

## RESEARCH PAPER

# Modulation of $K_{2P}2.1$ and $K_{2P}10.1$ $K^+$ channel sensitivity to carvedilol by alternative mRNA translation initiation

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## BACKGROUND AND PURPOSE

The  $\beta$ -receptor antagonist carvedilol blocks a range of ion channels.  $K_{2P}2.1$  (TREK1) and  $K_{2P}10.1$  (TREK2) channels are expressed in the heart and regulated by alternative translation initiation (ATI) of their mRNA, producing functionally distinct channel variants. The first objective was to investigate acute effects of carvedilol on human  $K_{2P}2.1$  and  $K_{2P}10.1$  channels. Second, we sought to study ATI-dependent modulation of  $K_{2P}$   $K^+$  current sensitivity to carvedilol.

## EXPERIMENTAL APPROACH

Using standard electrophysiological techniques, we recorded currents from wild-type and mutant  $K_{2P}2.1$  and  $K_{2P}10.1$  channels in *Xenopus* oocytes and HEK 293 cells.

## KEY RESULTS

Carvedilol concentration-dependently inhibited  $K_{2P}2.1$  channels ( $IC_{50, \text{oocytes}} = 20.3 \mu\text{M}$ ;  $IC_{50, \text{HEK}} = 1.6 \mu\text{M}$ ) and this inhibition was frequency-independent. When  $K_{2P}2.1$  isoforms generated by ATI were studied separately in oocytes, the  $IC_{50}$  value for carvedilol inhibition of full-length channels ( $16.5 \mu\text{M}$ ) was almost 5-fold less than that for the truncated channel variant ( $IC_{50} = 79.0 \mu\text{M}$ ). Similarly, the related  $K_{2P}10.1$  channels were blocked by carvedilol ( $IC_{50, \text{oocytes}} = 24.0 \mu\text{M}$ ;  $IC_{50, \text{HEK}} = 7.6 \mu\text{M}$ ) and subject to ATI-dependent modulation of drug sensitivity.

## CONCLUSIONS AND IMPLICATIONS

Carvedilol targets  $K_{2P}2.1$  and  $K_{2P}10.1$   $K^+$  channels. This previously unrecognized mechanism supports a general role of cardiac  $K_{2P}$  channels as antiarrhythmic drug targets. Furthermore, the work reveals that the sensitivity of the cardiac ion channels  $K_{2P}2.1$  and  $K_{2P}10.1$  to block was modulated by alternative mRNA translation initiation.

## Abbreviations

ATI, alternative translation initiation;  $K_{2P}$ , two-pore-domain  $K^+$  channel; TREK, TWIK-related  $K^+$  channel; TWIK, tandem of P domains in a weak inward rectifying  $K^+$  channel

## Introduction

Two-pore-domain potassium ( $K_{2P}$ ) channels stabilize the resting membrane potential and facilitate action potential

repolarization (channel nomenclature follows Alexander *et al.*, 2013). Dynamic and polymodal regulation of  $K_{2P}$  currents determines cellular excitability (Goldstein *et al.*, 2001; Gierten *et al.*, 2008; 2012; Thomas *et al.*, 2008; Sandoz *et al.*,

2011; Staudacher *et al.*, 2011a; Kisselbach *et al.*, 2012; Rahm *et al.*, 2012; 2014; Seyler *et al.*, 2012). In the heart, K<sub>2P</sub>3.1 inhibition or genetic inactivation of K<sub>2P</sub>3.1 currents results in action potential prolongation, suggesting that these channels might represent targets for antiarrhythmic therapy (Putzke *et al.*, 2007; Gierten *et al.*, 2010; Ravens, 2010; Decher *et al.*, 2011; Donner *et al.*, 2011; Limberg *et al.*, 2011; Staudacher *et al.*, 2011b; Petric *et al.*, 2012; Schmidt *et al.*, 2012). Prolongation of action potentials and cardiac effective refractory periods is a characteristic of class III antiarrhythmic drugs, reducing membrane excitability and decreasing arrhythmia susceptibility. In addition to K<sub>2P</sub>3.1, expression of K<sub>2P</sub>2.1 (TREK1, TWIK-related K<sup>+</sup> channel) channels has been shown in atria and ventricles at mRNA, protein and functional levels (Fink *et al.*, 1996; Reyes *et al.*, 1998; Aimond *et al.*, 2000; Medhurst *et al.*, 2001; Terrenoire *et al.*, 2001; Liu and Saint, 2004; Tan *et al.*, 2004; Li *et al.*, 2006; Zhang *et al.*, 2008; Decher *et al.*, 2011; Zhao *et al.*, 2011; Goonetilleke and Quayle, 2012; Schmidt *et al.*, 2014). The closely related K<sub>2P</sub>10.1 (TREK2) channels have been detected in the heart as well (Medhurst *et al.*, 2001; Liu and Saint, 2004; Li *et al.*, 2006; Staudacher *et al.*, 2011a; Gierten *et al.*, 2012). However, data delineating the cardiac functions of K<sub>2P</sub>2.1 and K<sub>2P</sub>10.1 channels are limited. Synthesis of different K<sub>2P</sub> proteins with different N terminal domains from a single mRNA by alternative mRNA translation initiation (ATI) was recently revealed as a novel mechanism to increase the number of functional TREK protein subunits, producing two (K<sub>2P</sub>2.1) or three (K<sub>2P</sub>10.1) endogenous variants (Simkin *et al.*, 2008; Thomas *et al.*, 2008). ATI regulates K<sub>2P</sub> current magnitude, ion selectivity and intersubunit interactions (Simkin *et al.*, 2008; Thomas *et al.*, 2008; Veale *et al.*, 2010; Eckert *et al.*, 2011). Effects of ATI on K<sub>2P</sub> channel sensitivity to antiarrhythmic drugs have not been investigated to date.

Carvedilol is a non-selective  $\beta$ -adrenoreceptor antagonist with a multichannel blocking pharmacological profile. Carvedilol targets a variety of ion currents including K<sup>+</sup> channels (Karle *et al.*, 2001; Staudacher *et al.*, 2011b). The clinical use of carvedilol is associated with reduced morbidity and mortality in heart failure patients (Poole-Wilson *et al.*, 2003; Remme *et al.*, 2007). In addition, suppression of atrial fibrillation and ventricular tachyarrhythmia as well as pronounced prolongation of atrial and ventricular effective refractory periods suggest specific antiarrhythmic effects of the drug (Senior *et al.*, 1992; Brunvand *et al.*, 1996; Cice *et al.*, 2000; Takusagawa *et al.*, 2000; Katritsis *et al.*, 2003; Merritt *et al.*, 2003; Ramaswamy, 2003; Acikel *et al.*, 2008; Kanoupakis *et al.*, 2008).

We hypothesized that inhibition of K<sub>2P</sub>2.1 and K<sub>2P</sub>10.1 background currents contributed to the antiarrhythmic effect of carvedilol. Furthermore, this study was designed to investigate modulation of the drug sensitivity of cardiac K<sub>2P</sub> channels by ATI.

## Methods

### Molecular biology

Human cDNAs encoding wild-type and mutant K<sub>2P</sub>2.1 (TREK1b) (GenBank accession number, EF165335) were

kindly provided by Dr. Steve Goldstein (Brandeis University, Waltham, MA, USA) (Thomas *et al.*, 2008). Human K<sub>2P</sub>2.1 was amplified from a heart cDNA library (Clontech, Palo Alto, CA, USA), inserted into the pCR2.1-TOPO vector (Invitrogen, Carlsbad, CA, USA) and subcloned into pMAX, a dual-purpose expression vector containing a CMV promoter for mammalian expression and a T7 promoter for cRNA synthesis. Human K<sub>2P</sub>10.1 (TREK2c) (GenBank accession number EU978939) was amplified from a brain cDNA library as described (Gierten *et al.*, 2008; Staudacher *et al.*, 2011a). Complementary DNA was inserted into the pCR2.1-TOPO vector and subcloned into pRAT, a dual-purpose expression vector containing a CMV promoter for mammalian expression and a T7 promoter for cRNA synthesis (Bockenbauer *et al.*, 2001). Generation of mutant K<sub>2P</sub>2.1 and K<sub>2P</sub>10.1 clones was previously reported (Thomas *et al.*, 2008; Staudacher *et al.*, 2011a). Briefly, K<sub>2P</sub>2.1 and K<sub>2P</sub>10.1 mutations described in the text were made with the QuikChange Site-Directed Mutagenesis kit (Stratagene, La Jolla, CA, USA). N terminal channel truncation was achieved using PCR. All cDNA constructs were confirmed by DNA sequencing. Complementary RNAs were transcribed after vector linearization using T7 RNA polymerase and the mMessage mMachine kit (Ambion, Austin, TX, USA). Transcripts were quantified by spectrophotometry and cRNA integrity was assessed by agarose gel electrophoresis.

