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# Integrin β1D Deficiency-mediated RyR2 Dysfunction Contributes to Catecholamine-Sensitive Ventricular Tachycardia in ARVC

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### Abstract

**Background:** Arrhythmogenic right ventricular cardiomyopathy (ARVC) is a hereditary heart disease characterized by fatty infiltration, life-threatening arrhythmias and increased risk of

DISCLOSURES: None.

SUPPLEMENTAL MATERIALS

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sudden cardiac death (SCD). The guideline for management of ARVC patients is to improve quality of life by reducing arrhythmic symptoms and to prevent SCD. However, the mechanism underlying ARVC-associated cardiac arrhythmias remains poorly understood.

**Methods:** Using protein mass spectrometry analyses, we identified integrin  $\beta 1$  is down-regulated in ARVC hearts without changes to Ca<sup>2+</sup>-handling proteins. As adult cardiomyocytes express only the  $\beta 1D$  isoform, we generated a cardiac specific  $\beta 1D$  knockout ( $\beta 1D^{-/-}$ ) mouse model, and performed functional imaging and biochemical analyses to determine the consequences from integrin  $\beta 1D$  loss of function in the heart in vivo and in vitro.

**Results:** Integrin  $\beta$ 1D deficiency and RyR2 Ser-2030 hyper-phosphorylation were detected by western blotting in left ventricular tissues from patients with ARVC but not in patients with ischemic or hypertrophic cardiomyopathy. Using lipid bilayer patch clamp single channel recordings, we found purified integrin  $\beta$ 1D protein could stabilize RyR2 function by decreasing RyR2 open probability (Po), mean open time (To), and increasing mean close time (Tc).  $\beta$ 1D<sup>-/-</sup> mice exhibited normal cardiac function and morphology, but presented with catecholamine-sensitive polymorphic ventricular tachycardia, consistent with increased RyR2 Ser-2030 phosphorylation and aberrant Ca<sup>2+</sup> handling in  $\beta$ 1D<sup>-/-</sup> cardiomyocytes. Mechanistically, we revealed that loss of desmoplakin induces integrin  $\beta$ 1D deficiency in ARVC mediated through an ERK1/2 – fibronectin – ubiquitin/lysosome pathway.

**Conclusions:** Our data suggest that integrin  $\beta$ 1D deficiency represents a novel mechanism underlying the increased risk of ventricular arrhythmias in patients with ARVC.

#### Keywords

ARVC; catecholamine-sensitive ventricular tachycardia; mechanism; integrin  $\beta$ 1D; RyR2 phosphorylation

### INTRODUCTION

Arrhythmogenic right ventricular cardiomyopathy (ARVC) is a rare hereditary progressive heart disease characterized by fibrous or fatty infiltration of the heart muscle with predominantly right ventricular (RV) dysfunction, life-threatening ventricular arrhythmias and increased risk of sudden cardiac death (SCD).<sup>1</sup> As left ventricular (LV) involvement is found more commonly than previously appreciated in patients with ARVC, ARVC is increasingly recognized as a biventricular entity.<sup>2, 3</sup> A recent study indicates that LV involvement in ARVC may even precede the onset of significant RV dysfunction and is associated with a higher prevalence of ventricular arrhythmia.<sup>3</sup> In adolescents and young adults with ARVC, especially in athletic individuals in the absence of previous reported cardiac symptoms, exercise or competitive sports is a critical factor that may trigger fatal ventricular arrhythmias and increase the risk of SCD.<sup>3, 4</sup> Exercise-induced catecholamine release is a major trigger linking the onset of ventricular arrhythmias and SCD in athletes with ARVC.<sup>5, 6</sup>

ARVC is associated with mutations in the genes encoding desmosomal proteins including desmoplakin (*DSP*), plakophilin 2 (*PKP2*), desmoglein 2 (*DSG2*), and desmocollin 2 (*DSC2*).<sup>7–9</sup> Mutation-associated electrical uncoupling between cardiomyocytes from

fibrofatty tissue replacement of the myocardium in ARVC is thought to be the main mechanism underlying ventricular arrhythmias. The fibro-fatty tissues in affected ventricles may provide a substrate for ventricular tachycardia through a re-entry mechanism.<sup>9–11</sup>

ARVC is distinct from catecholaminergic polymorphic ventricular tachycardia (or CPVT), another type of familial hereditary arrhythmogenic disease. Although both are life threatening, CPVT occurs in individuals with structurally normal hearts, characterized by exercise induced bidirectional/polymorphic ventricular tachycardia.<sup>12</sup> CPVT originates from dysregulation of intracellular calcium homeostasis. The pathogenic mediators of CPVT include mutations in genes associated with intracellular calcium release from the sarcoplasmic reticulum (SR) such as the ryanodine receptor type 2 (*RyR2*), calsequestrin-2 (*CASQ2*), triadin (*TRDN*) and calmodulin (*CALM1*).<sup>13–18</sup> Mutations in these genes cause defects in RyR2 activities, abnormal Ca<sup>2+</sup> handling and triggered activities, which are exacerbated under sympathetic stress or exercise.<sup>13, 17, 19</sup> It is generally agreed that ARVC and CPVT are caused by mutations of different sets of genes involved in distinct molecular and cellular functions within cardiomyocytes.

Interestingly, new lines of evidence from both clinical and animal experimental data indicate there is some degree of convergence (overlap) between ARVC and CPVT. First, mutations in the RyR2 gene were found in a well clinically characterized cohort of ARVC patients without mutations in desmosomal genes with a frequency of 9%.<sup>20</sup> Second, *PKP2* mutations were identified in a significant percentage (5 out of 18, 27.7%) of a clinically diagnosed CPVT patient cohort whose genetic testing panels reported negative on mutations of conventional CPVT genes.<sup>4</sup> In these patients, cardiac imaging or autopsy demonstrated structurally normal hearts at the time of diagnosis or death. Third, in a case report of a young patient with a clinical diagnosis of typical ARVC but negative for mutations of known genes associated with both CPVT and ARVC, showed classic bi-directional and polymorphic ventricular tachycardia during exercise stress testing.<sup>21</sup> Fourth, like CPVT, exercise or exertion is a risk factor of ventricular arrhythmias and SCD in ARVC patients with more clinical studies supporting the diagnostic value of isoproterenol testing in ARVC.<sup>22, 23</sup> Fifth, in an animal model, Cerrone et al. reported that cardiac-specific deletion of Pkp2 in mice leads to reduced expression of genes related to intracellular Ca<sup>2+</sup> handling, i.e., Ryr2 and Trdn, and higher vulnerability to isoproterenol-triggered polymorphic ventricular arrhythmias that mimics CPVT.<sup>24</sup> These findings collectively suggest that CPVT-like ventricular arrhythmias are often present in ARVC. However, the molecular mechanism by which ARVC-associated mutations give rise to CPVT-like ventricular arrhythmias (i.e., catecholamine-sensitive ventricular tachycardia), in the absence of canonical CPVT gene mutations, has been a longstanding mystery.

