

ω -3 and ω -6 polyunsaturated fatty acids block *HERG* channels

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Guizy, Miriam, Cristina Arias, Miren David, Teresa González, and Carmen Valenzuela. ω -3 and ω -6 polyunsaturated fatty acids block *HERG* channels. *Am J Physiol Cell Physiol* 289: C1251–C1260, 2005. First published June 29, 2005; doi:10.1152/ajpcell.00036.2005.—Dietary polyunsaturated fatty acids (PUFAs) have been reported to exhibit antiarrhythmic properties, which have been attributed to their availability to modulate Na^+ , Ca^{2+} , and several K^+ channels. However, their effects on human *ether-a-go-go*-related gene (*HERG*) channels are unknown. In this study we have analyzed the effects of arachidonic acid (AA, ω -6) and docosahexaenoic acid (DHA, ω -3) on *HERG* channels stably expressed in Chinese hamster ovary cells by using the whole cell patch-clamp technique. At 10 μM , AA and DHA blocked *HERG* channels, at the end of 5-s pulses to -10 mV, to a similar extent ($37.7 \pm 2.4\%$ vs. $50.2 \pm 8.1\%$, $n = 7$ – 10 , $P > 0.05$). 5,6,11,14-Eicosatetrayenoic acid, a nonmetabolizable AA analog, induced effects similar to those of AA on *HERG* current. Both PUFAs shifted the midpoint of activation curves of *HERG* channels by -5.1 ± 1.8 mV ($n = 10$, $P < 0.05$) and -11.2 ± 1.1 mV ($n = 7$, $P < 0.01$). Also, AA and DHA shifted the midpoint of inactivation curves by $+12.0 \pm 3.9$ mV ($n = 4$; $P < 0.05$) and $+15.8 \pm 4.3$ mV ($n = 4$; $P < 0.05$), respectively. DHA and AA accelerated the deactivation kinetics and slowed the inactivation kinetics at potentials positive to $+40$ mV. Block induced by DHA, but not that produced by AA, was higher when measured after applying a pulse to -120 mV ($I \rightarrow O$). Finally, both AA and DHA induced a use-dependent inhibition of *HERG* channels. In summary, block induced by AA and DHA was time, voltage, and use dependent. The results obtained suggest that both PUFAs bind preferentially to the open state of the channel, although an interaction with inactivated *HERG* channels cannot be ruled out for AA.

K^+ channel; membrane currents; ion channels; arrhythmia; antiarrhythmics

POLYUNSATURATED FATTY ACIDS (PUFAs) present in nature belong to two main classes: ω -6 class, mostly present in vegetable oils; and ω -3 class, which comes mostly from fish. Both are “essential” because they are necessary for an optimal health status and cannot be synthesized *de novo*. In addition, mammals cannot convert ω -6 into ω -3 PUFAs. The dramatic increase in the ω -6/ ω -3 ratio in the diet of the population of Western countries after the industrial revolution has contributed to the rise in cardiovascular disease (7, 21). There is a growing body of evidence that dietary ω -3 PUFAs, especially docosahexaenoic acid (DHA), play an important role in the prevention of coronary heart disease, in decreasing the risk of sudden cardiac death, and, in particular, in preventing fatal ventricular arrhythmias (1, 5, 8, 10, 21, 37). The cardioprotective effects of ω -3 PUFAs have been attributed to their availability to modulate several ionic channels involved in the onset and maintenance of the cardiac action potential (21, 22).

PUFAs block Na^+ channels by binding to the receptor site that share local anesthetics and antiarrhythmic drugs (29, 32, 42). PUFAs also interact with other cardiac ion channels, inhibiting ultrarapid delayed-rectifier K^+ current ($I_{\text{K,ur}}$), transient outward current (I_{to}), and Ca^{2+} current (I_{Ca}) (14, 15, 22, 40) and enhancing long-activated delayed-rectifier K^+ current (I_{K1}) and slow-activated delayed-rectifier K^+ current (I_{Ks} ; the latter by their interaction with minK) (9, 24). Moreover, it has been demonstrated that membrane lipids can convert Kv A-type channels into delayed rectifiers and vice versa. Thus phosphoinositides remove N-type inactivation, whereas arachidonic acid (AA) converts Kv-delayed rectifiers into A-type rectifier channels (30). However, the effects of PUFAs on human *ether-a-go-go*-related gene (*HERG*) channels have not been studied yet. *HERG* channel activity determines the ventricular action potential duration and, therefore, the refractory period. Mutations in the gene encoding *HERG* channels are involved in the genotype of congenital long QT syndrome (36). Moreover, most drugs that induce acquired long QT syndrome block *HERG* channels (35), and thus, *HERG* channels can be considered a target of antiarrhythmic agents. The purpose of the present study is to analyze and compare the effects of ω -6 and ω -3 PUFAs (AA, precursor of the prostaglandins, leukotrienes, and thromboxane cascade; and DHA) on *HERG* channels, whose activity determines the action potential duration (23). A preliminary report of this study has been published in abstract form (12).