### Oocyte preparation and injection

All animal care and experimental procedures complied with the Directive 2010/63/EU of the European Parliament and the United States National Institutes of Health Guide for the Care and Use of Laboratory Animals (NIH Publication no. 85-23, revised 1996). Approval was granted by the local Animal Welfare Committee (reference number A-38/11). All studies involving animals are reported in accordance with the ARRIVE guidelines for reporting experiments involving animals (Kilkenny *et al.*, 2010; McGrath *et al.*, 2010). The preparation of oocytes has been reported earlier in detail (Gierten *et al.*, 2008; Staudacher *et al.*, 2011b; Seyler *et al.*, 2012). Briefly, oocytes were isolated from *Xenopus laevis* ovarian lobes after surgical removal under tricaine anaesthesia (1 g·L<sup>-1</sup>; pH 7.5). Oocyte collection was alternated between left and right ovaries, and a maximum of three surgeries were performed on one individual frog. After the final collection of oocytes, anaesthetized frogs were killed by decerebration and pithing. Complementary RNA (0.1–25 ng; 46 nl per oocyte) was injected into stage V–VI defolliculated *Xenopus* oocytes.

### Cell culture

HEK 293 cells were cultured in DMEM (Invitrogen) supplemented with 10% FBS, 2 mM L-glutamine, 100 000 U·L<sup>-1</sup> penicillin and 100 mg·L<sup>-1</sup> streptomycin in an atmosphere of 95% humidified air and 5% CO<sub>2</sub> at 37°C. Cells were passaged regularly. Transient transfections (5  $\mu$ g cDNA/35 mm dish) were performed using FuGENE HD transfection reagent (Promega, Madison, WI, USA) according to the manufacturer's instructions.

### Electrophysiology

Whole-cell currents were recorded from *Xenopus* oocytes using two-electrode voltage-clamp electrophysiology with an

Oocyte Clamp amplifier (Warner Instruments, Hamden, CT, USA) and pCLAMP9 (Axon Instruments, Foster City, CA, USA) software 2–3 days after injection, as described (Thomas *et al.*, 2008; Gierten *et al.*, 2012). Data were sampled at 2 kHz and filtered at 1 kHz. Electrodes were filled with 3 M KCl. The extracellular solution contained 96 mM NaCl, 4 mM KCl, 1.1 mM CaCl<sub>2</sub>, 1 mM MgCl<sub>2</sub>, 5 mM HEPES (pH 7.4, adjusted with NaOH).

Whole-cell patch-clamp recordings from HEK 293 cells were carried out using a RK-400 amplifier (Bio-Logic SAS, Claix, France) as reported (Gierten *et al.*, 2008; Schmidt *et al.*, 2012; Seyler *et al.*, 2012). Electrodes were filled with the following solution: 100 mM K-aspartate, 20 mM KCl, 2 mM MgCl<sub>2</sub>, 1 mM CaCl<sub>2</sub>, 10 mM EGTA, 10 mM HEPES, 40 mM glucose, 5 mM K-ATP (pH adjusted to 7.2 with KOH). The external solution for these experiments was composed of 140 mM NaCl, 5 mM KCl, 1 mM MgCl<sub>2</sub>, 1.8 mM CaCl<sub>2</sub>, 10 mM HEPES, 10 mM glucose (pH adjusted to 7.4 with NaOH). Identification of transfected cells was performed by requiring outward current amplitudes of at least 200 pA and leak currents of ≤10% of the initial maximum current. Recordings were carried out under constant perfusion at room temperature, and no leak subtraction was done during the experiments.

### Drug administration

Carvedilol, metoprolol and propranolol (all from Sigma-Aldrich, St. Louis, MO, USA) were prepared as 100 mM stock solutions in DMSO (carvedilol, propranolol) or H<sub>2</sub>O (metoprolol) and stored at –20°C. On the day of experiments, aliquots of the stock solutions were diluted to the desired concentrations with the bath solution. Wild-type and mutant K<sub>2P</sub>2.1 or K<sub>2P</sub>10.1 currents recorded from *Xenopus* oocytes were not significantly altered upon application of 0.1% DMSO ( $v^*v^{-1}$ ; maximum bath concentration) for 20 min ( $n = 4$  to 10 cells were studied for each channel,  $P > 0.05$ ).

### Biochemistry

Pooled tissue-specific (heart, brain, aorta, lung, skeletal muscle, stomach, small intestine, kidney, uterus and prostate) total protein samples ('Protein Medleys') were obtained commercially from Clontech (Mountain View, CA, USA). The samples (30 µg per well) were subjected to SDS-PAGE on precast 10% gels ('Ready Gels', Bio-Rad, Hercules, CA, USA), followed by wet-transfer onto nitrocellulose paper and immunoblot analysis using anti-K<sub>2P</sub>2.1 antibodies raised against recombinant C terminal rat K<sub>2P</sub>2.1 protein containing 68 amino acids (1:400; Thomas *et al.*, 2008). Successful detection of human K<sub>2P</sub>2.1 protein using this antibody has been demonstrated previously (Thomas *et al.*, 2008). Secondary fluorescence-labelled goat anti-rabbit antibodies were used at 1:5000 dilutions, and secondary labelling of fluorescent conjugates was documented with an Odyssey scanner (Li-Cor Biosciences, Lincoln, NE, USA).

### Data analysis

Origin (OriginLab, Northampton, MA, USA) software was used for data analysis. Data are expressed as mean ± SEM. Concentration-response relationships for drug-induced block were fit with a Hill equation of the following form:  $I_{\text{drug}}/I_{\text{control}}$

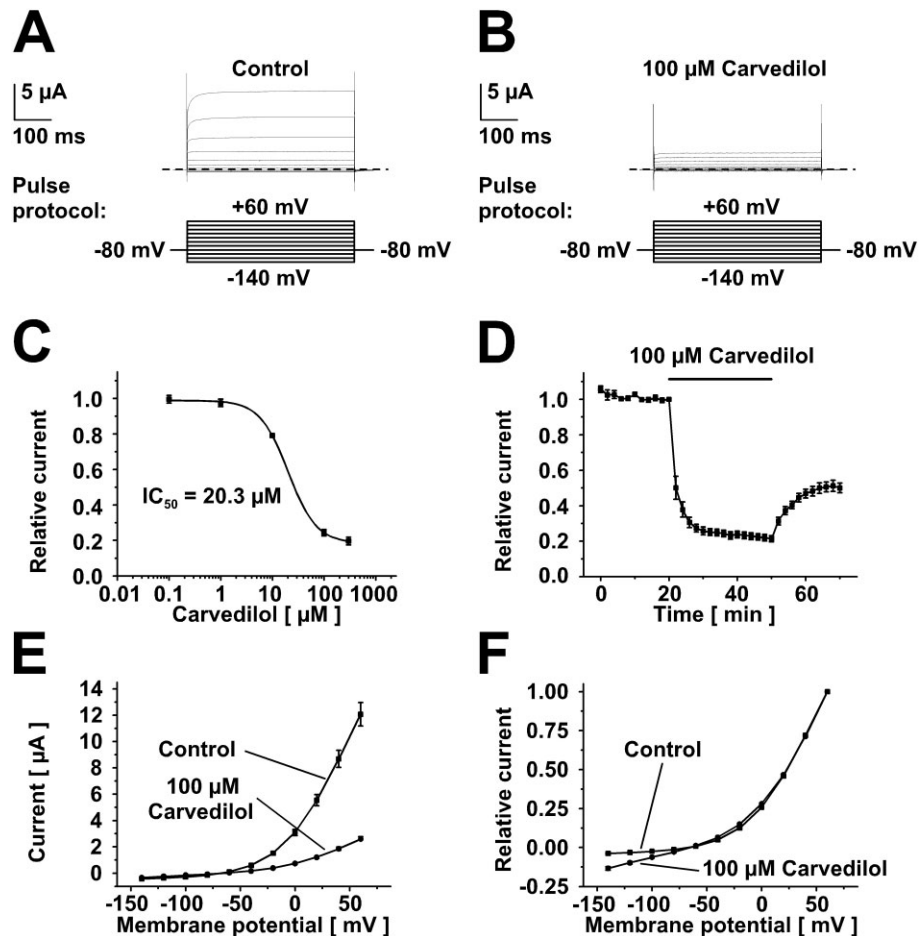
$= 1/[1 + (D/IC_{50})^n]$ , where  $I$  indicates current,  $D$  is the drug concentration,  $n$  is the Hill coefficient and  $IC_{50}$  is the concentration necessary for 50% block. We applied paired and unpaired Student's  $t$ -tests (two-tailed tests) to compare statistical significance of the results:  $P < 0.05$  was considered statistically significant. The multiple comparisons shown in Figure 2F and in Table 3 were performed using one-way ANOVA. If the hypothesis of equal means could be rejected at the 0.05-level, pair-wise comparisons of groups were made and the probability values were adjusted for multiple comparisons using the Bonferroni correction.

