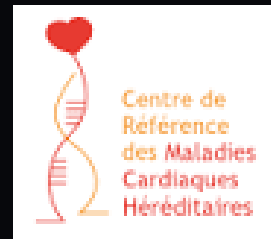


March 2-3

9th Congress Edition
Novotel PARIS Tour Eiffel



What genetic abnormalities should we search for after a sudden death?

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Rythmologie

Centre de référence pour les maladies cardiaques
héréditaires

Disclosure

I have the following potential conflicts of interest to report:

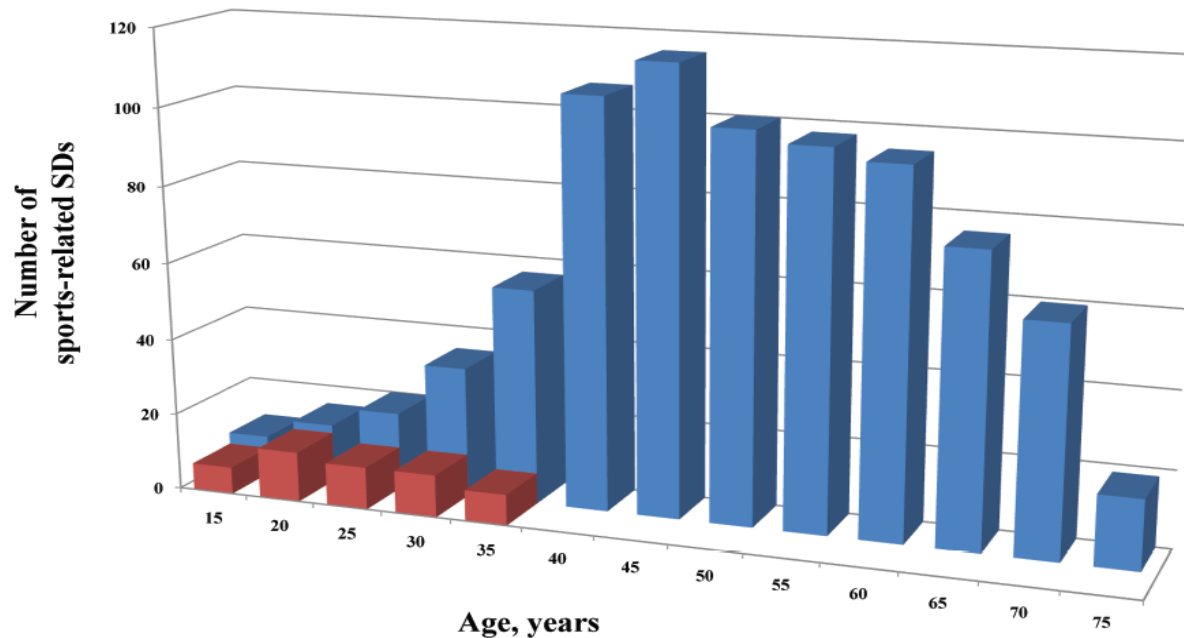
Consulting: Sorin, Medtronic, Boston



Sports-Related Sudden Death in the General Population

Eloi Marijon, Muriel Tafflet, David S. Celermajer, Florence Dumas, Marie-Cécile Perier, Hazrije Mustafic, Jean-François Toussaint, Michel Desnos, Michel Rieu, Nordine Benameur, Jean-Yves Le Heuzey, Jean-Philippe Empana and Xavier Jouven

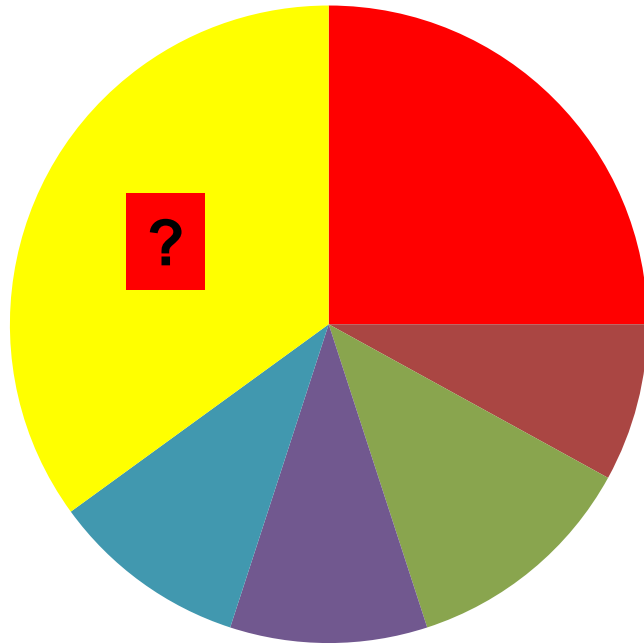
Circulation. 2011;124:672-681; originally published online July 25, 2011;



Distribution by age of sports-related sudden deaths (SDs) in the overall population (blue) and among young competitive athletes (red).