In the present study, using protein mass spectrometry analyses, we identified integrin  $\beta$ 1, but not other Ca<sup>2+</sup>-handling proteins, is significantly down-regulated in left ventricles of ARVC patients. Integrin  $\beta$ 1 is composed of 4 isoforms, including  $\beta$ 1A, B, C & D. Integrin  $\beta$ 1D is mainly expressed in striated muscles, including heart muscle, and is the only isoform expressed in adult cardiomyocytes.<sup>25, 26</sup> Using a cardiac specific  $\beta$ 1D knockout ( $\beta$ 1D<sup>-/-</sup>) mouse model, we further show that integrin  $\beta$ 1D deficiency represents an important

mechanism underlying  $Ca^{2+}$  disorders and fatal catecholaminergic-sensitive  $Ca^{2+}$ -dependent ventricular arrhythmias in ARVC.

### METHODS

The data, analytic methods, and study materials will be/have been made available to other researchers for purposes of reproducing the results or replicating the procedure. Requests for reagents and materials may be sent to the corresponding authors.

All studies involving human tissues strictly abided by the principles outlined in the World Medical Association Declaration of Helsinki.<sup>27</sup> All procedures involving human tissues were approved by Fuwai Hospital (Beijing, China). Informed consent was obtained from all subjects. All animal experiments were performed in accordance with the Guide for the Care and Use of Laboratory Animals (NIH publication No. 85 – 23, revised 1985) and were approved by the Institutional Animal Care and Use Committee of Fuwai Hospital.

#### Human heart samples

Human LV samples from patients with ARVC, ischemic cardiomyopathy (ICM) and hypertrophic cardiomyopathy (HCM) were obtained from explanted hearts through the Fuwai Hospital Heart Failure Transplant Program. A total of 4 non-failing healthy donor (HD) hearts without evidence of cardiac dysfunction were obtained through organ procurement agencies. Patient demographics, and corresponding clinic information of ARVC, ICM and HCM are detailed in Supplemental Table 1,2,3.

#### Generation of integrin $\beta$ 1D knockout mouse model

Itgb1 flox/flox (Itgb1<sup>f/f</sup>) mice with LoxP sites inserted between canonical exon 15 and exon 16 flanking the alternatively spliced and cardiomyocyte specific "D" exon<sup>26</sup> were generated by the Model Animal Research Center of Nanjing University. Tamoxifen-inducible Cre-Lox mice ( $\alpha$ MHC-MerCreMer, or  $\alpha$ MHC-MCM) were obtained from the same center. Itgb1<sup>f/f</sup> mice were bred with  $\alpha$ MHC-MCM mice to generate experimental cohorts including cardiac-specific Itgb1 homozygous knockout mice ( $\beta$ 1D<sup>-/-</sup>, Cre<sup>+</sup>/Itgb1<sup>f/f</sup>) and wildtype (WT, Cre<sup>-/</sup> Itgb1<sup>f/f</sup>) littermates. Mice were injected intraperitoneally with tamoxifen (40 mg/kg body weight, Sigma Aldrich, USA, dissolved in corn oil) or with corn oil only (control group) for 10 days. Expression levels of integrin  $\beta$ 1D were assessed by western blotting with heart tissues collected at 1 week and 2 weeks after the last dose of tamoxifen injection (~9–11 weeks old). The numbers of mice or myocytes for each experimental group are provided in the Figures and/or Figure Legends.

#### Statistical analyses

All data were analyzed with SPSS Statistics Version 22. Differences between groups were analyzed by Student's t-test or  $\chi 2$  test. The data represent the means  $\pm$  SEM. \*P <0.05, \*\*P <0.01 and \*\*\*P <0.001 were considered statistically significant.

All detailed experimental methods can be found in the online-only Supplemental Materials.

### RESULTS

### Integrin $\beta$ 1D expression is decreased together with RyR2 hyper-phosphorylation in left ventricle tissues from patients with ARVC, but not in patients with ICM and HCM.

We followed 39 patients who were diagnosed with ARVC according to the 2010 Task Force Criteria (TFC).<sup>28</sup> For the studies presented here, LV tissue samples were processed from the four patients who had successfully undergone cardiac transplantation (Supplemental Figure 1). All patients had severe, emotional excitement- or exercise-induced ventricular tachycardias that were not present at rest. (Figure 1A and Supplemental Figure 2). MRI reported fibrofatty infiltration of RV predominance with LV involvement in all four patients (Supplemental Figure 3). Whole exon sequencing found three patients had mutations in desmosomal genes including desmoglein-2 (DSG2 F531C; Case 1), plakophilin-2 (PKP2 C796R; Case 2) and desmoplakin (DSP S299R; Case 4) genes (Figure S3). Using protein mass spectrometry analyses, we found that integrin  $\beta 1$  is significantly down-regulated in LV hearts from patients with ARVC, but no change in other Ca<sup>2+</sup>-handling proteins including Amphyphisin-2 (Bin1), L-type Ca<sup>2+</sup> channel (Ca<sub>V</sub>1.2), calmodulin (CaM), IP3R, junctin (JCTN), junctophilin-2 (JP2), sodium-calcium exchanger (NCX1), phospholamban (PLB), RyR2, SR Ca<sup>2+</sup>-ATPase (SERCA2a) and triadin (TRDN), as compared to these of healthy donor hearts (HD) (Figure 1B). The expression levels of membrane ion channels such as sodium and potassium channels remained unaltered in LV hearts of ARVC, as compared to HDs (Supplemental Figure 4). Immunofluorescent confocal images of LV heart cryosections from healthy donors and ARVC patients demonstrate that integrin  $\beta$ 1D is regularly distributed in a striated pattern, like T-tubules, in control myocardium but markedly reduced in intensity in ARVC (Figures 1C–D). Previous work has suggested that integrin  $\beta$ 1D is present at intercalated discs and T-tubules in mouse and rat hearts.<sup>25, 26</sup> and co-localizes with and regulates RyR2 phosphorylation levels at its Serine 2808 (Ser-2808) site in cardiomyocytes.<sup>29</sup> These led us to hypothesize that integrin  $\beta 1D$  downregulation may contribute to RyR2 dysfunction and Ca<sup>2+</sup>-dependent arrhythmogenesis in ARVC.

To investigate the function of integrin  $\beta$ 1D and its potential interaction with RyR2 in cardiomyocytes, we first examined integrin  $\beta$ 1D protein expression levels and the relative phosphorylation status of RyR2 in LV heart tissues from the four ARVC patients and HDs. Consistent with protein mass spectrometry data, western blot analyses further confirmed that integrin  $\beta$ 1D expression was markedly decreased in patients with ARVC, as compared to HDs (Figures 1E–F). While the phosphorylation levels of RyR2 at S2814 and S2808 sites did not differ between patients and HDs, we did observe a hyper-phosphorylation state of RyR2 at S2030 site (Figures 1E–I). On the contrary, integrin  $\beta$ 1D is significantly upregulated in LV hearts of ICM and HCM, as compared to HDs, respectively (Supplemental Figure 5). Interestingly, ICM and HCM hearts showed increased phosphorylation of RyR2 at Ser-2808 and Ser-2814, but not at Ser-2030 site (Supplemental Figure 5). These results suggest integrin  $\beta$ 1D downregulation is a unique phenomenon of ARVC.

