

# 2023 ESC Guidelines for the management of cardiomyopathies Supplementary data

# Developed by the task force on the management of cardiomyopathies of the European Society of Cardiology (ESC)

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#### **Patient Forum**

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## Abbreviations and acronyms

3D	Three-dimensional
AFD	Anderson–Fabry disease
AlcCM	Alcoholic cardiomyopathy
AR	Autosomal recessive
ARVC	Arrhythmogenic right ventricular cardiomyopathy
ASA	Alcohol septal ablation
ATTR	Transthyretin amyloidosis
ATTRv	Hereditary transthyretin amyloidosis
ATTRwt	Wild-type OR Acquired transthyretin amyloidosis
AV	Atrioventricular
BNP	Brain natriuretic peptide
CCM	Cancer therapy-induced cardiomyopathy
CGU	Cardiogenetic unit
CMP	Cardiomyopathy
CMR	Cardiac magnetic resonance
CMU	Cardiomyopathy unit
СТ	Computed tomography
CV	Cardiovascular
DCM	Dilated cardiomyopathy
DIP	Drug-induced phospholipidosis
FCG	Electrocardiogram
ECV	Extracellular volume
FF	Election fraction
eGFR	Estimated glomerular filtration rate
FORP	EURObservational Research Programme
EU	European Union
FLC	Free light chain
HCM	Hypertrophic cardiomyopathy
HE	Heart failure
нн	Hereditary baemochromatosis
HRR	Heart rate reserve
	Inherited cardiac condition
	Implantable cardioverter defibrillator
IAD	l eft anterior descending
LGE	Late gadolinium enhancement
	Left ventricular ejection fraction
	Left ventricular bypertrophy
	Left ventricular hypertrophy
	Left ventricular outflow tract obstruction
	Left ventricular systolic dysfunction
MRI	Magnetic resonance imaging
	National amyloid centre
NCS	Non-cardiac surgery
	Non-sustained ventricular tachycardia
	N terminal pro brain patriuratic poptida
ΝΥΠΖ	New York Heart Association
DET	Positron omission tomography
	Pandiatric Cardiamyapathy Panistry
	Paripartum cardiomyopathy Registry
	renpartum cardiomyopathy

RCM	Restrictive cardiomyopathy
RPE	Rate of perceived exertion
SCD	Sudden cardiac death
SRT	Septal reduction therapy
SVD	Small vessel disease
TTNtv	Titin gene truncating variants
VF	Ventricular fibrillation
VO <sub>2</sub> max	Maximum oxygen consumption
VT	Ventricular tachycardia
VUS	Variant of uncertain significance

# 1. Epidemiology

### 1.1. General epidemiological data

Cardiomyopathies are in many cases inherited cardiac diseases with an identifiable or suspected genetic cause, with variable and often incomplete penetrance throughout life.<sup>1</sup> Geographical differences in the distribution of genetic variants may influence the estimated prevalence of different cardiomyopathy phenotypes in different populations, ethnicities, regions, and countries. The complexity of diagnostic criteria for some conditions, such as arrhythmogenic right ventricular cardiomyopathy (ARVC), limits the evaluation of the true prevalence of the disease in the general population. Moreover, epidemiological data are often not collected systematically at the population level, particularly in children.

Hypertrophic cardiomyopathy (HCM) is the most common cardiomyopathy, with an estimated prevalence taken from a pooled analysis of eight studies of 0.2% (95% confidence interval [CI], 1.44–2.71) or 1/460 population.<sup>2–9</sup> Data from cardiac magnetic resonance (CMR) analysis of a large adult cohort (>45 years of age) suggested that the prevalence may be higher than those derived from early echocardiography-based studies, with up to 1.4% prevalence, or 1/76 population.<sup>10</sup> In contrast, data from the UK Biobank reported a prevalence of 0.11% of left ventricular hypertrophy (LVH) above 15 mm in a population where arterial hypertension and aortic stenosis were excluded (0.22% in men and 0.04% in women).<sup>11</sup> Of note, (likely and definite) pathogenic gene variants have been detected in up to 60% of HCM.<sup>12</sup>

The prevalence of dilated cardiomyopathy (DCM) has been traditionally estimated at 0.036% (95% Cl, 0.023–0.050) or 1/2740 based on one large population study.<sup>13</sup> More recent data suggest that the prevalence of DCM may be almost 10 times higher, based on indirect assumptions of the prevalence of genetic variants associated with the disease in general populations<sup>14</sup> and with less stringent diagnostic criteria.<sup>15</sup> Likely and/or definitely pathogenic gene variants occur in up to 40% of DCM,<sup>16</sup> and the prevalence of (likely) pathogenic genetic variants is also over 10% in non-familial DCM.

The true prevalence of ARVC based on the pooled analysis of three relatively large series is estimated at 0.078% (95% CI, 0.077–0.078) or 1/1290.<sup>17–19</sup> However, substantial geographical differences in the estimated population prevalence have been reported.

Data from the EURObservational Research Programme (EORP) Cardiomyopathy and Myocarditis registry with consecutive inclusion of cardiomyopathy cases provide information on the relative burden of cardiomyopathies seen in specialized clinics across Europe.<sup>20</sup> Hypertrophic cardiomyopathy burden in inherited cardiac disease units is 1.4 times higher than DCM, 12.2 times higher than classical ARVC, and 26.3 times higher than restrictive cardiomyopathy (RCM). Overall, the annual incidence of paediatric cardiomyopathies is estimated to be about 1 per 100 000 children, with the highest incidence in the first year of life, based on population studies in Finland, the United States of America, and Australia.<sup>21–23</sup>

Paediatric cardiomyopathies form a diverse group of uncommon diseases, including genetic causes, metabolic or neuromuscular diseases, toxic causes, and infections. Dilated cardiomyopathy and HCM make up  $\sim$ 50% and 25% of all children with cardiomyopathy, respectively, whereas ARVC rarely presents in the first decade of life.<sup>21,23</sup> Restrictive cardiomyopathy is rare in childhood. The annual incidence of DCM is 0.34–0.58 per 100 000 children, with the highest incidence in infants (<1 year of age) of 3.80–4.58 per 100 000. $^{21-24}$  The annual incidence is higher in boys than in girls (0.66 vs. 0.47) and in Blacks than in Whites (0.98 vs. 0.46).<sup>25</sup> Data from the Paediatric Cardiomyopathy Registry (PMCR) suggest that 66% of children have idiopathic disease. Of known causes, myocarditis was the most common (46%), followed by neuromuscular disease (26%), familial DCM (14%), metabolic disease (11%), and malformation syndrome (3%).<sup>25</sup> In the current era, clinical genetic testing detects a genetic cause in about one-third of paediatric DCM cases, including a relatively high percentage of *de novo* mutations.<sup>26,27</sup> The overall annual incidence of HCM is 0.24–0.47 per 100 000 children.<sup>21–23</sup> A large peak in incidence in the first year of life is mainly caused by HCM due to metabolic disorders and malformation syndromes. Another smaller peak in the second decade is mainly due to sarcomeric diseases.<sup>28</sup> Arrhythmogenic right ventricular cardiomyopathy is very rare in early childhood; however, large ARVC registry data show that ~15% of ARVC patients present with paediatric onset disease with mean  $15.3 \pm 2.4$  years of age. Paediatric ARVC patients more often present with sudden cardiac death (SCD).<sup>29,30</sup> Restrictive cardiomyopathy is very rare, with an annual incidence of 0.03–0.04 per 100 000 children.<sup>21,23</sup> Age at diagnosis ranges from infancy to adolescence; however, in contrast with other paediatric cardiomyopathies, RCM shows no peak in infancy and the incidence increases with age.<sup>31</sup>

#### **1.3. Special populations**

Direct causes of cardiomyopathies include pathogenic gene variants, toxins (e.g. alcohol, chemotherapy), autoimmunity, storage diseases (e.g. Fabry disease), infiltrative diseases (e.g. amyloidosis), various stressors (e.g. pregnancy, tachyarrhythmias), or infections.