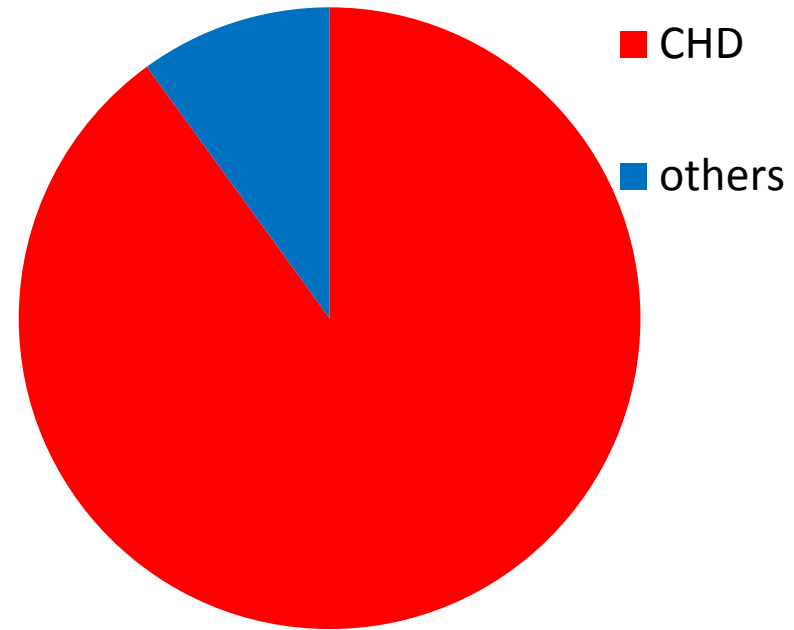
Necropsy study: SCD causes according to age

< 40 Y



- CHD
- myocarditis
- HCM
- ARVC/D
- other
- unknown (USCD)

