





Detrimental proarrhythmogenic interaction of Ca^{2+} /calmodulin-dependent protein kinase II and $\text{Na}_V1.8$ in heart failure

Philipp Bengel ^{1,2,7}, Nataliya Dybkova^{1,2,7}, Petros Tirilomis^{1,2,7}, Shakil Ahmad^{1,2,3}, Nico Hartmann^{1,2}, Belal A. Mohamed^{1,2}, Miriam Celine Krekeler^{1,2}, Wiebke Maurer^{1,2}, Steffen Pabel³, Maximilian Trum³, Julian Mustroph³, Jan Gummert⁴, Hendrik Milting⁴, Stefan Wagner³, Senka Ljubojevic-Holzer ⁵, Karl Toischer^{1,2}, Lars S. Maier³, Gerd Hasenfuss^{1,2}, Katrin Streckfuss-Bömeke ^{1,2,6,7} & Samuel Sossalla ^{1,2,3,7}✉

An interplay between Ca^{2+} /calmodulin-dependent protein kinase II δ c (CaMKII δ c) and late Na^+ current (I_{NaL}) is known to induce arrhythmias in the failing heart. Here, we elucidate the role of the sodium channel isoform $\text{Na}_V1.8$ for CaMKII δ c-dependent proarrhythmia. In a CRISPR-Cas9-generated human iPSC-cardiomyocyte homozygous knock-out of $\text{Na}_V1.8$, we demonstrate that $\text{Na}_V1.8$ contributes to I_{NaL} formation. In addition, we reveal a direct interaction between $\text{Na}_V1.8$ and CaMKII δ c in cardiomyocytes isolated from patients with heart failure (HF). Using specific blockers of $\text{Na}_V1.8$ and CaMKII δ c, we show that $\text{Na}_V1.8$ -driven I_{NaL} is CaMKII δ c-dependent and that $\text{Na}_V1.8$ -inhibition reduces diastolic SR- Ca^{2+} leak in human failing cardiomyocytes. Moreover, increased mortality of CaMKII δ c-overexpressing HF mice is reduced when a $\text{Na}_V1.8$ knock-out is introduced. Cellular and in vivo experiments reveal reduced ventricular arrhythmias without changes in HF progression. Our work therefore identifies a proarrhythmic CaMKII δ c downstream target which may constitute a prognostic and antiarrhythmic strategy.

¹Clinic for Cardiology & Pneumology, Georg-August University Göttingen, Göttingen, Germany. ²DZHK (German Centre for Cardiovascular Research), partner site Göttingen, Göttingen, Germany. ³Clinic and Polyclinic for Internal Medicine II, University Medical Centre Regensburg, Regensburg, Germany. ⁴Heart and Diabetes Centre North Rhine-Westphalia, Bad Oeynhausen, Germany. ⁵Department of Cardiology, Medical University of Graz, Graz, Austria. ⁶Institute of Pharmacology and Toxicology, University of Würzburg, Würzburg, Germany. ⁷These authors contributed equally: Philipp Bengel, Nataliya Dybkova, Petros Tirilomis, Katrin Streckfuss-Bömeke, Samuel Sossalla. ✉email: Samuel.Sossalla@klinik.uni-regensburg.de

Voltage-gated sodium channels (Na_V) play a critical role in physiological cardiac conduction. Na_V channels become inactive within a few milliseconds after activation under physiological conditions. However, in cardiac pathologies such as ischemia, hypoxia, oxidative stress, and heart failure (HF), some Na_V remain persistently open or reopen, generating a small but persistent Na^+ current, referred to as the late Na^+ current (I_{NaL})^{1–5}. This current slows the repolarisation rate and thereby prolongs the action potential duration (APD). Augmented I_{NaL} may additionally cause Na^+ -dependent Ca^{2+} overload in cardiomyocytes, thereby playing an essential role for arrhythmogenesis and diastolic dysfunction^{1,2,6,7}. Furthermore, $\text{Na}^+/\text{Ca}^{2+}$ overload caused by augmented I_{NaL} can give rise to early after-depolarizations (EADs), delayed afterdepolarizations (DADs), and hence sustained triggered arrhythmias^{8–11}. In the failing heart, increased I_{NaL} induces an influx of Na^+ into the cardiomyocyte, which in turn stimulates Ca^{2+} influx via the reverse mode of $\text{Na}^+/\text{Ca}^{2+}$ exchanger (NCX)^{12,13}. Cytosolic Ca^{2+} may now bind to calmodulin (CaM), forming a $\text{Ca}^{2+}/\text{CaM}$ complex, which activates $\text{Ca}^{2+}/\text{calmodulin}$ -dependent protein kinase II δ (CaMKII δ), a multifunctional serine/threonine protein kinase^{8,10,14,15}. CaMKII is expressed in four isoforms α , β , γ , and δ . CaMKII δ is the predominant isoform in heart¹⁶ while the δc isoform is mainly located in the cytosol. Once CaMKII δc is activated, it may cause hyperphosphorylation of the ryanodine receptor 2 (RyR2) residing within the sarcoplasmic reticulum (SR)-sarcolemma junction, leading to spontaneous proarrhythmic SR- Ca^{2+} release events in HF^{13,17–20}. Further, this augmented CaMKII δc activity can also induce I_{NaL} augmentation by phosphorylating Na_V channels^{10,14,21,22} leading to a vicious cycle between I_{NaL} and CaMKII δc .

Besides $\text{Na}_V1.5$ other Na_V isoforms have been reported to be present in the heart. Theoretically, they could also generate I_{NaL} , induce APD prolongation, and spontaneous SR- Ca^{2+} release. In the previous few years, different reports have been published on the existence of noncardiac Na_V in the heart. $\text{Na}_V1.8$, a noncardiac tetrodotoxin-resistant Na_V channel, is encoded by the *SCN10A* gene and was originally reported to be expressed in the dorsal root ganglion²³. Certain genome-wide association studies reported an association of *SCN10A* with changes in ECG parameters but most importantly with cardiac arrhythmias such as atrial fibrillation and sudden cardiac death^{24–27}. Later, the presence of $\text{Na}_V1.8$ was detected in atria and further evidence came from studies conducted in mouse and rabbit cardiomyocytes investigating its involvement in cardiac electrophysiology^{28–30}. Moreover, we recently reported that $\text{Na}_V1.8$ mRNA and protein expression are upregulated in tissue from human hypertrophied and failing ventricles and that $\text{Na}_V1.8$ contributes to I_{NaL} generation in the human heart under these pathological conditions^{31,32}. However, $\text{Na}_V1.8$ regulation, a potential interplay with pathologically increased CaMKII δc activity in HF, and the

role of $\text{Na}_V1.8$ on HF progression and arrhythmias in vivo and in vitro remain elusive.

In this work, we describe a detrimental interaction of $\text{Na}_V1.8$ with CaMKII δc in human and murine failing cardiomyocytes. Moreover, we investigate the contribution of $\text{Na}_V1.8$ to cellular electrophysiology in relation to enhanced CaMKII δc activity and consequently show a reduction of arrhythmias which is paralleled by an improved survival due to $\text{Na}_V1.8$ deletion in CaMKII δc transgenic HF mice.

Results

$\text{Na}_V1.8$ and CaMKII δc interaction in human HF. Since interaction between CaMKII δc and $\text{Na}_V1.5$ was shown previously, we hypothesized that CaMKII δc interacts also with $\text{Na}_V1.8$ and therefore performed co-immunoprecipitation using human ventricular tissue homogenates. Indeed, we found that CaMKII δc associates with $\text{Na}_V1.8$ in human non-failing as well as HF myocardium (Fig. 1a). Using immunofluorescence stainings, CaMKII δc and $\text{Na}_V1.8$ were found to be co-localized in isolated human cardiomyocytes (Fig. 1b). Of note, *SCN10A* mRNA expression in tissue from non-failing and HF hearts, as well as isolated cardiomyocytes from human HF hearts was found to be much lower than *SCN5A*. Further, RT-qPCR experiments revealed that a relevant part of *SCN10A* mRNA in the heart originates from cardiomyocytes (Supplementary information, Supplementary Figs. 1, 2).

$\text{Na}_V1.8$ inhibition reduces I_{NaL} in human failing, mouse CaMKII δc transgenic cardiomyocytes, and in *SCN10A* knockout iPSC-cardiomyocytes. In functional experiments, we could show that I_{NaL} , induced by increased CaMKII δc activity, was significantly reduced following pharmacological inhibition and genetical knockout of $\text{Na}_V1.8$ in human and murine cardiomyocytes. In isolated murine CaMKII $\delta\text{c}^{+/T}$ cardiomyocytes, I_{NaL} was augmented to ~150% compared to wildtype (WT), whereas the specific $\text{Na}_V1.8$ blockers A-806734 or PF-01247324 reduced I_{NaL} by ~40% in the same background (CaMKII $\delta\text{c}^{+/T}$) (Fig. 2a, b). These results illustrate that CaMKII δc -induced I_{NaL} can be clearly ameliorated by inhibiting $\text{Na}_V1.8$. A current–voltage relationship of $\text{Na}_V1.8$ -dependent I_{NaL} in isolated ventricular myocytes from CaMKII $\delta\text{c}^{+/T}$ mice is presented in the Supplementary information (Supplementary Fig. 3).

In the human failing heart, CaMKII δc and I_{NaL} are upregulated in parallel^{2,33}. Therefore, we inhibited $\text{Na}_V1.8$ by using PF-01247324 and compared its ability to reduce I_{NaL} to CaMKII δc inhibition using autocamtide-2-related inhibitory peptide (AIP) in human failing ventricular cardiomyocytes. In addition, we blocked $\text{Na}_V1.8$ and CaMKII δc in parallel by exposing human failing cells simultaneously to PF-01247324 and AIP. I_{NaL} measurements demonstrated that $\text{Na}_V1.8$ inhibition alone (PF-01247324) led to a ~40% decrease and CaMKII δc inhibition

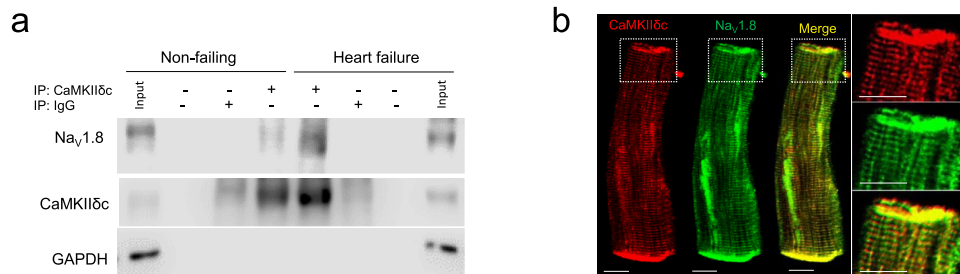


Fig. 1 CaMKII δc interacts with $\text{Na}_V1.8$ in human myocardium and isolated cardiomyocytes. **a** Co-immunoprecipitation of CaMKII δc and $\text{Na}_V1.8$ from left ventricular homogenates of human non-failing and failing hearts (NF: $n = 7$; HF: $n = 7$). **b** Co-localization of CaMKII δc and $\text{Na}_V1.8$ in human failing cardiomyocytes with immunofluorescence staining. Scale bar: 10 μm (staining was performed in cardiomyocytes isolated from five heart failure patients).

