Scientific letters

Phenotypic Patterns of Cardiomyopathy Caused by Mutations in the Desmin Gene. A Clinical and Genetic Study in Two Inherited Heart Disease Units



Patrón fenotípico de las miocardiopatías por mutaciones en el gen de la desmina. Estudio clínico y genético en dos unidades de cardiopatías familiares

To the Editor,

Desminopathies are a largely autosomal dominant group of rare diseases caused by mutations in the desmin gene. Because desmin is the main component of intermediate filaments in cardiac, skeletal, and smooth muscle and of Purkinje fibers, these conditions are characterized by skeletal myopathy and cardiomy-opathy (mainly restrictive) with arrhythmias or conduction disorders.^{1–3}

The aim of our present study was to analyze the genotype and phenotype of patients with desmin mutation-related cardiomyopathy. Because published series normally include few patients, any further contributions will boost our understanding of this disease.

In 2 centers with 819 studied families, we analyzed all individuals found to have a desmin mutation after a phenotypeguided genetic study (restrictive cardiomyopathy/dilated cardiomyopathy with a restrictive pattern, families with high rates of pacemaker implants, skeletal myopathy, and/or creatine kinase elevation). Gene sequencing was performed using Sanger or nextgeneration sequencing. A pathogenic mutation was defined as any mutation involving an amino acid change from the reference sequence that met 3 criteria: it segregated with affected members of the family, it was not present in 200 chromosomes of healthy unrelated individuals, and the affected residue was conserved among species and desmin isoforms.

We studied 20 patients from 4 families, identifying 3 pathogenic mutations: lle367Phe (2 families, Sanger), Pro419Ser (Sanger), and Arg415Gln (next-generation sequencing). Of these, 16 had desminopathy (including 2 obligate carriers who died of cardiomyopathy, without genetic confirmation) and 4 were young unaffected carriers (Table). The mean age at diagnosis was 35 ± 15 years. Two

Table

Clinical and Genetic Data of the Families and Patients Included in this Study

Family	Desmin mutation	Age/ sex	Cardiac phenotype	Age at diagnosis, y	FHSD	Presentation	NHYA	AF	VT	LVEF	Pacemaker/ age, y	ICD	CK elevation	Skeletal myopathy	Death
1/Proband	I367F	40/M	RM	36	Y	Dyspnea	3	Ν	Ν	25	Y/37	Y ^c	Y	Y	Y
1/Cousin	I367F	43/F	RM	22		Syncope	2	Y	Ν	60	Y/22	Ν	Ν	Ν	Ν
1/Mother	I367F	67/F	RM	49		Dyspnea	2	Y	Ν	58	Y/49	Ν	N	Ν	Ν
1/Brother		28/M	?	28		?	?	?	?	?	Ν	Ν	?	?	Y
1/Nephew ^a	I367F	21/M	Normal ^a				1	Ν	Ν	60	Ν	Ν	Ν	Ν	Ν
1/Cousin ^a	I367F	34/F	Normal ^a				1	Ν	Ν	60	Ν	Ν	Ν	Ν	Ν
2/Proband	I367F	34/M	RM	27	Y	Dyspnea	2	Ν	Ν	65	Y	Ν	Ν	Y	Ν
2/Brother	I367F	39/M	RM	37		Dyspnea	3	Ν	Ν	60	Y	Ν	N	Y	Ν
2/Aunt	I367F	53/F	Normal	50		Myopathy	2	Ν	Ν	60	Ν	Ν	N	Y	Ν
2/Great-aunt	I367F	73/F	RM	?		Myopathy	1	Ν	Ν	?	Ν	Ν	?	Y	Y
2/Father		48/M	?	?		?	?	?	?	?	Y	Ν	?	Y	Y
2/Half-sister	I367F	26/F	Normal ^a				1	Ν	Ν	60	Ν	Ν	Ν	Ν	Ν
2/Auncle	I367F	46/M	RM	?		?	2	?	?	?	Y	Ν	Y	Y	Ν
3/Proband	P419S	28/M	RM	24	Ν	Myopathy	2	Ν	Y	55	Y/26	Y ^d	Y	Y	Ν
3/Mother	P419S	56/F	RM	42		Myopathy	2	Ν	Ν	60	Y/48	Ν	Ν	Y	Ν
4/Proband	R415E	30/M	LVAC	30	N	Sport-related SCD	1	Ν	Y	?	Ν	N	?	Ν	Y
4/Father	R415E	69/M	DC	52		Palpitations	3 ^b	Y	Y	26	Y/52	Y ^e	N	N	Ν
4/Brother	R415E	26/M	NSC	26		Family screening	1	N	N	70	Ν	N	N	N	N
4/Uncle	R415E	66/M	RM	65		Family screening	2	Y	Y	50	Ν	N	N	N	Y
4/Cousin	R415E	37/F	Normal ^a	32		Family screening	1	N	N	70	Ν	N	Ν	Ν	N