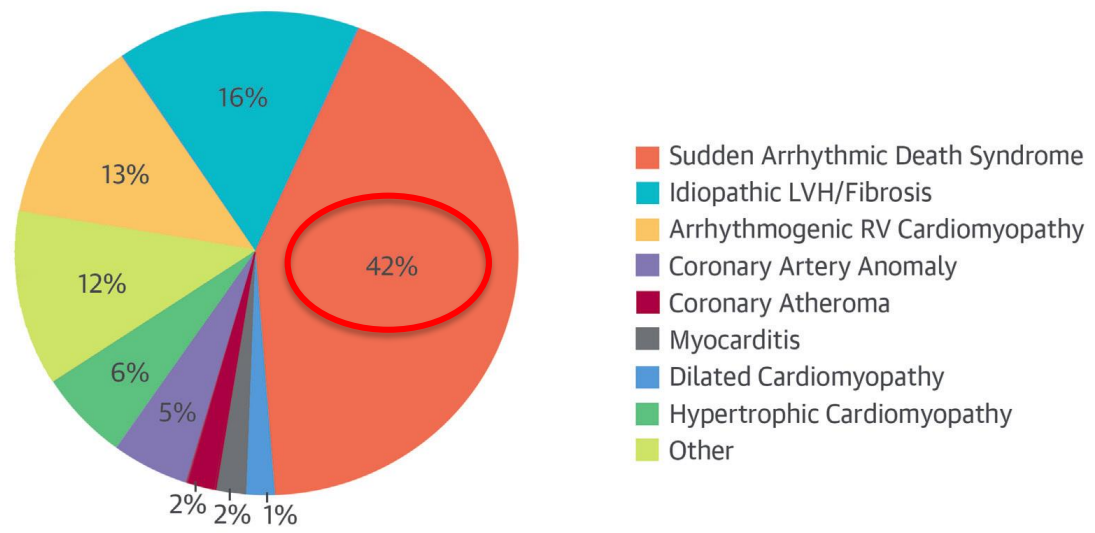
> 40 Y



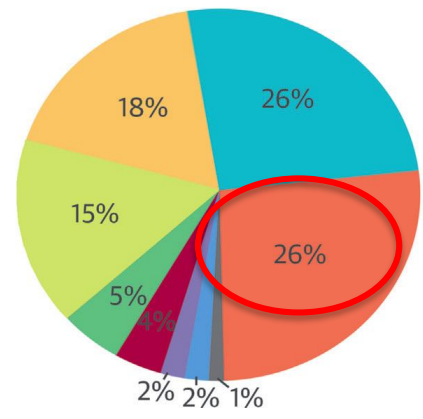
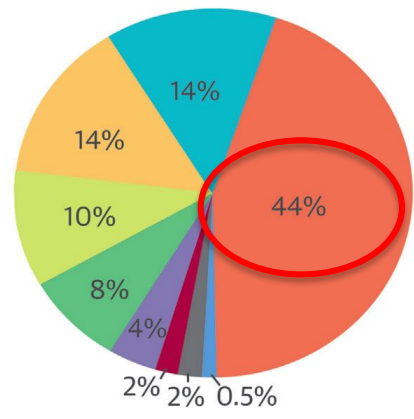
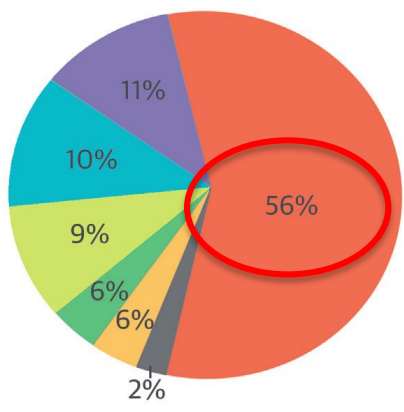
- CHD
- others

SCD in athletes (> 3 hours per week) : UK series

A. Sudden Death in Overall Population
 (n = 357)



B. Sudden Death <18 Years (n = 79) **C. Sudden Death 18-35 Years** (n = 179) **D. Sudden Death >35 Years** (n = 99)



Risk of cardiovascular disease in family members of young sudden cardiac death victims

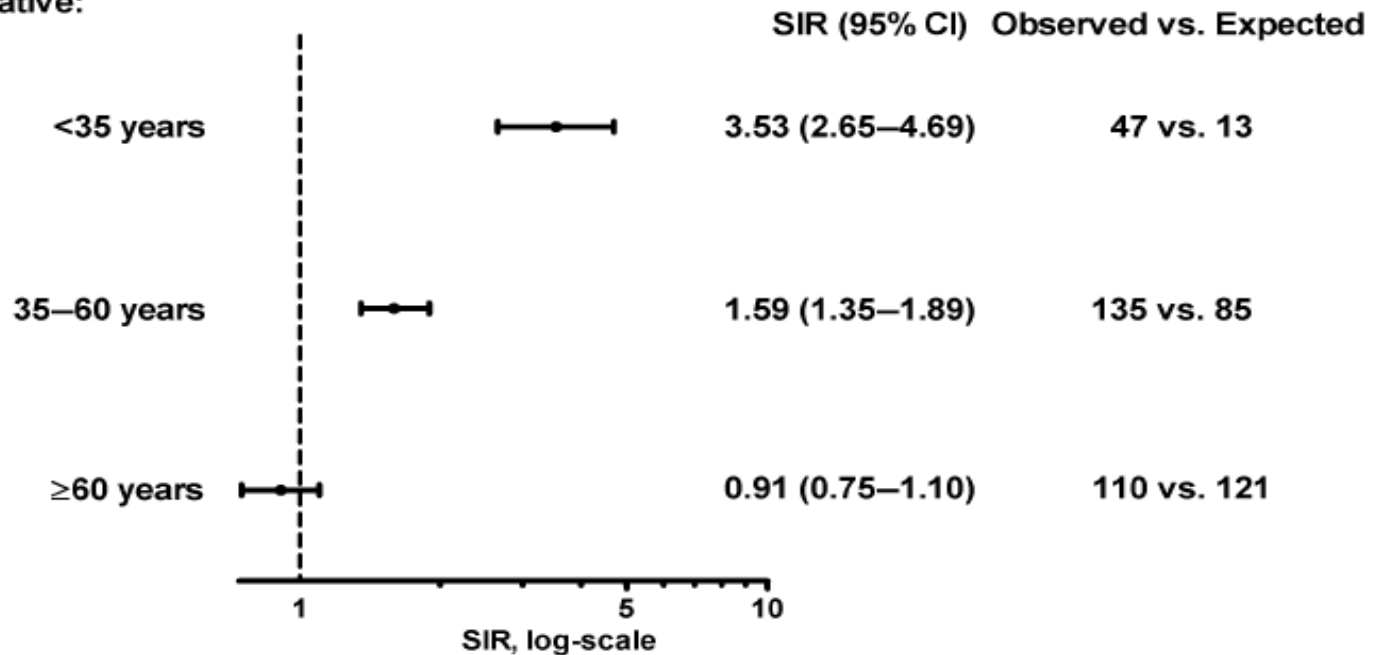
Denmark 2000-2006

n=470 SD

Screening and follow-up of 3073 relatives

Risk of any cardiovascular disease in relatives of young first degree relatives

Age of relative:



