



An autoantibody identifies arrhythmogenic right ventricular cardiomyopathy and participates in its pathogenesis

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Aims

Arrhythmogenic right ventricular cardiomyopathy (ARVC) is characterized by right ventricular myocardial replacement and life-threatening ventricular arrhythmias. Desmosomal gene mutations are sometimes identified, but clinical and genetic diagnosis remains challenging. Desmosomal skin disorders can be caused by desmosomal gene mutations or autoantibodies. We sought to determine if anti-desmosome antibodies are present in subjects with ARVC.

Methods and results

We evaluated ARVC subjects and controls for antibodies to cardiac desmosomal cadherin proteins. Desmoglein-2 (DSG2), desmocollin-2, and N-cadherin proteins on western blots were exposed to sera, in primary and validation cohorts of subjects and controls, as well as the naturally occurring Boxer dog model of ARVC. We identified anti-DSG2 antibodies in 12/12 and 25/25 definite ARVC cohorts and 7/8 borderline subjects. Antibody was absent in 11/12, faint in 1/12, and absent in 20/20 of two control cohorts. Anti-DSG2 antibodies were present in 10/10 Boxer dogs with ARVC, and absent in 18/18 without. In humans, the level of anti-DSG2 antibodies correlated with the burden of premature ventricular contractions ($r=0.70$), and antibodies caused gap junction dysfunction, a common feature of ARVC, *in vitro*. Anti-DSG2 antibodies were present in ARVC subjects regardless of whether an underlying mutation was identified, or which mutation was present. A disease-specific DSG2 epitope was identified.

Conclusion

Anti-DSG2 antibodies are a sensitive and specific biomarker for ARVC. The development of autoimmunity as a result of target-related mutations is unique. Anti-DSG2 antibodies likely explain the cardiac inflammation that is frequently identified in ARVC and may represent a new therapeutic target.

Keywords

Arrhythmogenic right ventricular cardiomyopathy • Biomarker • Autoantibody • Human desmoglein-2 • Boxer dog

Introduction

Arrhythmogenic right ventricular cardiomyopathy (ARVC) is a heritable heart muscle disease characterized by replacement of myocardium by fat and/or fibrosis, and arrhythmias arising usually from the right ventricular free wall.¹ It is a major contributor to sudden cardiac deaths (SCDs) in young adults,² athletes,³ and children and adolescents age 10–18 years,⁴ accounting for 11%, 22%, and 25% of SCD in these groups, respectively. Arrhythmogenic right ventricular cardiomyopathy is estimated to occur in one in 2000 individuals,⁵ and sudden deaths during sport are often attributed to this disorder.

Arrhythmogenic right ventricular cardiomyopathy is difficult to diagnose, but a set of Task Force criteria developed in 1994⁶ by the European Society of Cardiology, and revised in 2010,⁷ provides the current standard for identification. These criteria evaluate detailed imaging, pathology, electrocardiography, family history and genetics assessments, and the total number of major and minor criteria characterize patients as having definite, borderline, possible, or no ARVC. Task Force criteria assessments are complex and costly, must be repeated regularly and are still considered to miss 30% of affected individuals. Once diagnosed, risk stratification and therapy also pose clinical challenges that are addressed by additional guidelines.⁸

Mutations in 12 genes are associated with the development of ARVC or ARVC mimics^{9–19}; however, the genetic cause remains unidentified in approximately 50% of cases. Arrhythmogenic right ventricular cardiomyopathy is often characterized by incomplete penetrance and variable expression, with the disease appearing to skip generations or appear in isolated individuals. Activity and exercise clearly modify disease expression.²⁰ Where identified, genetic mutations most typically affect proteins of the desmosome, a structure that links heart muscle cells together at their intercalated discs. These abnormal desmosomes fail to mechanically link cardiomyocytes, resulting in myocardial replacement with fat and scar tissue either directly, or as a result of nuclear signalling altering cell fate.²¹ A common feature identified in ARVC, regardless of the gene cause, is secondary failure of gap junction electrical connections at the cardiomyocyte intercalated discs, resulting in cardiac conduction delay and ventricular arrhythmias.²² Frequently, there is also evidence for inflammation in the pathogenesis of ARVC. Lymphocytic infiltrates are frequently seen in myocardial specimens,²³ and the disease often demonstrates flares or hot phases of disease.²⁴

Outside of the heart, desmosomes also provide mechanical links between epithelial cells. Since skin desmosomal disease is caused by either mutations of desmosomal proteins or autoantibodies against these proteins,²⁵ we hypothesized that anti-desmosomal autoantibodies might contribute to ARVC as well.

In this study, we report the presence of anti-desmoglein-2 (DSG2) autoantibodies as a highly sensitive and specific biomarker identifying Task Force positive ARVC human subjects, as well as ARVC affected Boxer dogs. Our functional analyses suggest that anti-DSG2 autoantibodies are intrinsically involved in the pathogenesis of ARVC.

Materials and methods

Subjects and animals

The study protocol followed the ethical guidelines of the Declaration of Helsinki. The Research Ethics Board of the Hospital for Sick Children and

the Cantonal Ethical Committee Zurich, Switzerland, and all human participants gave written informed consent. Patients referred to these clinics for assessment of ARVC were evaluated using the 2010 ARVC Task Force criteria [including electrocardiograms (ECGs), signal-averaged ECGs (SAECGs), 24-h ambulatory monitoring, echocardiography, and magnetic resonance imaging] and only those meeting Definite or Borderline criteria (Hospital for Sick Children) or Definite criteria (Zurich validation cohort) were included. Sera were collected and frozen at -80°C until analysed. Sera from Zurich were transported on ice for evaluation. Normal Control sera were purchased from two commercial sources: Innovative Research Inc. (Novi, MI, USA) and from BBI Solutions (Cardiff, UK).

Optimization of western blot

We used a recombinant fusion protein of human DSG2 Fc Chimera (a portion of Fc region of human IgG attached to full DSG2 protein). This structure is commonly used in antibody studies to improve solubility and presentation of the protein. Though the attached Fc region is not the native Fc region of IgG, it may give some weak binding to the secondary antibody [Goat anti-human IgG-horseradish peroxidase (IgG-HRP) conjugate] and thus can appear as a false positive for negative samples. Therefore, we optimized the conditions for the western blot to avoid this weak positive reaction by testing different amounts of antigen, dilutions of Goat anti-human IgG-HRP, and exposure times for the enhanced chemiluminescence reaction (see [Supplementary material online](#)).

Western blot analysis

Western blot procedure was adopted from Chatterjee-Chakraborty and Chatterjee.²⁶ The analysis of serum from controls and patients were carried out using recombinant human DSG2, DSC2, and N-Cadherin proteins. The conditions used for our study to prevent false positive reactions for the attached Fc fragment in the fusion proteins were: (i) keeping the recombinant DSG2 fusion protein per lane to 100 ng, (ii) using the dilution of Goat anti-human IgG-HRP to 1:10 000, and (iii) maintaining the exposure time to X-ray film at 5 s. We have checked the specificity of the blotting method used by different prior control experiments as described (see [Supplementary material online](#)).

Desmoglein-2 antibody ELISA protocol

A direct ELISA was performed by first coating a micro-titre plate with recombinant human DSG2 protein and then exposing each well to diluted human sera (100 µL of 1:100 dilution) followed by washing. The resultant bound anti-DSG2 antibody was then assayed by anti-human IgG-HRP using tetramethylbenzidine chromophore to measure optical density (O.D.) at 450 nm wavelength. Details of the solutions, dilutions, incubations, and washing are provided in the [Supplementary material online](#).

