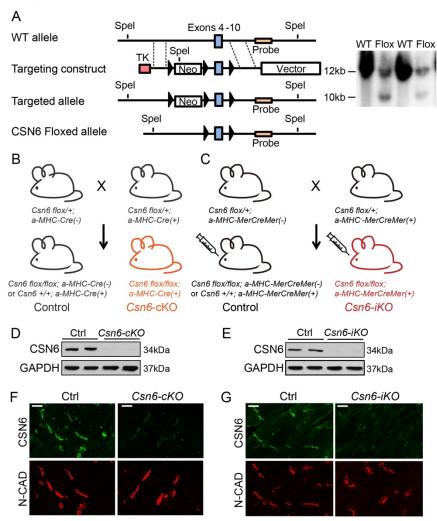
SUPPLEMENTAL DATA

Desmosomal COP9 regulates proteome degradation in arrhythmogenic right ventricular dysplasia/cardiomyopathy

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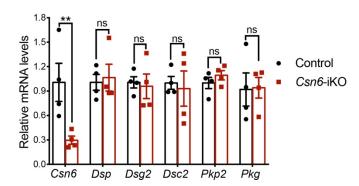
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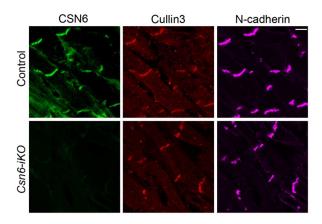


Supplemental Figure 1: Generation and validation of CSN6 loss in cardiac specific and cardiac inducible *Csn6* knockout mice. (A) Restriction map of the relevant genomic region of *Csn6*, targeting construct, the mutated locus following recombination and *Csn6* floxed allele (left). TK: thymidine kinase gene, Neo: neomycin resistance gene. Detection WT and mutant (target) alleles by southern blot analysis (right). (B) Schematic representation of breeding strategy to generate *Csn6*-cKO (*Csn6 flox/flox*; *a-MHC-Cre(+)*) and littermate control (*Csn6 flox/flox*; *a-MHC-Cre(-)*) or *Csn6+/+*; *a-MHC-Cre(+)*) mice. (C) Schematic representation of breeding strategy to generate *Csn6*-iKO (*Csn6 flox/flox*; *a-MHC-MerCreMer(+)*) and littermate control (*Csn6 flox/flox*; *a-MHC-MerCreMer(+)*) mice. To induce ablation of *Csn6*, 5- to 6-week-old mice were treated with daily intraperitoneal injection of tamoxifen (Sigma, 50µg/g per day) for 5 consecutive days. (D and E) Protein blot analysis of CSN6 in total

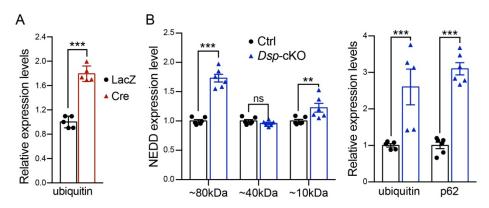
protein extracts from *Csn6*-cKO and control mice at 2 weeks of age (**D**) and *Csn6*-iKO mice at 2 weeks post tamoxifen injection (**E**) (n=4 mice). GAPDH is used as a loading control. (**F** and **G**) Immunofluorescence staining of CSN6 (green) and N-cadherin (red) in cardiac sections from *Csn6*-cKO and control mice at 2 weeks of age (**F**) and *Csn6*-iKO at 2 weeks post tamoxifen injection (**G**). Scale Bar: 10 μm.



Supplemental Figure 2: RNA levels of desmosomal genes are not impacted in *Csn6* knockout hearts. Quantitative RT-PCR analysis of *Csn6* and desmosomal gene expression in *Csn6*-iKO and control mouse hearts at 6 weeks post-tamoxifen injection (n=4 mice). *18S* and *Gapdh* RNA were used as loading controls and showed similar results. Data are mean \pm s.e.m; two-way ANOVA with Sidak multiple comparison test, **P<0.01, ns, not significant.

