A patient with atrioventricular block and a family history of dilated cardiomyopathy

Lamin A/C cardiomyopathy: case report and review of the literature

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Summary

A 45-year-old man known for bradycardia underwent cardiac evaluation because of complete atrioventricular block. His mother underwent heart transplantation at the age of 60 for dilated cardiomyopathy. Echocardiography revealed mild left ventricular dilatation, and a subnormal left ventricular ejection fraction (LVEF) of 50% due to septal hypokinesia. Cardiac magnetic resonance imaging showed linear mid-myocardial late gadolinium enhancement of the basal septum, indicating myocardial fibrosis. We decided to implant a dual chamber pacemaker programmed in DDD modality. On the basis of the clinical history we suspected a lamin A/C (LMNA) mutation cardiomyopathy. A known pathogenic heterozygote mutation c.1129C>T (p.Arg377His) in the LMNA gene was identified, confirming the diagnosis of LMNA cardiopathy. At this time we suggested an upgrade to an implantable cardioverter-defibrillator in order to prevent sudden cardiac death, but the patient refused. Eighteen months after the pacemaker implantation the patient is alive and well.

Key words: dilatative cardiomyopathy; lamin A/C

Figure 1: Sinus rhythm 78 bpm with 2:1 atrioventricular block (ventricular rate 39 bpm) and left anterior fascicular block.
ventricular dilatation (left ventricular end-diastolic diameter 62 mm), and a subnormal left ventricular ejection fraction (LVEF) of 50% due to septal hypokinesia. Cardiac magnetic resonance imaging (CMR) confirmed the mild left ventricular dilatation and showed linear mid-myocardial late gadolinium enhancement of the basal septum, indicating myocardial fibrosis (fig. 3). On the basis of the family history of DCM, the complete AVB, the mild left ventricular dilatation with sub-nor-
mal LVEF and linear mid-myocardial septal fibrosis, we suspected cardiomyopathy due to a lamin A/C (LMNA) gene mutation.

We decided to implant a dual chamber pacemaker pro-
grammed in DDD modality (fig. 4) and performed ge-
netic testing using the TruSight Cardio Panel (Illumina, San Diego, USA), which includes 176 genes. All associ-
ated mutations/variants were confirmed by direct Sanger sequencing. A known disease-causing hete-
rozygote missense mutation c.1129C>T (p.Arg377His) in the LMNA gene was identified, confirming the diagno-
sis of LMNA cardiopathy. Six months after pacemaker implantation the patient remained asymptomatic. Left ventricular dimension and function were unchanged. The pacemaker memory showed an episode of atrial arrhythmia (atrial flutter and atrial fibrillation) lasting 15 days, and several short episodes of relatively slow

Figure 2: Sinus rhythm with complete atrioventricular block and ventricular escape rhythm 39 bpm.

Figure 3: Post-contrast inversion recovery gradient echo images (late gadolinium enhancement) in a four-chamber (A) and short axis view (B), showing linear, mid-myocardial enhancement of the basal ventricular septum, consistent with fibrosis.
(150–180 bpm) nonsustained ventricular tachycardia (fig. 5). Beta-blocker therapy was initiated but after a few weeks was stopped because of side effects. At this time the question was: “What to do? Upgrade to a defibrillator (ICD)? To a resynchronisation system (CRT)? Close clinical follow-up?”

After extensive discussion with the patient, and considering that he had several risk factors for malignant ventricular arrhythmia, we suggested an upgrade to an ICD in order to prevent sudden cardiac death (SCD). Nevertheless, the patient refused our proposal. Eighteen months after the pacemaker implantation, the patient is alive and well. He has two sisters, who are asymptomatic with a normal ECG and echocardiogram. A genetic analysis has been planned.

**Discussion**

DCM is characterised by a dilated left ventricle with systolic dysfunction that is not caused by ischaemic or valvular heart disease [1]. The prevalence is about 1 DCM case in 2500 individuals [2]. DCM is currently responsible for approximately 10 000 deaths and 46 000 hospitalisations each year in the United States [3]. Despite a comprehensive evaluation, about 50% of cases lack an underlying diagnosis and are classified as idiopathic DCM [4]. Familial DCM accounts for up to 50% of all cases of idiopathic DCM [5]. LMNA gene mutations are one of the most frequent genetic abnormalities involved in DCM and it has been estimated that LMNA mutations cause up to 10% of familial DCM [6]. Lamin A and C are both encoded by the LMNA gene, which is localised to chromosome 1q21.2-q21.3 [7]. Lamins are type V intermediate filament proteins that are able to polymerise and form the nuclear lamina, an organised meshwork that lies between the inner nuclear membrane and the chromatin [8]. Mutations in LMNA are highly penetrant and may cause severe and progressive cardiopathy in a relevant proportion of patients. Furthermore, patients carrying a mutation in the LMNA gene may have one of several forms of muscular dystrophy such as Emery-Dreifuss muscular dystrophy [9], autosomal dominant limb girdle muscular dystrophy [10], or sensory and motor axonal neuropathy Charcot-Marie-Tooth type 2 [11].