Western blot analysis of synthetic desmoglein-2 oligopeptide library

Western blots with serum from control samples and patient samples were carried out using synthetic oligopeptides, made on the basis of the known sequence of human DSG2 protein. We used 112 sequential peptides of 15 amino acid (a.a.) residues (overlapping by five residues) to completely cover the DSG2 sequence from the N-terminal to C-terminal, and characterized the resultant binding as strong, moderate, weak, or faint (see also [Supplementary material online](#)).

Statistical analysis

To correlate the phenotype to the band intensities of the patients, the ambulatory burden of premature ventricular contractions (PVCs) was

Table 1 Phenotypic and genetic data for patients included in the original and validation cohorts

Patients	Age ^a	Sex	I ^b	II ^c	III ^d	IV ^e	V ^f	VI ^g	Total (major, minor)	Diagnosis	Biomarker	Therapy ^h	PVC ⁱ	Genetics ^j
HCG001	15	M	1,0	0,0	0,0	0,1	0,1	0,0	1,2	ARVC	Yes	Nadolol	2600 ^k	No Mut ^l
HRC0168	12	M	0,1	0,0	0,0	0,0	0,1	1,0	1,2	ARVC	Yes	Levothyroxin	921	DSC2
HRC0160	34	F	1,0	0,0	0,0	0,0	0,0	1,0	2,0	ARVC	Yes	Bisoprolol	0 ^k	No Mut
HRC0118	34	F	1,0	0,0	1,0	—	—	1,0	3,0	ARVC	Yes	Metoprolol, Sotalol	854	PKP2
HRC0101	37	F	1,0	0,0	1,0	—	0,0	0,1	2,1	ARVC	Yes	Bisoprolol	114 ^k	No Mut
HRC0099	17	F	1,0	0,0	1,0	0,0	—,1	1,0	3,1	ARVC	Yes	Nadolol	10 266	PKP2
HRC0098	54	F	1,0	0,0	0,0	0,0	0,0	1,0	2,0	ARVC	Yes	—	—	PKP2
HRC0085	13	M	0,1	0,0	0,0	0,1	0,0	1,0	1,2	ARVC	Yes	—	—	PKP2
HRC0084	16	F	1,0	0,0	1,0	0,0	0,1	1,0	3,1	ARVC	Yes	Atenolol	3091	PKP2
HRC0083	37	M	—	0,0	0,1	0,1	—	1,0	1,2	ARVC	Yes	Sotalol, L-thyroxin	—	DSP
HRC0179	46	F	1,0	—	1,0	0,1	0,1	1,0	3,2	ARVC	Yes	—	667	PKP2
HRC0066	42	F	1,0	—	1,0	0,0	0,1	—	2,1	ARVC	Yes	None	9126	—
HRC0159	9	M	0,0	0,0	0,0	0,1	0,0	1,0	1,1	Bord. ^m ARVC	Yes	None	0	No Mut
HRC0086	37	F	—	0,0	0,0	1,0	0,0	1,0	1,1	Bord. ARVC	No	—	—	PKP2
HRC0090	11	F	0,0	0,0	0,0	0,1	—	1,0	1,1	Bord. ARVC	Yes	None	0	DSG2
HRC0174	44	M	0,0	0,0	0,0	0,1	0,0	1,0	1,1	Bord. ARVC	Yes	None	26	PKP2
HRC0189	12	M	0,1	—	0,0	0,0	0,0	1,0	1,1	Bord. ARVC	Yes	None	1	No Mut
HRC0109	15	M	0,0	0,0	0,0	0,1	0,0	1,0	1,1	Bord. ARVC	Yes	None	0	No Mut
HRC0108	40	M	0,0	0,0	0,1	0,0	0,0	1,0	1,1	Bord. ARVC	Yes	None	—	No Mut
HRC0087	36	F	—	—	—	0,1	0,0	1,0	1,1	Bord. ARVC	Yes	—	0	No Mut
ZH-001	60	M	1,0	0,0	1,0	0,1	0,1	1,0	3,2	ARVC	Yes	Amiodarone	124 ^k	DSG2
ZH-007	64	M	1,0	0,0	1,0	0,1	1,0	0,0	3,1	ARVC	Yes	Betablocker, Amiodarone	238 ^k	No Mut
ZH-013	50	M	1,0	0,0	1,0	1,0	0,1	1,0	4,1	ARVC	Yes	Betablocker, Amiodarone	1673 ^k	PKP2
ZH-015	26	M	1,0	0,0	1,0	0,0	0,1	1,0	3,1	ARVC	Yes	Sotalol	2511 ^k	PKP2
ZH-016	46	M	1,0	1,0	0,1	1,0	0,1	0,0	3,2	ARVC	Yes	Sotalol, ACE-I/ARB, Aldosterone antagonist	90	TTN
ZH-017	52	M	1,0	0,0	0,1	0,1	1,0	0,0	2,2	ARVC	Yes	Betablocker	13 ^k	No Mut
ZH-019	28	M	1,0	0,0	1,0	0,0	0,1	0,0	2,1	ARVC	Yes	Betablocker	7899 ^k	DSG2
ZH-025	60	M	1,0	1,0	0,1	0,0	0,0	1,0	3,1	ARVC	Yes	Betablocker; Amiodarone	<500	DSP
ZH-027	53	F	1,0	0,0	0,0	0,1	0,1	1,0	2,2	ARVC	Yes	N/A	8769 ^k	PKP2
ZH-028	59	M	1,0	0,0	0,0	0,1	0,1	1,0	2,2	ARVC	Yes	Betablocker; ACE-I/ ARB, Aldosterone antagonist	92 ^k	DSP
ZH-038	54	M	1,0	0,0	0,1	1,0	0,1	0,0	2,2	ARVC	Yes	None	138	No Mut
ZH-051	43	M	1,0	0,0	1,0	1,0	1,0	0,0	4,0	ARVC	Yes	None	2203	TTN, SCN5A
ZH-053	44	M	1,0	0,0	0,1	0,0	1,0	0,0	2,1	ARVC	Yes	None	6275 ^k	LMNA
ZH-081	66	F	1,0	0,0	1,0	0,0	0,0	1,0	3,0	ARVC	Yes	Betablocker, ACE-I/ ARB	<500 ^k	DSC2
ZH-087	62	F	1,0	0,0	1,0	0,0	0,1	1,0	3,1	ARVC	Yes	Betablocker	1063 ^k	No Mut
ZH-092	52	F	1,0	0,0	1,0	0,0	1,0	1,0	4,0	ARVC	Yes	Sotalol	901 ^k	PKP2
ZH-101	26	M	1,0	0,0	0,0	0,0	0,1	1,0	2,1	ARVC	Yes	None	1373 ^k	PKP2
ZH-102	20	F	1,0	0,0	1,0	0,0	1,0	1,0	4,0	ARVC	Yes	None	2005 ^k	DSG2
ZH-107	74	M	1,0	0,0	1,0	0,0	0,1	0,0	2,1	ARVC	Yes	Amiodarone, ACE-I/ ARB, Aldosterone antagonist	209 ^k	No Mut
ZH-121	45	F	1,0	0,0	1,0	1,0	1,0	0,0	4,0	ARVC	Yes	—	26 ^k	No Mut

Continued

Table 1 Continued

Patients	Age ^a	Sex	I ^b	II ^c	III ^d	IV ^e	V ^f	VI ^g	Total (major, minor)	Diagnosis	Biomarker	Therapy ^h	PVC ⁱ	Genetics ^j
ZH-123	51	M	1,0	0,0	0,1	0,0	1,0	0,0	1,2	ARVC	Yes	Betablocker. Amiodarone	>500 ^k	RYR2
ZH-148	19	F	1,0	0,0	1,0	0,0	0,1	0,0	2,1	ARVC	Yes	Amiodarone	4499 ^k	SCN5A
ZH-206	45	M	1,0	0,0	1,0	0,1	0,1	0,0	2,2	ARVC	Yes	Betablocker, Aldosterone antagonist	11 875 ^k	DSG2
ZH-213	38	M	1,0	0,0	1,0	0,0	1,0	1,0	4,0	ARVC	Yes	Sotalol	<500 ^k	DSG2
ZH-222	37	F	1,0	0,0	1,0	0,0	1,0	1,0	4,0	ARVC	Yes	Amiodarone	>500 ^k	PKP2

^aAge at time of blood draw.