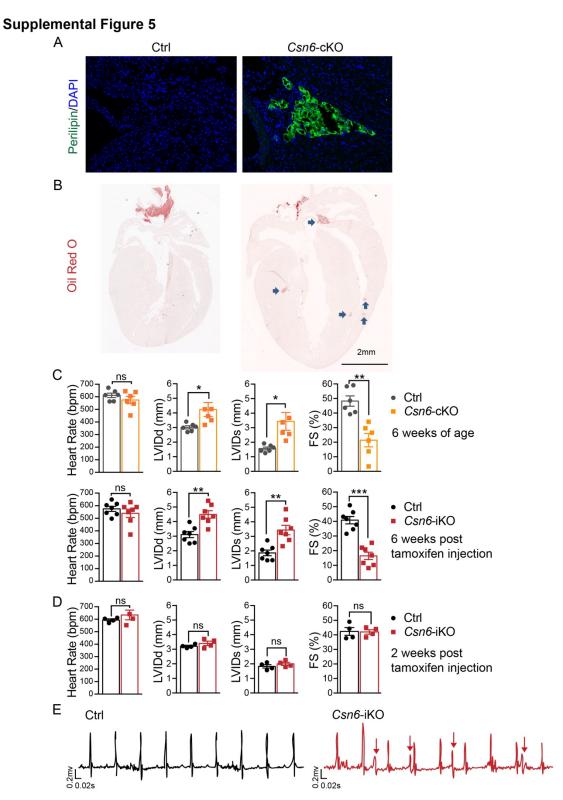


Supplemental Figure 3: Cullin 3 is localized to the cardiac cell-cell junction in adult WT and *Csn6*-iKO mouse hearts. Immunofluorescence staining of Cullin 3 (red), CSN6 (green) and N-cadherin (magenta) in control and *Csn6*-iKO mouse hearts at 6 weeks post tamoxifen injection (n=3 per group). Scale bar: 10μm.



Supplemental Figure 4: Quantification of protein expression levels in Figure 4 and Figure 5.

(A) Quantification of protein expression levels in Figure 4A. The expression levels of the proteins were normalized to the loading control GAPDH. Data are mean \pm s.e.m; two-tailed Student's t-test. ***P<0.001. (B) Quantification of protein expression levels in Figure 5A. The expression levels of the proteins were normalized to the loading control GAPDH. Data are mean \pm s.e.m;, two-way ANOVA with Sidak multiple comparison test, **P<0.01, ***P<0.001, ns, not significant.



Supplemental Figure 5: Characterization of cardiac specific *Csn6* knockout mice. (A) Immunofluorescence staining of perilipin in triangular area of cardiac sections in *Csn6*-cKO and control mice at 6 weeks of age. Cardiac sections were counterstained with DAPI nuclear stain

(blue). Scale Bar: 100 µm. (**B**) Representative Oil Red O stained whole cardiac sections from Csn6-cKO and control mice at 6 weeks of age (n=4 per group). Scale bar: 2mm.(**C**) Echocardiographic M-mode analysis of heart function in Csn6 deficient and control mice at 6 weeks of age and post-tamoxifen injection (n=6 mice). FS %: percent fractional shortening, LVIDd: left ventricular internal dimension at end-diastole. LVIDs: left ventricular internal dimension at end-systole. Data are mean \pm s.e.m.; two-tailed Student's t-test, *P<0.05; **P<0.01; ***P<0.001; ns, not significant. (**D**) Echocardiographic M-mode analysis of heart function in Csn6-iKO and control mice at 2 weeks post tamoxifen injection (n=4 mice). Data are mean \pm s.e.m.; two-tailed Student's t-test, ns, not significant. (**E**) Representative conscious ECG tracings from Csn6-iKO (red) and control mice (black) at 2 weeks post tamoxifen injection (n=4 mice). Arrows denotes ectopic beats.