## Results

### Inhibition of human cardiac K<sub>2P</sub>2.1 (TREK1) channels by carvedilol

Carvedilol blockade of hK<sub>2P</sub>2.1 channels was investigated in *Xenopus laevis* oocytes. From a holding potential of –80 mV, pulses were applied for 500 ms to voltages between –140 and +60 mV in 20 mV increments (0.5 Hz). Representative families of current traces from one cell are shown for control conditions (Figure 1A) and after application of 100 µM carvedilol for 20 min (Figure 1B). Current amplitudes were analysed at the end of the 0 mV test pulse. After a control period (20 min) with no significant amplitude changes ( $I_{0 \text{ min}}$ ,  $3.17 \pm 0.4 \mu\text{A}$ ;  $I_{20 \text{ min}}$ ,  $3.08 \pm 0.4 \mu\text{A}$ ;  $n = 14$ ;  $P = 0.18$ ), currents decreased concentration-dependently after administration of carvedilol (Figure 1A–D) and the derived  $IC_{50}$  value is shown in Table 1, with a Hill coefficient,  $n_H$ , of  $1.6 \pm 0.1$  ( $n = 12$  to 15 cells were studied at each concentration). Blockade of K<sub>2P</sub>2.1 channels in oocytes was incomplete with maximum current reduction of  $80.3 \pm 2.2\%$  by 300 µM (20 min) carvedilol (Figure 1C), similar to dronedarone inhibition of K<sub>2P</sub>2.1 reported earlier (Schmidt *et al.*, 2012). The onset of block was rapid (Figure 1D) and 100 µM carvedilol inhibited K<sub>2P</sub>2.1 currents by  $76.3 \pm 1.9\%$  ( $n = 6$ ;  $P = 0.001$ ). Inhibitory effects were partially reversible, as current levels reached  $50.0 \pm 2.7\%$  of controls, 20 min after removal of the drug. We cannot exclude a minor contribution of the bath solution exchange (usually requiring ~0.5–1 min) to the kinetics of block and unblock shown in Figure 1D. Mean voltages required to achieve 50% of maximum K<sub>2P</sub> current amplitudes recorded at +60 mV ( $V_{50\%,K2P}$ ) yielded  $23.1 \pm 0.5$  mV under control conditions and  $22.8 \pm 0.4$  mV after application of 100 µM carvedilol ( $n = 15$ ;  $P = 0.36$ ; Figure 1E,F). We observed an apparent increase of relative (but not absolute) inward currents compared with outward currents in the presence of carvedilol (Figure 1E,F). As low levels of endogenous currents (at hyperpolarizing pulses) are not blocked by carvedilol, they are expected to become more apparent in relation to significantly blocked K<sub>2P</sub>2.1 currents at more depolarized potentials. Similar findings were obtained with K<sub>2P</sub>10.1 channels.

K<sub>2P</sub>2.1 currents show outward (or open) rectification. Linear ramp voltage protocols were applied between –140 and +60 mV (500 ms) before and after application of 100 µM carvedilol (20 min) to assess drug effects on current rectification (Figure 2A). The currents displayed outward rectification before and after carvedilol administration. The degree of blockade at +20 mV ramp potential was  $80.5 \pm 1.3\%$  ( $n = 4$ ;



**Figure 1**

Inhibition of human K<sub>2P</sub>.1 (TREK1) channels by carvedilol in *Xenopus laevis* oocytes. (A, B) Representative macroscopic currents recorded under control conditions and after application of 100  $\mu$ M carvedilol. Zero current levels are indicated by dashed lines. (C) Concentration-response relationships for the effect of carvedilol on K<sub>2P</sub>.1 currents measured at the end of the 0 mV voltage step ( $n = 12$  to 15 cells). (D) Time course of hK<sub>2P</sub>.1 current blockade by 100  $\mu$ M carvedilol ( $n = 6$ ). Panels E and F display  $I$ - $V$  relationships (i.e. mean current amplitudes as function of the test pulse potential) recorded under isochronal conditions (E, original current amplitudes; F, values normalized to maximum currents;  $n = 15$ ).

$P = 0.001$ ). Next, carvedilol block was investigated using extended voltage pulse durations. K<sub>2P</sub>.1 currents were recorded during a single depolarizing step to +20 mV for 7.5 s under control conditions and after application of 100  $\mu$ M carvedilol for 20 min while keeping the membrane potential at -80 mV (Figure 2B). Current inhibition during the first 400 ms of the protocol is displayed with linear and logarithmic time scales in Figure 2C and D. Analysis of channel block after carvedilol administration revealed that significant inhibition of hK<sub>2P</sub>.1 channels had occurred at -80 mV, indicated by the level of block at the beginning of the depolarizing pulse ( $60.2 \pm 2.3\%$ ;  $n = 8$ ;  $P < 0.0001$ ) (Figure 2C,D). A second, time-dependent component of block was observed during the +20 mV-pulse (block at 400 ms:  $72.2 \pm 1.0\%$ ;  $n = 8$ ;  $P < 0.0001$ ), similar to ~76% inhibition observed above with 500 ms-voltage steps.

To study frequency-dependence of block, human K<sub>2P</sub>.1 channels were rapidly activated by a depolarizing step to +20 mV (500 ms) at intervals of 2 or 10 s, respectively. Current reduction by 30  $\mu$ M carvedilol was plotted versus

time (Figure 2E). The degree of inhibition after 20 min was not significantly different ( $P = 0.079$ ) between 0.5 Hz ( $53.0 \pm 0.9\%$ ;  $n = 7$ ;  $P = 0.002$ ) and 0.1 Hz stimulation rate ( $58.2 \pm 2.6\%$ ;  $n = 7$ ;  $P = 0.001$ ). Finally, the specificity of K<sub>2P</sub>.1 current block was investigated. Compared with carvedilol, K<sub>2P</sub>.1 displayed reduced affinity to the selective  $\beta$ -adrenoceptor antagonist metoprolol (100  $\mu$ M; 20 min) and to the non-selective  $\beta_1$ - and  $\beta_2$ - adrenoceptor blocker propranolol (100  $\mu$ M; 20 min) under experimental conditions similar to those described above (Figure 1). Metoprolol reduced hK<sub>2P</sub>.1 currents by  $18.8 \pm 2.5\%$  ( $n = 13$ ;  $P = 0.0003$ ) and propranolol blocked the channels by  $31.2 \pm 3.7\%$  ( $n = 14$ ;  $P = 0.001$ ) (Figure 2F).

### Carvedilol targets hK<sub>2P</sub>10.1 (TREK2) channels

To further elucidate the pharmacological profile of carvedilol, its effects on human K<sub>2P</sub>10.1 channels were analysed in oocytes. Concentration-dependent inhibition of K<sub>2P</sub>10.1 current was revealed using the experimental approach described above (Figure 3A-D) and the derived  $IC_{50}$  value

**Table 1**

Sensitivity to carvedilol of wild-type  $K_{2P}$  channels and of subunits generated by alternative mRNA translation initiation

Subunit	IC <sub>50, oocytes</sub> (mean ± SEM)	Relative drug sensitivity (full-length channel/ truncated subunit)	IC <sub>50, HEK</sub> (mean ± SEM)
hK <sub>2P</sub> 2.1			
hK <sub>2P</sub> 2.1 WT	20.3 ± 1.5 μM	–	1.6 ± 0.9 μM
hK <sub>2P</sub> 2.1 M57I	16.5 ± 3.7 μM	–	NI
hK <sub>2P</sub> 2.1 Δ1–56	79.0 ± 33.8 μM	4.8	NI
hK <sub>2P</sub> 10.1			
hK <sub>2P</sub> 10.1 WT	24.0 ± 11.1 μM	–	7.6 ± 0.2 μM
hK <sub>2P</sub> 10.1 M60I M72I	17.5 ± 4.3 μM	–	NI
hK <sub>2P</sub> 10.1 Δ1–59 M72I	45.5 ± 34.2 μM	2.6	NI
hK <sub>2P</sub> 10.1 Δ1–71	87.0 ± 66.3 μM	5.0	NI

NI, not investigated; WT, wild type.