AF, atrial fibrillation; CK, creatine kinase; DC, dilated cardiomyopathy; F, female; FHSD, family history of sudden death; I367F, Ile367Phe; ICD, implantable cardioverterdefibrillator; LVAC, left ventricular arrhythmogenic cardiomyopathy; LVEF, left ventricular ejection fraction; M, male; MR, restrictive cardiomyopathy; NHYA, New York Heart Association classification of dyspnea; NSC, nonspecific cardiomyopathy; NSVT, nonsustained ventricular tachycardia; P419S, Pro419Ser; R415E, Arg415Glu; SCD, sudden cardiac death: VT, ventricular tachycardia.

^a Genotype⁺, phenotype⁻ carriers.

^b Heart transplant.

ICD indications.

^c NSVT + atrioventricular block.

^d NSVT + symptomatic trifascicular block.

^e NSVT + bradycardia-tachycardia syndrome.

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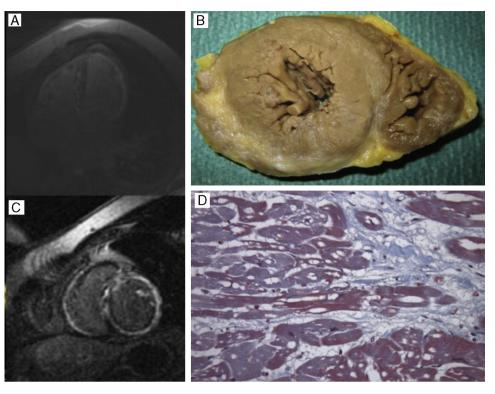


Figure. A and C: Cardiac magnetic resonance imaging of the proband of family 1 (*DES* Ile367Phe mutation) who shows biatrial and biventricular dilatation and extensive transmural fibrosis in the free wall of the left ventricle, septum, and right ventricle. B and D: Necropsy of the proband of family 4 (*DES* Arg415Glu mutation), revealing normal myocardial thicknesses and diameters, fiber hypertrophy, and severe subepicardial fibrosis, cytoplasmic vacuolization, and myofibrillar loss, which are all compatible with left ventricular arrhythmogenic cardiomyopathy.

families reported several relatives with pacemakers who died of cardiac arrest in their third or fourth decade of life. The most common symptoms were heart failure with an atrioventricular conduction disorder. The most common echocardiographic pattern was restrictive cardiomyopathy, with normal myocardial thickness or mild hypertrophy (11.4 ± 2.4 mm) and preserved left ventricular (LV) systolic function, except for 2 patients with severe LV dysfunction and another with LV arrhythmogenic cardiomyopathy (Figure). Cardiac magnetic resonance imaging was performed in 4 patients, who all had extensive transmural fibrosis; 7 of 11 men and 3 of 5 women required a pacemaker at early ages due to atrioventricular block, including 2 men who required ventricular resynchronization and 3 who required an implantable cardioverter-defibrillator due to nonsustained ventricular tachycardia.

The degree of skeletal myopathy was variable and mainly involved distal and progressive muscle weakness and atrophy of the lower limbs. Penetrance was high from the third decade of life, with more severe expression in men.

Six patients died during follow-up: 1 from cardiac arrest (a 30-year-old man with LV arrhythmogenic cardiomyopathy), 2 from heart failure (a 40-year-old man with severe LV dysfunction and a 66-year-old man with restrictive cardiomyopathy), and 3 from unknown causes (1 had a pacemaker).