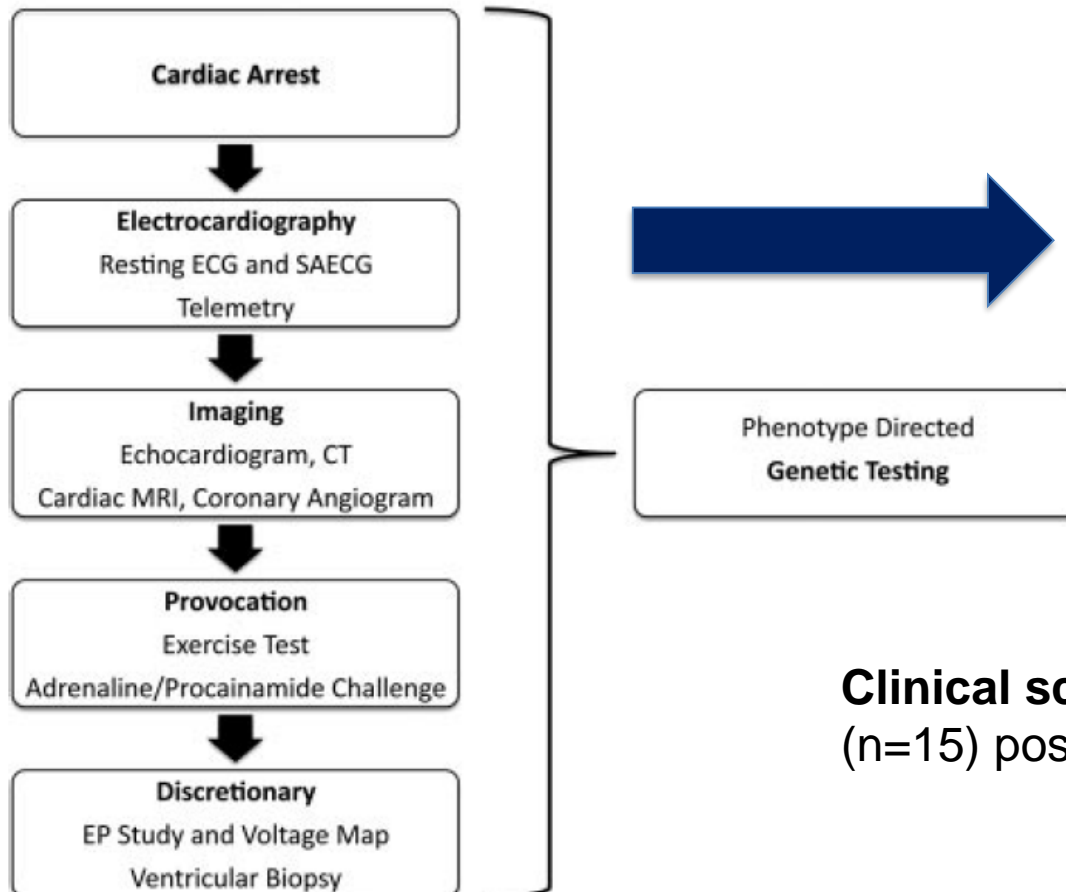
Risk increases were even more dramatic for first-degree relatives and persons <35 years of age

Importance of a complete clinical screening in patients with USCD (Casper study)

Unexplained SCD:

normal cardiac function on echocardiogram, no evidence of coronary artery disease, and a normal ECG

63 USCD (mean age=43 Y)



Diagnosis in 56% of patients

LQTs: 8 patients

CPVT: 8 patients

ARVC: 6 patients

Early repolarisation: 5 patients

Coronary Spasm: 4 patients

Brugada: 3 patients

Myocarditis: 1 patient

Clinical screening of 64 relatives → 24% (n=15) positives

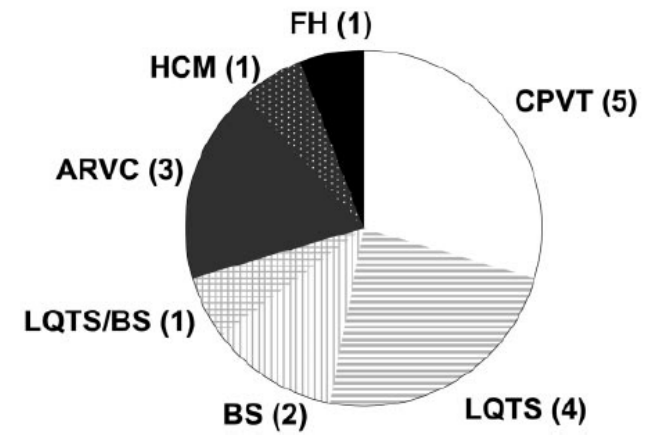
Diagnostic yield of clinical and genetic screening of relatives