### Integrin β1D stabilizes RyR2 by suppressing RyR2 phosphorylation at Ser-2030.

Previous studies suggest that integrin  $\beta$ 1D stabilizes RyR2 by binding to the interdomain of RyR2, but lack of evidence for the functional implications of this interaction.<sup>29</sup> As we found that integrin β1D protein expression is markedly and specifically down-regulated in ARVC hearts, we investigated the effect of exogenous integrin  $\beta$ 1D on RyR2 activity by fusing SR vesicles into planar lipid bilayers. We isolated single RyR2 channels from a freshly explanted ARVC heart (Case 1) and then performed single-channel recordings before and after the addition of GST- $\beta$ 1D (1  $\mu$ M) to the cytosolic side of the channel (Figures 2A and B). After a 30 min incubation, GST-β1D reduced RyR2 activity, which is reflected by a reduction in RyR2 open probability (Po) (0.002133±0.000096 vs. 0.001873±0.000033, P=0.053, n=3) and mean open time (To) (0.114043±0.002483 ms vs. 0.093333±0.005624 ms, P=0.093), and an increase in mean closed time (Tc) (144.467±5.284 ms vs. 169.667±7.323 ms, P=0.085) (Figures 2F-H). PKA treatment substantially increased RyR2 activity, an effect that was suppressed by GST-B1D (Po: 0.032401±0.002245 vs. 0.015177±0.001277, P<0.01; To: 1.766667±0.131852 ms vs. 0.630132 ±0.148399 ms, P<0.01; Tc: 44.376±6.244 ms vs. 70.014±3.300 ms, P<0.05, n=3) (Figures 2C-H). These findings suggest that integrin  $\beta$ 1D directly modulates RyR2 channel activity by suppressing channel open probability and time.

To investigate how integrin  $\beta$ 1D stabilizes RyR2 function, we analyzed RyR2 phosphorylation levels following the same treatments described above (Figures 2A–D) by western blotting with phospho-specific antibodies. PKA treatment increased the phosphorylation level of RyR2 at Ser-2030, but not Ser-2808 or Ser-2814 (Figures 2I–J). GST-fused Integrin  $\beta$ 1D markedly repressed PKA-induced RyR2 phosphorylation at Ser-2030. These results suggest that integrin  $\beta$ 1D can stabilize RyR2 function by suppressing PKA-dependent RyR2 Ser-2030 phosphorylation levels.

### CPVT-like ventricular arrhythmias in integrin β1D<sup>-/-</sup> mice

To reveal the in vivo effects of integrin  $\beta$ 1D deficiency on cardiac function and to validate our findings from human patients, we engineered a cardiac-specific  $\beta$ 1D<sup>-/-</sup> mouse model using the tamoxifen-inducible Cre-LoxP system (Figures 3A–B). We first confirmed the loss of integrin  $\beta$ 1D expression in left ventricle of  $\beta$ 1D<sup>-/-</sup> mice by immunofluorescence microscopy and immunoblotting. Integrin  $\beta$ 1D fluorescence was significantly diminished in  $\beta$ 1D<sup>-/-</sup> myocardium, as compared to WT control hearts (0.23±0.089 *vs.* 1.00±0.153, p<0.001, n=4) (Figure 3C). Western blotting data confirmed a 77% reduction in integrin  $\beta$ 1D in  $\beta$ 1D<sup>-/-</sup> hearts (Figures 3D–E and Supplemental Figure 6).

Next, we examined total and phosphorylated RyR2 levels in the left ventricle of  $\beta 1D^{-/-}$  mice two weeks after the last dose of tamoxifen injection. The phosphorylation of RyR2 at Ser-2030 was increased by 72.5% in  $\beta 1D^{-/-}$  mice as compared to WT littermate controls. However, the phosphorylation levels of RyR2 at Ser-2808 and Ser-2814 did not differ between  $\beta 1D^{-/-}$  and WT mice (Figures 3D and 3F–H). These data are consistent with the studies performed using human ARVC samples.

Having shown that the  $\beta 1D^{-/-}$  mouse model replicates the molecular findings in human ARVC patient hearts, we next characterized the in vivo effects of cardiac-specific integrin  $\beta 1D$  knockout on cardiac function two weeks after the last dose of tamoxifen injection. Compared with WT littermates,  $\beta 1D^{-/-}$  mice displayed no overt alterations in cardiac function in terms of echocardiographic parameters Supplemental Figure 7), histology or cardiomyocyte size (Masson and WGA staining, respectively) (Supplemental Figure 8). We also observed no differences in the distribution and expression levels of key excitation-contraction (E-C) coupling proteins or the organization of T-tubules in cardiomyocytes (Figure 3I–K, Supplemental Figure 9). Cardiac fibrosis was dramatically increased 3 months later in  $\beta 1D^{-/-}$  mice (Supplemental Figure 10). These data indicate that chronic integrin  $\beta 1D$  deficiency may contribute to cardiac fibrosis in ARVC.

Hyperactivation of RyR2 is linked to the increased propensity for stress-induced cardiac arrhythmias such as CPVT and atrial fibrillation.<sup>13–15, 30–33</sup> We then analyzed the electrophysiological properties of  $\beta 1D^{-/-}$  and WT mice under baseline conditions and cardiac stress. We found that  $\beta 1D^{-/-}$  mice have normal ECG recordings and regular heart rhythm at rest (Figures 4A–D, Supplemental Figure 11). However, upon sympathetic stress through administration of epinephrine (1.6 mg/kg) and caffeine (120 mg/kg),  $\beta 1D^{-/-}$  mice more readily developed ventricular arrhythmias than WT mice, characterized by irregular heart rates, sustained VT episodes, typical bidirectional ventricular arrhythmias and prolonged tachycardias (Figures 4B–H). These data clearly demonstrate that integrin  $\beta 1D$  deficiency predisposes the heart to stress-induced, CPVT-like ventricular arrhythmias.

# $\beta$ 1D<sup>-/-</sup> cardiomyocytes exhibit RyR2 Ser-2030 hyper-phosphorylation, delayed afterdepolarizations and spontaneous Ca<sup>2+</sup> release under $\beta$ -adrenergic stimulation.

Given that the  $\beta 1D^{-/-}$  mice exhibited CPVT-like ventricular arrhythmias under stress, we next tested if there is any change in RyR2 Ser-2030 and Ser-2808 phosphorylation levels in hearts from WT and  $\beta 1D^{-/-}$  mice under  $\beta$ -adrenergic stimulation. In addition to an increase in basal Ser-2030 phosphorylation (by 72%, Figure 5A–B), we found that ISO markedly increased RyR2 phosphorylation levels at Ser-2030 in both strains (by 78±7%, WT *vs.* by  $49\pm11\%$ ,  $\beta 1D^{-/-}$ , p<0.001). In contrast,  $\beta 1D$  deficiency only results into a small trend toward an increase in Ser-2808 phosphorylation level at baseline (by 19% compared to WT). ISO stimulation also induced no more than a 20% increase in Ser-2808 phosphorylation ( $20\pm8\%$ , WT, p=0.043 *vs.*  $16\pm9\%$ ,  $\beta 1D^{-/-}$ , p=0.057) (Figures 5A–C). In addition, using anti-phospho antibodies against Ser-2808 from different sources, we found similar moderate increases in RyR2 phosphorylation levels in WT hearts in response to ISO (Supplemental Figure 12). These results indicate that integrin  $\beta 1D$  regulates RyR2 phosphorylation primarily on Ser-2030, and RyR2 Ser-2030 hyper-phosphorylation may be responsible for adrenergic stimulation-induced VT.