Several forms of DCM previously considered secondary to external factors were recently proved to have genetic contributors, including alcoholic cardiomyopathy (AlcCM), cancer therapy-induced cardiomyopathy (CCM), and peripartum cardiomyopathy (PPCM). Titin gene truncating variants (*TTN*tv) represent a prevalent genetic predisposition for AlcCM (present in 13.5% vs. 2.9% in controls), as they are associated with a worse left ventricular ejection fraction (LVEF) in DCM patients who consume alcohol above recommended levels.<sup>32</sup> Unrecognized rare variants in cardiomyopathy-associated genes, particularly *TTN*tv (in 7.5% of cases), appeared to be associated with an increased risk of CCM in children and adults, and adverse cardiac events in adults.<sup>33</sup> Rare truncating variants in eight genes were found in 15% of women with PPCM, and two-thirds were *TTN*tv (10% of patients vs. 1.4% of the reference population).<sup>34,35</sup> Additionally, other truncating variants were identified in the genes *DSP* (1%), *FLNC* (1%), and *BAG3* (0.2%).<sup>35</sup>

Hypertrophic cardiomyopathy can represent the common phenotype for several phenocopies and genocopies.<sup>12</sup> Anderson–Fabry disease was found in 0.94% of males and 0.90% of females in cardiac screening programmes for LVH and HCM.<sup>36</sup> Screening with bone scintigraphy found a high prevalence of transthyretin cardiac amyloidosis in specific populations: 11.4% in aortic stenosis, 4.8% in heart failure with preserved ejection fraction, 12.9% in LVH/HCM, and 2.6% in carpal tunnel syndrome (more if it is bilateral).<sup>37</sup>

The concept that cardiomyopathies are more prevalent than initially thought has generally been accepted by the scientific community, but more work needs to be done to ensure that the classification and detection of these rarer cardiac conditions are carried out in a systematic manner.

## 2. The patient pathway

#### 2.1. Genetic testing

#### 2.1.1. Variant interpretation

If a variant is identified in a gene known to be associated with the presenting phenotype, the laboratory will classify it according to a 5-tier system. Where there is sufficient evidence to support causation, the variant will be classified as pathogenic or likely pathogenic (P/LP). Conversely, where there is sufficient evidence to support a variant as not being the cause of disease, it is classified as benign or likely benign. Where there is conflicting or insufficient evidence, this will be considered a variant of uncertain significance (VUS). Benign/likely benign variants may not be included in the report. Pathogenic and benign classifications convey high confidence regarding the causal significance of a variant. 'Likely' represents >90% likelihood that a variant is pathogenic or benign, usually sufficient confidence to act on a result. Often there are insufficient data to achieve 90% confidence for or against pathogenicity, and variants are reported as uncertain. There are many rare variants in the genomes of healthy individuals, and many of these, even in well-known disease-associated genes, are innocent bystanders and not causative for the observed disease. VUS should therefore not be used for cascade testing and are seldom actionable in the proband.

It is important for the clinician to recognize when they can contribute additional information that might allow a VUS to be re-interpreted with more confidence, and the laboratory will often flag in the report if extra information might be sufficient to reclassify. For example, if a variant is not inherited from either parent, then it has likely arisen *de novo* in the affected individual and is more likely to be pathogenic. Segregation analysis may reveal that the variant is shared by several affected relatives, providing additional evidence for a role in disease rather than a bystander role. Reclassification of variants can be challenging for patients and families,<sup>38,39</sup> but should be part of pre-test genetic counselling discussions.

## 2.2. Genetic counselling

#### 2.2.1. What is genetic counselling?

Genetic counselling is a process that aims to support patients and their families to understand and adapt to the medical, psychosocial, and familial impact of genetic diseases.<sup>40,41</sup> Genetic counselling can include discussion of inheritance risks, provide education, perform pre- and post-genetic test counselling and variant interpretation, obtain a three-generation family history, and provide psychosocial support.<sup>42</sup> The unique psychosocial needs of the inherited cardiomyopathy patient population have shaped the field of cardiac genetic counselling.<sup>43</sup> For those with a new diagnosis, there can be difficulty adjusting to life with an inherited cardiomyopathy, challenges living with an implantable

cardioverter defibrillator (ICD), and ongoing trauma and grief for those who have experienced a young SCD in their family. Attention to the psychological support needs of patients is therefore critical. Indeed, in the general setting, genetic counselling can improve knowledge, recall, and patient empowerment; increase satisfaction with decision-making; and reduce anxiety.<sup>44–47</sup>

#### 2.2.2. Who performs genetic counselling?

Genetic counselling is a process that is preferably performed by healthcare professionals with specific training, such as genetic counsellors, genetic nurses, or clinical/medical geneticists. Regardless of who takes on this role, it involves more than simply the provision of information: it requires careful attention to psychosocial needs. A systematic review of genetic counselling communication showed that higher levels of empathic responses, less verbal dominance (ratio of counsellor to patient talk), and the provision of a summary letter were associated with more positive outcomes.<sup>48</sup>

#### 2.2.3. Genetic counselling in paediatrics

Genetic counselling performed by trained healthcare professionals working within a multidisciplinary team is recommended for all children with all types of cardiomyopathies, regardless of whether genetic testing is being considered.

There are specific issues to consider when counselling children and their families and considering clinical screening and predictive genetic testing.<sup>49–51</sup> The guiding principle remains that any testing, clinical or genetic, should be in the best interests of the child and have an impact on management, lifestyle, and/or ongoing clinical testing.<sup>50</sup> The psychosocial impact should be considered and counselling should take place in the context of a multidisciplinary setting involving the child and their family.<sup>42,52–54</sup> With appropriate multidisciplinary support in a paediatric setting, psychosocial outcomes in children undergoing clinical screening and predictive genetic testing are no different than those of the general population.<sup>54</sup> Pre- and post-test counselling should be performed,<sup>42,53</sup> and the wishes of the child in terms of testing and disclosure of results should be at the forefront. Although the Task Force acknowledges that different healthcare systems may not allow for this, psychological support should be made available to all children and families with cardiomyopathy or undergoing clinical and/or genetic testing, with a particular focus on those with a recent diagnosis, a family history of SCD, and who are at important points during childhood, such as moving from primary to secondary education and transitioning from paediatric to adult services.<sup>52,54–56</sup>

# 2.2.4. Pre- and post-test genetic counselling (proband)

One critical role for genetic counselling is alongside genetic testing. This includes a discussion prior to a decision to undertake genetic testing (pre-test) and when the results are returned (post-test). Pre-test genetic counselling focuses on the collection of a detailed family history, providing genetic education about inheritance risks, the process and logistics of genetic testing and options for return of the results, an explanation of all possible outcomes, clinical and family implications, the risk of reclassification and identification of secondary genetic findings, insurance implications, and an exploration of feelings and understanding of the information provided. Post-test discussions include a review of the key points previously discussed in the pre-test session, return of the result, discussion of the implications of this result to the patient's care, and implications for the family. It can be helpful to provide tailored

information from the family history, explicitly noting every family member's inheritance risk, and clinical and genetic testing options. Where a suspicious VUS has been identified, specific explanation of what this means is important, including whether there is a plan for resolving it, such as segregation to other affected relatives in the family, and how often the classification will be reviewed. There should be ample opportunity for questions or clarification at all points in the process.