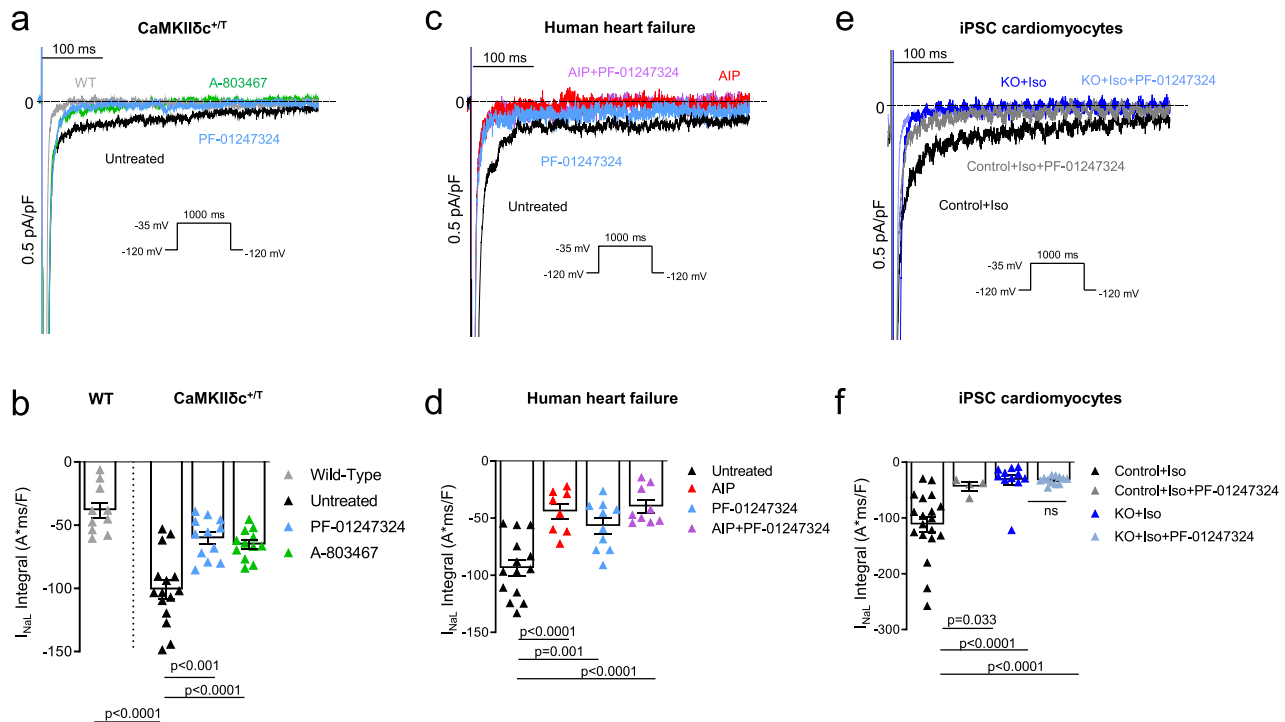


Fig. 2 Reduced I_{NaL} upon $Na_V1.8$ inhibition in human failing and mouse $CaMKII\delta c^{+/T}$ transgenic cardiomyocytes, and in $SCN10A$ knockout iPSC-cardiomyocytes. **a** Original traces of I_{NaL} in WT and $CaMKII\delta c^{+/T}$ mouse ventricular cardiomyocytes elicited using the protocol shown in the inset. **b** Mean data \pm SEM along with individual values shown in the graph plotting WT ($n = 10$ cells/5 mice) and $CaMKII\delta c^{+/T}$ (untreated: $n = 15$ cells/7 mice; A-806467: $n = 12$ cells/7 mice; PF-01247324: $n = 12$ cells/7 mice). Probability vs. untreated (One-way ANOVA with post hoc Bonferroni's correction). **c** Original traces of I_{NaL} from human failing ventricular cardiomyocytes elicited using the protocol shown in the inset. **d** Mean data \pm SEM along with individual values shown in the graph plotting (untreated: $n = 14$ cells/8 patients; AIP: $n = 8$ cells/5 patients; PF-01247324: $n = 10$ cells/6 patients; AIP + PF-01247324 = 9 cells/5 patients). Probability vs. untreated (One-way ANOVA with post hoc Bonferroni's correction). **e** Original traces of I_{NaL} from human ventricular $SCN10A$ knockout iPSC-cardiomyocytes elicited using the protocol shown in the inset. **f** Mean data \pm SEM along with individual values shown in the graph plotting (control + Iso: $n = 19$ cells/3 cardiac differentiations; control + Iso + PF: $n = 4$ cells/3 differentiations; $SCN10A$ knockout (KO) + Iso: $n = 11$ cells/3 differentiations; KO + Iso + PF-01247324 = 12 cells/3 differentiations). Probability vs. control + Iso (One-way ANOVA with post hoc Bonferroni's correction).

by AIP to a $\sim 53\%$ reduction of I_{NaL} in human failing cardiomyocytes (Fig. 2c, d). However, preincubation with AIP and PF-01247324 together decreased I_{NaL} comparable to AIP alone suggesting that $CaMKII\delta c$ inhibition might already suppress $Na_V1.8$ -driven I_{NaL} . Further, peak I_{Na} measurements revealed, that I_{NaL} reduction due to $Na_V1.8$ inhibition is not caused by a reduction of overall Na^+ current (Supplementary information, Supplementary Fig. 4).

As the existence of $Na_V1.8$ and its role in cardiomyocytes is still a matter of debate, we generated homozygous $Na_V1.8$ knockout (KO) lines by using CRISPR-Cas9 in human induced pluripotent stem cells (iPSC) and differentiated these cells into 2-month-old cardiomyocytes. Sanger sequencing demonstrated frameshifts and premature stop codons on both alleles (Supplementary information, Supplementary Fig. 5). As the amplitude of I_{NaL} is relatively small under healthy conditions we used isoproterenol (Iso, 50 nmol/l) to enhance I_{NaL} . Control iPSC-cardiomyocytes treated simultaneously with Iso and PF-01247324 exhibited $\sim 60\%$ less I_{NaL} compared to Iso alone, (Fig. 2e, f). Most importantly, in $Na_V1.8$ KO iPSC-cardiomyocytes I_{NaL} was reduced by $\sim 70\%$ compared to the control iPSC-cardiomyocytes. PF-01247324 did not exert any further impact on I_{NaL} in $Na_V1.8$ KO iPSC-cardiomyocytes compared to untreated $Na_V1.8$ KO cells, underlining the specificity of the drug.

$Na_V1.8$ modulates Ca^{2+} homeostasis under enhanced $CaMKII\delta c$ activity. In HF, enhanced I_{NaL} can potentially induce

proarrhythmic $SR-Ca^{2+}$ release events^{8,11}. Accordingly, we investigated whether inhibition of the $Na_V1.8$ -mediated I_{NaL} could attenuate the increase of the proarrhythmogenic $SR-Ca^{2+}$ spark frequency (CaSpF) caused by overexpression of $CaMKII\delta c$ (Fig. 3a, b). We incubated $CaMKII\delta c^{+/T}$ mouse cardiomyocytes with $Na_V1.8$ inhibitors and measured the CaSpF. A $\sim 50\%$ reduction of CaSpF was observed in both $Na_V1.8$ inhibitor groups compared to untreated cells (Fig. 3a, b). These results display that $SR-Ca^{2+}$ leak due to increased $CaMKII\delta c$ expression and activity can be reduced by inhibiting $Na_V1.8$.

It is well known that inhibition of $CaMKII\delta c$ can attenuate $SR-Ca^{2+}$ leak^{18,34}. However, therapeutic general inhibition of $CaMKII\delta c$ in humans may not be suitable because of its pivotal involvement in different vital pathways such as learning processes³⁵. We explored whether the $Na_V1.8$ inhibitor PF-01247324 exerts similar effects comparable to the inhibition of $CaMKII\delta c$. Incubation of human failing cardiomyocytes with either the $CaMKII\delta c$ inhibitor AIP or the $Na_V1.8$ inhibitor PF-01247324 resulted in a similar reduction of CaSpF compared to untreated cells (Fig. 3c, d). Furthermore, blocking $CaMKII\delta c$ and $Na_V1.8$ in parallel resulted in a significant reduction of CaSpF, comparable to AIP or PF-01247324 alone in human failing cardiomyocytes (Fig. 3c, d).

We further investigated whether $Na_V1.8$ inhibition modulates the Ca^{2+} transient amplitude and $SR-Ca^{2+}$ load in cardiomyocytes isolated from $CaMKII\delta c^{+/T}$ mice. $Na_V1.8$ inhibition using PF-01247324 did not pose any effect on either the Ca^{2+} transient

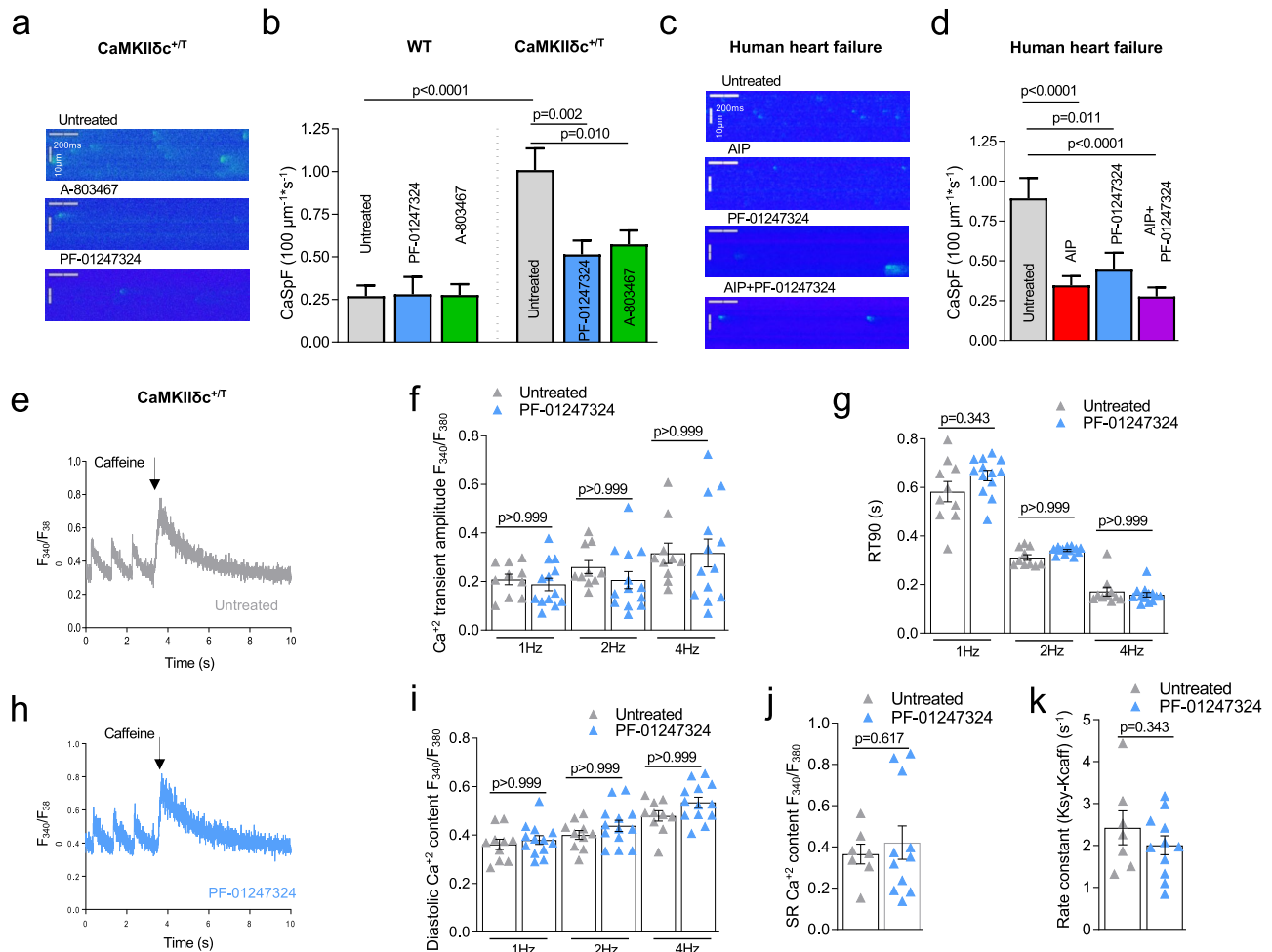


Fig. 3 Effects $\text{Na}_v1.8$ inhibition on intracellular Ca^{2+} handling. **a** Representative line scan images of $\text{CaMKII}\delta^{\text{c}/\text{T}}$ ventricular cardiomyocytes. **b** CaSpF data shown as mean \pm SEM for wildtype (WT) (untreated: $n = 58$ cells/4 mice; PF-01247324: $n = 41$ cells/4 mice; A-806467: $n = 41$ cells/4 mice) and $\text{CaMKII}\delta^{\text{c}/\text{T}}$ (untreated: $n = 122$ cells/8 mice; PF-01247324: $n = 105$ cells/7 mice; A-806467: $n = 101$ cells/8 mice). Data were analyzed by one-way ANOVA with post hoc Bonferroni's correction. **c** Representative line scan images of human failing ventricular cardiomyocytes. **d** Data shown as mean \pm SEM (untreated: $n = 123$ cells/14 patients; Autocamtide-2-related inhibitory peptide (AIP): $n = 105$ cells/15 patients; PF-01247324: $n = 59$ cells/10 patients; AIP + PF-01247324 = 89 cells/13 patients). Data were analyzed by one-way ANOVA with post hoc Bonferroni's correction, Probability vs. untreated. **e** Representative Ca^{2+} transients stimulated at 1 Hz and caffeine-induced Ca^{2+} transients in ventricular cardiomyocytes from $\text{CaMKII}\delta^{\text{c}/\text{T}}$ under untreated conditions. **f** Mean data \pm SEM show no effect of PF-01347324 treatment on Ca^{2+} transient amplitude at 1.0, 2.0, and 4.0 Hz stimulation ($n = 13$ cells/5 mice) compared to untreated ($n = 10$ cells/5 mice). **g** Ca^{2+} -transient decay (90% of Ca^{2+} -removal) RT90 was unchanged in PF-01347324-treated cells ($n = 11$ cells/5 mice) compared to untreated ($n = 8$ cells/5 mice). Data were presented as mean values \pm SEM. **h** Representative Ca^{2+} transients stimulated at 1 Hz and caffeine-induced Ca^{2+} transients in ventricular cardiomyocytes from $\text{CaMKII}\delta^{\text{c}/\text{T}}$ treated with PF-01247324. **i** Diastolic Ca^{2+} after addition of PF-01347324 ($n = 13$ cells/5 mice) compared to untreated cells ($n = 10$ cells/5 mice) at different stimulation frequencies was unchanged (one-way ANOVA with post hoc Bonferroni's correction, Fig. 3f, g, i). Data were presented as mean values \pm SEM. **j** Mean and individual values \pm SEM of caffeine-induced Ca^{2+} transients (untreated: $n = 7$ cells/4 mice, PF-01247324: $n = 11$ cells/5 mice) did not differ between the groups (Student's t -test). **k** Ca^{2+} -reuptake into the SR was not affected by inhibition of $\text{Na}_v1.8$ (untreated: $n = 7$ cells/4 mice, PF-01247324: $n = 11$ cells/5 mice), analyzed by Student's t -test. Data were presented as mean values \pm SEM.

amplitude or Ca^{2+} transient decay measured at different stimulation frequencies (Fig. 3e–h). Furthermore, diastolic Ca^{2+} , SR- Ca^{2+} content, and SERCA2a activity were not affected by PF-01247324 (Fig. 3i–k). Similar effects of PF-01247324 on Ca^{2+} transient kinetics and SR- Ca^{2+} content were observed in cardiomyocytes isolated from WT mice (Supplementary information, Supplementary Fig. 6).

Scn10a knockout improves survival in $\text{CaMKII}\delta^{\text{c}/\text{T}}$ mice in the absence of structural ventricular changes. To study whether inhibition of $\text{Na}_v1.8$ influences the development of HF, arrhythmogenesis, or survival in $\text{CaMKII}\delta^{\text{c}/\text{T}}$ mice, we crossbred these mice with $\text{Na}_v1.8$ knockout mice. Interestingly,

$\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{\text{c}/\text{T}}$ mice showed a significantly improved survival compared to $\text{CaMKII}\delta^{\text{c}/\text{T}}$ (median survival 98 vs. 72 days, 64 vs. 43 animals, Hazard Ratio 0.6) as assessed in blinded investigations. Specifically, $\text{CaMKII}\delta^{\text{c}/\text{T}}$ mice showed only a 37% survival at 12 weeks, whereas $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{\text{c}/\text{T}}$ died at a slower rate, with 67% survival at this age (Fig. 4a). These data illustrate that $\text{Na}_v1.8$ knockout is capable to counteract the lethal phenotype of $\text{CaMKII}\delta$ overexpression to a relevant extent.