^bGlobal or regional dysfunction and structural alterations.

^cTissue characterization of wall.

^dRepolarization abnormalities.

^eDepolarization/conduction abnormalities.

^fArrhythmias.

^gFamily history.

^hMedications/therapies patient was on during time of blood draw.

ⁱPremature ventricular contractions recorded from 24 h-Holter ECG closest to time of blood draw.

^jGene in which mutation has been identified for patient (according to www.arvcdatabase.info, the American College of Medical Genetics and Genomics, and previous literature).

^kPatient was on medical therapy during Holter from which PVC count was recorded.

^lNo mutations were identified in patient.

^mBorderline ARVC as defined by meeting one major and one minor criterion, or three minor criteria as defined by the 2010 Task Force Criteria for the diagnosis of ARVC.

plotted with the pixels calculated from the band intensities. Statistical calculations and graphs were completed with Prism 5.0 for Mac OS X Version 5.0 of 31 July 2012 (Graphpad Software, Inc.).

Results

Study population

We analysed sera from both definite and borderline affected adult and childhood cases of ARVC based on 2010 Task Force criteria followed in the inherited arrhythmia clinic of the Hospital for Sick Children. These include cases referred for ventricular arrhythmias, resuscitated sudden cardiac arrest or family history of ARVC. Subjects had undergone an ECG, SAECG, ambulatory ECG, echocardiogram, cardiac magnetic resonance imaging, and testing for genetic causes of ARVC. When these investigations did not yield a diagnosis of ARVC, an endomyocardial biopsy was also performed. Eleven of these subjects had mutations in known desmosomal genes (PKP2 in eight, DSC2 in one, DSG2 in one, and DSP in one). Eight had no pathogenic mutation detected and one patient did not have genetic testing performed. A validation cohort of adults with definite ARVC from the Zurich ARVC programme was also assessed. Nineteen of these subjects had mutations in known ARVC or ARVC mimic genes: PKP2 in six, DSG2 in five, DSP in two, SCN5A in two, TTN in two, DSC2 in one, LMNA in one, RYR2 in one. One subject had a digenic mutation, one subject had a homozygous DSG2 mutation and six had no pathogenic mutations detected. All subjects are summarized according to ARVC Task Force criteria in Table 1. Six subjects had neither a Family History criterion nor pathogenic ARVC mutation, although

two had suspicious deaths in second degree relatives (not meeting criteria) and one had a rare conserved DSC2 variant (DSC2 c.236T>G). Thus three subjects in total could be considered to have sporadic ARVC.

Normal control sera were obtained from two commercial sources: 12 from Innovative Research Inc. Novi, MI, USA and 20 from BBI Solutions, Cardiff, UK. In addition, a cohort of subjects from Mayo clinic with hypertrophic cardiomyopathy (see [Supplementary material online, Table S1](#)) and a cohort of subjects from Zurich with dilated cardiomyopathy (see [Supplementary material online, Table S2](#)) served as non-ARVC cardiomyopathic controls. Finally, as additional validation, we assessed sera from 10 Boxer dogs (a naturally occurring animal model of ARVC)²⁷ with manifest disease (see [Supplementary material online, Table S3](#)) compared with 18 unaffected dogs (2 mongrels and 16 Boxers) with no pedigree history of ARVC.

Autoantibody assessments

In 12 of 12 cases of definite ARVC in our human samples, and seven of eight cases of Borderline ARVC in our human samples, we found that autoantibodies to the desmosomal protein DSG2 were present on western blots, whereas these autoantibodies were absent (in 11) or very faint (in one) from sera of 12 control subjects from our first commercial source ([Figure 1](#)). No antibodies to desmocollin-2 or N-cadherin were identified in any sample.

In a subset of five subject and three control sera, we isolated the immunoglobulin G fraction, which also demonstrated similar antibody staining to DSG2 only (see [Supplementary material online, Figure S1](#)). A subset of sera was also assessed for antibodies to non-cadherin desmosomal proteins, as well as to skin cadherin proteins,