#### 2.2.5. Cascade genetic testing

Once a P/LP variant has been identified within an index patient following investigations of relevant disease genes associated with the specific phenotype, it is possible to offer cascade genetic testing of first-degree at-risk relatives, including pre-test genetic counselling. In a scenario where a first-degree relative has died, evaluation of close relatives of the deceased individual (i.e. second-degree relatives of the index patient) should also be considered. Genetic cascade screening should continue for all offspring of relatives who are shown to carry a P/LP variant.

Relatives who do not carry the variant can be dismissed from followup, while genotype-positive relatives should continue regular clinical evaluation.

The right assignment of the level of pathogenicity of a variant is crucial for cascade genetic testing. Inappropriate use of genetic testing in a family has the potential to introduce unnecessary worry and fear, as well as potential harm related to the misinterpretation of genetic variants. Therefore, it is advised that variants should be classified by a specialized multidisciplinary cardiac genetic team with an appropriate level of expertise. In this context, it is important to understand that up to 50% of published variants are wrongly classified. In addition, systematic reclassification of identified variants is crucial, including its communication to families, emphasizing the need for these specialized multidisciplinary cardiogenetic teams.

#### 2.2.6. Uptake of genetic counselling and testing

Several studies have evaluated the uptake of genetic testing and counselling and/or cardiological screening in inherited cardiomyopathies in first-degree relatives of index patients. Roughly 39–45% of at-risk relatives for an inherited cardiomyopathy come for screening within 1–2 years after the identification of the pathogenic variant in the family.<sup>57–60</sup> After a longer follow-up (median 16 years) this increased to 60%.<sup>60</sup> A first-intervention study (more intensive discussion of informing relatives at risk with the index patient, followed by a family letter sent directly to the relatives) did not increase the uptake.<sup>60</sup> This suboptimal uptake leaves room for improvement, and future studies will give guidance on how to do this. Currently, it is recommended to discuss with the index patient how they will reach out to first-degree relatives and how the healthcare provider can help and provide support.

#### 2.2.7. Approaching relatives of the index patient

At-risk relatives are generally informed by the index patient, who is supported with a family letter about the condition, the content of the family screening programme, and links to relevant websites with further information. However, the approach to relatives varies between countries due to differences in tradition and legislation.

The discussion of how to approach family members is an important part of the genetic counselling process with the index patient.

For a variety of reasons, the index patient may refuse to inform atrisk relatives of having inherited the condition and about the offer of family screening. This is a cause for concern, since most inherited cardiac conditions (ICCs) are associated with adverse complications and SCD, which may be prevented if diagnosed in due time. Depending on the legislation of each country, it may be possible to bypass the index patient and approach at-risk relatives directly. This may be done through a letter informing each relative that a hereditary condition is suspected in the family and that they should contact the ICC unit and obtain more information if they wish. Sometimes, affected relatives may refuse to have their minor children investigated; an issue which may be difficult to solve unless there is legislation in place that outlines how to deal with this issue.

When these kinds of difficult situations occur, it is important to be patient and try to understand the reasons of the affected individuals for withholding information from at-risk relatives. Offering repeated counselling sessions or suggesting a phone call with relatives at a later stage are two possible solutions. Usually, however, the problem resolves once the affected individual has had sufficient time to consider the implications.

#### 2.2.8. Inheritance and family communication

Knowing that one's family is at increased risk of inheriting a genetic condition can create uncertainty and worry for many index patients. Because most inherited cardiomyopathies follow an autosomal dominant inheritance pattern, there is a need to communicate the importance of both clinical and genetic testing of relatives. Conveying this information to at-risk relatives is typically reliant on the proband in the family understanding the information and passing it on to the appropriate relatives. Numerous studies show that effective family communication is an important challenge, and interventions to support communication have so far not yielded favourable results, highlighting the multidimensional nature of the issue.<sup>52,60,61</sup> Common barriers to communication may include poor family relationships; guilt about passing a causative variant on to children; psychosocial factors, including distress; and comprehension of the result.<sup>62,63</sup> A patient will often selectively communicate genetic information to relatives, assessing their ability to understand and cope with the information, their life stage, and risk status.<sup>61</sup> Health literacy is an important barrier to effectively communicating genetic risk information to relatives, highlighting the need for targeted resources and mechanisms for support.<sup>64</sup>

#### 2.2.9. Psychological support

Coming to terms with a diagnosis of an inherited cardiomyopathy can be difficult and have a significant impact on psychological well-being.  $^{43,65-70}$ 

As a patient, being at risk of hereditary cardiomyopathy is not only a matter of facts and test results, but also of perception: while one affected family member can have only mild symptoms that can be treated

with medication, others may have a different outcome with a greater impact in terms of the treatment needed (e.g. ICD or transplant). This fact should be considered when treating a family with a hereditary cause of cardiomyopathy, in aspects such as the communication of test results and targeting psychological needs. Not every patient within a family has the same burden of symptoms and/or disease outcome.

The 'estimate of risk' is not only a matter for the professional caretaker in terms of how to treat the patient: the impact on the patient themselves and the consequences of that treatment may necessitate a very thorough and professional need for psychological assistance. The psychological well-being of a patient with cardiomyopathy has a broad range of aspects to consider, from the very personal to specific and practical needs:

- Self-care should be clearly emphasized: from day-to-day blood pressure testing to the importance of a strict medication schedule. This contributes to a patient's feeling of self-control.
- The aspects of treatment (medication, ICD, transplant) that have an effect on daily life should be clearly communicated (e.g. medication schedule, side effects of medication, diet restrictions, exercise restrictions, etc.).
- The significance of the hereditary component of the condition should be made clear, as the possibility of having children with the defect (or the decision not to have children at all) can be a difficult burden to bear for many patients.
- Practical issues should be emphasized (these may vary according to local legislation); e.g. seeking a mortgage, getting a driver's licence, career planning, etc.

Psychological assistance should support and encourage the patient as well as the family, but also give guidance to the patient to try to find answers to the very specific challenges they will face.

# 3. Specific entities

#### **3.1. Hypertrophic cardiomyopathy** 3.1.1. Cardiac magnetic resonance imaging and sudden cardiac death risk in hypertrophic cardiomyopathy

Although current data suggest that the extent of late gadolinium enhancement (LGE) on CMR may be of use in predicting cardiovascular mortality, data in regards to the use of LGE in the prediction of SCD risk are conflicting (Table S1). Similarly, the roles of the presence of LV apical aneurysms (Table S2) and LV systolic dysfunction (Table S3) as additional independent predictors of SCD risk in HCM remain to be determined.

Study name		Details and quality	/ of evidence		Summary of key findings	
author	Study type	Number of patients and follow-up duration	Key inclusion and exclusion criteria	Relevant aims and outcome(s)	Key findings	Conclusions and limitations
Maron et <i>a</i> l., 2008∕1	Observational, prospective	n = 202 Two centres FU: 681 ± 249 days	Inclusion: consecutive HCM pts	<b>Aim:</b> describe clinical profile and relation of LGE to CV outcomes	<ul> <li>Adverse cardiovascular events in 11/202 pts.</li> <li>Among 7/11 with LGE:</li> <li>2 died suddenly;</li> <li>2 died suddenly;</li> <li>2 died suddenly;</li> <li>2 and</li> <li>3 with progressive HF symptoms vs. 4/11 pts without LGE:</li> <li>3 with sudden death; and</li> <li>1 with progressive HF.</li> <li>Annual cardiovascular event rate in HCM pts with LGE exceeded that of pts without LGE (5.5% vs. 3.3%), but not statistically significant (HR 1.45; 95% CI, 0.43–4.97; P = 0.5)</li> </ul>	<b>Conclusions:</b> LGE was an independent predictor of systolic dysfunction. Data insufficient to consider LGE as an independent risk factor for adverse prognosis
Bruder et <i>al.</i> , 2010 <sup>72</sup>	Observational, prospective	n = 243; 220 in analysis Single centre Mean FU: 1090 days	Inclusion: consecutive HCM pts Exclusion: Hx of previous septal ablation or myectomy, known coronary artery disease, aortic stenosis, amyloidosis, systemic hypertension, or contraindications to CMR	Primary endpoints: (i) all-cause death; and (ii) cardiac death (death from all cardiac causes, including SCD, heart failure, and aborted SCD)	All-cause mortality: 20/220 SCD: 11/220 among which 2 with ICD discharged Univariate analysis for SCD: Presence of LGE: OR 5.14; 95% CI, 0.65–41.0; $P = 0.10$ (non-significant) Multivariable Cox regression analysis, (presence of LGE, LVEF, and LV mass): LGE independent predictor of cardiac death (HR 4.81; $P = 0.035$ ) Not performed in the subgroup of SCD patients ( $n = 11$ , limited number of events)	<b>Conclusions:</b> LGE is a predictor of all-cause and cardiac mortality. Unable to demonstrate statistically significant relation to SCD <b>Limitations:</b> single centre, small size, limited number of events