Supplemental Figure 6 В C Mouse desmoplakin Fxon6 Exon8 Exon9 R sgRNA PAM WT ... ctttcttttcacagATACGCATGAGTCAACTGGAGGTC 243bp R327C ...ctttcttttcacagATATGCATGAGTCAACTGGAGGTC. D Ε WT Mut Mouse plakophilin Exon9 Exon11 PAM ... ggatgtgtctcacaacagATACCCACATTGGCTCGAA IVS10-1 G>C ... ggatgtgtctcacaacacATACCCACATTGGCTCGAA 9 1.2 1.0 DSP DSG2 1.2 1.0 WT Double Het CHX (h) 0 8 16 24 36 48 0 8 16 24 36 48 Relative protein le 0.8 DSP 280kDa Relative p. 0.0 WT Double Het DSG2 150kDa Double Het 16 24 36 48 16 24 36 48 PKP2 90kDa 1.2 1.0 PKG 1.2-1.0-PKP2 PKG 85kDa 0.8 0.6 0.8 **GAPDH** 0.6 Relative pr 0.4

800

700

600

200

3.5

3.0

(mm)

LVIDs

2.5-E) 2.0-

PO 1.5-

Supplemental Figure 6: Generation and characterization of knock-in mice harboring human equivalent DSP R315C and PKP2 IVS10-1 G>C mutations. (A and D) Schematic of the Cas9/sgRNA targeting site for the Dsp R327C (A) and Pkp2 IVS10-1 G>C (D) which are equivalent to the human R315C mutation and PKP2 IVS10-1 G>C respectively. The sgRNA coding sequence is underlined. The protospacer-adjacent motif (PAM) sequence is labeled in green. Uppercase: exon sequences. Lowercase: intron sequences. (B and E) Validation of Dsp R327C (**B**) and Pkp2 IVS10-1 G > C (**E**) allele by sequencing. (**C** and **F**) PCR using mutant-specific primers confirmed newborn mice contained the correct gene variant allele. (G and H) Representative protein blots (G) and quantification analyses (H) of desmosomal proteins in ventricular cardiomyocytes which isolated from Double Het and WT pups. The cardiomyocytes

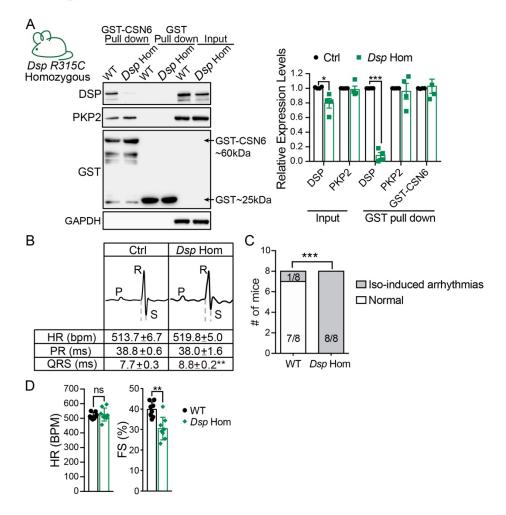
ഗ 30 34 % 20 Relative 0.2 0.0

16 24 36 48

WT

Double Het

were treated with cycloheximide (CHX, $10 \mu g/ml$) for 0, 8, 16, 24, 36 and 48 hours. The expression levels of proteins were normalized to the loading control GAPDH. Data are mean \pm s.e.m; two-way ANOVA with Sidak multiple comparison test, *P<0.05, **P<0.01. (I) Echocardiographic M-mode analysis of left ventricular dimensions and function in Double Het and WT mice at 4 months of age (n=6 mice). FS %: percent fractional shortening, LVIDd: left ventricular internal dimension at end-diastole. LVIDs: left ventricular internal dimension at end-systole. Data are mean \pm s.e.m.; two-tailed Student's t-test.