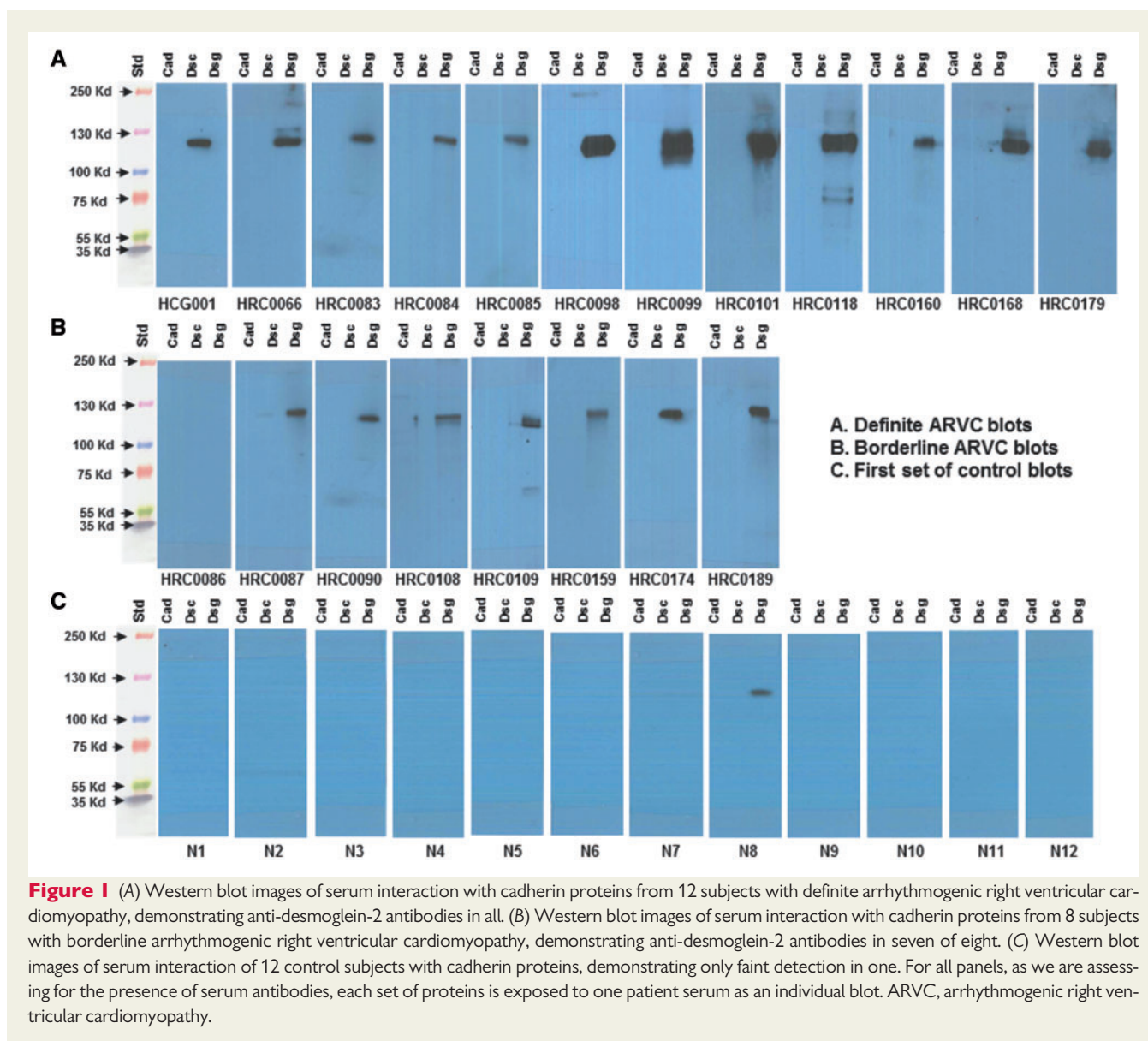


Figure 1 (A) Western blot images of serum interaction with cadherin proteins from 12 subjects with definite arrhythmogenic right ventricular cardiomyopathy, demonstrating anti-desmoglein-2 antibodies in all. (B) Western blot images of serum interaction with cadherin proteins from 8 subjects with borderline arrhythmogenic right ventricular cardiomyopathy, demonstrating anti-desmoglein-2 antibodies in seven of eight. (C) Western blot images of serum interaction of 12 control subjects with cadherin proteins, demonstrating only faint detection in one. For all panels, as we are assessing for the presence of serum antibodies, each set of proteins is exposed to one patient serum as an individual blot. ARVC, arrhythmogenic right ventricular cardiomyopathy.

with none found (see [Supplementary material online, Figure S2](#)). Signal levels for anti-DSG2 antibody varied between ARVC samples and were measured from western blots using quantitative analysis of the band densities as previously described.^{26,28}

Anti-DSG2 antibodies were absent on western blot of serum from one subject with asymptomatic borderline ARVC. This individual was the mother of an identified ARVC subject with one borderline abnormal measurement of high frequency, low amplitude signal duration on SAECG. She has a very rare novel missense variant of PKP2 that has been classified as pathogenic (www.arvcdatabase.info) based on moderately reduced co-localization of connexin-43 gap junction protein to the cardiac intercalated disk.¹⁹ Anti-DSG2 antibodies were weakly present in one control sample from a 53-year-old Hispanic female, but did not reach the level seen in the 12 definite ARVC subjects with antibodies based on quantitation of western blot density,

nor the seven of eight borderline ARVC cases that demonstrated antibody.

Within our validation cohort of 25 additional adult subjects with definite ARVC from the Zurich ARVC Program, we identified anti-DSG2 antibodies in all 25 ([Figure 2A](#)). We assessed a second set of 20 clinically well characterized control sera from another commercial source (BBI Solutions, Cardiff, UK) and did not identify anti-DSG2 antibodies in any ([Figure 2B](#)). In addition, no anti-DSG2 antibodies were identified among 10 subjects with hypertrophic cardiomyopathy or among six subjects with dilated cardiomyopathy (see [Supplementary material online, Figure S3](#)). Assessing the naturally occurring Boxer dog model of ARVC, we identified anti-DSG2 antibodies in 10 of 10 Boxers presenting with manifest disease, but in none of 18 healthy dogs with no pedigree history of ARVC (2 mongrel and 16 Boxer dogs; [Figure 3](#)).

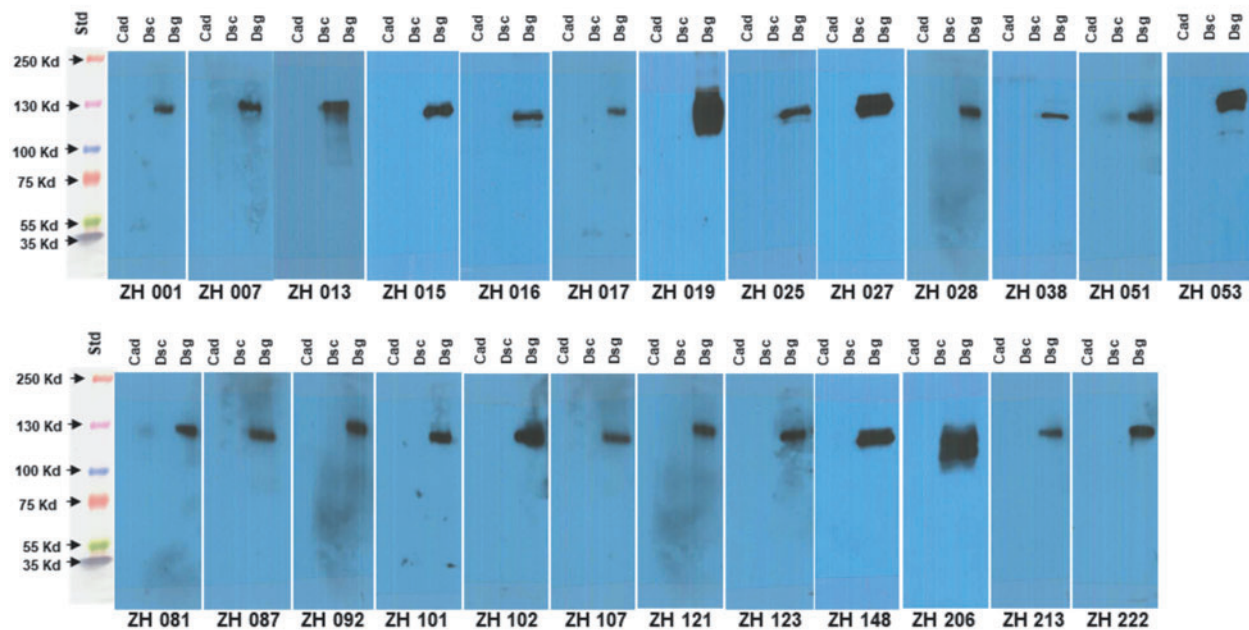
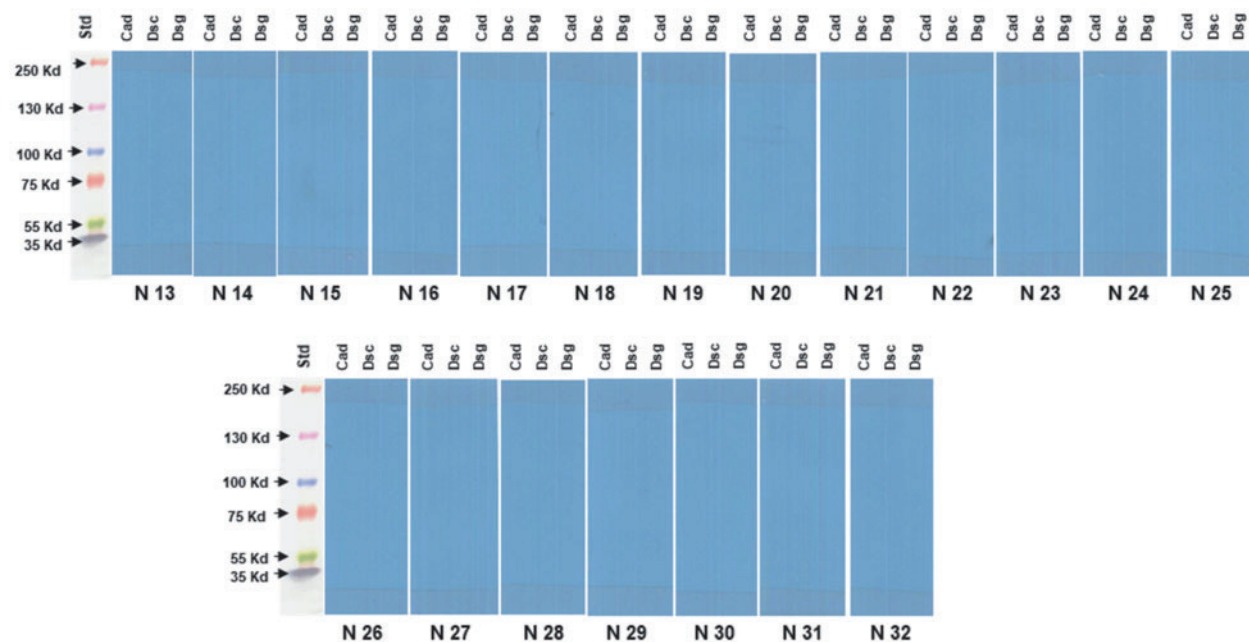
A Validation Cohort (Zurich)**B 20 Additional Controls**

