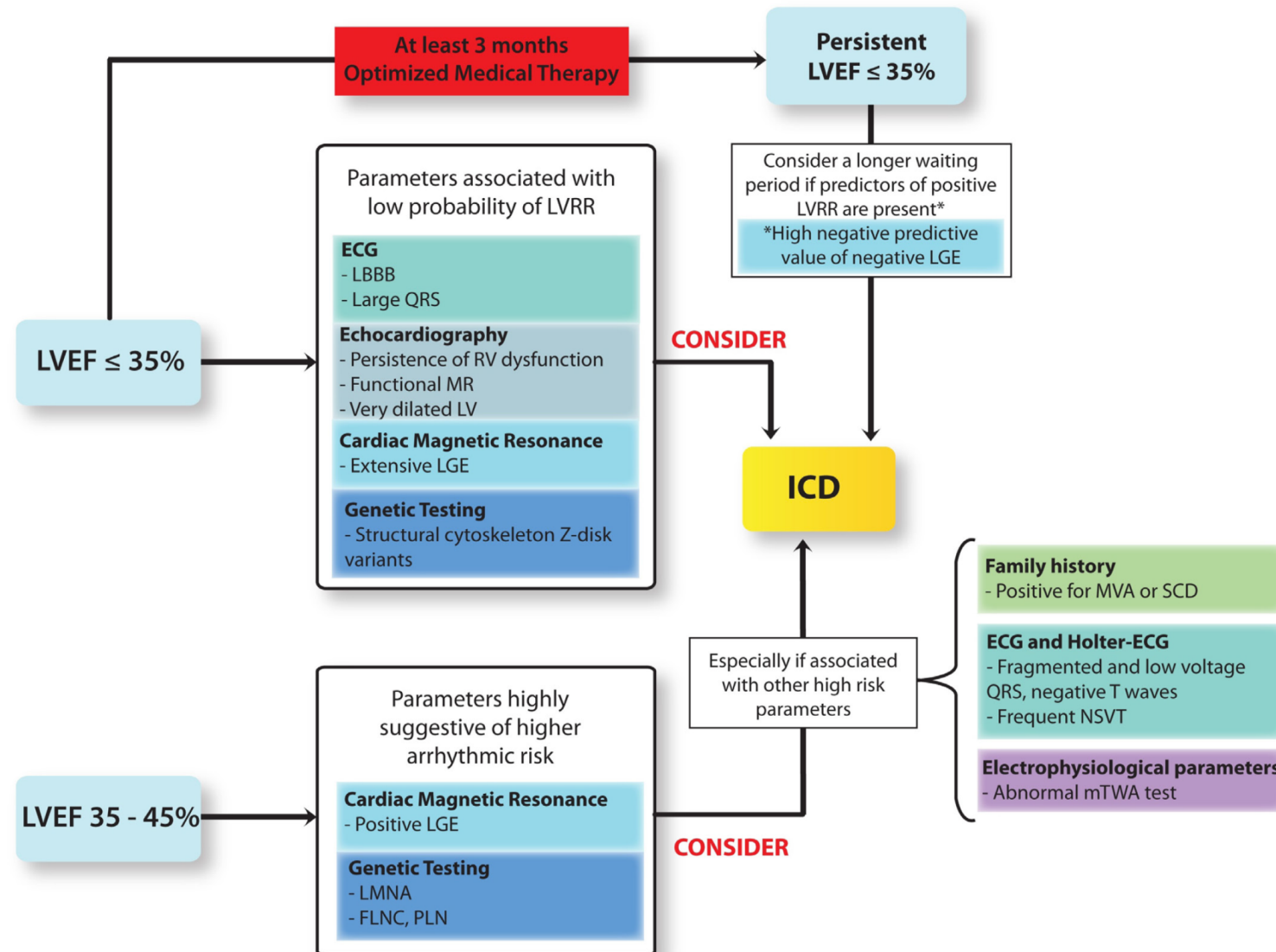
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September, 18th - 19th

Genetic basis of cardiomyopathy
and the genotypes involved in
prognosis and left ventricular
reverse remodeling



PERSONALIZED ARRHYTHMIC RISK ASSESSMENT

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| Risk Classification by Current Guidelines | + LGE Fibrosis Assessment | + Genetic Profiling | → Risk Reclassification |
|---|------------------------------------|--|---|
| <p>EF \leq 35%</p> <p>High risk of SCD</p> <p>ICD is recommended</p> | <p>Absence of Fibrosis on LGE</p> | <p>No familial DCM/SCD</p> | <p>Lower risk of SCD</p> <p>Is ICD necessary?</p> |
| <p>EF > 35% and \leq 49%</p> <p>Low risk of SCD</p> <p>ICD is NOT recommended</p> | <p>Presence of Fibrosis on LGE</p> | <p>And/or Presence of:</p> <ul style="list-style-type: none">– Familial DCM/SCD– Gene Mutations | <p>Higher risk of SCD</p> <p>Is ICD useful?</p> |

ARRHYTHMO "GENETIC" CARDIOMYOPATHIES

OVERLAPPING SYNDROMES WITH SPECIFIC SINGULARITIES DICTATED BY GENETICS

Several shared features

- Family history
- Genetic basis
- Histopathology
- Biventricular involvement
- ECG anomalies
- Arrhythmias

Personalized Medicine

- Sudden death risk stratification
- Therapeutic interventions
 - In asymptomatic pathogenic variant carriers
 - Target therapies
- Familial screening
 - Early detection of the disease
 - Management of incipient disease
 - Counselling (sports, reproductive issues...)

Vielen Dank ありがとう

хвала kiitos

merc

tack dank u

הדוּת 謝謝

thank you

ačiū

obrigado ευχαριστώ

dankie , اراك ش

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ARRHYTHMO "GENETIC" CARDIOMYOPATHIES: PERSONALIZING SUDDEN DEATH RISK STRATIFICATION

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Advances in Heart Failure