To investigate the role of integrin  $\beta$ 1D on myocyte electrophysiological properties, we recorded action potentials (APs) from WT and  $\beta$ 1D<sup>-/-</sup> cardiomyocytes in the absence or presence of 100 nM ISO during and after pacing stimulation. In the absence of ISO, spontaneous DADs were not found in WT but were present in 10% (3/30) of  $\beta$ 1D<sup>-/-</sup> cells. In the presence of ISO, 4% (1/25) WT cardiomyocytes showed spontaneous DADs which

increased to 66.67% (20/30) in  $\beta 1D^{-/-}$  cardiomyocytes (P<0.01) (Figures 5D–E). Moreover, triggered activities were not observed in WT cardiomyocytes but occurred in 50% (15/30) of  $\beta 1D^{-/-}$  cardiomyocytes (p<0.01) in the presence of ISO (Figure 5F).

It is well accepted that diastolic spontaneous Ca<sup>2+</sup> release from the SR underlies DAD and triggered activity.<sup>34</sup> To determine the contribution of RyR2-mediated SR Ca<sup>2+</sup> release in the increased arrhythmogenesis we saw in  $\beta 1D^{-/-}$  cardiomyocytes, we measured the occurrence of spontaneous Ca<sup>2+</sup> transients (SCaTs) after halting regular pacing (2 Hz) in Fluo-4AMloaded cardiomyocytes. In the absence of ISO stimulation, 13% (4/30) of  $\beta 1D^{-/-}$  cells exhibited SCaTs, while all WT cells remained quiescent after suspension of pacing. In the presence of ISO, however, 73% (22/30) of the  $\beta 1D^{-/-}$  cardiomyocytes developed SCaTs compared with only 4% (1/25) of the WT cardiomyocytes (P<0.05) (Figures 5G-H). Moreover, 47% (14/30) of the  $\beta 1D^{-/-}$  cells but no WT cells developed triggered events (P<0.01) (Figures 5G and I). To test whether spontaneous SR Ca<sup>2+</sup> release underlies the abnormal electric activities, we pretreated  $\beta 1D^{-/-}$  cardiomyocytes with ryanodine (1  $\mu$ M). Here, ryanodine significantly inhibited spontaneous Ca<sup>2+</sup> waves and, subsequently, DADs and triggered activities in  $\beta 1D^{-/-}$  cardiomyocytes under ISO stimulation (Supplemental Figure 13). Taken together, these data suggest that  $\beta 1D^{-/-}$  cardiomyocytes are more susceptible to  $Ca^{2+}$ -dependent arrhythmogenesis due most likely to abnormal  $Ca^{2+}$  handing caused by RyR2 hyperphosphorylation.

## Enhanced Ca<sup>2+</sup> spark activities and reduced SR Ca<sup>2+</sup> levels in $\beta$ 1D deficient cardiomyocytes

Ca<sup>2+</sup> sparks are elementary Ca<sup>2+</sup> release events mediated by Ca<sup>2+</sup> efflux through RyR2s, in cardiomyocytes, representing the gating properties of RyR2s.<sup>35</sup> We measured Ca<sup>2+</sup> sparks in quiescent Fluo-4–loaded cardiomyocytes. We found that the frequency of Ca<sup>2+</sup> sparks was significantly increased in  $\beta 1D^{-/-}$  cardiomyocytes compared to WT cardiomyocytes at baseline (0.98±0.064 *vs.* 2.45±0.472, P<0.05). In the presence of ISO, Ca<sup>2+</sup> spark frequency was further increased (2.23±0.054 *vs.* 5.96±0.39, P<0.01) (Figures 6A–B). Ca<sup>2+</sup> spark amplitude showed the same trend as the Ca<sup>2+</sup> spark frequency (Figure 6C). The full width or full duration at half-maximum (FWHM or FDHM) and was unchanged in  $\beta 1D^{-/-}$  cardiomyocytes (Figure 6D). In addition, we assessed SR Ca<sup>2+</sup> content by measuring caffeine-induced Ca<sup>2+</sup> transients after cessation of pacing followed by rapid application of caffeine (10 mM, Figures 6F–H). We found that  $\beta 1D^{-/-}$  cardiomyocytes showed significantly lower SR Ca<sup>2+</sup> levels than WT cardiomyocytes (3.43 ±0.013 *vs.* 2.78±0.025, P<0.01) in the presence of ISO. Collectively, these data implicate that  $\beta 1D$  deficiency leads to altered RyR2 function consistent with the biochemical evidence aforementioned, i.e., hyperphosphorylation of RyR2 at Ser-2030.

### DSP deficiency leads to elevated ubiquitination and degradation of integrin $\beta$ 1D through an ERK1/2 – Fibronectin pathway

Loss of desmosomal proteins is considered a molecular hallmark of ARVC.<sup>9</sup> We identified severe downregulation of desmoplakin (DSP) in human ARVC tissues by immunostaining (Figure 7A & B), western blots (Figures 7C & D,  $1.00 \pm 0.09 \text{ vs}$ .  $0.41 \pm 0.11$ , P<0.01) and protein mass spectrometry analyses (Supplemental Table 4). It has been suggested that DSP

loss triggers robust activation of ERK1/2 in cardiomyocytes.<sup>36</sup> Hyperactivation of ERK1/2 may boost the expression of fibronectin 1 (FN1),<sup>37</sup> an extracellular matrix ligand of integrin  $\beta$ 1, which promotes integrin  $\beta$ 1 ubiquitination and degradation.<sup>38</sup> Indeed, our protein mass spectrometry analyses also showed FN1 is significantly upregulated in ARVC patient hearts (Supplemental Table 4). Our western blotting analyses demonstrate 3-fold higher levels in FN1 and active phospho/total ERK1/2 ratios in ARVC heart samples, compared to those of healthy donors (Figures 7C, E–F & Supplemental Figure 14). We postulated that downregulation of integrin  $\beta$ 1D may result from a pathological ERK1/2-dependent fibronectin induction pathway in ARVC cardiomyocytes. To investigate whether DSP loss is responsible for ERK1/2 activation and fibronectin elevation, we used DSP siRNA to specifically knock down DSP in HL-1 cells and then quantified the expression levels of candidate downstream proteins. Consistent with our hypothesis, DSP knockdown (KD, by ~70%) resulted into a marked increase in ERK1/2 phosphorylation, FN1 protein expression and degradation of integrin  $\beta$ 1D protein (Figures 7G–K) comparable to that in ARVC hearts. Furthermore, treatment with an ERK specific inhibitor (FR180204) or knockdown of FN1 prevented β1D degradation in DSP-KD HL-1 cells (Supplemental Figure 15 & 16). Quantitative PCR (qPCR) analysis revealed that integrin  $\beta$ 1D transcript levels are similar between ARVC and HD hearts and between control and DSP-KD HL-1 cells, suggesting that DSP-dependent regulation of integrin B1D expression is not related to changes at the transcript level Supplemental Figure 7).

