Table S2-- Comparison of task force criteria and patients' conditions

2010 Task force criteria [1]	ARVC patient
I. Global or regional dysfunction and structural alterations	Major
Major	The patient had regional RV
By 2D echo:	dyskinesia and PLAX/BSA
• Regional RV akinesia, dyskinesia, or aneurysm and one of the following (end	RVOT ≥21 mm/m ² by 2D
diastole):	echo.
— PLAX RVOT ≥32 mm (corrected for body size [PLAX/BSA] ≥19 mm/m2)	
— PSAX RVOT ≥36 mm (corrected for body size [PSAX/BSA] ≥21 mm/m2)	
— or fractional area change ≤33%	
By MRI:	
Regional RV akinesia or dyskinesia or dyssynchronous RV contraction and	
one of the following:	
— Ratio of RV end-diastolic volume to BSA ≥110 mL/m2 (male) or ≥100	
mL/m2 (female)	
— or RV ejection fraction ≤40%	
By RV angiography:	
Regional RV akinesia, dyskinesia, or aneurysm	
Minor	
By 2D echo:	
• Regional RV akinesia or dyskinesia and one of the following (end diastole):	
— PLAX RVOT ≥29 to <32 mm (corrected for body size [PLAX/BSA] ≥16 to	
<19 mm/m2)	
— PSAX RVOT ≥32 to <36 mm (corrected for body size [PSAX/BSA] ≥18 to	
<21 mm/m2)	
— or fractional area change >33% to ≤40%	
By MRI:	

Regional RV akinesia or dyskinesia or dyssynchronous RV contraction and	
one of the following:	
− Ratio of RV end-diastolic volume to BSA ≥100 to <110 mL/m2 (male) or	
≥90 to <100 mL/m2 (female)	
— or RV ejection fraction >40% to ≤45%	
PLAX indicates parasternal long-axis view; RVOT, RV outflow tract; BSA, body	
surface area; PSAX, parasternal short-axis view; aVF, augmented voltage	
unipolar left foot lead; and aVL, augmented voltage unipolar left arm lead.	
II. Tissue characterization of wall	Undetected
Major	
 Fibrofatty replacement of myocardium on endomyocardial biopsy 	
• Residual myocytes <60% by morphometric analysis (or <50% if estimated),	
with the fibrous replacement of the RV free wall myocardium in \geq 1 sample, with	
or without fatty replacement of tissue on endomyocardial biopsy	
Minor	
\bullet Residual myocytes 60% to 75% by morphometric analysis (or 50% to 65% if	
estimated), with the fibrous replacement of the RV free wall myocardium in \geq 1	
sample, with or without fatty replacement of tissue on endomyocardial biopsy	
III. Repolarization abnormalities	No match
Major	
- Inverted T waves in the right precordial leads (V1, V2, and V3) or beyond in	Inverted T waves in leads V1,
individuals >14 years of age (in the absence of complete right bundle-branch	V2, V3, in individuals >14
block QRS ≥120 ms)	years of age in the presence
Minor	of complete right bundle-
- Inverted T waves in leads V_1 and V_2 in individuals >14 years of age (in the	branch block
absence of complete right bundle-branch block) or in V ₄ , V ₅ , or V ₆	
• Inverted T waves in leads V1, V2, V3, and V4 in individuals >14 years of age	
in the presence of complete right bundle-branch block	
IV. Depolarization/conduction abnormalities	Major
Major	Epsilon wave (reproducible
	low-amplitude signals

• Epsilon wave (reproducible low-amplitude signals between the end of QRS	between end of QRS complex
complex to the onset of the T wave) in the right precordial leads (V $_1$ to V $_3$)	to onset of the T wave) in the
Minor	right precordial leads (V $_1$ to
 Late potentials by SAECG in ≥1 of 3 parameters in the absence of a QRS 	V ₃)
duration of ≥110 ms on the standard ECG	
 Filtered QRS duration (fQRS) ≥114 ms 	
 Duration of terminal QRS <40 μV (low-amplitude signal duration) ≥38 ms 	
• Root-mean-square voltage of terminal 40 ms ≤20 μV	
 Terminal activation duration of QRS ≥55 ms measured from the nadir of the S 	
wave to the end of the QRS, including R', in V1, V2, or V3, in the absence of	
complete right bundle-branch block	
V. Arrhythmias	Minor
Major	Sustained ventricular
• Nonsustained or sustained ventricular tachycardia of left bundle-branch	tachycardia of unknown axis
morphology with a superior axis (negative or indeterminate QRS in leads II, III,	
and aVF and positive in lead aVL)	
Minor	
Nonsustained or sustained ventricular tachycardia of RV outflow configuration,	
left bundle-branch block morphology with the inferior axis (positive QRS in leads	
II, III, and aVF and negative in lead aVL) or of an unknown axis	
 > 500 ventricular extrasystoles per 24 hours (Holter) 	
VI. Family history	No match
Major	
• ARVC/D confirmed in a first-degree relative who met current Task Force	
criteria	
• ARVC/D confirmed pathologically at autopsy or surgery in a first-degree	
relative	
Identification of a pathogenic mutation† categorized as associated or probably	
associated with ARVC/D in the patient under evaluation	
Minor	