Detailed investigations of hearts from $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{\text{c}/\text{T}}$ double-mutant and $\text{CaMKII}\delta^{\text{c}/\text{T}}$ mice by the age of 12 weeks exhibited comparably enlarged heart chambers (Fig. 4b). Heart weight to tibia length ratio was similarly increased in double-mutant and $\text{CaMKII}\delta^{\text{c}/\text{T}}$ mice (Fig. 4c). To investigate whether $\text{Na}_v1.8$

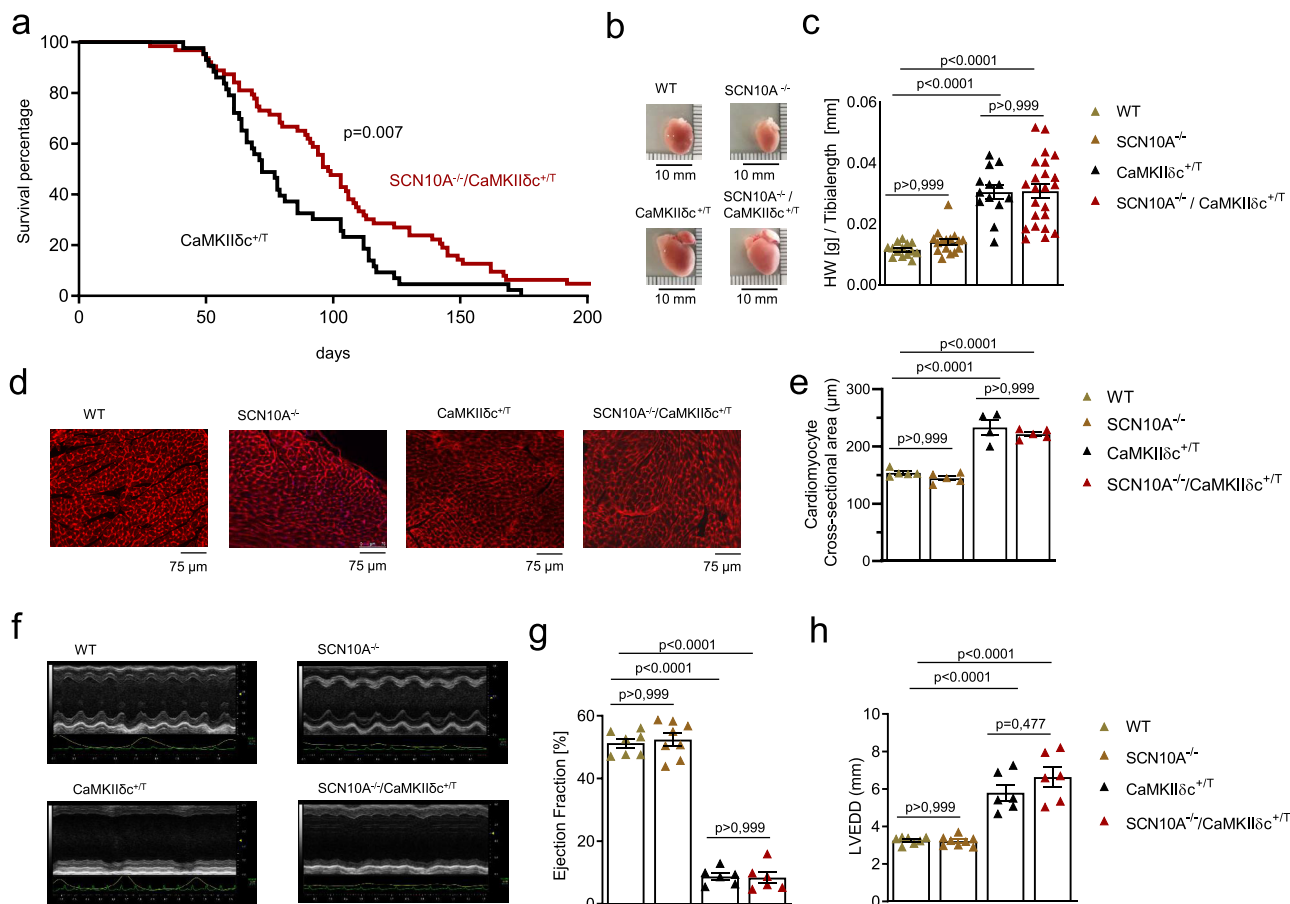


Fig. 4 Knockout of the *Scn10a* ($\text{Na}_V1.8$) gene in $\text{CaMKII}\delta^{\text{c}/\text{T}}$ mice improves survival. **a Survival curve of $\text{CaMKII}\delta^{\text{c}/\text{T}}$ and $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{\text{c}/\text{T}}$ (43 vs. 64 animals, median survival 72 vs. 98 days, blinded analysis). Log-rank (Mantel-Cox test and Gehan-Breslow-Wilcoxon test (two-tailed analysis) were performed to calculate the survival percentage of mice. Probability vs $\text{CaMKII}\delta^{\text{c}/\text{T}}$. **b** Hearts from WT, $\text{SCN10A}^{-/-}$, $\text{CaMKII}\delta^{\text{c}/\text{T}}$, and $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{\text{c}/\text{T}}$ mice. **c** Ratio of heart weight to tibia length as a parameter of cardiac hypertrophy. $\text{CaMKII}\delta^{\text{c}/\text{T}}$ and $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{\text{c}/\text{T}}$ showed a significant increase in this ratio compared to WT and $\text{SCN10A}^{-/-}$ mice. Data were analyzed by one-way ANOVA with post hoc Bonferroni's correction. (N = hearts studied, WT = 14, $\text{SCN10A}^{-/-}$ = 16, $\text{CaMKII}\delta^{\text{c}/\text{T}}$ = 13, and $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{\text{c}/\text{T}}$ = 23). Data were presented as mean values \pm SEM. **d** Original histological wheat germ agglutinin staining from WT, $\text{SCN10A}^{-/-}$, $\text{CaMKII}\delta^{\text{c}/\text{T}}$, and $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{\text{c}/\text{T}}$ mice. Scale bars = 75 μm . Stainings were produced from different sections and three different regions (basal, mid-ventricular, and apical) of each heart studied. **e** Cardiomyocyte cross-sectional-area (CSA) as a parameter for cellular hypertrophy. $\text{CaMKII}\delta^{\text{c}/\text{T}}$ and $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{\text{c}/\text{T}}$ showed a significant increase in CSA compared to WT and $\text{SCN10A}^{-/-}$ mice. CSA in $\text{CaMKII}\delta^{\text{c}/\text{T}}$ and $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{\text{c}/\text{T}}$ mice did not significantly differ. Data were analyzed by one-way ANOVA with post hoc Bonferroni's correction. N = hearts studied (>300 cardiomyocytes were studied per heart, from different sections and different regions (basal, mid-ventricular, apical), WT = 5 hearts, $\text{SCN10A}^{-/-}$ = 5 hearts, $\text{CaMKII}\delta^{\text{c}/\text{T}}$ = 4 hearts, $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{\text{c}/\text{T}}$ = 5 hearts). Data were presented as mean values \pm SEM. **f** Original echocardiography recordings from WT, $\text{SCN10A}^{-/-}$, $\text{CaMKII}\delta^{\text{c}/\text{T}}$, and $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{\text{c}/\text{T}}$ at M-mode in 12–13-week-old mice. **g** Echocardiography recordings revealed a decrease in left ventricular ejection fraction (EF) in $\text{CaMKII}\delta^{\text{c}/\text{T}}$ (six mice) and $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{\text{c}/\text{T}}$ (six mice) compared to WT (seven mice) or $\text{SCN10A}^{-/-}$ (eight mice) ($p < 0.0001$ (one-way ANOVA with post hoc Bonferroni's correction)). Data were presented as mean values \pm SEM. **h** Echocardiography recordings revealed a significant increase in left ventricular end-diastolic diameter (LVEDD) in $\text{CaMKII}\delta^{\text{c}/\text{T}}$ (six mice) and $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{\text{c}/\text{T}}$ (six mice) compared to WT (seven mice) or $\text{SCN10A}^{-/-}$ (eight mice) ($p < 0.0001$). LVEDD was not significantly different in WT vs $\text{SCN10A}^{-/-}$ or $\text{CaMKII}\delta^{\text{c}/\text{T}}$ vs $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{\text{c}/\text{T}}$ (one-way ANOVA with post hoc Bonferroni's correction)). Data were presented as mean values \pm SEM.**

knockout influences $\text{CaMKII}\delta$ -mediated hypertrophy we prepared histological heart stainings. $\text{CaMKII}\delta$ overexpression led to an increase of cardiomyocyte cross-sectional area (CSA) compared to WT and $\text{SCN10A}^{-/-}$ in mouse left ventricles. However, in double-mutant mice CSA did not differ from $\text{CaMKII}\delta^{\text{c}/\text{T}}$ alone (Fig. 4d, e). In addition, *Scn10a* knockout did not change the expression of $\text{Na}_V1.5$ (*Scn5a*) and $\text{CaMKII}\delta$ on protein and mRNA levels in hearts from WT and $\text{CaMKII}\delta^{\text{c}/\text{T}}$ mice (Supplementary information, Supplementary Figs. 7, 8).

Serial echocardiography revealed a severe HF phenotype in $\text{CaMKII}\delta^{\text{c}/\text{T}}$ and $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{\text{c}/\text{T}}$ with a significant reduction of left ventricular ejection fraction (EF) (Fig. 4f, g).

Moreover, significantly enlarged left ventricular end-diastolic diameters (LVEDD) were measured in $\text{CaMKII}\delta^{\text{c}/\text{T}}$ and $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{\text{c}/\text{T}}$ compared to WT or $\text{SCN10A}^{-/-}$ (Fig. 4f, h). There were no changes between WT and $\text{SCN10A}^{-/-}$ or $\text{CaMKII}\delta^{\text{c}/\text{T}}$ and $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{\text{c}/\text{T}}$ mice with respect to LVEDD and EF.

Reduction of proarrhythmic activity in $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{\text{c}/\text{T}}$ cardiomyocytes. To test whether enhanced I_{NaL} in $\text{CaMKII}\delta^{\text{c}/\text{T}}$ mice can be ameliorated by genetic knockout of $\text{Na}_V1.8$, we measured I_{NaL} in isolated mouse cardiomyocytes (Fig. 5a, b). While $\text{CaMKII}\delta^{\text{c}/\text{T}}$ cardiomyocytes showed significantly enhanced I_{NaL} ,