To determine whether integrin  $\beta$ 1D is ubiquitinated in response to DSP loss, immunoprecipitates of endogenous integrin  $\beta$ 1D from control and DSP-KD HL-1 cells were immunoblotted for endogenous ubiquitin. Using the mono/polyubiquitin antibody FK2, it was observed that integrin  $\beta$ 1D was ubiquitinated at a much higher level in DSP-KD samples compared with control group (Figures 7L–M). Further experiments demonstrated that inhibition of lysosomal degradation (with chloroquine), but not inhibition of proteasome function (with MG132) protects against DSP-KD induced Integrin  $\beta$ 1D downregulation (Figure 7N). These data suggest that integrin  $\beta$ 1D is not directly degraded by the ubiquitinproteasome system but targeted by ubiquitin for degradation in the lysosome.

### DISCUSSION

ARVC is an inherited desmosomal cardiomyopathy characterized by a high burden of ventricular arrhythmias and increased risk for SCD.<sup>8, 9</sup> Ventricular arrhythmias in ARVC are often related to exercise or emotional excitement, suggesting that they are sensitive to catecholamines.<sup>22, 23, 39</sup> Indeed, clinical studies have shown a high inducibility of polymorphic premature ventricular contractions or ventricular tachycardia during isoproterenol infusion testing in ARVC patients.<sup>22, 23</sup> However, the arrhythmogenic mechanism of ARVC remains elusive. Here, in the present study, we reveal for the first time a critical role of integrin  $\beta$ 1D-mediated Ca<sup>2+</sup> signaling in the arrhythmogenesis of ARVC.

In this study, using non-biased protein mass spectrometry, we first identified that integrin  $\beta$ 1D protein is significantly downregulated in ARVC patient heart tissue in parallel with reduction of desmosomal proteins such as DSP, DSC2 and DSG2. On the contrary, integrin  $\beta$ 1D is upregulated in human ICM and HCM, suggesting that integrin  $\beta$ 1D downregulation

is a unique change in ARVC. Functionally, integrin  $\beta 1D$  is linked to RyR2 phosphorylation at Ser-2030. We observed RyR2 hyper-phosphorylation at Ser-2030 in ARVC, but not in ICM or HCM heart samples. Exogenous integrin  $\beta 1D$  decreases PKA-induced RyR2 Ser-2030 phosphorylation and RyR2 channel open probability in samples isolated from hearts with ARVC. These results indicate that integrin  $\beta 1D$  stabilizes RyR2 via its influence on Ser-2030 phosphorylation. Importantly, we validated the in vivo function of integrin  $\beta 1D$ in cardiomyocytes using a cardiac inducible integrin  $\beta 1D$  specific knockout mouse model. We demonstrated that integrin  $\beta 1D$  deletion in mice for only two weeks, a time point preceding overt structural and functional phenotypes, results in CPVT-like ventricular arrhythmias, RyR2 Ser-2030 hyper-phosphorylation, SR Ca<sup>2+</sup> leak, DADs and triggered activities under isoproterenol stress. Lastly, we revealed that loss of desmoplakin causes integrin  $\beta 1D$  deficiency in ARVC, which is mediated through an ERK1/2 - fibronectin ubiquitin/lysosome pathway. Taken together, our study supports an important role for integrin  $\beta 1D$  deficiency and subsequent RyR2 Ser-2030 hyper-phosphorylation on mediating catecholamine-sensitive ventricular tachycardia in patients with ARVC.

Integrin  $\beta$ 1D deficiency is a unique molecular change observed in ARVC, which is not seen in other forms of human heart diseases including ischemic cardiomyopathy and hypertrophic cardiomyopathy. More interestingly, integrin  $\beta$ 1D is consistently downregulated among patients with distinct ARVC mutations, suggesting that integrin  $\beta$ 1D deficiency is a common consequence of ARVC, independent of heritable causation. ARVC is characterized by intercalated disk remodeling and loss of desmosomes.<sup>40</sup> Loss of desmosomal proteins is considered a molecular hallmark of ARVC. We did observe significant downregulation of multiple desmosomal proteins in human ARVC. This is confirmed by our data from protein mass spectrometry analyses. DSP knockdown in HL-1 cells triggered the robust activation of ERK1/2 and supports previous findings in neonatal rat ventricular myoyctes.<sup>36</sup> Furthermore, we were surprised to find ERK1/2 activation in DSP deficient cells is required for the induction of fibronectin 1 and subsequent loss of integrin  $\beta$ 1D through ubiquitination and lysosome-mediated degradation. These mechanistic studies are sufficient to explain how defects in desmosome function in ARVC patients produce similar reductions in integrin  $\beta$ 1D levels (Figure 8).

Previous studies have shown that in addition to localizing to intercalated discs, integrin  $\beta$ 1D also distributes to T-tubule regions of cardiomyocytes,<sup>25, 26</sup> where it regulates RyR2 phosphorylation levels at Ser-2808.<sup>29</sup> However, our combined results from ARVC heart tissue, isolated ARVC patient cardiomyocytes and our  $\beta$ 1D knockout mouse model strongly suggests that the level of integrin  $\beta$ 1D is linked to the status of RyR2 phosphorylation, particularly at Ser-2030, rather than at Ser-2808 or Ser-2814 sites (See Figures 1E–I, 2I–J, 3D–H, 5A–C). RyR2 phosphorylation is complex, and has been a controversial topic, especially regarding its role in mediating heart failure development.<sup>41–46</sup> Our results showing integrin  $\beta$ 1D antagonizes PKA-induced RyR2 phosphorylation at Ser-2030 (Figures 2I & 5A) are consistent with previous observations from Wayne Chen's group who reported that Ser-2030 is the major PKA site responding to  $\beta$ -adrenergic stimulation.<sup>47</sup> Our data also demonstrate that when integrin  $\beta$ 1D expression level is reduced,  $\beta$ -adrenergic stimulation induces stronger RyR2 phosphorylation at Ser-2030 (Figure 5A–C), leading to aberrant SR Ca<sup>2+</sup> activity and Ca<sup>2+</sup>-dependent arrhythmias. Taken together, we conclude

that loss of integrin  $\beta$ 1D results into exaggerated RyR2 phosphorylation at Ser-2030, thereby enhancing propensity to catecholamines-sensitive ventricular tachycardia in both mice and ARVC patients. On the other hand, we observed increased expression level of integrin  $\beta$ 1D in ICM and HCM. We postulate that upregulation of integrin  $\beta$ 1D may provide a protective mechanism prohibiting the incidence of cardiac arrhythmias in these diseases.