(Figure 3C), with  $n_H$  of  $2.2 \pm 1.0$ , from a series of 6–7 cells is shown in Table 1. Currents were stable during a 20-min control period ( $I_{0 \text{ min}}$ ,  $1.73 \pm 0.23 \mu\text{A}$ ;  $I_{20 \text{ min}}$ ,  $1.65 \pm 0.2 \mu\text{A}$ ;  $P = 0.12$ ). During carvedilol application (100 μM),  $K_{2P}10.1$  currents decreased by  $67.5 \pm 3.3\%$  ( $n = 7$ ;  $P = 0.0007$ ) (Figure 3D). Current inhibition was reduced to  $40.5 \pm 10.0\%$  during washout, indicating partial reversibility of block. Analysis of  $K_{2P}10.1$  current-voltage ( $I$ - $V$ ) relationships yielded a negligible change of 1.1 mV towards more negative potentials in the presence of carvedilol ( $V_{50\%, K_{2P}, \text{control}} = 25.5 \pm 1.2 \text{ mV}$ ;  $V_{50\%, K_{2P}, \text{carvedilol}} = 24.4 \pm 1.0 \text{ mV}$ ;  $n = 7$ ;  $P = 0.02$ ) (Figure 3E,F).

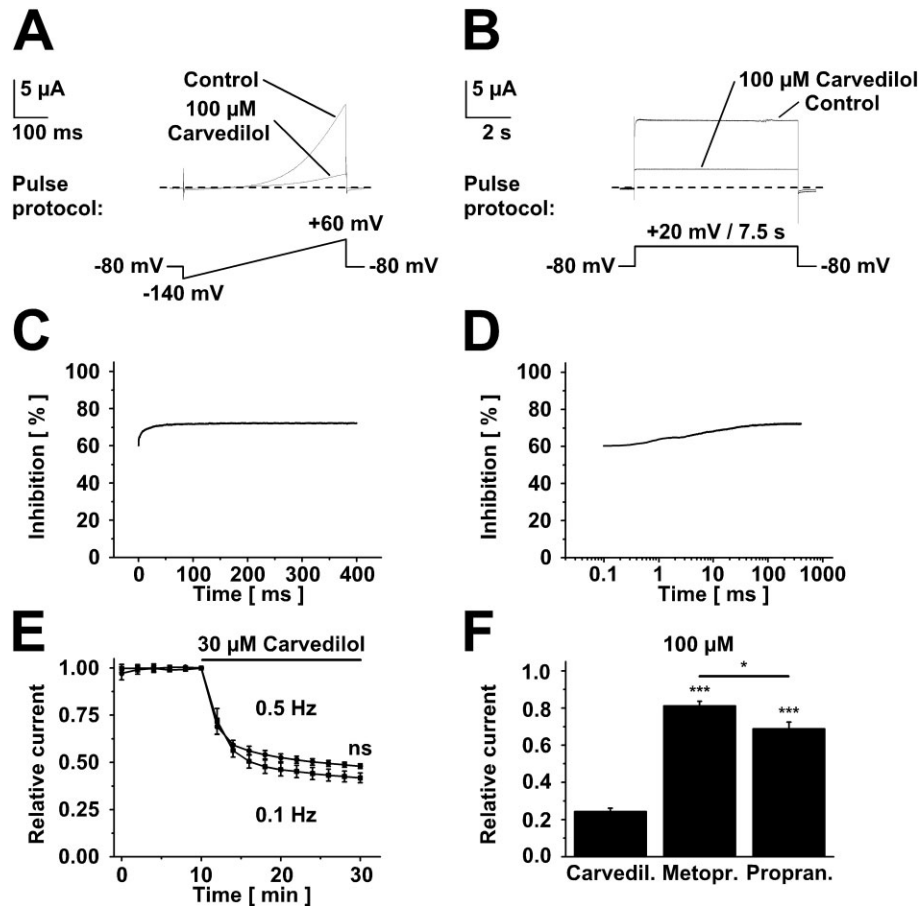
### Concentration-dependent blockade of hK<sub>2P</sub>2.1 and hK<sub>2P</sub>10.1 channels in mammalian cells

To evaluate the physiological significance of drug-receptor interactions, concentration-response relationships obtained from mammalian expression systems are required.  $K_{2P}2.1$  and  $K_{2P}10.1$  potassium channels were expressed in HEK 293 cells to analyse channel sensitivities to carvedilol (Figure 4). From a holding potential of  $-80 \text{ mV}$ , pulses were applied for 500 ms to voltages between  $-120$  and  $+80 \text{ mV}$  in 20 mV increments (0.2 Hz). Following current equilibration (10 min), the degree of inhibition was determined at  $+60 \text{ mV}$  10 min after carvedilol application (Figure 4A,B,D,E). For HEK 293 cells, the IC<sub>50</sub> values for blockade of hK<sub>2P</sub>2.1 channels (Hill coefficient  $n_H = 0.61 \pm 0.20$ ; Figure 4C) and for hK<sub>2P</sub>10.1 background channels (Hill coefficient  $n_H = 1.26 \pm 0.06$ ; Figure 4F) are given in Table 1. Four to eight cells were investigated at each concentration. Steady-state outward currents recorded in the presence of 100 μM carvedilol were blocked by  $89.4 \pm 4.0\%$  (hK<sub>2P</sub>2.1;  $n = 4$ ;  $P = 0.17$ ) and by  $96.6 \pm 1.5\%$  (hK<sub>2P</sub>10.1;  $n = 4$ ;  $P = 0.04$ ).

### Alternative translation initiation regulates $K_{2P}$ channel sensitivity to carvedilol

Alternative translation initiation generates two ( $K_{2P}2.1$ ) and three ( $K_{2P}10.1$ ) functionally different  $K_{2P}$  ion channel subunits respectively (Figure 5A–E) (Simkin *et al.*, 2008; Thomas *et al.*,

2008; Staudacher *et al.*, 2011a). Immunoblot analysis of tissue-specific total protein samples revealed significant  $K_{2P}2.1$  protein expression in adult human heart in addition to brain, aorta, skeletal muscle, stomach, small intestine, uterus and prostate (Figure 5C). Furthermore, the presence of a smaller  $K_{2P}2.1$  band (which may correspond to truncated  $K_{2P}2.1$  channels produced by ATI) was detected in the heart and other tissues (Figure 5C).  $K_{2P}10.1$  protein analysis was not performed as reliable cardiac immunoblot protocols and antibodies have not yet been established. To study differential pharmacological effects of carvedilol on  $K_{2P}$  isoforms, mutant subunits that specifically produce full-length channels by elimination of downstream start codons were used ( $K_{2P}2.1$  M57I;  $K_{2P}10.1$  M60I M72I). Drug effects on these channels were compared with truncated subunits associated with ATI ( $K_{2P}2.1$  Δ1–56;  $K_{2P}10.1$  Δ1–59 M72I;  $K_{2P}10.1$  Δ1–71). Figures 5B and E provide an overview of the channels investigated. Using the experimental approach described in Figures 1 and 3, marked differences in drug affinity were detected. Carvedilol (100 μM) reduced full-length hK<sub>2P</sub>2.1 M57I currents by  $72.6 \pm 3.3\%$  ( $n = 10$ ;  $P = 0.0002$ ; Figure 6A,B) compared with  $40.7 \pm 3.7\%$  inhibition of truncated hK<sub>2P</sub>2.1 Δ1–56 channels ( $n = 10$ ;  $P = 0.0001$ ; Figure 6F,G). The corresponding IC<sub>50</sub> values for blockade of hK<sub>2P</sub>2.1 M57I channels (Hill coefficient  $n_H = 1.4 \pm 0.5$ ; Figure 6C) and of hK<sub>2P</sub>2.1 Δ1–56 channels (Hill coefficient  $n_H = 1.5 \pm 0.6$ ; Figure 6H) are shown in Table 1. Four to 10 cells were investigated at each concentration. Carvedilol inhibition of full-length and truncated  $K_{2P}2.1$  variants ( $K_{2P}2.1$  M57I and  $K_{2P}2.1$  Δ1–56) was statistically compared. The resulting  $P$ -values (Table 2) illustrate statistically significant differences in the degree of blockade at concentrations  $>1 \mu\text{M}$ , providing statistical support for different drug affinities between full-length and truncated  $K_{2P}2.1$  channel variants. Please note that significant differences at 0.1 and 1 μM carvedilol are not expected as there was virtually no inhibition of either subunit at these concentrations (Figure 6C,H). Activation voltages of both  $K_{2P}2.1$  isoforms were virtually unchanged in the presence of 100 μM carvedilol (M57I;  $V_{50\%, K_{2P}, \text{control}} = 28.1 \pm 0.3 \text{ mV}$ ;  $V_{50\%, K_{2P}, \text{carvedilol}} = 26.0$