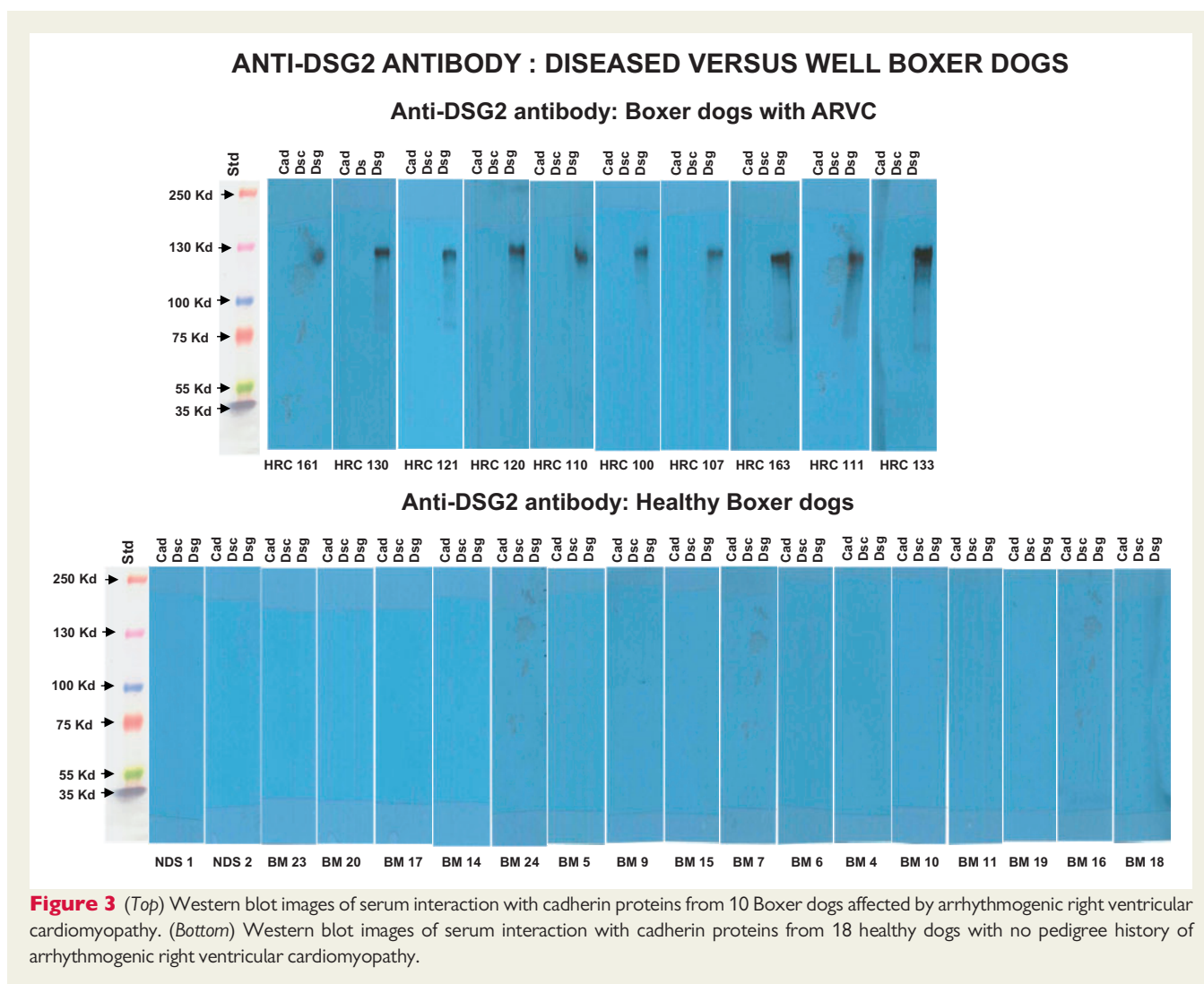
Figure 2 (A) Western blot images of serum interaction with cadherin proteins from 25 additional subjects with definite arrhythmogenic right ventricular cardiomyopathy from our validation cohort. (B) Western blot images of serum interaction with cadherin proteins from 20 additional control sera.

We also created an ELISA assay and characterized antibody concentration by O.D. in our control, borderline, and definite (primary and validation) subjects (Figure 4). Our ELISA assay provided comparable results, with a cut-point of O.D. = 0.1060 providing 97.78% sensitivity (95% confidence interval = 88.23–99.94%) and 96.88% specificity (95% confidence interval = 83.78–99.92%). A Receiver

Operator Characteristic (ROC) curve (see [Supplementary material online, Figure S4](#)) demonstrates an area under the curve of 0.99.

Correlation with disease severity

We assessed whether antibody density as measured by pixel count of the western blot or ELISA O.D. correlated with disease severity as



measured by 24-h burden of PVCs within the groups of definite ARVC. The Pearson correlation coefficient (r) was 0.70 ($P = 0.0001$) for PVCs vs. pixel count of the western band, indicating that 49% of the variation in PVC count could be accounted for by its linear relationship with antibody density (Figure 5).

Antibody effects on gap junction function

To assess whether anti-DSG2 antibody might cause gap junction dysfunction, a common feature of ARVC,¹⁹ we tested purified IgG from two randomly selected ARVC patients as well as commercial anti-DSG2 antibodies for effects on cell-to-cell dye transfer. Using automated microinjection of fluorescent dye molecules into single cells (Figure 6)²⁹ of confluent normal human iPSC-derived cardiomyocytes (i-Cell[®], Cellular Dynamics International), we identified reduced dye transfer distances from each injected cell to its adjacent cells in those cultures exposed to patient-derived purified antibodies or commercial anti-DSG2 antibody, compared with those exposed to purified antibody from control sera (Figure 7). We injected between 37 and 42 cells for each assessment.

Identification of the antigen epitope

To identify the DSG2 epitope bound by ARVC serum autoantibodies, we assessed two random patient sera against a peptide library (Sigma-Aldrich Canada) of sequential 15 a.a. peptides (with 5 a.a. overlap) derived from the human DSG2 protein sequence. Strong binding of serum autoantibodies was identified against a.a. 485–531 and a.a. 586–610, within the fourth extracellular domain and extracellular anchor regions, respectively (Figure 8).