Relatives of USCD victim (negative of no post mortem analysis) < 40 Y = 43 families

- ECG
- Exercise test
- TEE
- Lipid analysis
- Ajmaline challenge if suspect ECG
- MRI if suspect ARVC

+

Targeted genetic testing if positive phenotype



Identification of cause in 17/43 families (40%)

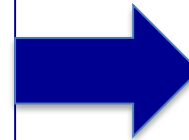
Exercise test ++

333 1st and 2nd-degree relatives screened

Diagnostic yield of clinical and genetic screening of relatives (CASPER)

**398 Relatives of 212 USCD victim
(negative of no post mortem analysis)**

- ECG
- Exercise test
- TEE
- SA ECG
- Ajmaline challenge if suspect BrS ECG/ER or nocturnal SCD
- epinephrine challenge if non diagnosis exercise test and exercise induced SCD
- MRI if suspect cardiomyopathy



**Identification of possible cause in
120 relatives (30%)
Including 17% with
probable/definite diagnosis**

**Mutation in only 5% relatives
(targeted testing)**

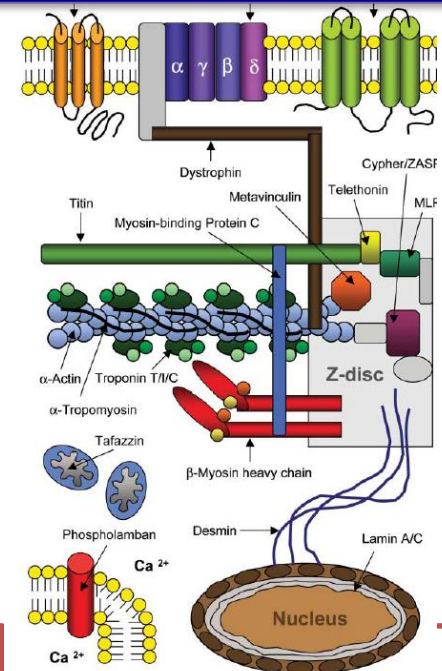
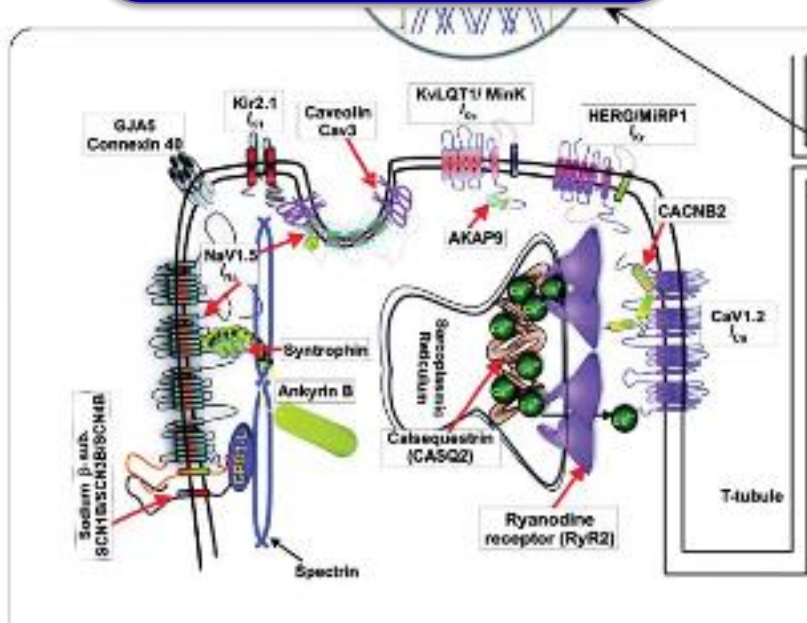


**Targeted genetic testing if positive phenotype
Bs: *SCN5A*, ARVC: *PKP2-DSP*, CPVT: *RYR2*, LQT: 5 genes**

CHANNELOPATHY

CARDIOMYOPATHY

Sarcomere,
cytoskeleton, nuclei



- Brugada: > 10 genes , modifier effect ++ (PGS 25%)
- LQT : >10 genes, PGS 60-70%
- SQT : >6 genes
- AVB, SSS :>5 genes
- CPVT : 4 genes
- ...