• History of ARVC/D in a first-degree relative in whom it was not possible or	
practical to determine whether the family member met current Task Force	
criteria	
• Premature sudden death (<35 years of age) due to suspected ARVC/D in a	
first-degree relative	
• ARVC/D confirmed pathologically or by current Task Force Criteria in a	
second-degree relative	
Definite diagnosis: 2 major, or 1 major and 2 minor, or 4 minor criteria from	2 major and 1 minor criteria
different categories; borderline diagnosis: 1 major and 1 minor, or 3 minor	
criteria from different categories; possible diagnosis: 1 major, or 2 minor criteria	
from different categories.	

References

1. Marcus FI, McKenna WJ, Sherrill D, Basso C, Bauce B, Bluemke DA, et al. Diagnosis of arrhythmogenic right ventricular cardiomyopathy/dysplasia: proposed modification of the Task Force Criteria. Eur Heart J. 2010; 31: 806-14.

Table S3. Primers for real-time PCR

Gene	Forward (5' to 3')	Reverse (5' to 3')
GAPDH	GGTGAAGGTCGGAGTCAACGGATTTGGTCG	GGATCTCGCTCCTGGAAGATGGTGATGGG
OBSCN	GCTGAGCTCCGTGTGGACT	CAGTGGTGGACTCTGTTCCCT
ANK1.5	CCAGATGAATGGTTACTCCTCAC	CAAGGGGATGGCGTCTAGG
POU5F1	CAGTGCCCGAAACCCACAC	GGAGACCCAGCAGCCTCAAA
NANOG	CAGAAGGCCTCAGCACCTAC	ATTGTTCCAGGTCTGGTTGC
α-actinin	TCCATCGGAGCCGAAGAAATC	GTGTCGGTGGATCAAAGCACA
plakoglobin	TCAGCAGCAAGGGCATCAT	TGGGTGTAAGTGGTGGTTTTCTT
plakophilin2	GCAAATGGTTTGCTCGATTT	GGCTGGTAATCTGCAATGGT
N-cadherin	CCTGCTTCAGGCGTCTGTAG	CTGCCTTTGTAGGTGGCCAC
connexin43	GGAATGCAAGAGAGGTTGAAAG	GGCATTTGGAGAAACTGGTAGA
desmoplakin	CAGTGGTGTCAGCGATGATGT	TGACGCTGGATATGGTGGAA
C/EBPa	CACGAAGCACGATCAGTCCAT	CGCACATTCACATTGCACAAG
FABP4	TGGTGGTGGAATGCGTCAT	GGTCAACGTCCCTTGGCTTA
PPARγ	AGGCCATTTTCTCAAACGAG	CCATTACGGAGAGATCCACG

Figure S1-Recording of implantable cardioverter defibrillator



Figure S2-- Alignment of sequences of the OBSCN gene from human, tupaia, vulpes, microtus,

nothoprocta, vulpes and zonotrichia.

*The asterisk represents conservative sequences. The red box is the location of the target gene mutation in the study.

Homo Tupaia Vulpes	GVSSTKAELRVDL TSTDYDTAADATESSSYFSAQGYLSSREQEGTESTTD GVSSTKAELRVDL TSTDYDTAADATETSSYFSAQGYLSSREQEGTESTTD GVSSTKAELRVDL TSTDYETAADATETSSYFSAQGYLSSREQEGTESTSE *********************************
	C L5218
Homo	SMGVSSTKAELRVDLTSTDYDTAADATESSSYFSAQGYLSSREQEGTEST
Microtus	SVGVSSTKAELRVELTSTDYDTAADATETSSYFSAQGYLSSREQEGTE
Nothoprocta	SMGVASTKAELRVDL
	*:**:********: <mark>**</mark> *****:*****:****:***:*
	L5218
Homo	INEDQQGGHQLIITAVVPADMGVYRCLAENSMGVSSTKAELRVDL
Vulpes	ISEDQQGGHQLIITAVVPADMGVYRCMAENSVGVSSTKAELRVDL
Zonotrichia	INEDQEGCHQLIITAVVPTDMGVYRCLAENNMGVASTKAELRVDL
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