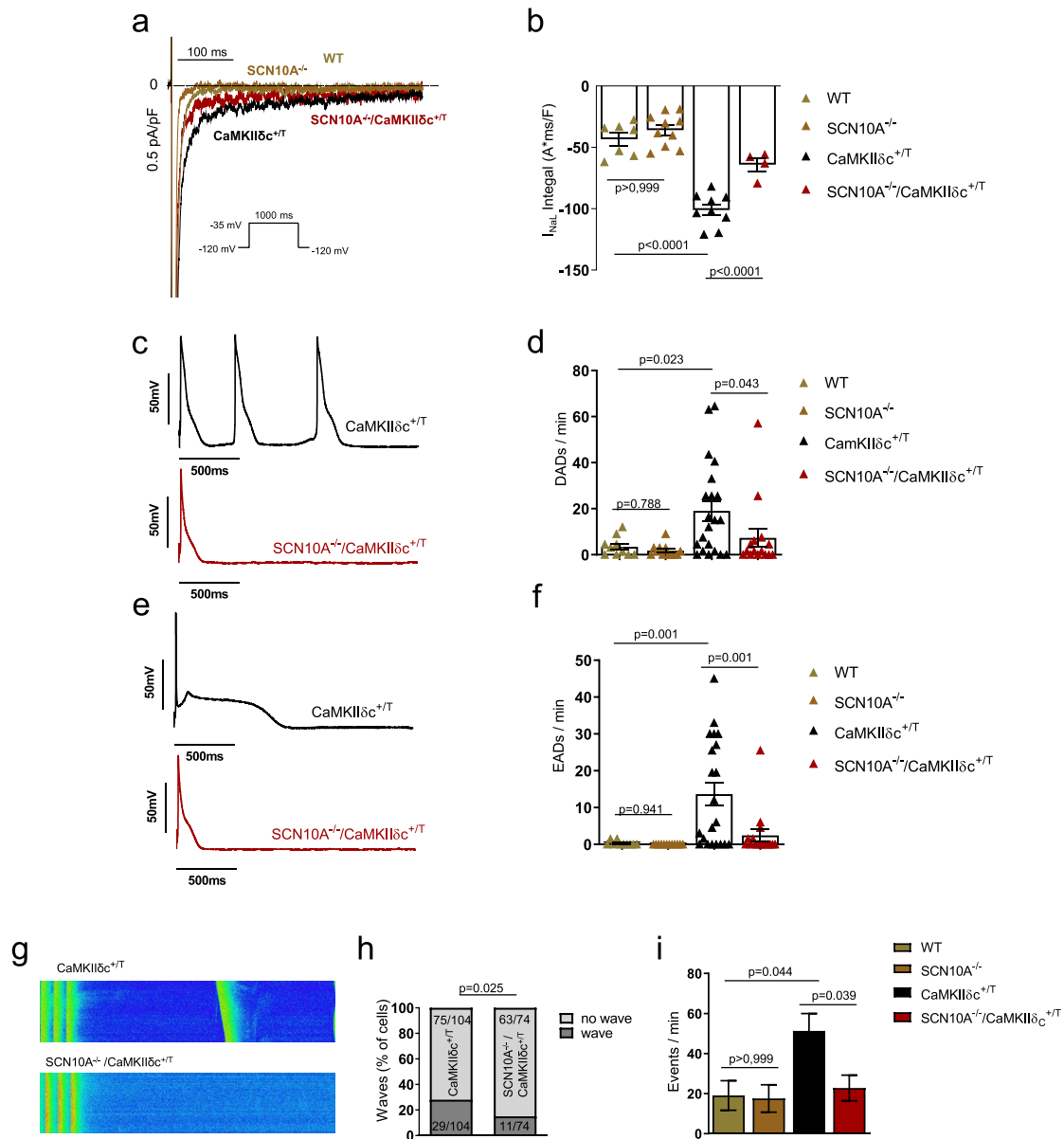


Fig. 5 Knockout of *Scn10a* ($Na_v1.8$) in $CaMKII\delta c^{+/T}$ mice ($SCN10A^{-/-}/CaMKII\delta c^{+/T}$) significantly reduces I_{NaL} and proarrhythmic triggers. **a** Original traces of I_{NaL} in WT, $SCN10A^{-/-}$, $CaMKII\delta c^{+/T}$, and $SCN10A^{-/-}/CaMKII\delta c^{+/T}$ mouse ventricular cardiomyocytes elicited using the protocol shown in the inset. **b** Mean data \pm SEM along with individual values shown in the graph plotting (WT: $n = 7$ cells/4 mice, $SCN10A^{-/-}$ $n = 10$ cells/5 mice, $CaMKII\delta c^{+/T}$: $n = 9$ cells/5 mice; $SCN10A^{-/-}/CaMKII\delta c^{+/T}$: $n = 4$ cells/4 mice), there was a significantly reduced I_{NaL} in $SCN10A^{-/-}/CaMKII\delta c^{+/T}$ cardiomyocytes compared to $CaMKII\delta c^{+/T}$. Data were analyzed by one-way ANOVA with post hoc Bonferroni's correction. **c** Original traces of action potentials showing triggered action potentials originating from delayed afterdepolarizations (DADs) in $CaMKII\delta c^{+/T}$ and $SCN10A^{-/-}/CaMKII\delta c^{+/T}$ cardiomyocytes. **d** Graph of mean data \pm SEM along with individual values showing DADs per minute in WT ($n = 10$ cells/5 mice), $SCN10A^{-/-}$ ($n = 12$ cells/5 mice), $CaMKII\delta c^{+/T}$ ($n = 21$ cells/5 mice) and $SCN10A^{-/-}/CaMKII\delta c^{+/T}$ ($n = 15$ cells/5 mice) cardiomyocytes. There were significantly less events of afterdepolarizations in $SCN10A^{-/-}/CaMKII\delta c^{+/T}$ compared to $CaMKII\delta c^{+/T}$ cardiomyocytes. Data were analyzed by one-way ANOVA with the post hoc two-stage step-up method of Benjamini, Krieger, and Yekutieli. **e** Original traces of action potential showing early afterdepolarizations (EADs) in $CaMKII\delta c^{+/T}$ and $SCN10A^{-/-}/CaMKII\delta c^{+/T}$ cardiomyocytes. **f** Graph of mean data \pm SEM along with individual values showing EADs per minute in WT ($n = 10$ cells/5 mice), $SCN10A^{-/-}$ ($n = 12$ cells/5 mice), $CaMKII\delta c^{+/T}$ ($n = 21$ cells/5 mice) and $SCN10A^{-/-}/CaMKII\delta c^{+/T}$ ($n = 16$ cells/5 mice) cardiomyocytes. There were significantly less events of afterdepolarizations in $SCN10A^{-/-}/CaMKII\delta c^{+/T}$ compared to $CaMKII\delta c^{+/T}$ cardiomyocytes. Data were analyzed by one-way ANOVA with the post hoc two-stage step-up method of Benjamini, Krieger, and Yekutieli. **g** Original confocal line scans images of $CaMKII\delta c^{+/T}$ and $SCN10A^{-/-}/CaMKII\delta c^{+/T}$ cardiomyocytes showing diastolic Ca^{2+} waves. **h** Percentage of cells exhibiting waves was significantly less in $SCN10A^{-/-}/CaMKII\delta c^{+/T}$ ($n = 74$ cells/7 mice) compared to $CaMKII\delta c^{+/T}$ ($n = 104$ cells/9 mice). Data were analyzed by Chi-square test, two-tailed analysis. **i** Significantly decreased number of Ca^{2+} waves per minute in $SCN10A^{-/-}/CaMKII\delta c^{+/T}$ compared to $CaMKII\delta c^{+/T}$. Data were analyzed by one-way ANOVA with post hoc Bonferroni's correction. Cells/mice studied, WT: $n = 48$ cells/5 mice, $SCN10A^{-/-}$: $n = 52$ cells/5 mice, $CaMKII\delta c^{+/T}$: $n = 104$ cells/9 mice; $SCN10A^{-/-}/CaMKII\delta c^{+/T}$: $n = 74$ cells/7 mice. Data were presented as mean values \pm SEM.

knockout of $\text{Na}_V1.8$ resulted in a ~45% decrease in I_{NaL} in $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{+/T}$ mice compared with $\text{CaMKII}\delta^{+/T}$. Of note, at basal/unstimulated conditions, I_{NaL} did not differ between WT and $\text{SCN10A}^{-/-}$.

To evaluate whether chronic ablation of $\text{Na}_V1.8$ in $\text{CaMKII}\delta^{+/T}$ mice may reduce proarrhythmic cellular activity, we performed electrophysiological measurements (Fig. 5c–f). Patch-clamp experiments revealed that $\text{CaMKII}\delta^{+/T}$ cardiomyocytes exhibited approximately fivefold more delayed afterdepolarizations/min (DADs) compared to WT and $\text{SCN10A}^{-/-}$. *Scn10a* knockout reduced the fraction of $\text{CaMKII}\delta$ -induced DADs/min by ~60% (Fig. 5c, d). Comparable observations were made regarding the occurrence of early afterdepolarizations (EADs, Fig. 5e, f). While WT and $\text{SCN10A}^{-/-}$ cardiomyocytes developed almost no EADs, 13.6 ± 3.2 EADs/min were recorded in $\text{CaMKII}\delta^{+/T}$. $\text{Na}_V1.8$ knockout caused an 80% reduction of EADs/min in these cells. A detailed description of the action potential characteristics of these measurements is provided in the Supplementary information (Supplementary Table 1).

Furthermore, to evaluate whether inhibition of $\text{Na}_V1.8$ may decrease the number of cardiomyocytes exhibiting Ca^{2+} -derived proarrhythmic events, we quantified the occurrence of Ca^{2+} waves in mouse ventricular cardiomyocytes. The fraction of cardiomyocytes developing Ca^{2+} waves was significantly less in $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{+/T}$ versus $\text{CaMKII}\delta^{+/T}$ (14.8 vs 27.8%, Fig. 5g, h). However, some cardiomyocytes showed more than one Ca^{2+} wave. Therefore, we calculated the occurring events per minute. In $\text{CaMKII}\delta^{+/T}$ the frequency of Ca^{2+} waves was ~2.5-fold higher compared to WT and $\text{SCN10A}^{-/-}$ and was reduced by ~55% in cardiomyocytes from $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{+/T}$ compared to $\text{CaMKII}\delta^{+/T}$ (Fig. 5i). Of note, Ca^{2+} transient amplitude and Ca^{2+} extrusion from cytosol was unchanged between $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{+/T}$ and $\text{CaMKII}\delta^{+/T}$ (Supplementary information, Supplementary Fig. 9). In summary, we describe a cellular rescue of the proarrhythmic $\text{CaMKII}\delta^{+/T}$ phenotype due to genetic ablation of $\text{Na}_V1.8$, which is associated with improved animal survival.

***Scn10a* knockout reduces ventricular arrhythmias in $\text{CaMKII}\delta^{+/T}$ mice.** To further investigate whether the proarrhythmic potential of *Scn10a* knockout on the cellular level is translatable into the in vivo situation, we implanted telemetric monitors into 8 week old $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{+/T}$ and $\text{CaMKII}\delta^{+/T}$ mice. After 10 days of recovery from surgery, telemetric measurements were performed twice a week for 24 h over a period of 2 weeks. Baseline ECG characteristics are presented in the Supplementary information (Supplementary Table 2). There was no relevant difference in overall physical animal activity between the groups (Fig. 6b). Blinded analysis revealed that $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{+/T}$ showed a strong trend towards a reduction of premature ventricular contractions (PVC) by ~93% compared to $\text{CaMKII}\delta^{+/T}$ (Fig. 6a, c, $p = 0.08$). Most importantly, the incidence of ventricular tachycardia (VT) was significantly lower (91%) in $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{+/T}$ (Fig. 6a, d). Therefore, the observed improved survival of $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{+/T}$ -animals is associated with reduced ventricular arrhythmias.

Discussion

In our present study, we demonstrate that $\text{CaMKII}\delta$ interacts with the neuronal sodium channel $\text{Na}_V1.8$ in human ventricular cardiomyocytes. Using different approaches in human and mouse cardiomyocytes, we demonstrated the relevance of $\text{Na}_V1.8$ for I_{NaL} generation in HF and that an enhanced $\text{CaMKII}\delta$, indeed, regulates this $\text{Na}_V1.8$ -driven I_{NaL} . Isolated cardiomyocytes from $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{+/T}$ compared to $\text{CaMKII}\delta^{+/T}$ mice

exhibit reduced cellular arrhythmic events. While there was no change with respect to HF progression, i.e., similar left ventricular ejection fraction and chamber diameters, we found reduced ventricular arrhythmias and an improved animal survival of $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{+/T}$ animals. Thus, we identified a modifiable proarrhythmic $\text{CaMKII}\delta$ downstream target in the failing heart.

We recently found that $\text{Na}_V1.8$ is upregulated and thereby contributes to I_{NaL} under conditions of HF and cardiac hypertrophy where $\text{CaMKII}\delta$ activity is known to be enhanced^{31,32}. Therefore, the aim of the current study was to investigate a potential crosstalk between $\text{Na}_V1.8$ and $\text{CaMKII}\delta$ and its consequences on I_{NaL} generation and cellular arrhythmogeny. Accordingly, we revealed an interaction of $\text{Na}_V1.8$ and $\text{CaMKII}\delta$ in human ventricular myocardium of both non-failing and HF samples. $\text{CaMKII}\delta$ is known to also interact with $\text{Na}_V1.5$ channels at the intercalated disc where $\text{Na}_V1.5$ and $\text{CaMKII}\delta$ are part of a macro complex with Ankyrin-G and βIV -spectrin³⁶. The interaction with $\text{CaMKII}\delta$ in this complex influences $\text{Na}_V1.5$ channel function in several cardiac disease states such as HF^{14,21}. As $\text{Na}_V1.8$ was previously shown to interact with Ankyrin-G in neurons³⁷ in a similar fashion as it is known for $\text{Na}_V1.5$ in cardiomyocytes, similar mechanisms of $\text{Na}_V1.8$ interaction with $\text{CaMKII}\delta$ like known for $\text{Na}_V1.5$ are conceivable. Several Serin residues as well as CaMKII -binding consensus motifs were found to be conserved between $\text{Na}_V1.5$ and $\text{Na}_V1.8$ (for details see Supplementary Fig. 10).

In a variety of cardiac pathologies, enhanced $\text{CaMKII}\delta$ activity and expression is a key contributor to maladaptive electrical remodeling and thereby promotes arrhythmias^{9,38–40}. $\text{CaMKII}\delta$ can influence I_{NaL} magnitude by phosphorylating Na_V , which has been exclusively investigated for cardiac $\text{Na}_V1.5$ before, whereas a possible $\text{CaMKII}\delta$ -dependent regulation of $\text{Na}_V1.8$ has not been investigated yet^{14,21,22}. In our study, we investigated the contribution of $\text{Na}_V1.8$ to I_{NaL} in an HF model that was exclusively induced by chronic $\text{CaMKII}\delta$ overexpression. As previously shown, I_{NaL} was augmented in cardiomyocytes from $\text{CaMKII}\delta^{+/T}$ compared to WT^{21,40}, while these effects were ameliorated by the application of specific $\text{Na}_V1.8$ blockers. Therefore, at least a relevant part of $\text{CaMKII}\delta$ -induced I_{NaL} appears to be driven by $\text{Na}_V1.8$. Additional support for this conclusion comes from our I_{NaL} measurements from human failing cardiomyocytes where $\text{CaMKII}\delta$ activity and I_{NaL} are both known to be increased in parallel^{2,13,16,21,41}. Inhibition of either $\text{CaMKII}\delta$ or $\text{Na}_V1.8$ reduced I_{NaL} as separately demonstrated before^{21,31}. However, simultaneous inhibition of $\text{Na}_V1.8$ and $\text{CaMKII}\delta$ had no additive effect compared to $\text{CaMKII}\delta$ inhibition alone. Therefore, it can be assumed that the majority of $\text{Na}_V1.8$ -driven I_{NaL} was already abolished by $\text{CaMKII}\delta$ inhibition and hence, seems to be $\text{CaMKII}\delta$ -dependent. In addition, *Scn10a* knockout in $\text{CaMKII}\delta^{+/T}$ mice resulted in a reduction in I_{NaL} comparable to $\text{Na}_V1.8$ inhibition upon specific blockers.

In previous studies, the functional relevance of $\text{Na}_V1.8$ expression in cardiomyocytes was questioned as an application of specific blockers had no effects on peak I_{Na} and I_{NaL} in healthy and unstimulated cardiomyocytes^{42,43}. These data are not in conflict with our findings, as we also did not observe differences in I_{NaL} magnitude in cardiomyocytes from healthy mice, while clear effects were evident under conditions of enhanced I_{NaL} either by chronic $\text{CaMKII}\delta$ overexpression or isoproterenol treatment of iPSC-cardiomyocytes. Therefore, the interaction of $\text{Na}_V1.8$ with enhanced $\text{CaMKII}\delta$ activity might be necessary to generate meaningful effects on cardiomyocyte electrophysiology while $\text{Na}_V1.8$ appears to play a negligible role in healthy cardiomyocytes. This establishes $\text{Na}_V1.8$ to be a disease-specific target.