- HCM >15 genes
- DCM > 30 gènes
- ARVC 10 genes
- LMNA
- LVNC >7 genes
- RCM... >10 genes

Large genetic heterogeneity++
Not all causative genes are known

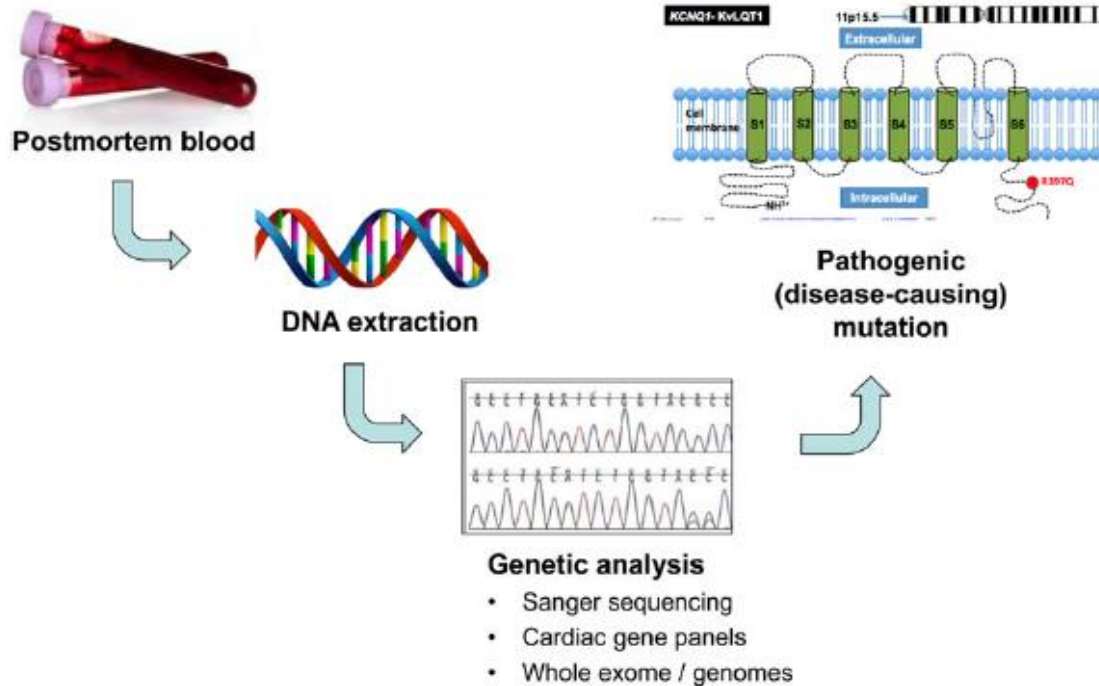
Molecular autopsy

=Post mortem genetic screening

*Analysis impossible from hair
Difficult from paraffin blocks*

Blood samples before death

Necropsy tissue (liver++)