**Figure 2**

Biophysical characteristics of hK<sub>2P</sub>2.1 blockade by carvedilol in oocytes. (A) Open rectification of K<sub>2P</sub>2.1 currents evoked by voltage ramps from -140 to +60 mV. Typical recordings in the absence of the drug and after superfusion with 100 μM carvedilol (20 min) are superimposed. (B–D) Blockade during extended voltage pulses. Currents activated by 7.5 s depolarizing voltage steps to +20 mV are shown under control conditions and after administration of 100 μM carvedilol (20 min). (C, D) The extent of current inhibition during the first 400 ms of the depolarizing voltage step is displayed as % (C, linear time scale; D, logarithmic time scale). (E) Carvedilol block of K<sub>2P</sub>2.1 is frequency-independent. Relative current amplitudes recorded at +20 mV membrane potential during carvedilol application (0.5 and 0.1 Hz stimulation rates) are plotted versus time ( $n = 7$  oocytes were studied at each rate). For the purpose of clear presentation, not all measurements are displayed. (F) Effects of metoprolol and propranolol on K<sub>2P</sub>2.1 channels. Mean relative current amplitudes after administration of 100 μM carvedilol (Carvedil;  $n = 15$ ), 100 μM metoprolol (Metopr;  $n = 13$ ) and 100 μM propranolol (Propan;  $n = 14$ ) are displayed. K<sub>2P</sub> current inhibition by carvedilol was significantly greater than those induced by metoprolol and propranolol ( $*P < 0.05$  vs. metoprolol;  $***P < 0.001$  vs. carvedilol). Dashed lines indicate zero current levels.

**Table 2**

Statistical comparisons of relative current inhibition by a range of carvedilol concentrations between full-length (K<sub>2P</sub>2.1 M57I) and truncated (K<sub>2P</sub>2.1 Δ1–56) hK<sub>2P</sub>2.1 channel variants

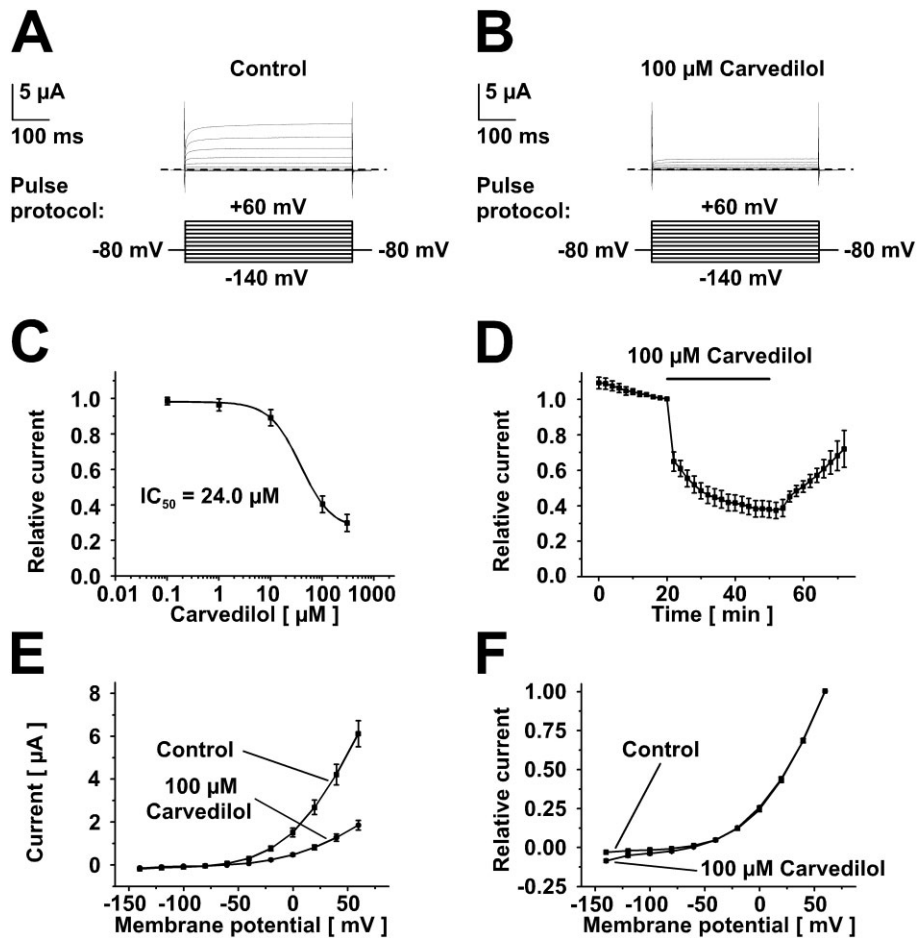
	0.1 μM	1 μM	10 μM	100 μM	300 μM
<i>P</i> -value ( $n$ , hK <sub>2P</sub> 2.1 M57I; $n$ , hK <sub>2P</sub> 2.1 Δ1–56)	0.19 <sup>ns</sup> (10; 8)	0.21 <sup>ns</sup> (10; 6)	0.002 <sup>**</sup> (9; 8)	<0.0001 <sup>***</sup> (10; 10)	0.014 <sup>*</sup> (10; 4)

\* $P < 0.05$ ; \*\* $P < 0.01$ ; \*\*\* $P < 0.001$ ; ns, not statistically significant.

$\pm 0.3$  mV;  $n = 10$ ;  $P < 0.001$ ; Δ1–56;  $V_{50\%,K2P,control} = 31.2 \pm 0.4$  mV;  $V_{50\%,K2P,carvedilol} = 29.7 \pm 0.7$  mV;  $n = 10$ ;  $P = 0.058$ ).

ATI-derived K<sub>2P</sub>10.1 channel isoforms were differentially regulated by carvedilol as well (Figure 7). Representative current traces are shown for control conditions

(Figure 7A,F,K) and after application of 100 μM carvedilol (Figure 7B,G,L) for indicated subunits. Currents were blocked by  $69.7 \pm 3.5\%$  (K<sub>2P</sub>10.1 M60I M72I;  $n = 7$ ;  $P = 0.001$ ), by  $45.3 \pm 4.6\%$  (K<sub>2P</sub>10.1 Δ1–59 M72I;  $n = 8$ ;  $P = 0.0002$ ) and by  $26.5 \pm 1.1\%$  (K<sub>2P</sub>10.1 Δ1–71;  $n = 7$ ;  $P = 0.002$ ) respectively. The IC<sub>50</sub>



**Figure 3**

Human  $K_{2P}10.1$  (TREK2) channels in oocytes are sensitive to carvedilol. Analyses performed as described in Figure 1 illustrate concentration-dependent  $K_{2P}10.1$  current inhibition (A–C;  $n = 6–7$  cells). Zero current levels are indicated by dashed lines. (D) Time course of blockade by 100  $\mu\text{M}$  carvedilol ( $n = 7$ ). Open rectification (E, F) and current-voltage relationships of  $K_{2P}10.1$  channels were not markedly affected by carvedilol (100  $\mu\text{M}$ ;  $n = 7$ ).

**Table 3**

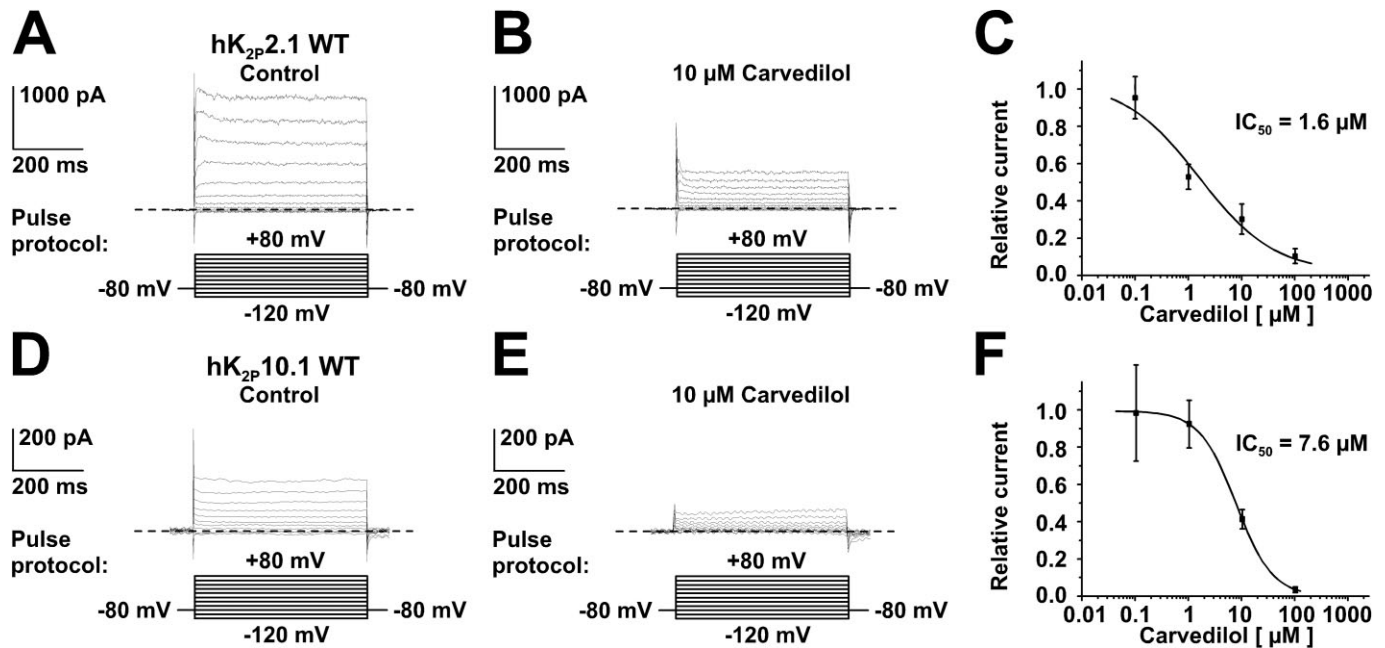
Statistical comparisons of relative current inhibition by indicated carvedilol concentrations between full-length (h $K_{2P}10.1$  M60I M72I) and truncated (h $K_{2P}10.1$   $\Delta 1–59$  M72I, h $K_{2P}10.1$   $\Delta 1–71$ ) h $K_{2P}10.1$  channel variants