Discussion

This study identified autoantibodies to the cardiac DSG2 protein as a common feature in the sera of patients with ARVC. These autoantibodies were specific to ARVC, as they were essentially absent in two independent sets of control sera, as well as sera from subjects with other forms of heritable cardiomyopathy. Assessment of anti-DSG2 antibodies may provide a reliable assay for detection of ARVC patients undergoing evaluation. Anti-DSG2 antibodies were confirmed within a validation cohort and were present independent of whether an underlying genetic defect had been identified, and

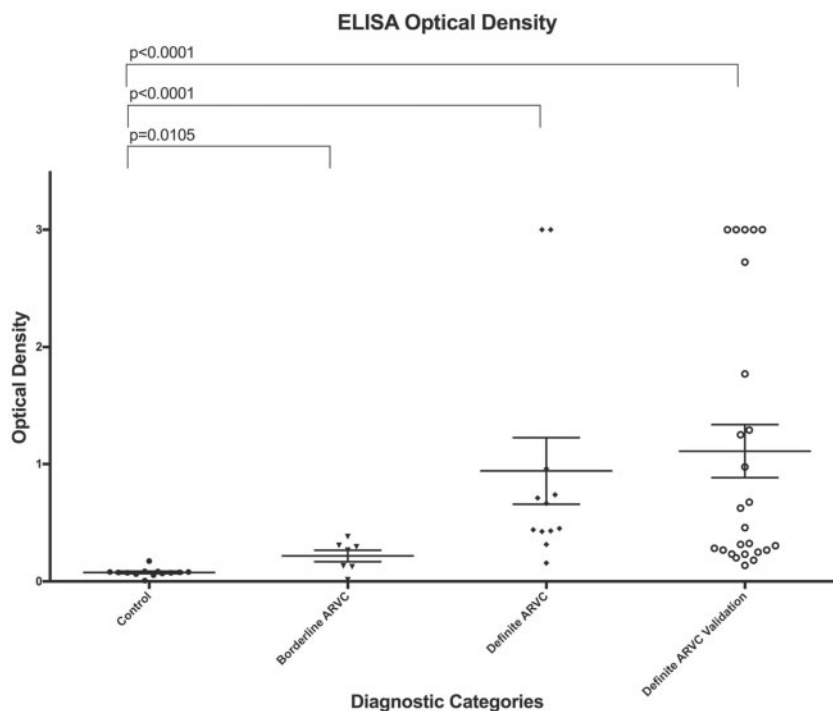


Figure 4 Dot plot of ELISA optical density for controls, borderline arrhythmogenic right ventricular cardiomyopathy subjects, and definite arrhythmogenic right ventricular cardiomyopathy subjects from our primary and validation cohorts. Error bars represent the mean value of each cohort (mid horizontal line) and standard deviation within each cohort. The two-tailed *P*-values between the indicated groups are exact values calculated using a Mann–Whitney *U*-test.

independent of the specific ARVC gene when known. Anti-DSG2 antibodies were also demonstrated to be specific and sensitive for disease in the naturally occurring Boxer dog model of ARVC, providing the potential for a veterinary test for this disease, as well as a potential large animal model for assessment of immune therapies for ARVC. In contrast to desmosomal disorders of the skin, which are caused by either mutations or autoantibodies, our ARVC subjects had anti-DSG2 antibodies usually in the presence of familial and/or genetic disease.

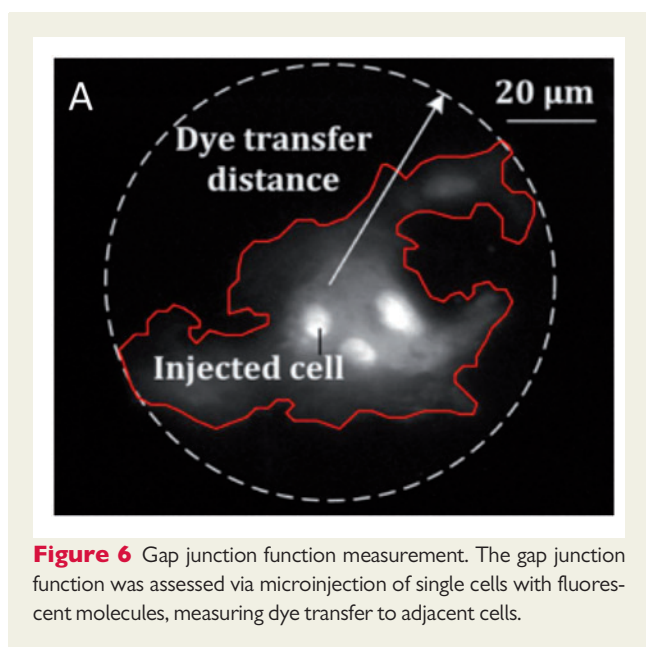
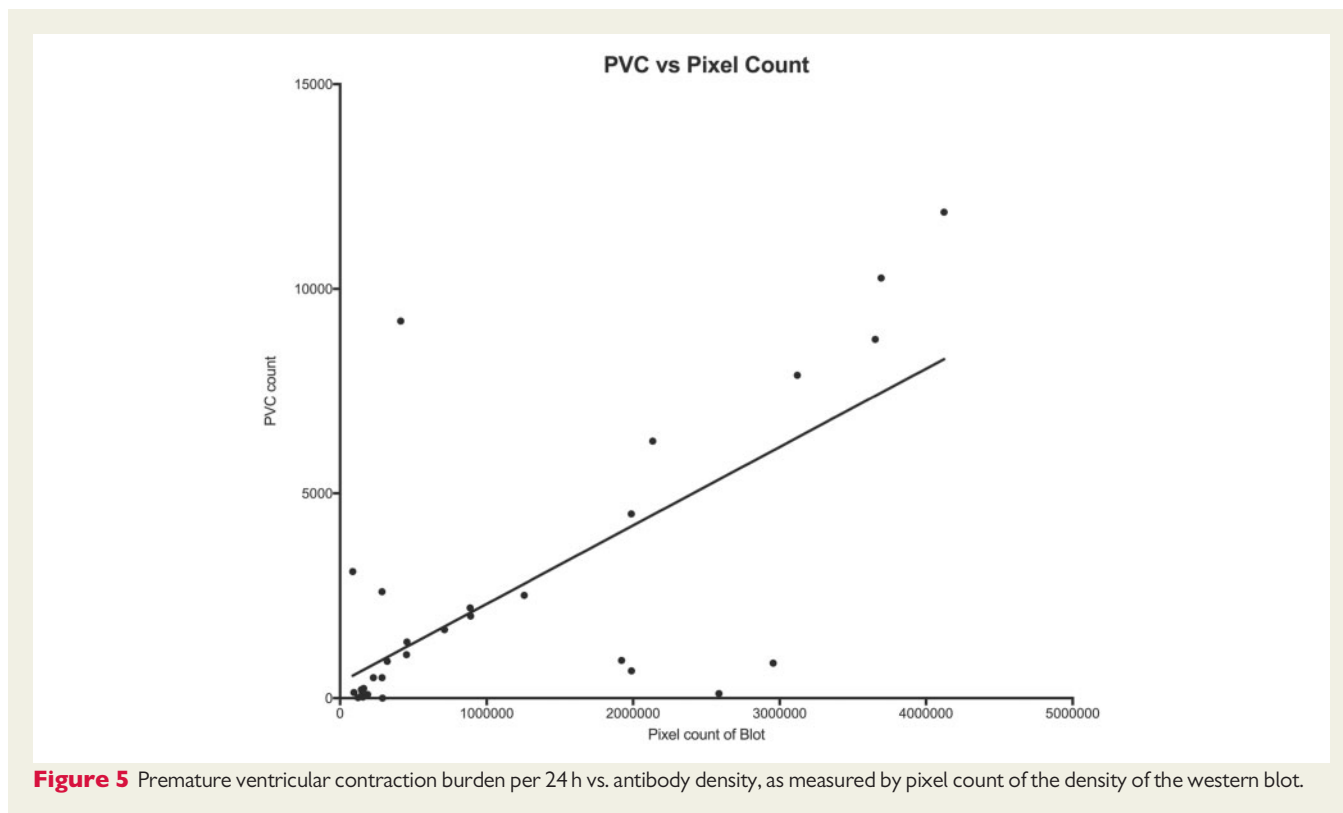
Prognostically, the level of anti-DSG2 antibodies correlated with disease burden as measure by PVC count, and purified antibodies resulted in reduced gap junction function, an identified common pathway in ARVC pathophysiology. This latter finding is consistent with a previous study that demonstrated that anti-N-cadherin antibody Fab fragments result in failure to form adherens junctions and inhibit gap junction dye transfer in an experimental cell model.³⁰ As such, our test provides a quantitative result that may contribute to the evaluation of risk within diagnosed ARVC patients, or predict hot phases of disease.