Table S1 Studies on the role of late gadolinium enhancement in predicting sudden cardiac death in patients with hypertrophic cardiomyopathy

<b>Conclusion:</b> LGE is an independent predictor of adverse outcomes. Longer follow-up in a larger cohort will clarify the role of LGE as a risk factor in SCD prediction <b>Limitations:</b> single referral centre, small cohort, underpowered, low event rate, selection bias (pts with ICDs excluded at baseline)	<b>Conclusion:</b> LGE remained a significant associate of subsequent SCD or appropriate ICD therapies after controlling for other factors <b>Limitations:</b> single centre, selection bias, low event rate, lack of multivariate Cox model, high number of SRT in FU
Primary endpoint: 40/217 (18.4%) pts • 6/81 (7.4%) in the LGE group (HR 3.4, 95% Cl, 1.4-8.1; P = 0.006) CV deaths: 9 LGE group: 8 Multivariable analysis: presence and amount of LGE remained independent predictors of the primary endpoint (HR 2.7; 95% Cl, 101–7.1; P = 0.046) Arrhythmic endpoint: 12/217 overall • LGE group: 10/136 (7.3%), and • non-LGE group: 2/81 (2.5%) (HR 3.15; 95% Cl, 0.69–14.4; P = 0.138) Univariate analysis: the amount of LGE was significantly associated with the outcome (HR 1.30; 95% Cl, 1.05–1.61; P = 0.014), but in the multivariable analysis: NSVT was the strongest predictor. No other variables were added to the model (low event rate)	Outcome occurred in 8/424 pts • SCD 4 pts; and • ICD discharge: 4 pts All 8 pts had LGE In addition to LGE, the presence of NSVT was another univariate associate of events (HR 6.9, 95% Cl, 1:14–52.4; <i>P</i> = 0.04) No multivariable Cox model analysis performed (low event rate). On bivariate analysis LGE remained associated with events after controlling for all other parameters
Combined primary endpoint: (i) CV death; (ii) unplanned CV admission; (iii) sustained VT/VF, or discharge Secondary outcomes: (i) HF; (i) arrhythmic	Aims: (i) evaluate the relation between LGE and HCM genes status, severity of symptoms, and the degree of ventricular ectopy on Holter ECG; (ii) Outcomes of SCD and appropriate ICD therapies were recorded
Inclusion: consecutive HCM pts	Inclusion: consecutive HCM pts
Exclusion: significant coronary	Exclusion: ICD <i>in situ</i> , previous hx
artery disease, or prior SRT	of SRT
n = 217	n = 424
Single centre	Single centre
Mean FU: 3.1 ± 1.7	Mean FU: $43 \pm 14$
years	months
Observational,	Observational,
prospective	retrospective
O'Hanlon	Rubinshtein
et <i>al.</i> , 2010 <sup>73</sup>	et <i>al.</i> , 2010 <sup>74</sup>

Green et <i>a</i> l., 2012 <sup>75</sup>	Systematic review and meta-analysis	n = 1063 Pooled HCM pts from 4 studies (1–4) Mean FU: 3.1 years	Inclusion: studies of LGE in HCM that reported outcomes of cardiovascular mortality, SCD, aborted SCD, and HF death	<b>Aim:</b> association of LGE with CV mortality, SCD, aborted SCD, and HF death	LGE correlated with: • cardiac death (OR 2.92; 95% Cl, 1.01–8.42; <i>P</i> = 0.047) • HF death (OR 5.68; 95% Cl, 1.04– 31.07; <i>P</i> = 0.045); and • all-cause mortality (OR 4.46; 95% Cl, 1.1.53–13.01; <i>P</i> = 0.006), but was not significant for SCD/ aborted SCD (OR 2.39; 95% Cl, 0.87–6.58; <i>P</i> = 0.091)	<b>Conclusion:</b> there are significant relationships between LGE and cardiovascular mortality, heart failure death, and all-cause mortality in HCM. Additionally, LGE and SCD/ aborted SCD displayed a trend toward significance
2014 <sup>76</sup> 2014 <sup>76</sup>	Observational, prospective	n = 1293 7 HCM centres Mean FU: 3.3 years	Inclusion: consecutive HCM patients with CMR at first evaluation Exclusion: prior ICD implantation or history of sustained VT/VF, claustrophobia, obstructive coronary artery disease, septal myectomy or alcohol ablation (before CMR), and incomplete follow-up	SCD composite endpoint: HCM-related sudden death, aborted arrest, ICD discharge	Cumulative SCD events incidence: 0.9%/year SCD events in 37/1293 patients (3%): • 14/1293 aborted cardiac arrest; • 17/1293 appropriate ICD therapy LIGE present in 548/1293 pts (42%): • ≤10% of the LV mass ( <i>n</i> = 381, 29%); • 11–19% ( <i>n</i> = 94, 7%); • 220% ( <i>n</i> = 73, 6%) Among 37 pts with SCD events, LIGE was present in 26 (70%), occupying 13 ± 14% of the LV myocardium Each 10% increase in LGE was associated with P0% increase in relative SCD events risk (adjusted HR 1.46 for 10% increase in LGE; 95% CI, 1.12–1.92; <i>P</i> = 0.002) Compared with patients without LGE, the adjusted (for conventional risk factors and EF) HR of SCD events related to %LGE: • 10%, 1.46; • 15%, 1.77; • 20%, 2.14 In low-risk pts SCD event risk increased in direct proportion to extent of LGE (HR 1.66/10% LGE; 95% CI, 1.24–2.23; <i>P</i> = 0.0007) The absence of LGE was associated with lower risk of SCD events (adjusted HR 0.39, 95% CI, 0.18–	<b>Conclusions:</b> extensive LGE provides additional information for assessing SCD event risk among HCM patients, particularly patients otherwise judged to be at low risk <b>Limitations:</b> selection bias, exclusion of high-risk pts with h CDs, low event rate in HCM. 17 out of 37 ICD shocks potentially non-fatal arrhythmias