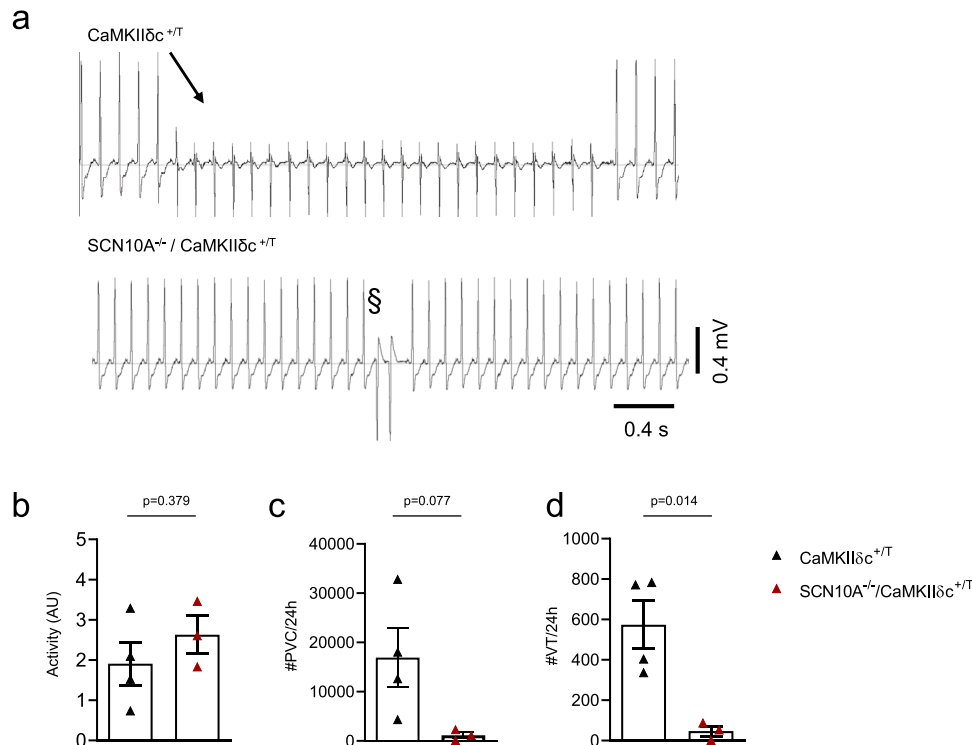


Fig. 6 *SCN10A*^{-/-}/*CaMKIIδc*^{+T} exhibit less in vivo arrhythmias compared to *CaMKIIδc*^{+T} mice. **a** Original ECG traces from telemetry recordings of 10-week-old *CaMKIIδc*^{+T} and *SCN10A*^{-/-}/*CaMKIIδc*^{+T} mice showing ventricular arrhythmias. **b** Unchanged activity levels in *SCN10A*^{-/-}/*CaMKIIδc*^{+T} compared to *CaMKIIδc*^{+T} (*CaMKIIδc*^{+T} (four mice); *SCN10A*^{-/-}/*CaMKIIδc*^{+T} (three mice), four individual recordings each). Data were presented as mean values ± SEM, analyzed by unpaired two-tailed Student's *t*-test. **c** Mean values of premature ventricular contractions (PVCs) in *SCN10A*^{-/-}/*CaMKIIδc*^{+T} ($p = 0.08$, Unpaired two-tailed Student's *t*-test), *CaMKIIδc*^{+T} (four mice); *SCN10A*^{-/-}/*CaMKIIδc*^{+T} (three mice), four individual recordings each. Data were presented as mean values ± SEM. **d** Reduction of ventricular tachycardia (VT) incidence in *SCN10A*^{-/-}/*CaMKIIδc*^{+T} ($p < 0.05$, Unpaired two-tailed Student's *t*-test), *CaMKIIδc*^{+T} (four mice); *SCN10A*^{-/-}/*CaMKIIδc*^{+T} (three mice), four individual recordings each. Data were presented as mean values ± SEM.

An augmentation of I_{NaL} was demonstrated to cause a Na^+ -dependent Ca^{2+} overload and spontaneous SR- Ca^{2+} release providing a substrate for cellular proarrhythmia^{8,15,30,44}. We previously reported that inhibition of I_{NaL} by specifically targeting $Na_v1.8$ is potent enough to reduce NCX reverse mode and thereby diastolic SR- Ca^{2+} leak^{11,31,32}. In the present work, we inhibited either *CaMKIIδc* or/and $Na_v1.8$ in failing humans and *CaMKIIδc*^{+T} mouse ventricular cardiomyocytes and correspondingly observed a decrease in I_{NaL} and diastolic SR- Ca^{2+} leak. A significant reduction of diastolic SR- Ca^{2+} release events was already prominent after $Na_v1.8$ inhibition alone in failing human cardiomyocytes which was almost comparable to the effect caused by *CaMKIIδc* inhibition. Moreover, we did not observe a further reduction in SR- Ca^{2+} leak when AIP was co-administrated with the $Na_v1.8$ blocker suggesting that effects of $Na_v1.8$ inhibition on SR- Ca^{2+} leak act via indirect inhibition of *CaMKIIδc* as it was previously shown for I_{NaL} inhibition with TTX or Ranolazine^{8,15}. The present findings, therefore confirm that $Na_v1.8$ inhibition abrogates the vicious proarrhythmic cycle between enhanced I_{NaL} and *CaMKIIδc*.

Increased expression and enhanced activity of *CaMKIIδc* is not only known to increase proarrhythmic triggers in HF but is also associated with a strong HF phenotype and increased mortality in mice^{9,13,16,45}. Our current findings illustrate that $Na_v1.8$ plays a relevant role for arrhythmogenesis under conditions of enhanced *CaMKIIδc* activity. For deeper mechanistic analysis, we crossbred our *CaMKIIδc*^{+T} mice with *Scn10a* knockout mice. In fact, we observed an improved survival in these *SCN10A*^{-/-}/*CaMKIIδc*^{+T} mice but still with an unchanged typical phenotype of dilated

cardiomyopathy in these blinded investigations. There may be three potential explanations for the improved survival, which were either reduced lethal arrhythmias, pump failure, or a combination of both. As we could demonstrate that induction and progression of HF is not influenced by the additional *Scn10a* knockout we propose a reduction of augmented I_{NaL} with subsequent lower proarrhythmic activity to constitute the underlying mechanism of this improved survival. The link between enhanced I_{NaL} and increased arrhythmia risk is rather complex. On the one hand, enhanced I_{NaL} prolongs APD and may therefore give rise to the formation of EADs. On the other hand, enhanced I_{NaL} causes Na^+ overload of the cardiomyocyte subsequently leading to Ca^{2+} overload by activating NCX reverse mode^{6,8,11,15}. This may trigger the occurrence of diastolic SR- Ca^{2+} release and DADs due to enhanced *CaMKIIδc*-dependent RyR2 phosphorylation^{8,10,15}. In our experiments, a reduction of I_{NaL} in *SCN10A*^{-/-}/*CaMKIIδc*^{+T} compared to *CaMKIIδc*^{+T} cardiomyocytes was observed. This was clearly associated with a reduction in the occurrence of EADs, DADs and diastolic SR- Ca^{2+} -release events, thereby illustrating that the *CaMKIIδc*-induced proarrhythmic phenotype can be ameliorated on the cellular level by $Na_v1.8$ knockout.

In a recent study by our groups, we could demonstrate that selective inhibition of diastolic SR- Ca^{2+} leak by the compound Rycal S36 improves survival in an HF mouse model, where I_{NaL} and *CaMKIIδc* activity were also described to be enhanced^{10,46}. As in our study, improved survival was caused by a significant reduction of malignant arrhythmias, while the severity of HF was unchanged. Likewise, we observed less ventricular arrhythmias in *SCN10A*^{-/-}/*CaMKIIδc*^{+T} mice in vivo. It is well known that

individuals with structural heart disease are at increased risk for sustained ventricular tachycardia and ventricular fibrillation⁴⁷. Moreover, sustained ventricular tachycardia may degenerate to ventricular fibrillation⁴⁷. In addition, clinical trials have shown an association of PVCs and ventricular tachycardia with adverse outcomes in patients with dilated cardiomyopathy^{47–49}. The $\text{Na}_V1.8$ knockout significantly reversed a relevant part of the arrhythmogenic CaMKII δ transgenic substrate in vitro and in vivo which is known to be associated with sudden cardiac death. However, due to the sporadic nature of sudden cardiac death, we were not able to correlate mortality with ventricular arrhythmias in these CaMKII δ transgenic mice as this is technically and ethically not feasible. We are also not aware of any other experimental trial that investigates such a high number of HF mice with implanted telemetric recorders during death struggle. Finally, a chicken or the egg question would remain, as detected rhythm disorders cannot be differentiated to be either caused primarily by arrhythmia or secondary due to hypoxia.

Persistent CaMKII δ hyperactivity in hearts of HF patients has several detrimental effects such as influencing myocardial remodeling and promoting arrhythmias^{33,39}. Despite the fact that several CaMKII δ inhibitors have been demonstrated to effectively reduce arrhythmogenesis in HF in vitro^{46,50}, CaMKII δ inhibition to manage cardiac arrhythmias has not been established in patients so far. This can be explained by the vast signaling network of this enzyme, which is required to keep a balance between pivotal off-target effects and the therapeutic benefits. Any other alternative downstream or upstream target of the CaMKII δ pathway, which is involved in promoting arrhythmias may be a more specific therapeutic option. As it has consistently been shown that activation of CaMKII δ is involved in enhancement of I_{NaL} and vice versa^{10,11,21,34,40} our proposed crosstalk between CaMKII δ and $\text{Na}_V1.8$ causing enhanced I_{NaL} in the presence of hyperactive CaMKII δ may be a novel strategy to prevent detrimental CaMKII δ -induced effects on cellular electrophysiology. Most importantly, we and others have shown that selective $\text{Na}_V1.8$ inhibition does not influence either peak Na^+ current or Na^+ current kinetics and thereby cardiac conduction as another critical regulator of proarrhythmia^{11,29,31,42,51}.

In summary, the results of our current study demonstrate that increased CaMKII δ activity in HF contributes to enhanced I_{NaL} via interaction with $\text{Na}_V1.8$. This augmented I_{NaL} generated by $\text{Na}_V1.8$ detrimentally influences cellular electrophysiology by increasing RyR2-leakiness and can give rise to cellular proarrhythmic events in HF cardiomyocytes. Importantly, the results of the present study demonstrate that pharmacological inhibition and genetic ablation of $\text{Na}_V1.8$ can specifically reverse these detrimental proarrhythmic effects in the human HF heart and in our $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{+/T}$ mouse model. This is associated with improved animal survival. Therefore, targeting $\text{Na}_V1.8$ as a specific substrate of increased CaMKII δ activity may constitute a promising antiarrhythmic approach in HF which merits further translational investigation.

Methods

Human tissue samples. The study conforms to the declaration of Helsinki and was approved by the local ethics committee. All participants were informed about the study prior to inclusion. All patients signed informed consent. We obtained left ventricular tissues from explanted hearts of patients with end-stage HF (NYHA HF classification IV) who were undergoing heart transplantation. After explantation, the whole heart or myocardial tissues were immediately placed in a container having precooled cardioplegic solution containing (in mmol/L): NaCl 110, KCl 16, MgCl₂ 16, NaHCO₃ 16, CaCl₂ 1.2, and glucose 11. Myocardial samples for Western blot and co-immunoprecipitation were immediately frozen in liquid nitrogen and stored at -80°C . The heart tissue for cell isolation was stored in cooled cardioprotective solution containing (in mmol/L): 156 Na^+ , 3.6 K^+ , 135 Cl^- , 25 HCO_3^- , 0.6 Mg^{2+} , 1.3 H_2PO_4^- , 0.6 SO_4^{2-} , 2.5 Ca^{2+} , 11.2 glucose, and 10 2,3-butanedionmonoxime (BDM) aerated with 95% O_2 and 5% CO_2 . We used healthy

myocardium from healthy donor hearts that could not be transplanted for technical reasons as controls co-immunoprecipitation. Patient characteristics are presented in the Supplementary information (Supplementary Table 3). The study was approved by the local ethics committee of the University Medicine of Goettingen.

CaMKII δ transgenic mice. CaMKII δ transgenic ($\text{CaMKII}\delta^{+/T}$) mice were generated using an α -MHC promoter¹³. We used 12- to 14-week-old mice for electrophysiology experiments. The animal investigations conform to the “Guide for the Care and Use of Laboratory Animals” published by the US National Institutes of Health (8th edition, revised 2011) and the guidelines from Directive 2010/63/EU of the European Parliament on the protection of animals used for scientific purposes. The mouse study was approved by the local ethics committee of the University Medicine of Goettingen and the public authority on animal welfare.

Generation of $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{+/T}$ mice. To generate the $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{+/T}$ mouse model we crossed $\text{CaMKII}\delta^{+/T}$ mice from a Black-Swiss strain with $\text{Na}_V1.8$ knockout mice ($\text{SCN10A}^{-/-}$) from a C57BL/6 strain. Eight weeks old male $\text{CaMKII}\delta^{+/T}$ mice were mated with 8 weeks old female $\text{SCN10A}^{-/-}$ mice. This breeding resulted in a genotype carrying a WT or transgenic (TG) allele of the CaMKII δ gene and a heterozygote knockout of *Scn10a*. From this first generation, 8 weeks old male mice carrying a TG CaMKII δ allele ($\text{CaMKII}\delta^{+/T}$) were mated with 8 weeks old female mice carrying homozygous WT CaMKII δ alleles and heterozygous for *Scn10a* gene. This breeding produced 25% mice with the desired genotype having heterozygous CaMKII δ gene ($\text{CaMKII}\delta^{+/T}$) and homozygous *Scn10a* knockout ($\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{+/T}$). Mice carrying a heterozygote *Scn10a* gene were excluded from survival analysis and experimental studies. A detailed scheme of the crossbreeding is displayed in the Supplementary information (Supplementary Fig. 11). The breeding rooms were maintained at $20\text{--}22^\circ\text{C}$ with 50–60% humidity. Mice were housed in a room with a 12-h light/dark cycle with ad libitum access to water and food.