Desmoglein-2 is one of several cardiac cadherin proteins. Cadherins are calcium-dependent adhering molecules providing mechanical attachment between cells in multiple tissues.³¹ Typically, three calcium ions (12 in total) are pocketed into binding motifs present between each pair of five successive extracellular cadherin (EC) domains, providing the crescent shape required to bring cadherin domains from opposing cells into a 90° conformation for *trans*

binding. This 90° conformation is critical for binding as it allows for tryptophan residues on each EC1 domain to be inserted simultaneously into the hydrophobic pocket of the alternate cadherin. Interfering with this orientation, such as through missense mutations, or antibody binding to proximal extracellular DSG2 domains, might be expected to reduce binding and adhesion.

Recently, it has been recognized that cell-to-cell binding at desmosomal structures of the cardiac intercalated disk is heterophilic, with desmogleins typically forming adhesive dimers with desmocollins.³² The ontogeny of the expression of intercalated disk structures from 15 weeks gestation through 11 years postnatal has been assessed, with structural (adherens junction and desmosomal) proteins localizing to the intercalated disk at 1 year after birth, followed by the cardiac sodium channel at 2 years and gap junction connexin 43 protein at 7 years.³³ Within proteins of the intercalated disk, DSG2 does not appear to localize to the intercalated disk until 6 weeks of postnatal life, whereas DSC2 is present in the intercalated disk prenatally.³⁴

We do not yet fully understand the pathobiology responsible for an autoimmune response to DSG2 in ARVC patients. We also do not know to what degree autoimmunity participates in the pathophysiology of ARVC. It is plausible that just as catenins are released into the sarcoplasm in the face of mutations that disrupt the desmosome, DSG2 protein may include ‘cryptic’ epitopes,³⁵ which are also exposed or released into the intercellular space and/or circulation and as a result of desmosomal mutations. Such released DSG2 proteins may link with an antigen-presenting cell to stimulate a T-cell



response, generating the observed autoantibodies. Unmasking of cryptic epitopes by gene mutations could contribute to other forms of autoimmunity. Alternately, the failure of tolerance for the DSG2 epitope may be related to its relatively late expression at or around 6 weeks of infancy.³⁴ Although three subjects had sporadic ARVC, it would be premature to suggest that autoantibodies cause ARVC in

isolation, as ARVC is recognized to be incompletely penetrant, variably expressed and often gene-elusive.

Our results are promising for the development of a serological test for the diagnosis of ARVC. The high sensitivity and specificity of this assay is anticipated to improve the diagnosis of ARVC, possibly even in the pre-symptomatic period. Cardiac tissue sections have been used by others to assess for the presence of anti-heart or anti-intercalated disk antibodies in myocarditis and dilated cardiomyopathy,³⁶ as well as ARVC.³⁷ The frequencies of anti-heart or anti-intercalated disk antibodies were higher (31%; 15%) in ARVC, than in non-*ischaemic* cardiomyopathy, *ischaemic* heart failure, or normal subjects.³⁸ Antibodies were associated with PKP2 and DSP mutation-positive individuals, and with those with reduced left ventricular function. However, the intercalated disk is known to contain more than 700 different proteins based on Human Protein Atlas.³⁹ Our technique assays autoantibodies to a single-specific isolated protein, and provides very high sensitivity for the diagnosis of ARVC. A diagnostic test for ARVC based on reduced intercalated disc plakoglobin immunofluorescence in endomyocardial biopsies has been previously proposed⁴⁰ and critiqued^{41,42} but has not been adopted in practice. Our serum blood test provides a more practical and accurate alternative.

The identification of an autoantibody associated with ARVC may open new therapeutic avenues, similar to those proposed for autoimmune pemphigus vulgaris.²⁵ Immunosuppression is often effective for disorders resulting from autoimmunity and is usually the standard to which other therapies are compared. Newer therapies include co-stimulation blockade, regulatory T-cell therapy, antigen-specific immunotherapy, and manipulation of the interleukin-2 pathway.⁴³ Each

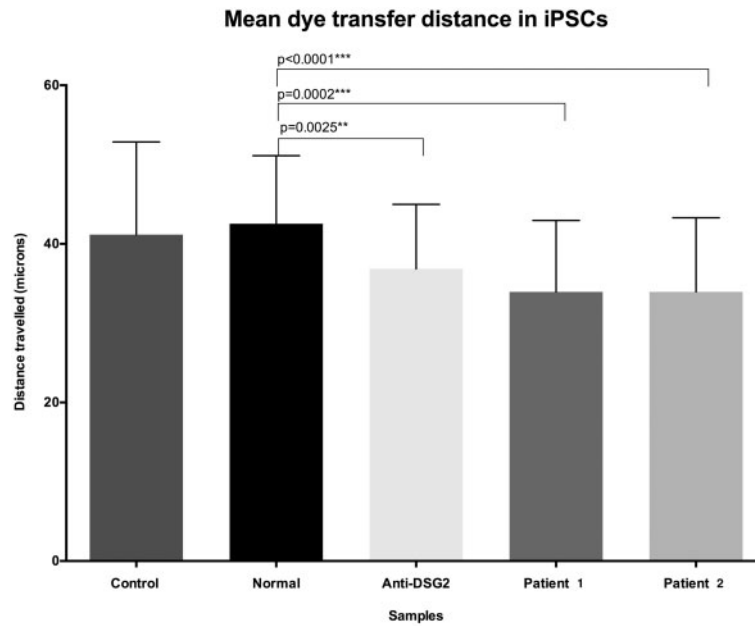


Figure 7 Purified IgG antibodies from the serum of two patients with arrhythmogenic right ventricular cardiomyopathy, as well as commercial anti-DSG-2 IgG demonstrate reduced cell-to-cell dye transfer distance compared with cells exposed to IgG from normal human serum ($P \leq 0.0001$ and $P = 0.0025$, respectively). The two-tailed P -values between the indicated groups are exact values calculated using a Mann–Whitney U -test. Error bars indicate standard deviation.