The contribution of dysfunctional Ca<sup>2+</sup> handling to the arrhythmogenic mechanism of ARVC has begun to be appreciated.<sup>24, 48</sup> Recent studies in PKP2 knockout mice from the Delmar group reported that loss of PKP2 leads to reduced expression of a group of proteins related to Ca<sup>2+</sup> handling, i.e., RyR2, Ankyrin-B, Ca<sub>v</sub>1.2 Ca<sup>2+</sup> channel, triadin and calsequestrin-2. These factors together contribute to intracellular Ca<sup>2+</sup> dysregulation and therefore the Ca<sup>2+</sup>-dependent arrhythmia phenotype in this mouse model.<sup>24, 48</sup> However, our protein mass spectrometry analyses failed to identify any significant changes in these Ca<sup>2+</sup> handling proteins in human ARVC (Figure 1B). Thus, whether findings from this study can be translated to human ARVC remains for to be investigated.

Our study has significant implications for current clinical practice. Although it is generally agreed that ARVC and CPVT are caused by mutations of different sets of genes involved in different functions at molecular and cellular levels in cardiomyocytes, we now recognize that there is some degree of convergence between ARVC and CPVT. This indeed has raised the question how clinicians and invasive electrophysiologists are to be informed on differential diagnoses of the two diseases. This is particularly more critical and difficult at the early stage of the disease when no structural findings are present in patients. In addition to medical history, treadmill test, electrophysiological examination with isoproterenol infusion, genetic test, careful examinations of cardiac MRI may provide valuable information in detecting subtle early changes in cardiac structure. For patient management, a beta blocker is strongly recommended as long as patients have CPVT-like ventricular arrhythmias. Our previous study has suggested that carvedilol is the only beta blocker with the additional benefit of suppressing RyR2 activities,<sup>49</sup> which may provide better outcomes than other beta blockers in the treatment of ARVC.

At present, there is still no effective treatment for ARVC, except for cardiac transplantation, due to a lack of mechanistic understanding of the ventricular arrhythmias and SCD in ARVC.<sup>50</sup> Here, we identify a novel pathway that integrin  $\beta$ 1D deficiency mediates RyR2 dysfunction and contributes to catecholamine-sensitive ventricular tachycardia in ARVC. Interventions targeting the signaling pathway that leads to integrin  $\beta$ 1D degradation or stabilizing RyR2 function could be promising approaches for preventing ventricular arrhythmias and SCD in ARVC patients.

In summary, we identified downregulation of integrin  $\beta$ 1D is a novel cause of catecholamine-sensitive ventricular tachycardia in ARVC. Our data implicate integrin  $\beta$ 1D deficiency as a mechanism leading to hyperphosphorylation of RyR2 at Ser-2030, aberrant Ca<sup>2+</sup> handling, DAD and triggered ventricular arrhythmia in ARVC, particularly under sympathetic stress. Taken together, these findings provide novel insights into the role of integrin  $\beta$ 1D deficiency in the pathogenesis of cardiac arrhythmias in ARVC, which may

have important implications for future development of new therapeutic strategies to ameliorate this condition.

### **Supplementary Material**

Refer to Web version on PubMed Central for supplementary material.

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### Nonstandard Abbreviations and Acronyms:

ARVC	Arrhythmogenic right ventricular cardiomyopathy
Bin1	Amphyphisin-2
BVT	Bidirectional ventricular tachycardia
CaM	Calmodulin
CaV1.2	L-type Ca <sup>2+</sup> channel
CPVT	Catecholaminergic polymorphic ventricular tachycardia
CSAQ2	Calsequestrin
CX43	Connexin-43
DADs	Delayed afterdepolarizations
DHPR	Dihydropyridine
DSC2	Desmocollin 2
DSG2	Desmoglein 2
DSP-KD	Desmoplakin knockdown
DSP	Desmoplakin
ECG	Electrocardiogram
ERK1/2	Extracellular signal-regulated kinase 1 and 2
FN1	Fibronectin1
GAPDH	Glyceraldehyde 3-phosphate dehydrogenase

GST-β1D	GST-tagged integrin $\beta$ 1D fusion protein
GST	Glutathione S-transferase
НСМ	Hypertrophic cardiomyopathy
HD	Healthy donor
ICM	Ischemic cardiomyopathy
IP3R	Inositol 1,4,5-trisphosphate receptor
ISO	Isoproterenol
JCTN	Junctin
JP2	Junctophilin-2
MVT	Monomorphic broad complex VT
NCX	Sodium-calcium exchanger
PBS	Phosphate-buffered saline
РКА	Protein kinase A
РКР2	Plakophilin 2
PLB	Phospholamban
PVC	Premature ventricular contraction
PVT	Polymorphic VT
RT-PCR	Reverse transcriptase coupled polymerase chain reaction
RyR2	Ryanodine Receptor 2
SCaTs	Spontaneous Ca <sup>2+</sup> transients
SCD	Sudden cardiac death
SERCA2a	SR Ca <sup>2+</sup> -ATPase
SR	Sarcoplasmic reticulum
ST	Sinus tachycardia
TRDN	Triadin
VT	Ventricular tachycardia
β1D <sup>-/-</sup>	Cardiac specific $\beta$ 1D knockout

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### **CLINICAL PERSPECTIVE**

### What Is New?

- We identify that integrin  $\beta$ 1D is uniquely downregulated in ARVC, but not in HCM or ICM.
- Exogenous integrin β1D decreases PKA-induced RyR2 Ser-2030 phosphorylation and stabilizes RyR2 channels in samples isolated from hearts with ARVC.
- Integrin β1D deletion in mice results in CPVT-like ventricular arrhythmias, RyR2 Ser-2030 hyper-phosphorylation, SR Ca<sup>2+</sup> leak, DADs and triggered activities under isoproterenol stress.
- We reveal that loss of desmoplakin mechanistically contributes to integrin β1D deficiency in ARVC, which is mediated through an ERK1/2 - fibronectin
  ubiquitin/lysosome pathway.

### What Are the Clinical Implications?

- Our study indicates that loss of integrin β1D may be responsible for ventricular arrhythmias in ARVC disease.
- Interventions that stabilize integrin  $\beta$ 1D levels through inhibition of its signalspecific degradation or by integrin  $\beta$ 1D overexpression could reduce ventricular arrhythmias and SCD in ARVC patients.





Figure 1. Left ventricular tissue from ARVC patients is characterized by integrin  $\beta 1D$  down-regulation and RyR2 Ser-2030 hyper-phosphorylation.