Survival of $\text{CaMKII}\delta^{+/T}$ mice was documented in a database of the local facility of animal experiments. A survival curve was prepared from this database and a comparison of survival was made between the groups $\text{CaMKII}\delta^{+/T}$ and $\text{SCN10A}^{-/-}/\text{CaMKII}\delta^{+/T}$. Animals used for experiments were excluded from survival analysis. Before sacrificing animals, body weight was recorded while after explantation heart weight and tibia length were measured to tabulate animal size. Some of the animals showed signs of heart failure with fluid retention while some had signs of terminal illness therefore, heart weight to tibia length ratio was analyzed instead of heart weight to body weight ratio. The mouse study was approved by the local ethics committee of the University Medicine of Goettingen and the public authority on animal welfare.

Generation of homozygous knockout iPSCs using CRISPR/Cas9 and directed differentiation into iPSC-cardiomyocytes. All procedures were performed according to the Declaration of Helsinki and were approved by the local ethics committee. Informed consent was signed by all tissue donors. Gene editing was conducted using two CRISPR guide RNAs (gRNA1: GTGACTCCGAG-TAAAGCGACGG and gRNA2: ACGGAAGTTGTTAGTTTCGAGG, designed with IDTdna.com design tool) targeting *SCN10A* exon1. About 2×10^6 wild-type iPSCs were electroporated with 2.5 μl (100 μM) of each gRNA, 5 μl tracrRNA (100 μM), 2 μl Cas9 protein (10 ng/ μl , IDT), and 1 μl electroporation enhancer (100 μM , IDT) using the Amaxa Nucleofection II Device (Lonza, program B-016) and the corresponding Human Stem Cell Nucleofection Kit (Amaxa, VPH-5022). Electroporated iPSC cells were expanded and analyzed by Sanger sequencing (Microsynth). After additional singularization two identical homozygous *SCN10A* knockout clones (K62.1 and K62.4) were chosen for pluripotency analysis and cardiac differentiation. Two $\text{Na}_V1.8$ knockout iPSC-lines (K62.1 and K62.4) and the corresponding isogenic control iPSC-line⁵² were cultured feeder-free and adherent by cultivation on Geltrex-coated cell culture dishes in the presence of chemically defined medium E8 (Life Technologies).

Directed cardiac differentiation of iPSCs was done by manipulation of the Wnt signaling pathway⁵². Briefly, iPSCs were cultured in E8 medium as a monolayer on Geltrex-coated 12-well plates to a confluence of 85–95%. Wnt signaling was initiated by a medium change to RPMI1640 GlutaMAX (Thermo Fisher Scientific) supplemented with human recombinant albumin, L-ascorbic acid 2-phosphate including the GSK3 β inhibitor CHIR99021 (4 $\mu\text{mol/L}$, Millipore). After 48 h, cells were treated with fresh medium supplemented with the inhibitor of Wnt production 2 (IWP2) (5 $\mu\text{mol/L}$, Millipore) for 48 h. On day 6, the medium was changed to RPMI1640 GlutaMAX with 1x B27 with insulin (Thermo Fisher Scientific) with a medium change every 2–3 days. Cardiomyocytes were purified by using a metabolic selection for 2–4 days with 4 mmol/L lactate as carbon source after 15–20 days of differentiation⁵³. Cells were cultured for 60 days and passaged onto glass-bottom FluoroDishes (WPI, 30 K/dish) by trypsinizing for 3 min at 37°C . Cells settled for 7 days prior to further measurements with medium change every 2 days. iPSC-cardiomyocytes were analyzed 8–10 weeks after initiation of differentiation except when mentioned otherwise. Following differentiation, purity of iPSC-CM was determined by flow analysis (>90% cardiac TNT+) and qPCR of ventricular cardiomyocytes marker MLC2v. Four to five differentiation

experiments into ventricular iPSC-CMs of two $N_{AV}1.8$ knockout lines and the corresponding healthy isogenic control line were used.

Cardiomyocyte isolation

Human. Human myocardium was rinsed, cut into small pieces, and incubated at 37 °C in a spinner flask filled with Joklik-MEM solution (JMEM; AppliChem, Darmstadt, Germany) containing 1.0 mg/mL collagenase (type 1, 185 U/mg; Worthington Biochem, New Jersey, USA) and trypsin (2.5 g/L, Life Technologies, Carlsbad, California, USA). After 45 min of digestion, the supernatant was discarded, and the tissue was incubated with fresh JMEM solution containing only collagenase. The solution was incubated for 10 to 20 min until cardiomyocytes were disaggregated using a Pasteur pipette. The supernatant containing disaggregated cells was removed and centrifuged (700 r.p.m., 5 min). Fresh JMEM with collagenase was added to the remaining tissue. This procedure was repeated 4 to 7 times. After every step, the centrifuged cells were resuspended in JMEM medium containing (in mmol/L): 10 taurine, 70 glutamic acid, 25 KCl, 10 KH_2PO_4 , 22 dextrose, 0.5 EGTA, and 10% bovine calf serum (pH 7.4 adjusted with KOH at room temperature). Only cell solutions containing elongated, non-granulated cardiomyocytes with cross-striations were selected for experiments, plated on laminin-coated recording chambers, and allowed to settle for 30 min.

Mouse. Explanted mouse hearts from male and female wild-type (WT), $SCN10A^{-/-}$, $CaMKII\delta^{+/T}$, and $SCN10A^{-/-}/CaMKII\delta^{+/T}$ mice were retrogradely perfused with a nominally Ca^{2+} -free Tyrode's solution containing (in mmol/L): 113 NaCl, 4.7 KCl, 0.6 KH_2PO_4 , 0.6 $Na_2HPO_4 \times 2H_2O$, 1.2 $MgSO_4 \times 7H_2O$, 12 $NaHCO_3$, 10 $KHCO_3$, 10 HEPES, 30 taurine, 10 BDM, 5.5 glucose, and 0.032 phenol red (pH 7.4, with NaOH at 37 °C) using a Langendorff perfusion apparatus. Then, 0.05 mg/mL liberase TM (Roche Diagnostics, Mannheim, Germany), 0.014% trypsin, and 0.1 mmol/L $CaCl_2$ were added to the perfusion solution. Once the heart became flaccid, ventricular tissue was removed, cut into small pieces, and dispersed. Ca^{2+} reintroductions were performed carefully by increasing Ca^{2+} stepwise from 0.1 to 0.4 mmol/L for the patch-clamp and to 1.6 mmol/L for Ca^{2+} spark experiments. Cells were plated on laminin-coated chambers, allowed to settle for 15 min, and then used for measurements.

Co-immunoprecipitation. Tissue homogenates from human ventricular myocardium were suspended in lysis buffer containing (1% CHAPS in RIPA buffer containing (in mmol/L): 50 Tris-HCl, 120 NaCl, 200 NaF, 1 Na_3VO_4 , 1 DTT, (pH 7.4), and complete protease and phosphatase inhibitor cocktails (Roche Diagnostics). Protein concentration was determined by a BCA assay. $CaMKII\delta$ was immunoprecipitated with rabbit polyclonal anti- $CaMKII\delta$ antibody (3 μ g/500 μ g of protein; preincubation at 4 °C overnight, a gift from Prof. Donald M. Bers, Department of Pharmacology, University of California Davis, USA) and protein G Plus Agarose (prewashed and blocked with 5% BSA overnight at 4 °C; Pierce, #22851) for 2 h and 30 min at 4 °C. As a control, rabbit anti-control IgG (3 μ g/500 μ g protein Santa Cruz, sc-2027) was used and a negative control without antibody. The pellets were collected and washed with the RIPA buffer without CHAPS 3 times. The immunoprecipitated proteins were eluted in 2x Laemmli sample buffer containing β -mercaptoethanol (10 min at 70 °C). Supernatants were subjected to Western blotting to detect $N_{AV}1.8$ (mouse monoclonal, LSBio, LS-C109037), $CaMKII\delta$ (rabbit polyclonal, Thermo Scientific, PA5-22168), and GAPDH (mouse monoclonal, Biotrend, BTMC-A437-9).

Immunofluorescence staining of human isolated cardiomyocytes. Isolated human ventricular cardiomyocytes were allowed to attach the poly-L-lysine coated glass chambers for 30 to 45 min at room temperature. Cardiomyocytes were fixed in ice-cold 99.2% ethanol for 20 min at -20 °C. After fixation, cardiomyocytes were washed with phosphate buffer saline (1X PBS) and subsequently blocked and permeabilized with 0.5% Triton X-100 and 5% BSA (bovine serum albumin) in PBS at room temperature.

After blocking, cells were washed with PBS (3x 5 min) and incubated with a primary antibody: mouse monoclonal anti- $N_{AV}1.8$ (1:50, LSBio, LS-C109037). The antibody was diluted in antibody diluent (Dako) containing 0.5% Triton X-100 and incubated overnight at 4 °C. On the next day, cells were washed with PBS (3x 10 min). After washing, cells were incubated for 1 h at room temperature in darkness with the fluorescent-conjugated secondary antibody: Alexa Fluor-488 goat anti-mouse (1:200, Life Technologies, A-11029). In the next step, cells were again incubated overnight at 4 °C with a primary antibody: rabbit polyclonal anti- $CaMKII\delta$ (1:100, gift from D. M. Bers, Department of Pharmacology, University of California Davis, USA). After washing with PBS (3x 5 min) cells were incubated for 1 h at room temperature in darkness with a fluorescent-conjugated secondary antibody: Alexa Fluor-555 goat anti-rabbit (1:200, Life Technologies, A-21424). After washing, cells were covered with Vectashield HardSet Mounting Medium (Vector Laboratories) and viewed using an LSM 5 Pascal confocal microscope (Zeiss) with a 40x oil immersion objective.

Immunofluorescence of iPSCs and iPSC-cardiomyocytes. $N_{AV}1.8$ KO-iPSCs and iPSC-CMs were fixated with 4% Histofix solution (Sigma) for 20 min at room temperature and subsequently blocked in 1% BSA overnight at 4 °C. The primary antibodies mouse anti-SSEA4 (1:200, Abcam), mouse anti- α -actinin (1:750, Sigma),

rabbit anti-Titin-M8/M9 (1:750, MyoMedix) were incubated overnight at 4 °C. The fluorescently labeled secondary antibodies were added for 1 h at 37 °C (AF488 donkey anti-mouse IgG (1:1000, Invitrogen); AF555 donkey anti-rabbit IgG (1:750, Invitrogen). Images were acquired with the Axiovert 200 fluorescence microscope (Zeiss) and the Axiovision software (Rel 4.8).

Patch-clamp experiments

I_{NaL} measurements. To measure I_{NaL} in human and mouse ventricular cardiomyocytes, a ruptured whole-cell patch-clamp was performed at room temperature. The resistance of the pipette was between 2 and 3 mega-Ohm ($M\Omega$) when filled with pipette solution containing (in mmol/L): 95 CsCl, 40 Cs-glutamate, 10 NaCl, 0.92 $MgCl_2$, 5 Mg-ATP, 0.3 Li-GTP, 5 HEPES, 0.03 niflumic acid (to block Ca^{2+} -activated chloride current), 0.02 nifedipine (to block Ca^{2+} current), 0.004 strophanthidin (to block Na^+/K^+ ATPase current) 1 EGTA, and 0.36 $CaCl_2$ (free $[Ca^{2+}]_i$, 100 nmol/L) (pH 7.2 with CsOH at room temperature). Cardiomyocytes were maintained in the bath solution containing (in mmol/L): 135 NaCl, 5 tetramethylammonium chloride, 4 CsCl, 2 $MgCl_2$, 10 glucose, 10 HEPES (pH 7.4 with CsOH at room temperature). To minimize contaminating Ca^{2+} currents during I_{NaL} measurements, Ca^{2+} was omitted from the bath solution. I_{NaL} was measured only in those cardiomyocytes where a seal of more than 1 giga-Ohm was achieved and the access resistance remained <7 $M\Omega$. When whole-cell patch configuration was achieved, cardiomyocytes were given a period of 3 min to be stabilized before conducting measurements. Then cardiomyocytes were held at -120 mV before depolarization to -35 mV for a duration of 1000 ms with 10 pulses and a basic cycle length of 2 s. I_{NaL} was measured as integral current amplitude between 100 and 500 ms and was normalized to the membrane capacitance. Mouse cardiomyocytes were incubated with either A-803467 (30 nmol/L, Sigma) or PF-01247324 (1 μ mol/L, Sigma) for 15 min before starting I_{NaL} measurements. Human cardiomyocytes were incubated with autocomptide-2-related inhibitor peptide (AIP, 1 μ mol/L), PF-01247324 (1 μ mol/L, Sigma), or PF-01247324+AIP for 15 min. For control groups, cardiomyocytes were incubated for 15 min in a normal bath solution to equilibrate the measurements.