LMNA mutations are associated with cardiac abnormalities characterised by arrhythmias (sinus node dysfunction, AVB, atrial and ventricular arrhythmias) and DCM leading to heart failure [12, 13]. Fatkin et al. [14] evaluated the clinical features of LMNA mutations in 39 affected patients with cardiac involvement. The study showed that the onset of disease occurred in middle age (mean age 38 years, range 19–53 years). Eighty-seven percent had sinus-node dysfunction or atrioventricular disturbances (sinus bradycardia, or first-, second- or third-degree heart block). Atrial fibrillation or atrial flutter were present in 59% of affected people and 64% had DCM. Pacemakers were implanted, owing to high-grade AVB or brady-arrhythmias, in 54% of affected patients. Often in LMNA cardiopathy, conduction disturbances or arrhythmia anticipate left ventricular dysfunction [12–15]. It was the case in our
patient with AVB, atrial arrhythmia and nonsustained ventricular tachycardia, but preserved LVEF. In addition to history, ECG and echocardiography, CMR plays an important role in determining cardiac involvement in LMNA cardiomyopathy. Holmström et al. [16] showed that 88% of patients with LMNA cardiomyopathy had left ventricular myocardial fibrosis. The pattern of enhancement was typically linear and less than 50% of the area of the segment. In all the patients, late gadolinium enhancement occurred in the basal or mid-ventricular septum, which strongly correlated with segmental wall motion abnormalities. Our patient had exactly this typical pattern. Patient with LMNA cardiopathy have a significantly worse prognosis than other patients with idiopathic DCM [12], with a high incidence of phenotypic progression and adverse arrhythmic and nonarrhythmic events during long-term follow-up. Besides heart failure due to left ventricular dysfunction, arrhythmias are a major threat, and cardiac arrest due to bradyarrhythmia or ventricular arrhythmia has been reported in up to 50% of patients [12–15, 17–19]. Life expectancy is around 50 to 60 years [15, 19]. There is consensus regarding the efficacy of ICD in the primary and secondary prevention of SCD in patients with cardiovascular diseases [20]. In LMNA cardiomyopathy, the risk factors of SCD are incompletely elucidated and have been correlated to several clinical and genetic factors. In a study including 94 patients, New York Heart Association (NYHA) functional class, competitive sporting activity and type of mutation predicted the incidence of heart failure and ventricular arrhythmia [17]. In a multicentre European registry of 269 patients, four independent risk factors for malignant ventricular arrhythmia were identified: nonsustained ventricular tachycardia, LVEF <45%, male sex and non-missense mutations of the LMNA gene [19]. In this study, malignant ventricular arrhythmias occurred only in subjects with at least two of these risk factors and there was a cumulative risk per additional risk factor. Male sex, non-missense mutations and LVEF ≤50% were also associated with malignant ventricular arrhythmias in a recently published multicentre study of 122 patients [15]. In two studies of 47 and 41 patients, malignant ventricular arrhythmias were correlated with atrioventricular conduction disorders even when LVEF

Figure 5: Electrograms from the pacemaker memory. A: Nonsustained ventricular tachycardia (6 complexes, 160–180 bpm). B: Beginning of the episode of atrial arrhythmia. A = atrial electrogram; V = ventricular electrogram; MC = marker channel
was preserved [21, 22]. It was suggested that implantation of an ICD in patients requiring a pacemaker should be considered [21]. The presence of myocardial fibrosis in the interventricular septum may be the mechanism of the relationship between atrioventricular conduction disease and ventricular arrhythmias [21, 22]. No study so far has evaluated the role of electrophysiological testing with programmed electrical stimulation or other testing (e.g., stress testing, etc.) in the risk stratification of patients with lamin A/C cardiopathy. According to the 2015 European Society of Cardiology Guidelines for the prevention of SCD [20], the implantation of an ICD is a class IIa recommendation in patients with DCM due to LMNA mutations and clinical risk factors (male sex, LVEF <45%, nonsustained ventricular tachycardia, non-missense mutations), and a class IIb recommendation in patients requiring a pacemaker. Our patient had several risk factors for malignant ventricular arrhythmias: male gender, advanced AVB and nonsustained ventricular tachycardia. Therefore, we proposed an upgrade to an ICD.

In patients with LMNA cardiomyopathy needing a pacemaker, implanting a CRT should be considered, even with preserved LVEF, because the progressive nature of the disease leads in a substantial number of patients to left ventricular dysfunction and heart failure. However, no study has addressed the preventive role of CRT in these patients so far.

We have described a patient with LMNA cardiomyopathy discovered through a high grade AVB. This case highlights several important elements. First, in a young patient with advanced AVB accompanied by left ventricular dysfunction or family history of DCM, one should think of LMNA cardiopathy; genetic analysis can confirm the disease. Second, LMNA cardiopathy has a poor prognosis, and risk stratification should be individualised on the basis of several clinical, instrumental and genetic factors. In patients deemed at risk of SCD, an ICD should be considered. Finally, genetic screening of family members should be offered. Mutation-positive, phenotype-negative patients should have close electrical and functional follow-up, given the highly penetrant nature of the disease.

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References
The full list of references is included in the online version of the article at www.cardiovascmed.ch.