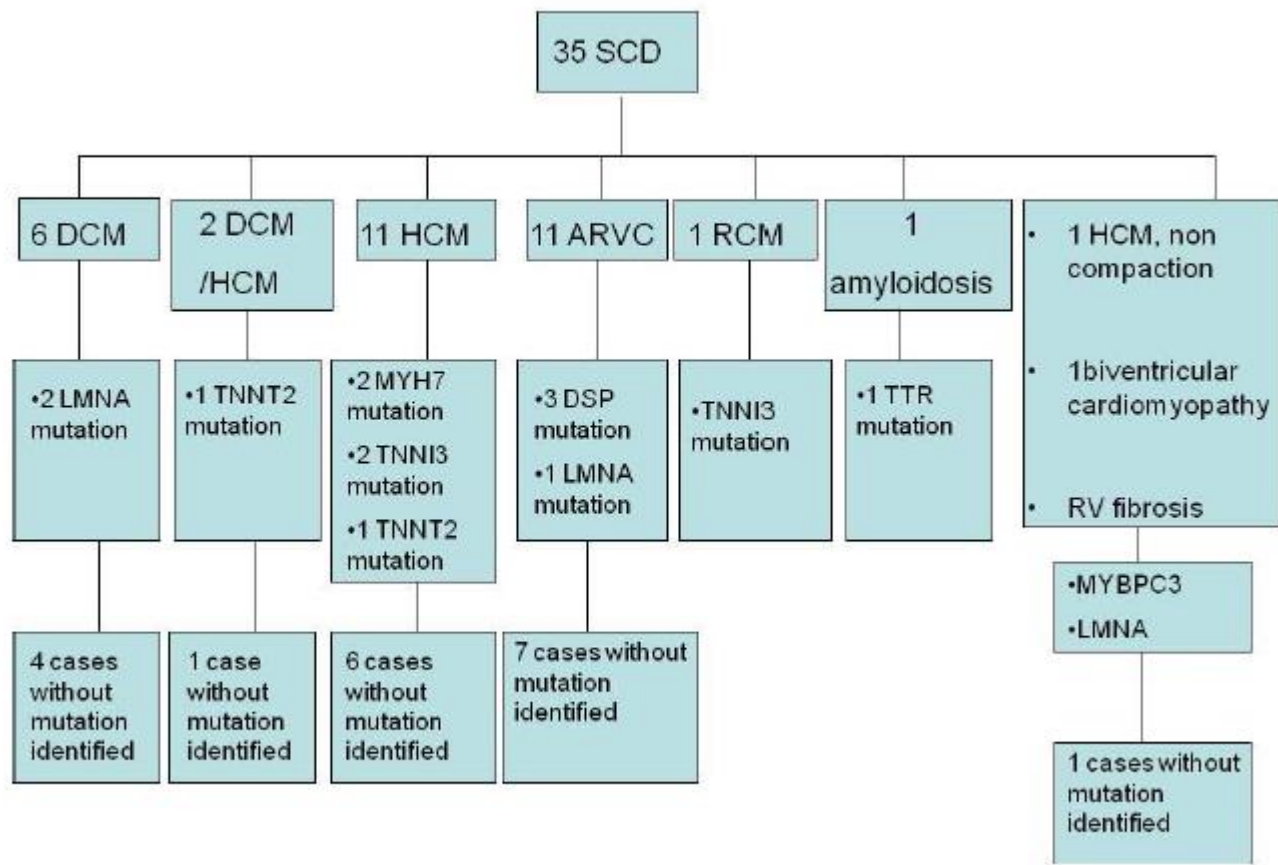


Legal issues

- Rightholder consent ++
- Result to right holder

Targeted post mortem genetic screening in cardiomyopathy: Case series from GHPS

Global results in the cohort of patients



→ 15 mutations in 15 patients were identified out of 35 patients (43%)

Post mortem genetic screening in USCD

Post mortem screening of *KCNQ1*, *KNCH2*, *SCN5A*, *RYR2* in unexplained SCD → mutation yield: 15% of cases

Table 3 Current 4-gene molecular autopsy

Gene Name	Encoded protein	Disease	% of disease	% of SADS ^a
<i>KCNQ1</i>	<i>I_{Ks}</i> K ⁺ channel α-subunit	LQTS1	35–40	10–15
<i>KCNH2</i>	<i>I_{Kr}</i> K ⁺ channel α-subunit	LQTS2	30–35	1–5
<i>SCN5A</i>	<i>I_{Na}</i> Na ⁺ channel α-subunit	LQTS3 BrS	5–10 15–25	<1 <1
<i>RYR2</i>	Ryanodine receptor	CPVT1	60–65	10–15

32 patients with exercise related USCD

KCNQ1, *KNCH2*, *SCN5A*, *RYR2*

Mutation in 11

Negative in 21

100 genes (exome)

Mutation in 3 cases:
CALM2 (TVC)
PKP2 (DVDA)
GVUS in 7 cases

Post mortem genetic screening in USCD

Genetic screening identifies a high proportion of mutations in patients with idiopathic ventricular fibrillation and sudden cardiac death

Vincent Probst (1), Solena Le Scouarnec (2), Florence Kyndt (3), Jean-Jacques Schott (2), Jean-Baptiste Gourraud (1), Frederic Sacher (4), Philippe Mabo (5), Matilde Karakachoff (2), Stéphanie Bonnaud (2), Jade Violleau (2), Eloi Marijon (6), Florence Dumas (6), Alain Cariou (6), Estelle Baron (2), Pierre Lindenbaum (2), Xavier Jouven (6), Richard Redon (2)

(1) CHU Nantes, Nantes, France – (2) CHU Nantes, Institut du Thorax, INSERM UMR 1087, Nantes, France – (3) CHU Nantes, Institut du Thorax, Nantes, France – (4) CHU Bordeaux, Bordeaux, France – (5) CHU Rennes, Rennes, France – (6) Paris Sudden Death Expertise Center INSERM U970, Paris, France