Current-voltage relationship of I_{NaL} . Ruptured whole-cell patch-clamp experiments were applied to establish the current-voltage relationship of I_{NaL} . The pipette resistance ranged between 1.5 and 3.5 $M\Omega$ when filled with the pipette solution containing (in mmol/L): 95 CsCl, 40 Cs-glutamate, 10 NaCl, 0.92 $MgCl_2$, 5 Mg-ATP, 0.3 Li-GTP, 5 HEPES, 0.03 niflumic acid (to block Ca^{2+} -activated chloride current), 0.02 nifedipine (to block Ca^{2+} current), 0.004 strophanthidin (to block Na^+/K^+ ATPase current), 1 EGTA, and 0.36 $CaCl_2$ (free $[Ca^{2+}]_i$, 100 nmol/L) (pH 7.2 with CsOH at room temperature). The bath solution contained (in mmol/L): 135 NaCl, 5 tetramethylammonium chloride, 4 CsCl, 2 $MgCl_2$, 10 glucose, 10 HEPES (pH 7.4 with CsOH at room temperature). In some experiments, cardiomyocytes were incubated with PF-01247324 (1 μ mol/L, Sigma) for 15 min before initiation of I_{NaL} measurements. After patch rupture cardiomyocytes were allowed to stabilize for at least 3 min. Currents were elicited using a voltage step protocol with 10 mV increments ranging from -120 to +30 mV (step duration: 1 s, holding potential: -120 mV). I_{NaL} was determined as the mean current density between 180 to 190 ms of each depolarization step. PF-sensitive I_{NaL} was calculated as the difference between the mean I_{NaL} densities of the vehicle and PF-treated cardiomyocytes at each membrane potential.

Peak Na^+ current measurements. Ruptured-patch whole-cell voltage-clamp was used to measure I_{Na} in mouse cardiomyocytes with microelectrodes (2-3 $M\Omega$, room temperature). Pipettes were filled with (in mmol/L): 80 CsCl, 40 Cs-glutamate, 5 NaCl, 0.92 $MgCl_2$, 5 Mg-ATP, 0.3 Li-GTP, 5 HEPES, 0.03 niflumic acid (to block Ca^{2+} -activated chloride current), 0.02 nifedipine (to block Ca^{2+} current), 0.004 strophanthidin (to block Na^+/K^+ ATPase current), 1 EGTA, and 0.36 $CaCl_2$ (free $[Ca^{2+}]_i$, 100 nmol/L) (pH 7.2 with CsOH at room temperature). Cardiomyocytes were maintained in the bath solution containing (in mmol/L): 5 NaCl, 135 tetramethylammonium chloride, 4 CsCl, 2 $MgCl_2$, 10 glucose, 10 HEPES, 0.4 $CaCl_2$ (pH 7.4 with KOH at room temperature). I_{Na} was measured only in those cardiomyocytes where a seal of more than 1 Giga-Ohm was achieved and the access resistance remained <7 $M\Omega$. Mouse cardiomyocytes were incubated with PF-01247324 (1 μ mol/L, Sigma) for 15 min before starting I_{Na} measurements. In all experiments, myocytes were mounted on the stage of a microscope (Nikon Eclipse TE2000-U). Liquid junction potentials were corrected with the pipette in the bath. Membrane capacitance and series resistance were compensated after patch rupture. Data were collected using Patchmaster 2.0 (HEKA Elektronik).

Membrane potential recordings. The whole-cell patch-clamp technique in current-clamp configuration was used to measure membrane potential in single isolated cardiomyocytes. Microelectrodes (3-5 $M\Omega$, room temperature) were filled with (in mmol/L): 92 K-Aspartate, 48 KCl, 1 Mg-ATP, 10 HEPES, 0.02 EGTA, 0.1 GTP-Tris, and 4 Na_2 -ATP (pH 7.2, KOH). The bath solution contained (in mmol/L): 140 NaCl, 4 KCl, 1 $MgCl_2$, 2 $CaCl_2$, 10 glucose, and 10 HEPES (pH 7.4, NaOH). Action potentials were continuously elicited by square current pulses of 1-2 nA amplitude and 1-5 ms duration at increasing stimulation frequency (0.5-3 Hz). Access resistance was typically ~5-15 $M\Omega$ after patch rupture. Fast capacitance was compensated for in a cell-attached configuration. Recordings were commenced

after cell stabilization, which was ~5 min after rupture. Data were collected using Patchmaster 2.0 (HEKA Elektronik) and was analyzed using LabChart 7.37 (ADInstruments). Afterdepolarizations such as EADs and DADs were counted under continuous stimulation with 0.5 Hz. An EAD was counted as an EAD when a renewed depolarization of at least 1 mV occurred before complete membrane repolarization. Criteria for counting DADs were similar, but after achieving complete membrane repolarization.

Ca²⁺ spark and Ca²⁺ wave measurement. Human and mouse ventricular cardiomyocytes were loaded with a Fluo-4AM (10 μmol/L for 15 min, Molecular Probes) at room temperature. For some experiments, either A-803467 (30 nmol/L for 15 min, Sigma) or PF-01247324 (1 μmol/L for 15 min, Sigma) was added to the loading buffer to inhibit Na_v1.8. Cells were perfused with Tyrode's solution containing (in mmol/L): 136 NaCl, 4 KCl, 0.33 NaH₂PO₄, 4 NaHCO₃, 2 CaCl₂, 1.6 MgCl₂, 10 HEPES, 10 glucose (pH 7.4 adjusted with NaOH at room temperature) and the respective active agent. The cardiomyocytes were washed with Tyrode's solution for 15 min for de-esterification of the Ca²⁺ dye. Ca²⁺ spark measurements were performed with a laser scanning confocal microscope (LSM 5 Pascal, Zeiss) using a 40× oil immersion objective. Fluo-4AM was excited by an argon ion laser (488 nm), and emitted fluorescence was collected through a 505-nm long-pass emission filter. Fluorescence images were recorded in the line scan mode with 512 pixels per line (width of each scanline: 38.4 μm) and a pixel time of 0.64 μs. One image consists of 10,000 unidirectional line scans, equating to a measurement period of 7.68 s. Experiments were conducted under resting conditions after loading the sarcoplasmic reticulum with Ca²⁺ by field stimulation at 1 Hz and 20 V. Ca²⁺ sparks were analyzed with the program SparkMaster for ImageJ. The mean spark frequency of the respective cell (CaSpF) resulted from the number of sparks normalized to cell width and scan rate (100 μm/s). Cardiomyocytes exhibiting major arrhythmic events (Ca²⁺ waves, spontaneous Ca²⁺ transients, and spark clouds) were excluded from the quantification of the Ca²⁺ spark. Cardiomyocytes showing Ca²⁺ waves were analyzed separately. The percentage of cardiomyocytes exhibiting Ca²⁺ waves was calculated and compared between the groups. Some cardiomyocytes showed more than one event therefore we calculated the events per time and represented this ratio as waves per minute. The time to the first arrhythmic event was measured in seconds.

Ca²⁺ transient measurements. Cardiomyocytes were isolated from CaMKIIδ^{+T} mouse ventricles, plated as described above, and incubated with a Fura-2AM loading buffer (10 μmol/L, Molecular Probes, Eugene, Oregon, USA) for 20 min. After staining, the cardiomyocytes were washed with experimental solution (as described above in the Ca²⁺ sparks measurement section, at room temperature) for 15 min before measurements were started to enable complete de-esterification of intracellular Fura-2AM and to allow cellular rebalance of Ca²⁺ cycling properties. During measurements, cardiomyocytes were continuously superfused with an experimental solution. Measurements were performed with a Motic AE31 microscope provided with a fluorescence detection system (ION OPTIX Corp.) at room temperature. Cells were excited at 340 and 380 nm and the emitted fluorescence was collected at 510 nm. The intracellular Ca²⁺ level was measured as the ratio of fluorescence at 340 and 380 nm (F340/F380 nm in ratio units, r.u.). Ca²⁺ transients were recorded at steady-state conditions under constant field stimulation (1.0, 2.0, and 4.0 Hz, 20 V). The recorded Ca²⁺ transients were analyzed with the software IONWizard^R (ION OPTIX Corp.).

Analysis of Ca²⁺-transients from confocal line scans. From confocal line scans that were used for quantification of diastolic Ca²⁺ waves in cardiomyocytes from CaMKIIδ^{+T} and SCN10A^{-/-}/CaMKIIδ^{+T} mice Ca²⁺-kinetics were analyzed from the Ca²⁺-transients at the beginning of each line scan before 3 Hz steady-state stimulation was paused. Ca²⁺-transient amplitude (F, normalized to diastole, F0) and time to half-maximal decay (RT50), and time to 90% of maximal decay were calculated from confocal line scan images using ImageJ software.

Quantitative real-time PCR (RT-qPCR). Human or mouse cardiac tissues, isolated human cardiomyocytes or iPSC-cardiomyocytes were snap-frozen in liquid nitrogen and stored at -80 °C. RNA was isolated by use of the ReliaPrepTM RNA Tissue Miniprep System (Promega). About 200 ng RNA was reverse transcribed into cDNA using standard protocols (QuantiNova Reverse Transcription Kit (QIAGEN) for mouse tissue and iScript cDNA Synthesis Kit (Bio-Rad) for human samples). For qPCR, 10 μL SYBR Green PCR Master Mix (Bio-Rad), 7 μL nuclease-free water, 1 μL forward and 1 μL reverse Primer, and 1 μL of cDNA were mixed. Q-PCR was carried out using the CFX ConnectTM Real-Time System (Bio-Rad). Forty cycles of 15 s at 95 °C followed by 1 min of 60 °C were used and fluorescence was measured after each cycle. After 40 cycles melt curve analysis was performed to ensure the specificity of the products. Thresholds cycles were evaluated and normalized to housekeeping genes and controls. A list of all primers used is presented in Supplementary Table 4.

Western blots. Ventricular tissue from WT, SCN10A^{-/-}, CaMKIIδ^{+T}, and SCN10A^{-/-}/CaMKIIδ^{+T} mice or pellets of iPSC-cardiomyocytes from WT and homozygous Na_v1.8 knockout (KO) lines were homogenized in Tris buffer

containing (in mmol/L): 20 Tris-HCl, 200 NaCl, 20 NaF, 1 Na₃VO₄, 1 DTT, 1% Triton X-100 (pH 7.4), and complete protease and phosphatase inhibitor cocktails (Roche Diagnostics). Protein concentration was determined by BCA assay (Pierce Biotechnology). Denatured tissue homogenates (10 min, 70 °C in 2% beta-mercaptoethanol) were separated on 8% SDS-polyacrylamide gels, then transferred to a nitrocellulose membrane, and incubated with the following primary antibodies: rabbit polyclonal anti-CaMKIIδ (1:5000, Thermo Scientific, PA5-22168), rabbit polyclonal anti-Na_v1.5 (1:2000, Alomone labs, ASC-005), and mouse monoclonal anti-GAPDH (1:20000, BIOTREND, BTMC-A473-9) at 4 °C overnight. Secondary antibodies included HRP-conjugated goat anti-rabbit and goat anti-mouse (1:20000, Jackson Immunoresearch, 111-035-144 and 115-035-062, respectively). The membrane was incubated with secondary antibodies for 1 h at RT. ImmobilonTM Western Chemiluminescent HRP Substrate (Millipore) was used for chemiluminescent detection. Analysis was performed using Image Studio Lite Ver 5.2. The full scan blots for Na_v1.5 and CaMKIIδ are shown in the Data Availability file.

Histology. Freshly explanted hearts were fixed in 4% buffered formaldehyde at 4 °C for 24 h. LV tissue was harvested, fixed in 4% buffered formaldehyde overnight, paraffin-embedded, sectioned (5 μm), and stained with fluorescein-conjugated wheat germ agglutinin (WGA-Alexa Fluor 594; Invitrogen, USA) for cross-sectional area (CSA) assessment. At least 300 randomly selected cardiomyocytes per animal from different sections and different regions (basal, mid-ventricular, apical) of each heart were measured using the ImageJ software (Bethesda, USA).

Mouse echocardiography. For echocardiography, mice were anesthetized using 1.5% isoflurane, and echocardiography was performed using a VS-VEVO 660/230 (VisualSonics, Canada). During this procedure, the core temperature was maintained at 37 °C, and heart rates were kept consistent between experimental groups at 400–500 b.p.m. Electrocardiogram monitoring was obtained using hind limb electrodes. LV geometry and systolic function were assessed by using standard 2D parasternal long and short-axis views. The examiner was blinded towards group assignment.

Telemetric ECG recordings in mice. Mice were implanted with an intraperitoneal telemetric ECG transmitter (TA11ETA-F10, Data Sciences International) with its Cable in lead II configuration. After a postoperative recovery period of 10 days, we recorded ECGs during periods of normal activity (24 h continuous recording, twice a week for 2 weeks). ECG parameters were analyzed, and the QT interval was corrected (QTc) using Bazett's formula, QTc = QT/(RR/100)^{1/2} established for mice with QT and RR expressed in ms. Premature ventricular contractions and ventricular tachycardia (>3 beats) were counted per 24 h and a medium value was per animal was calculated from all measurements. Further, physical activity was recorded by the telemetric monitors.

Statistics. All data were expressed as the mean ± standard error of the mean (SEM). Where appropriate, one-way ANOVA with multiple comparison tests (post hoc Bonferroni's and two-stage step-up methods of Benjamini, Krieger, and Yekutieli correction) was used. Mixed-effects analysis with Holm-Sidak's post hoc test was used to analyze I-V curves. Otherwise, Student's unpaired *t*-tests were used. The Chi-Square test was used to compare the occurrence of diastolic Ca²⁺ waves. Log-rank (Mantel-Cox) test was used to compare survival between CaMKIIδ^{+T} and SCN10A^{-/-}/CaMKIIδ^{+T} mice. Two-sided *p* ≤ 0.05 was considered significant.

Reporting Summary. Further information on research design is available in the Nature Research Reporting Summary linked to this article.

Data availability

All data generated or analyzed during this study are available within the Article and its Supplementary Information. All raw data supporting the findings from this study are available from the corresponding author upon reasonable request. Source data are provided with this paper. Any remaining raw data will be available from the corresponding author upon reasonable request.