2014 <sup>77</sup> Observ 2014 <sup>77</sup> prospec	Briasoulis Meta-ar et al., 2014 <sup>78</sup>
ctive ctive	nalysis
Size: 711 Median FU: 3.5 years Single HCM centre	n = 3067 Pooled pts from 6 HCM studies. Mean FU: 3.05 years
Inclusion: consecutive HCM pts Exclusion: previous myectomy or alcohol septal ablation; previous myocardial infarction; or contraindications to CMR (including prior device implantation) and gadolinium-based contrast agents	Inclusion: prospective HCM studies reporting the effects of LGE on clinical outcomes (SCD/aborted SCD, all-cause mortality, cardiac and heart failure death)
Primary endpoint: SCD or aborted SCD	<b>Aim:</b> assess the utility of CMR on LGE as a prognostic factor of SCD in HCM
SCD composite endpoint 22/711 pts (3.1%), in detail: • 18 (3.8%) pts with LGE and 4 (1.7%) with no LGE (HR LGE, 2.69, 95% CI, 0.91–7.97, $P = 0.073$ ) Patients with LGE were more likely to have extreme hypertrophy (LV wall thickness $\geq$ 30 mm) and non-sustained VT at baseline. On univariable analysis, the amount of fibrosis was a significant predictor of outcome: (HR per 5% LGE 1.24; 95% CI, 1.06–1.45, $P = 0.007$ ) On multivariable analysis, only LVFF was an independent predictor (HR 0.92; 95% CI, 0.89–0.95, $P \leq 0.001$ ) Increasing %LGE was associated with increased risk. However, not statistically significant after adjusting for EF	SCD/aborted SCD in HCM pts with LGE was significantly increased as compared with pts without LGE (OR 2.52; 95% Cl, 1.44-4.4; P = 0.001) The extent of LGE was not significantly related to the risk of SCD
<b>Conclusions:</b> the amount of myocardial fibrosis is an important predictor of outcome in HCM but may not provide information incremental to EF. <b>Limitations:</b> single centre. Small size. Low event rate. Selection bias. High-risk pts with ICDs excluded. Methodology of LGE quantification	<b>Conclusion:</b> LGE is significantly associated with SCD risk, cardiac mortality, and all-cause mortality in patients with non-high-risk HCM according to conventional risk factors

<pre> € ESC 2033 € ESC 203 € ESC 203 € ESC 203 € ESC 203 € ESC 2033 € ESC 203 €</pre>
S0/1423 (4%) composite primaryConceaseVents:- 40 actual SCD; $\otimes$ compared and SCD; $\otimes$ compared by LGE on CMR: 717 (50%) pts $\otimes$ LGEAny LGE on CMR: 717 (50%) pts $\otimes$ not minary $\otimes$ compared by the other and solved by the other and
Primary composite endpoint: actual SCD, successful arrest or appropriate ICD shocks P F F F F F F F F F F F F F F F F F F F
Inclusion: consecutive low-/ intermediate-risk HCM pts with preserved LVEF undergoing CMR. Exclusion: prior device implantation; claustrophobia; high SCD risk pts; ≥moderate aortic/mitral stenosis; subaortic membrane; LGE pattern consistent with ischaemic myocardial damage; LVEF <50%; prior history of alcohol septal ablation or surgical myectomy; and prior mitral and/or aortic valve replacement
n = 1423 (458 non-obstructive, 965 obstructive) Single HCM centre Median FU: 4.5 years
Observational, retrospective
Mentias et <i>al.</i> , 2018 <sup>79</sup>

Cl, confidence interval; CMR, cardiac magnetic resonance; CV, cardiovascular; ECG, electrocardiogram; EF, ejection fraction; ESC, European Society of Cardiology; FU, follow-up; HCN, hypertrophic cardiomyopathy; HF, heart failure; HR, hazard ratio; Hx, history; ICD, implantable cardioverter defibrillator; LGE, late gadolinium enhancement; LV, left ventricular; LVEF, left ventricular ejection fraction; NSVT, non-sustained ventricular tachycardia; OR, odds ratio; pts, patients; SCD, sudden cardiac death; SRT, septal reduction therapy; VF, ventricular fibrillation; VT, ventricular tachycardia.

Summary of key findings	Key findings Conclusion(s)	site endpoint event rate ar vs. 0.9%/year in ig HCM cohort. PHCM patients with LV apical aneurysms are at high risk of arrhythmic sudden death mposite endpoint 21/93 \$%), in detail: 5%), in detail: 5%), in detail: 5%), in detail: 15CD: 11% (1/93 pts) ed SCD: 2.2% (2/93 pts) priate ICD therapy (18/93 pts). Rhythm nonomorphic VT in 89% 3 pts)	site endpoint event rateConclusion:arararDenotype, associated with increased risk ofmposite endpoint 14/Phenotype, associated with increased risk of(8.8%), in detail:adverse cardiovascular events, includingSCD: noneadverse cardiovascular arrhythmiased SCD: 4.4% (7/160)Limitations:selection bias, high prevalence of confounderspriate ICD therapyin patients with events (previous VT/VF and LV(7/160 pts). Rhythm notsystolic function)
	Relevant outcome(s)	SCD composite endpoint Compos (actual SCD + resuscitated 4.7%/yes cardiac arrest + appropriate remainir ICD therapy) pts (22.6 pts (22.6 pts (22.4	SCD composite endpoint Compos (actual SCD + resuscitated 1.8%/yet cardiac arrest + appropriate 7.8% or co ICD therapy) 160 pts actual • abort pts) • appro 4.4% ( specifi
vidence	Key inclusion and exclusion criteria	<b>Inclusion:</b> diagnosis of HCM and apical aneurysm on cardiac MRI and/or echo <b>Exclusion:</b> known obstructive coronary artery disease	Inclusion: diagnosis of HCM and apical aneurysm on cardiac MRI and/or echo Exclusion: history of anterior infarction or significant LAD stenosis
Details and quality of ev	Number of patients and intervention(s)	93 pts from 2 centres	160 pts from a single centre
	Study type	Retrospective observational	Retrospective observational
Study	first author	Rowin et al., 2017 <sup>80</sup>	Lee et <i>al.</i> , 2022 <sup>81</sup>

 Table S2
 Apical aneurysms and sudden death in hypertrophic cardiomyopathy

HCM, hypertrophic cardiomyopathy; ICD, implantable cardioverter defibrillator; LAD, left anterior descending; LV, left ventricular; MRI, magnetic resonance imaging; pts, patients; SCD, sudden cardiac death; VT, ventricular tachycardia; VF, ventricular fibrillaton.

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Table S3 Studie:	s on the pr	evalence of and predic	tors for left ve	entricular systolic e	dysfunction in patients with hyper	trophic cardiomyopathy
Study name or first author	Number	Specific subset	LVSD prevalence	Annual incidence of HCM-LVSD	Main endpoints	Multivariable analysis for prediction of SCD events
Wasserstrum et al. <sup>82</sup>	1328	None	2.8%	0.39%	All-cause mortality or ventricular assist device or transplant	Not available
Biagini et <i>al.</i> <sup>83</sup>	222	Both paediatric and adult patients	4.9%	0.53%	Cardiovascular death (SCD, HF-related death, stroke-related death)	LV wall thickness (HR 1.07; 95% Cl, 1.01–1.14; $P = 0.03$ ) and LV end-diastolic diameter (HR 1.08; 95% Cl, 1.04–1.11; $P = 0.0001$ ) <sup>a</sup>
Thaman et al. <sup>84</sup>	1080	Serial echocardiography substudy ( <i>n</i> = 462)	2.4%	0.87%	Composite of SCD and ICD firing: composite of all-cause death, ICD firing, and heart transplantation	Not available
Harris et al. <sup>85</sup>	44	None	3.5%	1.12%	Death, SCD, ICD firing, heart transplantation	Not available
Kawarai et al. <sup>86</sup>	43	Incident HCM-LVSD	8.7%	0.73%	Sudden death including also non-fatal cardiac arrest and appropriate ICD intervention	Among HCM-LVSD the presence of syncope resulted as an independent predictor of sudden death (HR 6.15, 95% Cl, 2.40–15.75; $P < 0.001)^{\rm b}$
Rowin et al. <sup>87</sup>	2447	None	4.8%	па	SCD events, heart transplant listing, HCM-related death, all-cause death	Not available
SHaRe Consortium <sup>88</sup>	6793	None	8.1%	а	All-cause death, heart transplantation, sudden death, ICD activation	Adjusted <sup>c</sup> HCM-LVSD HR for SCD, 3.9; 95% Cl, 2.6– 2001; adjusted HCM-LVSD HR for ICD activation, 1.6; 95% Cl, 1.4–1.8; P < 0.001

Cl, confidence interval; HCM, hypertrophic cardiomyopathy; HF, heart failure; HR, hazard ratio; ICD, implantable cardioverter defibrillator; LV, left ventricular, LVSD, left ventricular systolic dysfunction; na, not available; pts, patients; SCD, sudden cardiac death; VT, ventricular tachycardia.