Supplemental Figure 7: Characterization of *Dsp R315C* homozygous mice. (A) Protein blot analyses and quantification of protein expression levels following GST pulldown assay. GST (~25kDa) and GST-CSN6 (~60kDa) protein were incubated with the heart extracts from WT and Dsp R315C homozygous mice (Dsp Hom) at 1 year of age (n=4 per group). The expression levels of input proteins were normalized to the loading control GAPDH. Data are mean \pm s.e.m; two-way ANOVA with Sidak multiple comparison test, *P<0.05, ***P<0.001. (B) Representative surface ECG tracings in Dsp R315C homozygous and WT mice at 1 year of age (n=8 per group). Quantification of heart rate (HR), PR and QRS intervals from ECG tracings was performed. Data from 100 ECG tracings per mouse were averaged for analysis. Data are mean \pm s.e.m; Student's two-tailed t-test, **P<0.01. QRS duration is indicated by the dotted line. (C) Quantification of number of Dsp R315C homozygous and WT mice which displayed arrhythmia with 2.5mg/kg

isoproterenol treatment during ECG recording at 1 year of age (n=8 per group). Chi-square test, ***P<0.05. (**D**) Echocardiographic M-mode analysis of left ventricular dimensions and function in *Dsp R315C* homozygous and WT mice at 1 year of age (n=8 per group). HR: heart rate; FS %: percent fractional shortening. Data are mean \pm s.e.m.; Student's two-tailed t-test. **P<0.01, no significant differences.

Supplemental Table 1: *Dsp R315C* and *Pkp2 IVS10-1 G>C* crRNA, trans-activating crRNA (tracrRNA), and single-strand oligodeoxynucleotides sequences used to generate knockin mice.

Oligo Name	Sequence (5' to 3')
Dsp R315C	acagauacgcaugagucaacGUUUUAGAGCUAUGCUGUUUUG
crRNA	
Pkp2 IVS10-1	cucacaacagauacccacauGUUUUAGAGCUAUGCUGUUUUG
$G > C \operatorname{crRNA}$	
tracrRNA	AAACAGCAUAGCAAGUUAAAAUAAGGCUAGUCCGUUAUCAAC
	UUGAAAAGUGGCACCGAGUCGGUGCU
Dsp R315C	AACAGAAATCTAACTGTGACTTGCTGTATGGACTGGTCCTTTCTTT
DNA	ACAGATATGCATGAGTCAACTGGAGGTCAAGGAAAAGGAACTCAAT
	AAGCTTAAACAAGAAAG
Pkp2 IVS10-1	AGAGAACTTCTCTGGTAGCAAATGTGATAGCATTTACAGGATGTGTC
DNA	TCACAACACATACCCACATTGGTGGCTCGAATGGTTGTCCAAAAGGA
	AAATGGTCTTCAGCATA

Supplemental Table 2: Primer sequences for real-time PCR analyses.

Gene	Primer 5'-3'
Csn6-F	GAGCTGGAGTTTCTGGGTTG
Csn6-R	GATCCGTTCAGCTTCCTCAG
Dsp-F	GCTCCATTACCAAGACTTCATC
Dsp-R	TGTCGTCGTCTCCAAACATCT
Dsg2-F	GAGGAATTGAGTGCAGCACATAC
Dsg2-R	CTTGCTTCCACCGTCAAGG
Dsc2-F	ATGCAGATGGGAGAAGCTGT
Dsc2-R	TGCAACAATTTCAGCAGAGG
<i>Pkp2-</i> F	GGCTCTCCAGAACCTCACAG
Pkp2-R	GGGAAAGATTCCGTGACAAA
Pkg-F1	CCTGTGGACTCTGCGCAAT
Pkg-R1	GACCAGGATCTTCAGCACACTCT