	0.1 $\mu\text{M}$	1 $\mu\text{M}$	10 $\mu\text{M}$	100 $\mu\text{M}$	300 $\mu\text{M}$
$P$ -value ( $n$ , h $K_{2P}10.1$ M60I M72I; $n$ , h $K_{2P}10.1$ $\Delta 1–59$ M72I)	0.013* (7;6)	0.032 <sup>ns</sup> (6;7)	0.001** (6;6)	0.001** (7;8)	0.086 <sup>ns</sup> (7;7)
$P$ -value ( $n$ , h $K_{2P}10.1$ M60I M72I; $n$ , h $K_{2P}10.1$ $\Delta 1–71$ )	0.87 <sup>ns</sup> (7;6)	0.20 <sup>ns</sup> (6;7)	0.001** (6;8)	<0.0001*** (7;7)	<0.0001*** (7;7)
$P$ -value ( $n$ , h $K_{2P}10.1$ $\Delta 1–59$ M72I; $n$ , h $K_{2P}10.1$ $\Delta 1–71$ )	0.18 <sup>ns</sup> (6;6)	0.26 <sup>ns</sup> (7;7)	0.60 <sup>ns</sup> (6;8)	0.002** (8;7)	0.006* (7;7)

\* $P < 0.05$  (adjusted to 0.05/3 = 0.017); \*\* $P < 0.01$  (adjusted to 0.01/3 = 0.003); \*\*\* $P < 0.001$  (adjusted to 0.001/3 = 0.0003); ns, not statistically significant; ANOVA with Bonferroni adjustment.

values for block of h $K_{2P}10.1$  M60I M72I ( $n_H = 1.3 \pm 0.4$ ;  $n = 6–7$ ) and of h $K_{2P}10.1$   $\Delta 1–59$  M72I ( $n_H = 0.97 \pm 0.3$ ;  $n = 6–8$ ) and of h $K_{2P}10.1$   $\Delta 1–71$  ( $n_H = 0.8 \pm 0.3$ ;  $n = 6–8$ ) channels are shown in Table 1. Statistically significant differences between inhi-

tion of full-length channels ( $K_{2P}10.1$  M60I M72I) and both truncated variants ( $K_{2P}10.1$   $\Delta 1–59$  M72I,  $K_{2P}10.1$   $\Delta 1–71$ ) were observed at 100  $\mu\text{M}$  carvedilol (Table 3), confirming different drug sensitivity at concentrations close to the respective  $IC_{50}$



**Figure 4**

Inhibition of hK<sub>2P</sub>.1 and hK<sub>2P</sub>.10.1 currents expressed in HEK293 cells. Representative K<sub>2P</sub>.1 and K<sub>2P</sub>.10.1 measurements before (A, D) and after exposure to 10 μM carvedilol (B, E) are displayed. (C, F) Concentration-response relationships indicating IC<sub>50</sub> values of respective K<sub>2P</sub> channels ( $n = 4-8$  cells).

values. *I-V* relationships of all K<sub>2P</sub>.10.1 variants tested were not markedly modified by 100 μM carvedilol (K<sub>2P</sub>.10.1 M60I M72I;  $V_{50\%,K2P,control} = 25.9 \pm 0.6$  mV;  $V_{50\%,K2P,carvedilol} = 25.0 \pm 0.5$  mV;  $n = 7$ ;  $P = 0.04$ ; K<sub>2P</sub>.10.1 Δ1-59 M72I;  $V_{50\%,K2P,control} = 30.3 \pm 0.4$  mV;  $V_{50\%,K2P,carvedilol} = 27.8 \pm 0.4$  mV;  $n = 8$ ;  $P = 0.001$ ; K<sub>2P</sub>.10.1 Δ1-71;  $V_{50\%,K2P,control} = 30.7 \pm 0.4$  mV;  $V_{50\%,K2P,carvedilol} = 29.3 \pm 0.5$  mV;  $n = 7$ ;  $P = 0.005$ ; Figure 7D,I,N).

## Discussion

### Carvedilol inhibits hK<sub>2P</sub>.1 (TREK1) and hK<sub>2P</sub>.10.1 (TREK2) K<sub>2P</sub> channels

We report concentration-dependent inhibition of hK<sub>2P</sub>.1 and hK<sub>2P</sub>.10.1 currents by carvedilol with IC<sub>50</sub> values of 20.3 and 24.0 μM obtained from *Xenopus laevis* oocytes (Table 1). An extension of these findings to *I*<sub>TREK1</sub> and *I*<sub>TREK2</sub> recorded from mammalian HEK 293 cells revealed IC<sub>50</sub> values of 1.6 and 7.6 μM respectively. Differences in IC<sub>50</sub> values between mammalian cells and oocytes are commonly observed and attributed to specific properties of oocytes (e.g. the vitelline membrane and the yolk) that reduce drug affinities. Different IC<sub>50,oocytes</sub>/IC<sub>50,mammalian cells</sub> ratios are primarily determined by physicochemical drug characteristics. Here, these ratios yield 12.7 (hK<sub>2P</sub>.1) and 3.2 (hK<sub>2P</sub>.10.1) for the same drug, carvedilol, suggesting that channel-specific mechanisms of drug access to the putative receptor site within the pore are affected by oocyte properties as well. During therapeutic application of the drug, total plasma levels between 47 and 615 nM have been measured (McPhillips *et al.*, 1988; Morgan *et al.*, 1990; Gehr *et al.*, 1999; Karle *et al.*, 2001). Carvedilol is

approximately 98% bound to plasma proteins, reducing effective free drug concentrations to 0.94–12.3 nM, which would suggest a minimal effect of K<sub>2P</sub> channel blockade on cardiac electrophysiology under normal pharmacokinetic conditions.

### Biophysical properties of hK<sub>2P</sub>.1 blockade by carvedilol

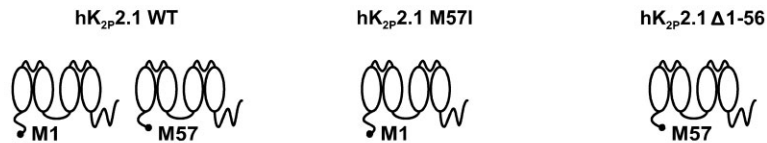
The biophysical mechanism of K<sub>2P</sub>.1 inhibition by carvedilol was analysed in detail. A rapid onset of block supports a direct drug-channel interaction as opposed to increased protein turnover or accelerated channel degradation as a molecular mechanism of action. In addition, carvedilol might indirectly affect K<sub>2P</sub>.1 or K<sub>2P</sub>.10.1 function through inhibition of β-adrenoreceptor signalling in oocytes or HEK 293 cells. However, this mechanism appears unlikely as activation of the cAMP/PKA system caused K<sub>2P</sub>.1 and K<sub>2P</sub>.10.1 current reduction, rather than enhancement (Gu *et al.*, 2002; Honoré *et al.*, 2002; Murbartian *et al.*, 2005). Furthermore, functional data suggest that levels of endogenous β-adrenoreceptors are negligible (Kathöfer *et al.*, 2000). The lack of complete K<sub>2P</sub>.1 and K<sub>2P</sub>.10.1 current inhibition in oocytes compared with mammalian cells may be attributed to specific lipophilic properties of the oocyte expression system. Open rectification that is characteristic to K<sub>2P</sub> channel function in physiological ionic conditions was observed before and during drug block. Finally, the lack of frequency-dependence of K<sub>2P</sub>.1 channel blockade supports continuous accessibility of the putative drug binding site inside the channel. It is likely that kinetics of drug block (Figure 2) were largely independent of channel opening as the cytosolic channel gate is constitutively open