Few data are available on these 3 mutations because they have only been described in the literature in 4 families: 1 family with lle367Phe,^{1,2} 3 with Pro419Ser,^{1,2,4} and none with Arg415Glu.

The lle367Phe mutation has previously been described as being pathogenic.² Segregation of the mutation with the disease was checked in our 2 families and the phenotype was similar to that already described: restrictive cardiomyopathy, atrioventricular block, and skeletal myopathy.

The Pro419Ser mutation has also been described and shows a more variable phenotype: marked neurological involvement with predominantly distal muscle weakness and a nasal voice, restrictive cardiomyopathy, and atrioventricular block and, in the other family, right ventricular arrhythmogenic cardiomyopathy.^{2,4} This mutation is located in the tail of the gene, and cardiomyopathy typically appears alone or precedes a skeletal condition. In contrast, the Ile367Phe mutation is located in a hot spot and is the most severe mutation.²

The Arg415Glu mutation in exon 6 has not been described before and affects splicing. Cosegregation was seen in our family. The proband died of sudden cardiac death, with histopathology findings of LV arrhythmogenic cardiomyopathy, and is the first published patient with this association. This mutation was seen in most of our patients with severe cardiomyopathy without skeletal myopathy.

Notably, these patients visited the clinic with other diagnoses, such as dilated hypertensive heart disease, or hypertrophic cardiomyopathy. The fourth proband was diagnosed after sudden cardiac death. The cause was suggested by a specialized study and confirmed by genetic analysis. A subsequent family study diagnosed 14 carriers, 10 with restrictive cardiomyopathy and/ or skeletal myopathy and 4 unaffected young carriers. The study also allowed 16 families to be excluded from follow-up.

In conclusion, desminopathies commonly present as restrictive cardiomyopathy with heart failure in the third or fourth decade of life or early advanced atrioventricular block requiring pacemaker implantation and, in some patients, additionally, implantable cardioverter-defibrillator due to nonsustained ventricular tachycardia.

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The Descending Septal Artery: Description of This Infrequent Coronary Anatomical Variant in Three Different Clinical Scenarios

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La arteria septal descendente: descripción de esta variante anatómica coronaria poco frecuente en tres escenarios clínicos diferentes

To the Editor,

Septal coronary branches arising from the right coronary artery (RCA) or the right coronary sinus, known as descending septal artery (DSA)¹ or Bonapace's branch,² have rarely been described. However, the DSA might play an important role in certain situations, highlighting the need for its proper identification and evaluation. The present series reviews 3 cases of DSA identified during coronary angiography in different clinical scenarios.

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A 60-year-old man was referred for coronary angiography due to exertional chest pain and a significant lesion in the RCA detected by coronary computed tomography angiography. Through the right femoral artery, a multipurpose catheter was used to cannulate the RCA. A DSA emerging from a common ostium with the RCA was visualized (Figure 1A). The RCA lesion was successfully stented.

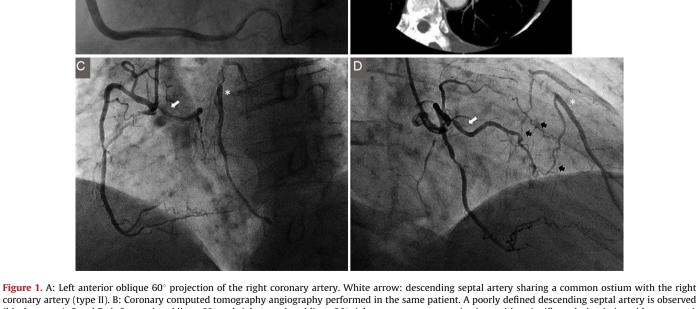


Figure 1. A: Left anterior oblique 60° projection of the right coronary artery. White arrow: descending septal artery sharing a common ostium with the right coronary artery (type II). B: Coronary computed tomography angiography performed in the same patient. A poorly defined descending septal artery is observed (black arrows). C and D: Left anterior oblique 60° and right anterior oblique 30° right coronary artery projections with a significant lesion in its mid-segment. A descending septal artery originating from the proximal segment (type I, white arrows) provides collaterals to a chronically occluded left anterior descending (asterisk) through several septal branches (black arrows).