<sup>a</sup>Gender, age at first evaluation, family history of sudden death, New York Heart Association (NYHA) functional class I to II to IV, medical treatment at first evaluation, non-sustained ventricular tachycardia, LV outflow obstruction, maximum LV wall thickness, posterior wall thickness, LV end-diastolic diameter, end-systolic left atrial diameter, and (as a time-dependant covariate) dilated-hypokinetic evolution.

<sup>b</sup>Multivariable Cox regression including: positive family history, maximum LV wall thickness >30 mm at the initial diagnosis of HCM, non-sustained VT at the diagnosis of end-stage HCM, and unexplained syncope at the diagnosis of end-stage HCM.

3.2. Restrictive cardiomyopathy

The systematic approach to the diagnosis of RCM should include clinical examination, ECG, advance cardiac imaging and genetic testing. Table S4 summarizes the spectrum of restrictive heart disease.

# Table S4 Spectrum of restrictive heart disease

		Myocardium			Endocardium	Pericardium
	Myocytes		EC	>		
RCM, primary, genetic	Genocopies	Phenocopies, non-genetic	Infiltration	Fibrosis/ECV remodelling		
With or without	Storage diseases with	Toxicity:	Extramyocyte	Non-specific	Diseases affecting	Pericardium diseases
aggregates of	accumulation:	intraniyocyte accumulation in acquired lysosomal diseases:	infiltration:	<ul> <li>diabetic heart</li> </ul>	enuoniyocar ulai layers:	anecung pericardium and
misfolded, mutated	Lysosomal diseases <sup>a</sup>	DIP by cationic amphiphilic drugs:	<ul> <li>amyloidosis, genetic</li> </ul>	disease (SVD,	<ul> <li>endomyocardial</li> </ul>	epicardial fat tissue:
proteins:	<ul> <li>glycogenoses (pompe,</li> </ul>	e.g.	and non-genetic <sup>a</sup> (+	fibrosis and	fibrosis	<ul> <li>constrictive pericarditis</li> </ul>
<ul> <li>troponinopathies (1)</li> </ul>	Mcardle, Danon, etc.)	chloroquine-hydroxychloroquine	myocyte toxicity in	myocyte damage)	<ul> <li>eosinophilic: tropical</li> </ul>	(panpericardial or
most common; T; C)	<ul> <li>Glycosphingolipidoses</li> </ul>	(other cationic amphiphilic drugs)	primary	<ul> <li>autoimmune</li> </ul>	and non-tropical forms	segmental; rare,
<ul> <li>Myosinopathies (MYH7)</li> </ul>	(AFD)	Less common:	amyloidosis)	diseases (fibrosis,	(myeloproliferative,	post-pericardiotomy)
<ul> <li>Desminopathies (DES)</li> </ul>	Iron storage diseases, genetic	serotonin	<ul> <li>cystinosis, genetic,</li> </ul>	large and small	with hypereosinophilia,	<ul> <li>pericardial tumours,<sup>a</sup></li> </ul>
BAG30pathies (BAG3)	forms:	<ul> <li>methysergide</li> </ul>	AR (CTNS gene)	vessel	Löffler endocarditis,	benign (cysts and
<ul> <li>Myotilinopathies</li> </ul>	<ul> <li>HH (myocytes)</li> </ul>	<ul> <li>ergotamine</li> </ul>	<ul> <li>hyperoxaluria:</li> </ul>	involvement) e.g.	endocardial	lipomas); malignant
(MYOT) Filaminopathies	Friedreich ataxia	<ul> <li>mercurial agents, busulfan</li> </ul>	primary, genetic:	scleroderma	thrombosis)	(primary
(FLNC)	(mitochondria)	<ul> <li>heavy metals</li> </ul>	AGXT and GRHPR	<ul> <li>inflammatory,</li> </ul>	<ul> <li>Endinger syndrome</li> </ul>	mesothelioma, most
<ul> <li>Ab-crystallinopathies</li> </ul>	Heritable	<ul> <li>potentially reversible</li> </ul>	genes; secondary	granulomatous	(carcinoid heart	common)
(CRYAB)	haemoglobinopaties and		(jejunoileal bypass,	(e.g. sarcoidosis:	disease)	<ul> <li>radiation therapy</li> </ul>
<ul> <li>Titinopathies (TTN)</li> </ul>	other haematologic diseases		oxalate poisoning,	sporadic; rare	<ul> <li>iatrogenic/drug toxicity</li> </ul>	<ul> <li>epicardial</li> </ul>
(rare)	may cause myocardial iron		drug toxicity)	genetic early-onset	<ul> <li>radiation therapy</li> </ul>	steatonecrosis in heart
<ul> <li>Less common: ACTCI,</li> </ul>	storage			sarcoidosis:	<ul> <li>endocardial neoplasms<sup>a</sup></li> </ul>	transplantation
MYBPC3, MYL3, TPMI,				CARD15/NOD2		. ጋՏ:
MYL3, MYL2				gene		

AFD, Anderson–Fabry disease; AR, autosomal recessive; DIP, drug-induced phospholipidosis; ECV, extracellular volume; HH, hereditary haemochromatosis; RCM, restrictive cardiomyopathy; SVD, small vessel disease. <sup>a</sup>Tumours: rhabdomyomas isolated or in tuberous sclerosis 1 and 2, fibromas; angiosarcomas, fibrosarcomas, liposarcomas, liposarcomas

## 3.3. Amyloid

Prognosis in cardiac amyloidosis was originally considered poor but this can be further stratified (Table S5). Several multiparametric biomarkerbased staging systems have been developed for  $AL^{89,90}$  and ATTR cardiac amyloidosis.<sup>91–93</sup>

Table S5	Prognostic staging scores i	n light chain amyloi	idosis and transthy	retin amyloidosis/

Kumar et al. <sup>8</sup> AL	9	Lillenes e AL	t al. <sup>90</sup>	Grogan e ATTR	et al. <sup>93</sup> Rwt	Gillmore of ATTRv and a	et al. <sup>92</sup> ATTRwt	C ATT	heng et al. <sup>91</sup> Rv and ATTRwt
Staging paramet FLC-diff >18 mg Troponin T > 0 NT-proBNP > 1	ers: g/dL .025 ng/mL 1800 pg/mL	Staging paramet troponin l > 0.1 BNP > 81 pg/ml	ers: ng/mL -	Staging paramet troponin T > 0.9 NT-proBNP > 30	ers: 5 ng/mL 000 pg/mL	Staging paramet eGFR <45 mL/r NT-proBNP > 30	ers: nin 000 pg/mL	Scoring parameters: Mayo or NAC Score (0–2 points) Daily dose of furosemide or equivalent: 0 mg/kg (0 points), >0–0.5 mg/kg (1 point), >0.5–1 mg/kg (2 points), and >1 mg/kg (3 points) NYHA class I–IV (1–4 points)	
Stage	5-year	Stage	Median	Stage	4-year	Stage	Median	Score	Median survival