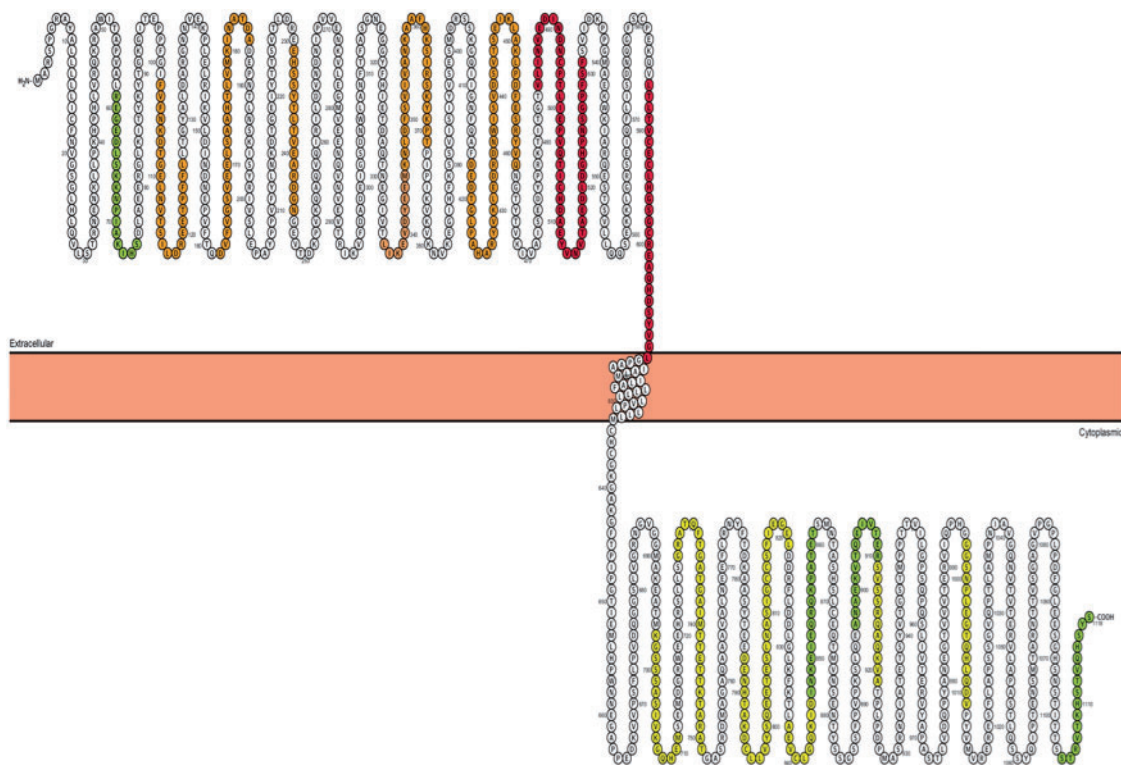
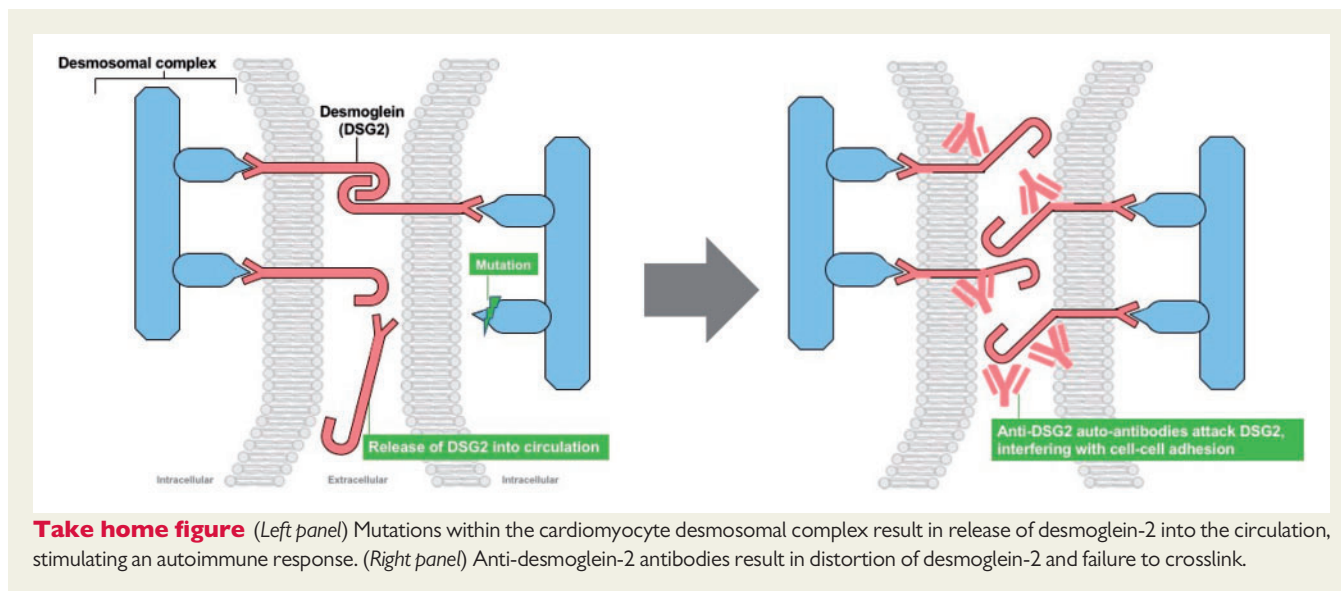


Figure 8 Serum binding intensity to desmoglein-2 oligopeptides is presented as colour grades onto a Protter (<http://wlab.ethz.ch/protter>) plot of the desmoglein-2 protein (red, strong binding; orange, moderate binding; yellow, weak binding; green, faint binding). Strongest binding is in regions of the fourth extracellular cadherin and extracellular anchor domains.



of these may provide an avenue for therapy in ARVC, once studies of the mechanism of autoimmunity have been completed. Of particular interest is a recent report of reengineered chimeric antigen receptor T-cells to target autoimmunity specifically, without off-target immuno-suppression.⁴⁴ This technique was applied successfully to treat a murine model of pemphigus vulgaris due to anti-desmoglein-3 antibodies, and may hold promise for an effective targeted therapy in ARVC. Finally, a 'negative vaccination' approach exposing at-risk subjects to the DSG2 antigen early in life might provide a specific and potentially safe approach to preventing autoimmunity in ARVC, similar to therapies under study for Type 1 diabetes.⁴⁵

Limitations

Our study evaluated a limited number of commercial control sera from ostensibly normal subjects for which we had limited clinical information, as well as a limited number of control sera from two other specific forms of cardiomyopathy: hypertrophic cardiomyopathy and dilated cardiomyopathy. Other cardiomyopathic processes including myocarditis and sarcoidosis should be assessed to determine if anti-DSG2 antibodies will also be detected in these disorders. Viral myocarditis has been associated with ARVC in some patients as an acquired cause of ARVC disease.³⁷ We have not yet assessed these disorders to further assess the specificity of our test.

For this initial study, we have not tested individuals in whom the disease has yet to manifest, and therefore cannot characterize the ability of this test to detect silent mutation carriers at this time.

Conclusion

Our discovery will enable the development of a new serological test for ARVC that can easily be adapted to clinical diagnosis and the prediction of disease. The assay is negative in normal control individuals and also negative in two other forms of cardiomyopathy, but further

assessment is required among other myocardial diseases to determine its specificity to ARVC. Further research is required to elucidate the pathophysiology of this autoimmune response, its sensitivity in predicting disease risk or hot phases of disease, and its utility in directing or monitoring new therapies for ARVC.

Supplementary material

Supplementary material is available at *European Heart Journal* online.

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Conflict of interest: M.J.A. is a consultant for Audentes Therapeutics, Boston Scientific, Gilead Sciences, Invitae, Medtronic, MyoKardia, and St. Jude Medical. M.J.A. and Mayo Clinic have an equity/royalty relationship with AliveCor, Blue Ox Health, and StemoniX. However, none of these

entities had any involvement with this study. All other authors have declared no conflict of interest.

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