(A) Representative ECG recordings from four patients with ARVC obtained during hospitalization but before cardiac transplantation. (B) Quantification of proteins associated with intracellular calcium release from the sarcoplasmic reticulum from protein mass spectrometry analyses. (C) Representative immunofluorescence images obtained using antiserum against integrin β1D in fixed left ventricle tissue sections from healthy donor (HD, left) or ARVC patients (right). (D) Quantification of integrin β1D Immunofluorescence intensity (n=4 per group; Each group includes 5 slides). (E) Representative immunoblots for integrin β1D, total RyR2, pRyR2 Ser-2808, pRyR2 Ser-2814, pRyR2 Ser-2030 and GAPDH protein expression in left ventricular tissue homogenates from HDs or patients with ARVC. (F) Quantification of integrin β1D immunoblots shown in (E) relative to GAPDH expression (n=4 per group). Note that both mature and precursor forms of β1D integrin subunit were reduced. (G-I) Quantification of immunoblots showing increased phosphorylation of RyR2

at Ser-2030 in left ventricle tissue from patients with ARVC compared to HD tissue HD. There were no significant differences between the groups in RyR2 phosphorylation at Ser-2808 or Ser-2814. (n=4 per group). The data represent the means  $\pm$  SEM; \*P<0.05; \*\*\*P<0.001; n.s., not significant; Student's t test. Abbreviations: ARVC, Arrhythmogenic right ventricular cardiomyopathy; HD, Healthy Donors;  $\beta$ 1D, integrin  $\beta$ 1D; RyR2, Ryanodine Receptor 2; BIN1, Bridging integrator 1; Ca<sub>v</sub>1.2, L-type Ca<sub>v</sub>1.2 calcium channel; CaM, Calmodulin; CASQ2, Calsequestrin 2; IP3R2; Inositol 1,4,5-trisphosphate receptor, type 2; ITGB1, Integrin beta-1; JCTN, Junctin; JP2, Junctophilin-2; NCX1, Na<sup>+</sup>/Ca<sup>2+</sup> exchanger, isoform 1; PLB, phospholamban; SERCA2a; Sarcoplasmic / endoplasmic reticulum Ca<sup>2+</sup> ATPase 2a; TRDN, Triadin; GAPDH, glyceraldehyde 3-phosphate dehydrogenase.



#### Figure 2. Integrin β1D attenuates PKA-mediated RyR2 activation.

(A-E) Single channel activity of RyR2 from SR vesicles isolated from an ARVC heart (case 1). RyR2 channels were recorded in a solution containing 250 mM KCl and 25 mM Hepes (pH 7. 4) 1 mM ATP, 5 mM MgCl<sub>2</sub>, and 260 nM Ca<sup>2+</sup> (cytosolic) and 2.5 mM Ca<sup>2+</sup> (luminal). Single RyR2-channel activity after pre-incubation with: 1  $\mu$ M GST for 30 min followed by 15 min treatment with 1  $\mu$ l of PBS (A), 1 $\mu$ M GST- $\beta$ 1D for 30 min followed by 15 min treatment with 1  $\mu$ l of PBS (B), 1 $\mu$ M GST for 30 min followed by 15 min treatment with 1  $\mu$ M GST- $\beta$ 1D for 30 min followed by 15 min treatment with 1  $\mu$ M GST- $\beta$ 1D for 30 min followed by 15 min treatment with 1 $\mu$ M PKA (C), or 1 $\mu$ M GST- $\beta$ 1D for 30 min followed by 15 min treatment with 1 $\mu$ M PKA (D); or 10 $\mu$ M ryanodine to confirm RyR2 channel identity (E). Recording potentials, -20 mV. Zero-current baselines are indicated (short bar). (F-H) Summary data from RyR2 single channel recordings determining mean open time (To, F), open probability (Po, G) and mean closed time (CT, H). (A-H), n=3 independent experiments for each group. (I) Representative immunoblots for total RyR2, pRyR2 Ser-2808, pRyR2 Ser-2814, and pRyR2

Ser-2030 protein levels in samples with the treatments described in (A-D). (J), Quantification of pRyR2 Ser-2808, pRyR2 Ser-2814 and pRyR2 Ser-2030 phosphorylation levels normalized to total RyR2 protein expression. n = 4 independent experiments for each group. The data represent the means  $\pm$  SEM; \*P<0.05; \*\*P<0.01; n. s., not significant; Student's t test. Abbreviations: PKA, Protein kinase A; PBS, Phosphate-buffered saline; GST, glutathione S-transferase; GST- $\beta$ 1D, GST-tagged integrin  $\beta$ 1D fusion protein;  $\beta$ 1D, integrin  $\beta$ 1D; RyR2, Ryanodine Receptor 2.



**Figure 3. Generation and evaluation of the \beta 1D^{-/-} mouse model under basal conditions.** (A) Schematic of the targeting vector and generation of  $\beta 1D^{-/-}$  mice. (B) RT-PCR analysis of integrin  $\beta 1D$  mRNA in ventricle samples from WT (Cre<sup>-</sup>/*Itgb1*<sup>f/f</sup>) and  $\beta 1D^{-/-}$  (Cre<sup>+</sup>/ *Itgb1*<sup>f/f</sup>) mice. (C) Representative immunostaining for integrin  $\beta 1D$  and nuclei in left ventricular tissue sections from  $\beta 1D^{-/-}$  and WT mice (Blue, DAPI stained nuclei; Red,  $\beta 1D$ ). Heart tissues were collected from mice two weeks after the last dose of tamoxifen injection. (D) Representative immunoblots for integrin  $\beta 1D$ , total RyR2, pRyR2 Ser-2030, pRyR2 Ser-2808, pRyR2 Ser-2814 and GAPDH protein levels in left ventricular tissue homogenates from WT and  $\beta 1D^{-/-}$  mice. (E-H) Quantitative assessment of the protein levels represented in (D). n=4 hearts/genotype. (I) Representative DHPR and RyR2 immunostaining of WT and  $\beta 1D^{-/-}$  isolated cardiomyocytes. (J) Representative Di-8-ANEPPS immunostaining of WT and  $\beta 1D^{-/-}$  isolated cardiomyocytes. (K) Mean TT power values, where TT power indicates the strength of the regularity of the T-tubule network.

n=25–30 cells from 3 hearts/genotype. The data represent the means  $\pm$  SEM; \*\*P<0.01; \*\*\*P<0.001; n. s., not significant; Student's t test. Abbreviations:  $\beta$ 1D, integrin  $\beta$ 1D;  $\beta$ 1D <sup>-/-</sup>, integrin  $\beta$ 1D knockout; GAPDH, glyceraldehyde 3-phosphate dehydrogenase; RT-PCR, reverse transcriptase coupled polymerase chain reaction; DHPR, Dihydropyridine (L-type Ca<sup>2+</sup> channel); RyR2, Ryanodine Receptor 2.