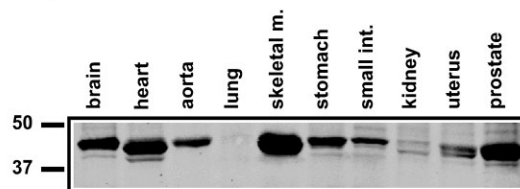
### A hK<sub>2p</sub>2.1 amino acid alignment

	M1		M57	
hK <sub>2p</sub> 2.1 WT	↓	MLPSASRERPGYRAGVAAPDLLDPKSAAQNSKPRLSFSTKPTVLASRVESDTTINVMKWK	↓	60
hK <sub>2p</sub> 2.1 M57I		MLPSASRERPGYRAGVAAPDLLDPKSAAQNSKPRLSFSTKPTVLASRVESDTTINVIKWK		60
hK <sub>2p</sub> 2.1 Δ1-56		-----MKWK		60

### B Protein subunits



### C K<sub>2p</sub>2.1 protein expression in humans



### D hK<sub>2p</sub>10.1 amino acid alignment

	M1		M60	M72	
hK <sub>2p</sub> 10.1 WT	↓	MKFFIETPRKQVNWDPKVAVPAAAPVCQPKSATNGQPPAPAPTPTPRLSISRATVVARMEGTSQGGGLQTVMKWK	↓	↓	75
hK <sub>2p</sub> 10.1 M60I M72I		MKFFIETPRKQVNWDPKVAVPAAAPVCQPKSATNGQPPAPAPTPTPRLSISRATVVARIEGTSQGGGLQTVIKWK			75
hK <sub>2p</sub> 10.1 Δ1-59 M72I		-----MEGTSQGGGLQTVIKWK			75
hK <sub>2p</sub> 10.1 Δ1-71		-----MKWK			75

### E Protein subunits



**Figure 5**

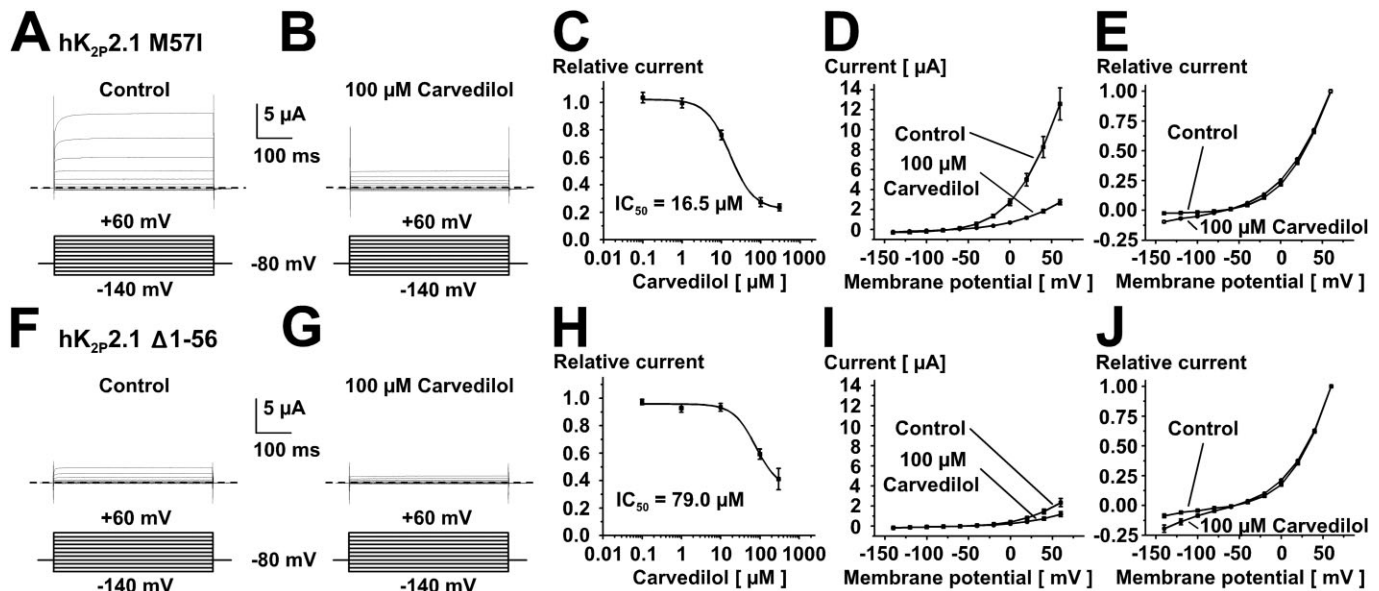
Alternative translation initiation of hK<sub>2p</sub>2.1 and hK<sub>2p</sub>10.1 mRNA produces multiple channel isoforms. (A, D) N terminal amino acid sequence alignments of wild-type (WT) K<sub>2p</sub>2.1 (A) and K<sub>2p</sub>10.1 channels (D). Alternative translation initiation sites are indicated by arrows. In addition, alignments of mutated channels used to specifically produce full-length or truncated K<sub>2p</sub> proteins in isolation are shown. (B, E) The topologies of K<sub>2p</sub>2.1 (B) and K<sub>2p</sub>10.1 (E) WT channels (left) and of protein variants arising from alternative mRNA translation initiation studied in this work (right) are displayed. (C) Expression of two K<sub>2p</sub>2.1 protein isoforms in indicated human tissues (skeletal m., skeletal muscle; small int., small intestine) visualized by Western blot analysis using an antibody to the K<sub>2p</sub>2.1 C terminus. Markers correspond to apparent molecular weights: 50, 50 kDa; 37, 37 kDa.

(Piechotta *et al.*, 2011; Brohawn *et al.*, 2012; Miller and Long, 2012), allowing for drug binding within the channel pore in state-independent manner.

### ATI determines drug sensitivity of K<sub>2p</sub> channels

This work identifies ATI as a genetic mechanism regulating sensitivity of K<sub>2p</sub>2.1 and K<sub>2p</sub>10.1 channels to carvedilol. Com-

pared with full-length K<sub>2p</sub> channels, truncated isoforms produced by ATI were 2.6 to 5.0-fold less sensitive to the drug (Table 1). These findings are in line with a previous report describing decreased sensitivity (by 70%) of truncated K<sub>2p</sub>2.1 channels to the antidepressant drug fluoxetine (Eckert *et al.*, 2011). The present study extends regulation of K<sup>+</sup> channel drug sensitivity by ATI to cardiac K<sub>2p</sub> channels and to a cardiovascular drug, suggesting a broader role of this



**Figure 6**

Human K<sub>2P</sub>2.1 variants produced by ATI exhibit different carvedilol affinity. Channels were studied in *Xenopus* oocytes using indicated voltage protocols. (A, B, F, G) Representative current recordings under control conditions (A, F) and after 20 min perfusion with 100 μM carvedilol (B, G) are illustrated for indicated isoforms. Concentration-response relationships show a 4.8-fold higher IC<sub>50</sub> value for K<sub>2P</sub>2.1 Δ1-56 (H) compared with K<sub>2P</sub>2.1 M57I (C) ( $n = 6$  to 10 cells analysed at each concentration). (D, E, I, J) Voltage-dependence of activation before and after carvedilol application (D, I, original current amplitudes; E, J, values normalized to maximum currents;  $n = 10$ ). Dashed lines indicate zero current level.

pharmacological mechanism. Several studies have identified the C-terminus of K<sub>2P</sub>2.1 channels as a critical region for polymodal channel regulation (Honoré, 2007). Reduced carvedilol affinity observed in N-terminally truncated isoforms now indicates a significant contribution of the N-terminus of K<sub>2P</sub>2.1 and K<sub>2P</sub>10.1 channels. Truncated K<sub>2P</sub>2.1 ATI variants may exhibit reduced carvedilol sensitivity, because of the additional Na<sup>+</sup> conductance of the channel pore (Thomas *et al.*, 2008). However, this mechanism could not explain reduced sensitivity among K<sub>2P</sub>10.1 ATI variants, as loss of K<sup>+</sup> selectivity has not been demonstrated for this channel.