0.000	survival	ougo	survival	04460	survival/ median survival	04460	survival		
Stage I (0 parameters)	68%	Stage I (0 parameters)	Not reached	Stage I (0 parameters)	57% 66 months	Stage I (0 parameters)	69.2 months	Score 1–3	90.5 months
Stage II (1 parameter)	60%	Stage II (1 parameter)	112.8 months	Stage II (1 parameter)	42% 40 months	Stage II (1 parameter)	46.7 months	Score 4–6	38.5 months (Mayo) 36 months (NAC)
Stage III (2 parameters)	28%	Stage III (2 parameters)	51.6 months	Stage III (2 parameters)	18% 20 months	Stage III (2 parameters)	24.1 months	Score 7–9	20.3 months (Mayo) 19.8 months (NAC)
Stage IV (3 parameters)	14%	Stage IIIb (2 parameters and BNP >700 pg/mL)	12 months						

AL, Monoclonal immunoglobulin light chains amyloidosis; ATTRv, Hereditary transthyretin amyloidosis; ATTRvt, wild-type transthyretin amyloidosis; BNP, brain natriuretic peptide; eGFR, estimated glomerular filtration rate; FLC, free light chain; NAC, national amyloid centre; NT-proBNP, N-terminal pro-brain natriuretic peptide; NYHA, New York Heart Association.

# 4. Sports

Exercise recommendations for the different cardiomyopathies are referred to intensity levels that are provided in Table S6.

Intensity	VO <sub>2</sub> max (%)	HRmax (%)	HRR (%)	RPE scale	Training zone
Low-intensity, light exercise	<40	<55	<40	10–11	Aerobic
Moderate-intensity exercise	40–69	55–74	40–69	12–13	Aerobic
High-intensity exercise	70–85	75–90	70–85	14–16	Aerobic + lactate
Very high-intensity exercise	>85	>90	>85	17–19	Aerobic + lactate + anaerobic

 Table S6
 Indices for exercise intensity during sport

HRmax, maximum heart rate; HRR, heart rate reserve; RPE, rate of perceived exertion;  $VO_2max$ , maximum oxygen consumption. Adapted from Vanhees et al.<sup>94</sup>

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# 5. Recommendations for non-cardiac interventions

Management of patients with HCM and LVOTO deserves careful evaluation. Objectives and specific actions are summarized in Table S7.

# Table S7 Management of a patient with hypertrophic cardiomyopathy and left ventricular outflow tract obstruction for non-cardiac interventions

Objectives	<ul> <li>To maintain normal to high cardiac pre-load.</li> <li>Afterload control.</li> <li>To maintain sinus rhythm/AV synchrony at low heart rates (60–65 b.p.m.).</li> <li>Medication to treat LVOTO should remain on board during NCS.</li> </ul>
To avoid	Diuretics and inotropic agents.
If hypotension appears	<ul> <li>Pre-load should be corrected (fluids, Trendelenburg position to favour venous return).</li> <li>Use drugs that increase systemic vascular resistance without causing greater ventricular obstruction and without increasing contractility and heart rate as phenylephrine or vasopressin.</li> </ul>
If LVOTO appears	Consider use of intravenous beta-blockers such as short-acting esmolol, metoprolol, or labetalol.
lf rapid atrial fibrillation appears	Rate control with beta-blockers or calcium blockers or electric cardioversion if haemodynamic instability.

AV, atrioventricular; b.p.m., beats per minute; LVOTO, left ventricular outflow tract obstruction; NCS, non-cardiac surgery.

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Adapted from Sahoo et al., Dhillon, et al., and Hensley et al.<sup>95–97</sup>

# 6. Requirements for specialized cardiomyopathy units

#### 6.1. Requirements and skills

A cardiogenetic unit/cardiomyopathy unit (CGU/CMU) should have access to both outpatient and inpatient facilities within a healthcare provider structure.<sup>98–101</sup> The activity of a CGU/CMU is based on the availability of multimodality cardiac imaging facilities (echocardiography

with advanced analysis, CMR, nuclear medicine), electrocardiographic (including Holter monitoring), and functional evaluation (cardiopulmonary test, electrocardiogram [ECG] exercise testing). Close collaboration with interventional cardiology, electrophysiology, and cardiac surgery departments is necessary. A genetic team with expertise in cardiomyopathies and links to a clinical genetic testing laboratory is also required. These units should also provide a framework for discussing the diagnosis, therapeutic options, and prognosis. Clinical psychologists and genetic counsellors assist with the integration of information by the patients and their families. These units should also collaborate in the management of relatives across different geographic areas. These requirements are summarized in *Table S8*.

# 7. Living with cardiomyopathy: patient and family education

Our knowledge of both the clinical and genetic basis of cardiomyopathies has been a rapidly evolving area of cardiology over the last 30 years. This poses significant challenges to healthcare professionals who seek to stay abreast of new knowledge and ensure that their patients are receiving the highest standard of care. This can also be difficult for patients who seek to better understand the condition they are being assessed or managed for. For a patient and their at-risk family members, resources to enable understanding and make sense of the tangible impact a diagnosis will have on them should be made available. Patient support groups may also play an important role in supporting patient and family education.

Patient education has been defined as a 'process of assisting consumers of healthcare to learn how to incorporate health-related behaviours (knowledge, skill, attitude) into everyday life with the purpose of achieving the goal of optimal health'.<sup>103</sup> Incorporating patient education into the design and development of healthcare services, as well as clinical guideline documents, can enable greater awareness and appreciation of the need for clear and understandable patient education resources.

Patient education reduces the uncertainty regarding symptoms, management, and prognosis; provides appropriate risk information; and promotes patient empowerment and self-efficacy, whereby the patient plays a leading role in their own health management. This is particularly relevant in the case of paediatric cardiomyopathy patients in which the parent's and child's environment should be appropriately addressed.

The process of patient education can be classified into four components: assessment, planning, implementation, and evaluation.<sup>104</sup> Assessment of the patient, including literacy levels, concerns, priorities, learning preferences, and potential barriers to learning, is crucial to personalizing education. The planning of patient education strategies involves setting mutual goals for the healthcare provider and the patient, and developing an education plan that uses evidence-based teaching strategies with a focus on patient's needs and priorities. Implementation of the plan should be adjusted according to how the patient responds and their changing learning needs. Finally, the plan should be evaluated to determine where learning has been achieved. The patient education process should be repeated accordingly. A comprehensive approach is shown in *Table S9*.