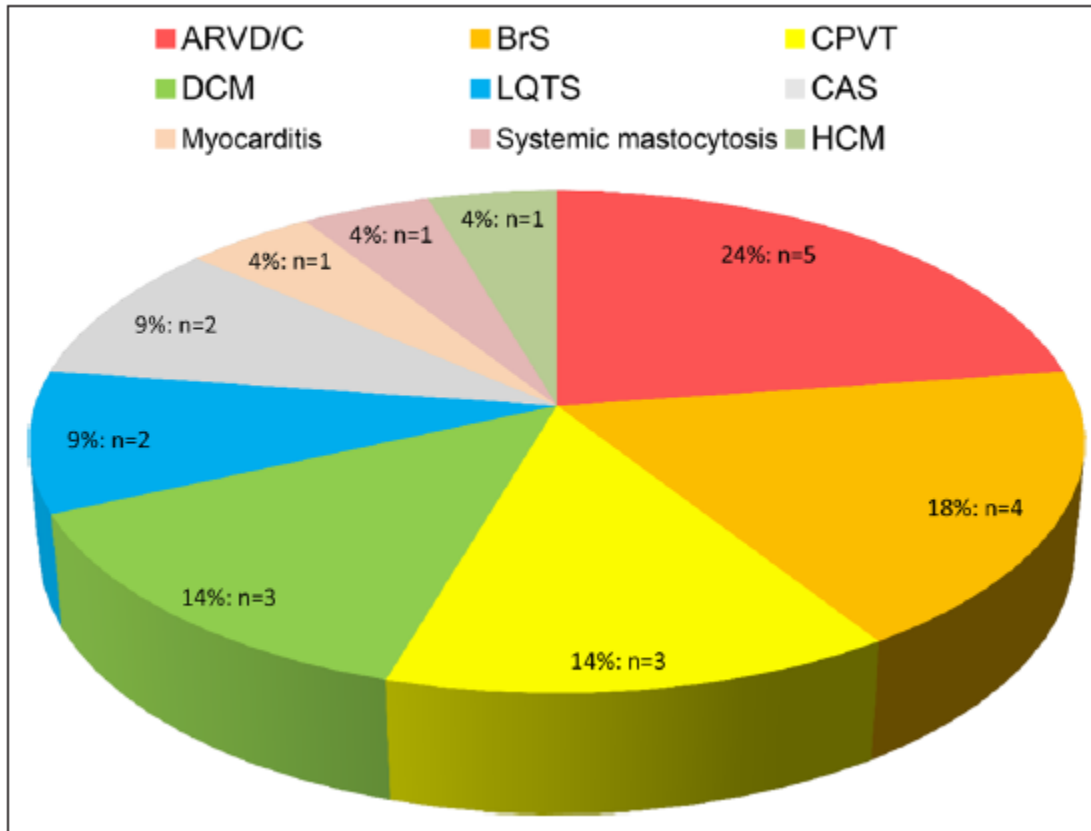
Abstract 0185-Table

	Cohort 1 (n=75)	Cohort 2 (n=99)
SCN5A	7 (9%)	0
HCM and DCM genes	22 (29%)	17 (17%)
ARVD/C genes	4 (5%)	3 (3%)
LQTS genes (excluding SCN5A)	4 (5%)	4 (4%)
CPVT genes (excluding SCN5A)	7 (9%)	3 (3%)
BRS genes	3 (4%)	1 (1%)
Other	3 (4%)	2 (2%)

Limit of large genetic screening: interpretation of rare missense variants?
→ EXAC database, prediction software, Study of familial segregation +++

Long term follow-up after USCD

79 patients with USCD → median FU = 10 Y
 Identification of cause in 20% of cases



<i>AKAP9</i>	<i>CASQ2</i>	<i>DSP</i>	<i>KCNE2</i>
<i>KCNJ8</i>	<i>RYR2</i>	<i>SNTA1</i>	<i>ANK2</i>
<i>CAV3</i>	<i>GPD1L</i>	<i>KCNE3</i>	<i>KCNQ1</i>
<i>SCN1B</i>	<i>TGFB3</i>	<i>CACNA1C</i>	<i>DES</i>
<i>HCN4</i>	<i>KCNH2</i>	<i>LMNA</i>	<i>SCN3B</i>
<i>TMEM43</i>	<i>DSC2</i>	<i>JUP</i>	<i>KCNJ2</i>
<i>DSG2</i>	<i>KCNE1</i>	<i>KCNJ5</i>	<i>PLN</i>
<i>CACNA2D1</i>	<i>PKP2</i>	<i>SCN4B</i>	<i>CACNB2</i>
<i>SCN5A</i>	<i>DPP6</i> (VF risk haplotype only)		



Pathogenic mutation in 15% (n=12)
 GVUS in 13 patients

Familial screening and long term follow-up ++

Index case with SCD

In all cases, long term follow-up of probands and relatives ++