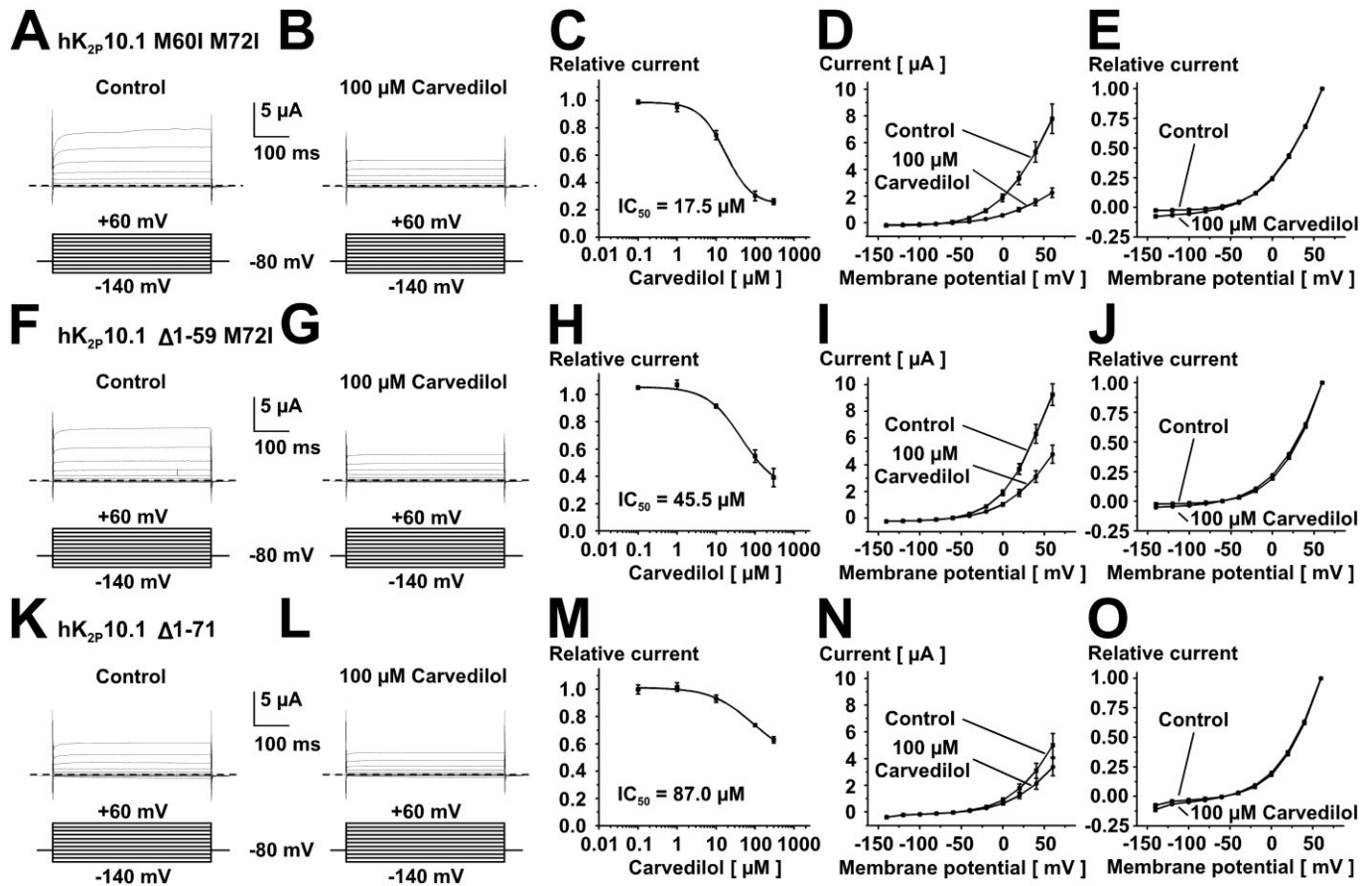
The apparent role of the N-terminus in K<sub>2P</sub> channel regulation is in line with data obtained from human ether-a-go-go-related gene (hERG) K<sup>+</sup> channels. The length of the hERG channel N-terminus is determined by mRNA splicing (as opposed to ATI-dependent regulation of the K<sub>2P</sub>2.1 N-terminus reported in the present work). The transcript variant hERG 1b is characterized by a shortened N-terminus and displays reduced affinity to the class III compounds dofetilide and E-4031 (Abi-Gerges *et al.*, 2011). It is reasonable to assume that reduced length of the N-terminus modulates drug sensitivity through alterations of the channel pore structure that contains the putative binding site. As the N-terminus is not detailed in the crystal structures of K<sub>2P</sub>1.1 and mechanosensitive K<sub>2P</sub>4.1 channels, the precise mechanism by which the N-terminus regulates K<sub>2P</sub> channel pharmacology remains to be elucidated (Brohawn *et al.*, 2012; Miller and Long, 2012).

### Clinical implications

Class III antiarrhythmic drugs suppress atrial fibrillation and ventricular tachyarrhythmia through K<sup>+</sup> channel block,

resulting in prevention of electrical reentry via prolongation of action potential duration and cardiac refractoriness. Outward potassium currents mediated by K<sub>2P</sub> channels contribute to cardiac repolarization. Thus, inhibition of cardiac K<sub>2P</sub> channels represents a novel concept in antiarrhythmic therapy, with carvedilol blockade of hK<sub>2P</sub>2.1 and hK<sub>2P</sub>10.1 K<sup>+</sup> currents observed in the current study and inhibition of hK<sub>2P</sub>3.1 channels reported previously (Staudacher *et al.*, 2011b) serving as proof of this principle. The specific contribution of K<sub>2P</sub> channel blockade to the antiarrhythmic efficacy of carvedilol in heart failure and atrial fibrillation patients (Senior *et al.*, 1992; Brunvand *et al.*, 1996; Cice *et al.*, 2000; Takusagawa *et al.*, 2000; Katritsis *et al.*, 2003; Merritt *et al.*, 2003; Ramaswamy, 2003; Haghjoo *et al.*, 2007; Acikel *et al.*, 2008), however, is expected to be low, as the IC<sub>50</sub> values determined *in vitro* in the low micromolar range were much higher than the nanomolar free therapeutic drug concentrations found in clinical practice. Compared with K<sub>2P</sub>2.1 (IC<sub>50</sub> = 1.6 μM) and K<sub>2P</sub>10.1 channels (IC<sub>50</sub> = 7.6 μM), higher sensitivity implying greater clinical effect was demonstrated for the K<sub>2P</sub>3.1 channels (IC<sub>50</sub> = 0.8 μM; Staudacher *et al.*, 2011b). The overall antiarrhythmic efficacy of the drug is likely to result primarily from its anti-adrenergic properties, supported by multichannel blocking effects on potassium currents ( $I_{Kr}$ ,  $I_{Ks}$ ,  $I_{to}$ ,  $I_{Kur}$ ,  $I_{TASK1}$ ,  $I_{TREK1}$ ,  $I_{TREK2}$ ,  $I_{K(ATP)}$ ), pacemaker current ( $I_h$ ), L-type calcium current and the cardiac ryanodine receptor with different affinities (Cheng *et al.*, 1999; Kawakami *et al.*, 2006; Kikuta *et al.*, 2006; Deng *et al.*, 2007; Yokoyama *et al.*, 2007; Staudacher *et al.*, 2011b; Zhou *et al.*, 2011).

In conclusion, this investigation showed inhibition of K<sub>2P</sub>2.1 and K<sub>2P</sub>10.1 channels by carvedilol, providing proof-of-concept to support a pharmacological role of cardiac K<sub>2P</sub>



**Figure 7**

ATI affects inhibition of hK<sub>2P</sub>10.1 channels by carvedilol. Representative macroscopic recordings from oocytes and activation curves reflect blockade of full-length (K<sub>2P</sub>10.1 M60I M72I; A, B, D, E) and of two truncated isoforms (K<sub>2P</sub>10.1 Δ1–59 M72I; F, G, I, J; K<sub>2P</sub>10.1 Δ1–71; K, L, N, O) by 100 μM carvedilol. Zero current levels are marked by dashed lines. Corresponding concentration-response curves demonstrate decreased carvedilol sensitivity of truncated isoforms (H, M) compared with full-length channels (C) (*n* = 7 to 8 cells were studied).

channels as antiarrhythmic drug targets. The therapeutic significance of K<sub>2P</sub> channel blockade in heart rhythm disorders requires validation in translational and clinical studies. At the molecular level, antiarrhythmic drug sensitivity is determined by alternative translation initiation of K<sub>2P</sub>2.1 and K<sub>2P</sub>10.1 mRNA that produces subunits with different carvedilol affinity. Differential expression of ATI-dependent K<sub>2P</sub> channel variants may result in spatiotemporal variation of electrophysiological carvedilol effects in the heart. In addition, the effect of ATI on antiarrhythmic drug efficacy has broad and general implications for cardiac drug discovery.

## Acknowledgements

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## Conflict of interest

None declared.

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