Received: 21 May 2019; Accepted: 14 October 2021;

Published online: 15 November 2021

References

- Despa, S., Islam, M. A., Weber, C. R., Pogwizd, S. M. & Bers, D. M. Intracellular Na(+) concentration is elevated in heart failure but Na/K pump function is unchanged. *Circulation* **105**, 2543–2548 (2002).
- Maltsev, V. A. et al. Novel, ultraslow inactivating sodium current in human ventricular cardiomyocytes. *Circulation* **98**, 2545–2552 (1998).

3. Maltsev, V. A., Silverman, N., Sabbah, H. N. & Undrovinas, A. I. Chronic heart failure slows late sodium current in human and canine ventricular myocytes: implications for repolarization variability. *Eur. J. Heart Fail.* **9**, 219–227 (2007).
4. Song, Y., Shryock, J. C. & Belardinelli, L. An increase of late sodium current induces delayed afterdepolarizations and sustained triggered activity in atrial myocytes. *AJP Hear. Circ. Physiol.* **294**, H2031–H2039 (2008).
5. Valdivia, C. R. et al. Increased late sodium current in myocytes from a canine heart failure model and from failing human heart. *J. Mol. Cell. Cardiol.* **38**, 475–483 (2005).
6. Sossalla, S. et al. Ranolazine improves diastolic dysfunction in isolated myocardium from failing human hearts — Role of late sodium current and intracellular ion accumulation. *J. Mol. Cell. Cardiol.* **45**, 32–43 (2008).
7. Undrovinas, A. I., Maltsev, V. A. & Sabbah, H. N. Repolarization abnormalities in cardiomyocytes of dogs with chronic heart failure: Role of sustained inward current. *Cell. Mol. Life Sci.* **55**, 494–505 (1999).
8. Fischer, T. H. et al. Late INa increases diastolic SR-Ca²⁺-leak in atrial myocardium by activating PKA and CaMKII. *Cardiovasc. Res.* **107**, 184–196 (2015).
9. Sag, C. M. et al. Calcium/calmodulin-dependent protein kinase II contributes to cardiac arrhythmogenesis in heart failure. *Circ. Hear. Fail.* **2**, 664–675 (2009).
10. Toischer, K. et al. Role of late sodium current as a potential arrhythmogenic mechanism in the progression of pressure-induced heart disease. *J. Mol. Cell. Cardiol.* **61**, 111–122 (2013).
11. Bengel, P. et al. Contribution of the neuronal sodium channel NaV1.8 to sodium- and calcium-dependent cellular proarrhythmia. *J. Mol. Cell. Cardiol.* **144**, 35–46 (2020).
12. Larbig, R., Torres, N., Bridge, J. H. B., Goldhaber, J. I. & Philipson, K. D. Activation of reverse Na⁺-Ca²⁺ exchange by the Na⁺ current augments the cardiac Ca²⁺ transient: evidence from NCX knockout mice. *J. Physiol.* **588**, 3267–3276 (2010).
13. Maier, L. S. et al. Transgenic CaMKII δ overexpression uniquely alters cardiac myocyte Ca²⁺ handling: reduced SR Ca²⁺ load and activated SR Ca²⁺ release. *Circ. Res.* **92**, 904–911 (2003).
14. Ashpole, N. M. et al. Ca²⁺/calmodulin-dependent protein kinase II (CaMKII) regulates cardiac sodium channel NaV1.5 gating by multiple phosphorylation sites. *J. Biol. Chem.* **287**, 19856–19869 (2012).
15. Sag, C. M. et al. Enhanced late INa induces proarrhythmic SR Ca leak in a CaMKII-dependent manner. *J. Mol. Cell. Cardiol.* **76**, 94–105 (2014).
16. Zhang, T. et al. The δ C isoform of CaMKII is activated in cardiac hypertrophy and induces dilated cardiomyopathy and heart failure. *Circ. Res.* **92**, 912–919 (2003).
17. Wehrens, X. H. T., Lehman, S. E., Reiken, S. R. & Marks, A. R. Ca²⁺/calmodulin-dependent protein kinase II phosphorylation regulates the cardiac ryanodine receptor. *Circ. Res.* **94**, e61–e70 (2004).
18. Kohlhaas, M. et al. Increased sarcoplasmic reticulum calcium leak but unaltered contractility by acute CaMKII overexpression in isolated rabbit cardiac myocytes. *Circ. Res.* **98**, 235–244 (2006).
19. Ai, X., Curran, J. W., Shannon, T. R., Bers, D. M. & Pogwizd, S. M. Ca²⁺/calmodulin-dependent protein kinase modulates cardiac ryanodine receptor phosphorylation and sarcoplasmic reticulum Ca²⁺ leak in heart failure. *Circ. Res.* **97**, 1314–1322 (2005).
20. Fischer, T. H. et al. Ca²⁺/calmodulin-dependent protein kinase II and protein kinase A differentially regulate sarcoplasmic reticulum Ca²⁺ leak in human cardiac pathology. *Circulation* **128**, 970–981 (2013).
21. Wagner, S. et al. Ca²⁺/calmodulin-dependent protein kinase II regulates cardiac Na⁺ channels. *J. Clin. Invest.* **116**, 3127–3138 (2006).
22. Glynn, P. et al. Voltage-gated sodium channel phosphorylation at Ser571 regulates late current, arrhythmia, and cardiac function in vivo. *Circulation* **132**, 567–577 (2015).
23. Akopian, A. N., Sivilotti, L. & Wood, J. N. A tetrodotoxin-resistant voltage-gated sodium channel expressed by sensory neurons. *Nature* **379**, 257–262 (1996).
24. Ritchie, M. D. et al. Genome- and phenome-wide analyses of cardiac conduction identifies markers of arrhythmia risk. *Circulation* **127**, 1377–1385 (2013).
25. Jabbari, J. et al. Common and rare variants in *SCN10A* modulate the risk of atrial fibrillation. *Circ. Cardiovasc. Genet.* **8**, 64–73 (2015).
26. Savio-Galimberti, E. et al. *SCN10A*/Nav1.8 modulation of peak and late sodium currents in patients with early onset atrial fibrillation. *Cardiovasc. Res.* **104**, 355–363 (2014).
27. Gando, I. et al. Functional characterization of *SCN10A* variants in several cases of sudden unexplained death. *Forensic Sci. Int.* **301**, 289–298 (2019).
28. Facer, P. et al. Localisation of *SCN10A* gene product Nav1.8 and novel pain-related ion channels in human heart. *Int. Heart J.* **52**, 146–152 (2011).
29. Yang, T. et al. Blocking *Scn10a* channels in heart reduces late sodium current and is antiarrhythmic. *Circ. Res.* **111**, 322–332 (2012).
30. Pabel, S. et al. Inhibition of NaV1.8 prevents atrial arrhythmogenesis in human and mice. *Basic Res. Cardiol.* **115**, 20 (2020).
31. Dybkova, N. et al. Differential regulation of sodium channels as a novel proarrhythmic mechanism in the human failing heart. *Cardiovasc. Res.* **114**, 1728–1737 (2018).
32. Ahmad, S. et al. The functional consequences of sodium channel NaV1.8 in human left ventricular hypertrophy. *ESC Hear. Fail.* **6**, 154–163 (2019).
33. Kirchhefer, U., Schmitz, W., Scholz, H. & Neumann, J. Activity of cAMP-dependent protein kinase and Ca²⁺/calmodulin-dependent protein kinase in failing and nonfailing human hearts. *Cardiovasc. Res.* **42**, 254–261 (1999).
34. Neef, S. et al. CaMKII-dependent diastolic SR Ca²⁺ leak and elevated diastolic Ca²⁺ levels in right atrial myocardium of patients with atrial fibrillation. *Circ. Res.* **106**, 1134–1144 (2010).
35. Zalcman, G., Federman, N. & Romano, A. CaMKII isoforms in learning and memory: Localization and function. *Front. Mol. Neurosci.* **11**, 445 (2018).
36. Hund, T. J. et al. A β IV-spectrin/CaMKII signaling complex is essential for membrane excitability in mice. *J. Clin. Invest.* **120**, 3508–3519 (2010).
37. Montersino, A. et al. Tetrodotoxin-resistant voltage-gated sodium channel Na_v1.8 constitutively interacts with ankyrin G. *J. Neurochem.* **131**, 33–41 (2014).
38. Koval, O. M. et al. Ca²⁺/calmodulin-dependent protein kinase ii-based regulation of voltage-gated na⁺ channel in cardiac disease. *Circulation* **126**, 2084–2094 (2012).
39. Sossalla, S. et al. Inhibition of elevated Ca²⁺/calmodulin-dependent protein kinase II improves contractility in human failing myocardium. *Circ. Res.* **107**, 1150–1161 (2010).
40. Sossalla, S. et al. Diastolic dysfunction and arrhythmias caused by overexpression of CaMKII δ C can be reversed by inhibition of late Na⁺ current. *Basic Res. Cardiol.* **106**, 263–272 (2011).
41. Undrovinas, A. I., Maltsev, V. A., Kyle, J. W., Silverman, N. & Sabbah, H. N. Gating of the late Na⁺ channel in normal and failing human myocardium. *J. Mol. Cell. Cardiol.* **34**, 1477–1489 (2002).
42. Casini, S. et al. Absence of functional Nav1.8 channels in non-diseased atrial and ventricular cardiomyocytes. *Cardiovasc. Drugs Ther.* **33**, 649–660 (2019).
43. Stroud, D. M. et al. Contrasting Nav1.8 activity in *Scn10a*^{-/-} ventricular myocytes and the intact heart. *J. Am. Heart Assoc.* **5**, e002946 (2016).
44. Bengel, P., Ahmad, S. & Sossalla, S. Inhibition of late sodium current as an innovative antiarrhythmic strategy. *Curr. Heart Fail. Rep.* **14**, 179–186 (2017).
45. Dybkova, N. et al. Overexpression of CaMKII δ C in RyR2R4496C^{+/−} knock-in mice leads to altered intracellular Ca²⁺ handling and increased mortality. *J. Am. Coll. Cardiol.* **57**, 469–479 (2011).
46. Mohamed, B. A. et al. Sarcoplasmic reticulum calcium leak contributes to arrhythmia but not to heart failure progression. *Sci. Transl. Med.* **10**, ean0724 (2018).
47. Al-Khatib, S. M. et al. 2017 AHA/ACC/HRS guideline for management of patients with ventricular arrhythmias and the prevention of sudden cardiac death. *Circulation* **138**, e272–e391 (2018).
48. Meinertz, T. et al. Significance of ventricular arrhythmias in idiopathic dilated cardiomyopathy. *Am. J. Cardiol.* **53**, 902–907 (1984).
49. Hofmann, T. et al. Mode of death in idiopathic dilated cardiomyopathy: a multivariate analysis of prognostic determinants. *Am. Heart J.* **116**, 1455–1463 (1988).
50. Neef, S., Mann, C., Zwenger, A., Dybkova, N. & Maier, L. S. Reduction of SR Ca²⁺ leak and arrhythmogenic cellular correlates by SMP-114, a novel CaMKII inhibitor with oral bioavailability. *Basic Res. Cardiol.* **112**, 45 (2017).
51. Payne, C. E. et al. A novel selective and orally bioavailable Nav1.8 channel blocker, PF-01247324, attenuates nociception and sensory neuron excitability. *Br. J. Pharmacol.* **172**, 2654–2670 (2015).
52. Borchert, T. et al. Catecholamine-dependent β -adrenergic signaling in a pluripotent stem cell model of Takotsubo cardiomyopathy. *J. Am. Coll. Cardiol.* **70**, 975–991 (2017).
53. Tohyama, S. et al. Distinct metabolic flow enables large-scale purification of mouse and human pluripotent stem cell-derived cardiomyocytes. *Cell Stem Cell* **12**, 127–137 (2013).

Acknowledgements

We thank Prof. John Wood (University College London) for providing us a pair of *SCN10A*^{+/−} mice. We thank Timo Schulte, Yvonne Metz, Johanna Heine, Sabrina Kozzawa, Manar El Kenani, and Sarah Zafar for technical assistance. This work was supported by the German Heart Foundation/German Foundation of Heart Research (to P.B., P.T., and S.S.); by the University Hospital Regensburg (ReForM C program) (to L.S.M. and S.S.); by the Marga und Walter Boll-Stiftung (to S.A. and S.S.); the College of Translational Medicine by the Ministry of Culture and Science, State of Lower Saxony (P.B.), the German Center for Cardiovascular Research (DZHK; to K.S.-B.), the Else Kröner-Fresenius Stiftung (N.H. and M.T.) and the Deutsche Forschungsgemeinschaft

(DFG) through the International Research Training Group Award (IRTG) 1816 (to K.S.-B.; W.M. is a fellow under IRTG 1816) and SFB 1002 (to K.T., N.D., and G.H.).

Author contributions

P.B., S.A., and S.S. conceived the study. K.S.-B., P.B., and S.S. designed the experiments and wrote the manuscript. P.B., N.D., P.T., S.A., B.M., N.H., M.C.K., W.M., S.P., and M.T. carried out experimental work and analyzed the data. K.S.-B. designed the CRISPR experiments. J.G., H.M., and S.L.-H. acquired and provided human tissue. K.T. applied for the authorization to carry out animal experiments and helped to analyze in vivo data. J.M., S.W., L.S.M., and G.H. provided expertise and feedback. All authors discussed the results and had the opportunity to comment on the manuscript.

Funding

Open Access funding enabled and organized by Projekt DEAL.

Competing interests

The authors declare no competing interests.

Additional information

Supplementary information The online version contains supplementary material available at <https://doi.org/10.1038/s41467-021-26690-1>.

Correspondence and requests for materials should be addressed to Samuel Sossalla.

Peer review information *Nature Communications* thanks Robin Shaw and the other, anonymous, reviewer(s) for their contribution to the peer review of this work. Peer reviewer reports are available.

Reprints and permission information is available at <http://www.nature.com/reprints>

Publisher's note Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.



Open Access This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons license, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons license, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons license and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this license, visit <http://creativecommons.org/licenses/by/4.0/>.

© The Author(s) 2021