**Figure 4.**  $\beta$ **1D**<sup>-/-</sup> **mice develop cardiac arrhythmias in response to sympathetic stress.** (A-B) Representative ECG recordings from WT and  $\beta$ 1D<sup>-/-</sup> mice before (A) and 3–6 min after (B) intraperitoneal injection of epi/caff. (C) PR interval from WT and  $\beta$ 1D<sup>-/-</sup> mice before and after intraperitoneal injection of epi/caff. (D) Incidence of different types of VT (ST, PVC, MVT and BVT) calculated from  $\beta$ 1D<sup>-/-</sup> mice in response to epi/caff. (E) Percentage of VT type occurring in individual  $\beta$ 1D<sup>-/-</sup> mice. (F) VT occurred intermittently and the percentage (%) time in VT (VT duration) was measured within each of 10 consecutive 3-min periods immediately after epi/caff injection. (G) VT duration over the entire 30-min recording period. The data represent the means ± SEM for WT (n=11) and  $\beta$ 1D<sup>-/-</sup> (n=12) mice; Student t-test. \*\*P<0.01, \*\*\*P<0.001 vs. WT. Abbreviations: ECG=electrocardiogram; WT, Cre<sup>-</sup>/Itgb1<sup>f/f</sup> mice;  $\beta$ 1D<sup>-/-</sup>, integrin  $\beta$ 1D knockout; epi/caff= epinephrine (1.6mg/kg) and caffeine (120mg/kg). VT= Ventricular tachycardia; ST= Sinus tachycardia; PVC= Premature ventricular contraction; MVT= Monomorphic broad complex

VT; BVT= Bidirectional ventricular tachycardia; PVT= Polymorphic VT. \* / # = Different morphology of PVCs.



Figure 5. Integrin  $\beta$ 1D deficiency induces RyR2 Ser-2030, but not Ser-2808 hyperphosphorylation and promotes stress-induced DAD and spontaneous Ca<sup>2+</sup> release. (A-C) RyR2 Ser-2030 and Ser-2808 phosphorylation levels at baseline or in response to ISO stimulation (100nM perfusion for 30 min) in WT and  $\beta$ 1D<sup>-/-</sup> cardiomyocytes. (A) Representative immunoblots of RyR2 Ser-2030 and Ser-2808 phosphorylation levels. (A-B) The mean ratios of RyR2 phosphorylated at Ser-2030 (B) and Ser-2808 (C) to total RyR2 protein levels at baseline or under ISO treatment. n=4 hearts for each group. Data represent means ± SEM; \*\*\*P<0.001; Student's t test. (D) Representative traces of action potentials in WT and  $\beta$ 1D<sup>-/-</sup> cardiomyocytes ±100nM ISO. Small DADs (yellows arrows) and triggered activities (red arrows) were recorded after a pause following 2-Hz field stimulation. (E-F) Cumulative incidence of DADs (E) and triggered activities (F) in WT and  $\beta$ 1D<sup>-/-</sup> mice (n=25–30 cells per group from four hearts/genotype). (G) Representative traces showing SCaTs (yellow arrows) and triggered beats (red arrows) after a pause following 2-Hz field stimulation in WT and  $\beta$ 1D<sup>-/-</sup> cardiomyocytes ±100nM ISO. (H-I) Cumulative incidence of

SCaTs (H) and triggered beats (I) in WT and  $\beta 1D^{-/-}$  cardiomyocytes (n=25–30 cells per group from 4 hearts/genotype). \*\*P<0.01;  $\chi 2$  test. Abbreviations: DADs, delayed afterdepolarizations; SCaTs, spontaneous Ca<sup>2+</sup> transients.



### Figure 6. $\beta 1D^{-/-}$ cardiomyocytes display increased $\mathrm{Ca}^{2+}$ spark frequency and reduced SR $\mathrm{Ca}^{2+}$ content.

(A) Representative  $Ca^{2+}$  spark images in myocytes from WT and  $\beta 1D^{-/-}$  mice at baseline or under 100nM ISO treatment. (B) Mean frequency of  $Ca^{2+}$  sparks. (C) Mean amplitude of  $Ca^{2+}$  sparks. (D) Mean full width at half-maximum (FWHM) of the  $Ca^{2+}$  sparks. (E) Mean duration at half-peak amplitude (FDHM) of  $Ca^{2+}$  sparks. n=20-30 cells from 4 hearts/ genotype per group. (F-H) SR content was measured by measuring caffeine-induced  $Ca^{2+}$ release. (F) Representative traces of 1-Hz field stimulation-triggered  $Ca^{2+}$  transients and caffeine-induced  $Ca^{2+}$  release (SR content) from WT and  $\beta 1D^{-/-}$  cardiomyocytes with or without 100nM ISO treatment. (G) Average data of the amplitude of  $Ca^{2+}$  transients. (H) Cumulative data on the amplitudes of caffeine-induced SR  $Ca^{2+}$  release (i.e., SR  $Ca^{2+}$ content). n= 20–30 cells from 4 hearts/genotype for each group; The data represent the means  $\pm$  SEM; \*P<0.05; \*\*P<0.01; n.s., not significant; Student's t test.



**Figure 7. Downregulation of DSP promotes ubiquitination and degradation of integrin β1D.** (A) Representative immunofluorescent images of DSP in HD and ARVC heart sections (Blue, nucleus; Red, DSP; Green, actin). (B) Image quantification showing significantly lower DSP in patients with ARVC compared to HD (n=4 per group; 5 slides for each group; \*\*p<0.05, Student's t-test). (C) Representative immunoblots for DSP, ERK1/2, pERK1/2, FN1, integrin β1D and GAPDH protein expression in left ventricle tissue homogenates from HDs and patients with ARVC (n = 4 per group). (D-F) Quantification of immunoblots shown in (B) demonstrating reduced DSP protein, increased phosphorylation of ERK1/2 and FN1 in left ventricle tissues from patients with ARVC than in tissues from HD. (G) Representative immunoblots for DSP, ERK1/2, pERK1/2, FN1, integrin  $\beta$ 1D and GAPDH protein expression in the control and DSP-knockdown (DSP-KD) HL-1 cells (n=3). (H-K) Quantification of immunoblots of DSP, pERK1/2 / ERK1/2, FN1, and integrin  $\beta$ 1D protein expression in the control and DSP-KD HL-1 cells. (L) integrin  $\beta$ 1D was immunoprecipitated

from control and DSP-KD HL-1 cells. Samples were subsequently blotted for integrin  $\beta$ 1D and ubiquitin. (M) Whole cell lysate immunoblots from samples used for immunoprecipitation in (L) (n=4). (N) The lysosomal inhibitor Chloroquine was able to restore  $\beta$ 1D expression levels in response to DSP-KD in HL cells (left). MG132, a potent proteasome inhibitor, was used to assess role of proteasomal degradation of  $\beta$ 1D caused by DSP loss in HL-1 cells (right) (n = 3 independent experiments, respectively). The data represent means ± SEM; \*\*P<0.01; \*\*\*P<0.001; n. s., not significant; Student's t test. Abbreviations: DSP, Desmoplakin; ERK1/2, extracellular signal-regulated kinase 1 and 2; FN1, Fibronectin1; DSP-KD, Desmoplakin knockdown; GAPDH, glyceraldehyde 3-phosphate dehydrogenase.



### Figure 8. Schematic to illustrate the proposed mechanism of catecholamine-sensitive VT in ARVC.

ARVC from loss of function mutations in desmosomal genes results in robust increase in pERK1/2 and the induction of fibronectin expression. Subsequent fibronectin secretion leads to the ubiquitination and degradation of integrin  $\beta$ 1D. Decreased integrin  $\beta$ 1D increases Ser-2030 phosphorylation on RyR2 and impairs RyR2 stability, thereby causing enhanced Ca<sup>2+</sup> leak that promotes VT under ISO stress.