#### Table S8 Cardiomyopathy units: requirements and skills

Requirements	Skills
<b>Integrated cardiology/paediatric cardiology unit</b> (should include adult and paediatric cardiovascular specialists with expertise in cardiac genetics and cardiomyopathies)	Ability to perform a family history and draw and interpret a family pedigree Ability to perform or interpret appropriate diagnostic tests and examine diagnostic clues to define the nature of the cardiomyopathy and its underlying aetiology Clinical Nurse Specialists and clinic co-ordinators to facilitate scheduling of new patients and families and arrange appropriate follow-up
Dedicated outpatient clinic	Capacity to see and follow-up cardiomyopathy patients and their families in a dedicated outpatient clinic
Family screening	Ability to carry out a complete clinical screening of relatives (adults and children)
<b>Diagnostic imaging unit</b> (with experience in and capacity to carry out studies on cardiomyopathies, including basal, stress, and transoesophageal echocardiography; CMR; and cardiac CT [whether integrated or not into the cardiology unit])	Performance and interpretation of echocardiography with focus on the evaluation of myocardial morphology, outflow geometry, septum morphology, valve apparatus anatomy, presence of LVOTO, mitral valve and/or papillary muscle anomalies, right ventricle myocardium, Doppler and strain studies, etc. Performance and/or interpretation of CMR with special focus on the characterization of myocardial tissue with parametric mapping methods (quantification of T1, T2, and extracellular volume fraction), LGE, etc. Performance and/or interpretation of cardiac CT scans
<b>Nuclear imaging unit</b> (with experience in cardiac involvement of systemic diseases [amyloidosis, sarcoidosis, etc.]; may not be co-located)	Ability to interpret cardiac scintigraphy studies and PET scans
<b>Genetics unit</b> (with experience in cardiomyopathies and links to a clinical genetic testing laboratory, with access to genetic counselling)	Ability to discuss genetic testing options with patients and their families, as well as the impact on the patient and/or family Access to clinical genetic testing encompassing genes with definitive gene- disease association with the condition tested for Ability to interpret the genetic result and differentiate the clinical implications of different variant classification for the patient and their family Ability to perform cascade genetic testing of at-risk relatives Access to genetic counselling for the patient and all families seen with an inherited cardiomyopathy Access to pre- and post-test genetic counselling for all patients undergoing genetic testing
<b>Catheterization and interventional cardiology unit</b> (diagnostic and therapeutic procedures including structural procedures, with experience in ASA and an adequate number of procedures according to guidelines)	Ability to perform and interpret invasive haemodynamic studies to assess cardiac performance and intracardiac pressures in cardiomyopathies and valve disease as well as coronary angiography to detect haemodynamically significant epicardial coronary disease as contributors to myocardial ischaemia or heart failure Performance and/or interpretation of coronary angiograms of HCM patients that allow for correct identification of a septal perforator branch with compatible anatomy for ASA Ability to contextualize the clinical condition of patients, determine the need for interventions, and the likelihood of safe and effective septal reduction with ASA
<b>Electrophysiology and arrhythmia unit</b> (with experience in electrophysiological studies, ventricular tachycardia and atrial fibrillation ablation, and device implantation [pacemakers, resynchronization devices, and ICDs])	Ability to interpret other arrhythmia screening tools (e.g. implantable loop recorders), perform basic device troubleshooting, and deactivate defibrillator therapy and pacemaker functions Accurate selection of suitable patients for defibrillator therapy/ resynchronization based on guidelines

Cardiac surgery unit (with experience in myectomies and mitral valve repair for the management of obstructive HCM with an adequate number of procedures according to guidelines; may be in another reference centre) Active heart transplant and ventricular assist device programme that allows for a response to and continuity for needs derived from the treatment of patients with cardiomyopathies (may be in another reference centre)	Ability to determine the need for effective septal reduction with myectomy and or mitral valve repair Ability to detect patient deterioration and offer referral for heart transplantation when necessary Ability to offer ventricular assist device therapy or access to such a centre
Pathological anatomy unit (with experience in cardiomyopathies) Forensic pathologist unit (directly involved in the study of SCDs in the community)	Ability to understand the microscopic study of patient tissue samples in order to arrive at a diagnosis of the disease Ability to develop protocols between forensics and the cardiomyopathy unit for study of SCDs that occur in the community in an out-of-hospital setting
<b>Clinical psychological support</b> (from healthcare professionals with appropriate qualifications in the psychotherapeutic treatment of emotional and behavioural disorders caused by inherited conditions and SCD in relatives)	Psychological support should be offered to the patient or the family, with referral to an appropriately qualified healthcare professional Access to clinical psychological expertise for families with psychological difficulties, e.g. following a young SCD in the family; and patients adjusting to a new diagnosis, ICD therapy, possible medical restrictions on their daily routine, and the psychological aspects of living with a condition for the foreseeable future (aided by medication, ICD, transplant, etc.)
Research, teaching, and educational capacity	Research capacity, and demonstrated research experience in the field of cardiomyopathies, at national and international levels Capacity to carry out teaching and educational activities related to the area of cardiomyopathies, aimed at improving the knowledge and technical capacity of healthcare providers involved in the same chain of care within and outside of the provider facility, such as continuing medical education and distance learning
Telemedicine and other e-health tools	Ability to exchange expertise with other healthcare providers and support them Established procedures and a framework for ensuring the management, safeguarding, and exchange of medical data, including established outcomes, process indicators, and patient registers for the specific area of expertise in accordance with EU data protection legislation Ability to foster the use of telemedicine and other e-health tools within and outside their facilities, by fulfilling the minimum interoperability requirements and, when possible, using agreed standards and recommendations <sup>102</sup>

ASA, alcohol septal ablation; CMR, cardiac magnetic resonance; CT, computed tomography; EU, European Union; HCM, hypertrophic cardiomyopathy; ICD, implantable cardioverter defibrillator; LGE, late gadolinium enhancement; LVOTO, left ventricular outflow tract obstruction; PET, positron emission tomography; SCD, sudden cardiac death.

#### Table S9 Educational steps

Step	Examples					
Assessment						
Demographic data	• Enquire about the patient's level of education, and cultural and religious beliefs and values.					
Patient's concerns and priorities	<ul> <li>Enquire what the patient is most concerned about.</li> <li>Enquire what knowledge the patient has regarding the cardiomyopathy they are being assessed for or affected by.</li> <li>Ask the patient about level of knowledge of the condition in their close social environment.</li> </ul>					
Patient's interests	• What does the patient want to understand?					
Learning preferences	• Enquire about the patient's preferred mode of learning (e.g. reading, video, verbal, demonstration, app).					
Assessment tools	• Questionnaires or other assessment tools can be used to establish the level of knowledge pre- and post-patient education activity.					
Planning						
Set mutual goals	<ul> <li>Set goals according to the context. For example, educate the patient so that they are able to inform family members about the condition.</li> <li>Aim to address individual health literacy needs and consider disabilities that might affect this.</li> <li>Offer education according to the patient's interests and learning preferences.</li> </ul>					

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Develop an educational plan	<ul> <li>Avoid medical jargon.</li> <li>Use simple, direct messaging.</li> <li>Create objectives and a step-wise approach.</li> <li>Create communities that promote patient education.</li> <li>Provide both written material and videos, but also engagement events that allow faster feedback in learning.</li> </ul>
Create resources	<ul> <li>Documents.</li> <li>Videos to watch asynchronously.</li> <li>Creation of mobile apps, if relevant.</li> <li>Decision aids to be used in shared decision-making (e.g. decision on genetic testing or ICD implantation).</li> <li>Illustrations, 3D models.</li> <li>Accommodate patients who are visually or hearing impaired.</li> <li>Make available in languages other than English, depending on the location.</li> </ul>
Implementation	
Focus on the patient	<ul> <li>Plain language.</li> <li>Active listening.</li> <li>Allow for interactive learning (e.g. encourage patients to ask questions).</li> <li>Be respectful and empathetic.</li> </ul>
Follow key educational principles	<ul> <li>Frame the objectives to be clear from the beginning.</li> <li>Create small educational snippets/segments that are easily understood and digestible.</li> <li>Create clear summaries and review key points.</li> </ul>
Adjust teaching	<ul> <li>Use analogies that make sense in terms of the context and the patient's profile.</li> <li>Allow enough time to reiterate information on more challenging topics.</li> <li>Allow for hands-on practice (e.g. have a model of an ICD to demonstrate, and allow the patient to manipulate when discussing ICDs).</li> <li>Use multiple teaching methods depending on the context.</li> </ul>
Evaluation	
Direct patient input	• Have direct conversation with patients to get feedback for any educational activity.
Patient outcomes	<ul> <li>Measure adherence to therapy plans.</li> <li>Assess re-admission rates.</li> <li>Assess engagement with genetic counselling.</li> <li>Assess adherence to lifestyle changes.</li> </ul>

3D, three-dimensional; ICD, implantable cardioverter defibrillator.

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