METHODS AND MATERIALS

Cell culture. Stably transfected Chinese hamster ovary cells with the gene encoding *HERG* channels (a gift of Drs. S. Nattel, T. E. Hébert, and W. Weerapura) were cultured at 37°C in Ham's F-12 medium supplemented with geneticine (600 $\mu\text{g}/\text{ml}$), penicillin-streptomycin (800 IU and 200 $\mu\text{g}/\text{ml}$, respectively), and 10% bovine serum, in a 5% CO_2 atmosphere (2, 11). Cultures were passaged every 3–5 days with the use of a brief trypsin treatment. Before experimental use, the cells were removed from the dish with a rubber policeman, a procedure that left the majority of the cells intact. The cell suspension was stored at room temperature (21 – 23°C) and used within 12 h for all the experiments reported.

Electrophysiological recording. The intracellular pipette-filling solution contained (in mM) 80 K-aspartate, 50 KCl, 3 phosphocreatine, 10 KH_2PO_4 , 3 MgATP, 10 HEPES-K, and 5 EGTA and was adjusted to pH 7.25 with KOH. The bath solution contained (in mM) 130 NaCl, 4 KCl, 1.8 CaCl_2 , 1 MgCl_2 , 10 HEPES-Na, and 10 glucose, and was adjusted to pH 7.4 with NaOH. AA, DHA, and 5,8,11,14-eicosatetrayenoic acid (ETYA) (Sigma, St. Louis, MO) were dissolved in ethanol at concentrations of 56.5, 52.5, and 10 mM, respectively. Experiments were performed to test the potential effects of ethanol

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(30 μM /100 ml) on *HERG* channels. Ethanol, at this concentration, did not modify the outward maximum current (56.3 ± 10.3 vs. 52.7 ± 11.2 pA; $n = 7$, $P > 0.05$) nor the maximum peak tail current (68.7 ± 16.4 to 63.6 ± 16.7 pA; $n = 7$, $P > 0.05$). AA and DHA were stored under argon atmosphere and maintained in sealed ampoules protected from light at -40°C to prevent oxidation. *HERG* currents were recorded at room temperature (21 – 23°C) using the whole cell patch-clamp technique (13) with an Axopatch 200A patch-clamp amplifier (Axon Instruments, Foster City, CA). Micropipettes were pulled from borosilicate glass capillary tubes (GD-1; Narishige, Tokyo, Japan) on a programmable horizontal puller (Sutter Instrument, San Rafael, CA) and heat polished with a microforge (Narishige). Micropipette resistance was 1–3 M Ω . Maximum *HERG* tail-current amplitudes averaged 717 ± 123 pA, mean uncompensated access resistance was 1.5 ± 0.5 M Ω , and cell capacitance 29.8 ± 2.0 pF ($n = 11$). Thus no significant voltage errors (<5 mV) were expected. *HERG* currents were filtered at 100 Hz and sampled at 200 Hz. Cells were held at -80 mV. After control data were obtained, bath perfusion was switched to fatty acid-containing solution. The effects of drug infusion were monitored with test pulses to -10 mV applied every 30 s until steady state was obtained. Steady-state current-voltage relationships (*I-V*) were obtained by averaging the current over a small window (2–5 ms) at the end of 5-s depolarizing pulses. Between -80 and -50 mV only passive linear leak was observed and least squares fit to these data were used for passive leak correction. Deactivating tail currents were recorded at -60 mV. The activation curves were obtained from the tail current amplitude measured at the maximum peak value. The inactivation curves were obtained from the maximum current amplitude measured at a test pulse at $+30$ mV applied after a two-pulse protocol that consisted of a 1-s depolarizing pulse from -80 mV to $+30$ mV followed by a second pulse of 20 ms in duration to different membrane potentials between -120 mV and $+10$ mV. Other pulse protocols are described in RESULTS. Command potentials and data acquisition were generated by using the CLAMPEX program of pCLAMP 6.0.1, 9.0.1 (Axon Instruments). Data analysis was performed using the CLAMPfit program of pCLAMP 9.0.1, Origin 7.0.3 (Microcal Software, Northampton, MA), and other custom-made analysis programs. Deactivation was fitted to a biexponential process

$$y = C + A_1 \exp(-t/\tau_1) + A_2 \exp(-t/\tau_2) \quad (1)$$

where τ_1 and τ_2 are the system time constants, A_1 and A_2 are the amplitudes of each component of the exponential, and C is the baseline value. Half-maximal voltages ($E_{1/2}$) and slope factors (s) of activation and inactivation were determined by fitting data with a Boltzmann equation $y = 1/[1 + \exp[-(E - E_{1/2})/s]]$. The curve-fitting procedure used a nonlinear least-squares (Gauss-Newton) algorithm; results were displayed in linear and semilogarithmic format, together with the difference plot. Goodness of fit was judged by the χ^2 criterion and by inspection for systematic nonrandom trends in the difference plot.

Statistical methods. Results are expressed as means \pm SE. Direct comparisons between mean values in control conditions and in the presence of drug for a single variable were performed by paired Student's *t*-test. Differences were considered significant if $P < 0.05$. Comparisons between the three groups were performed by a one-way ANOVA, with a posterior Newman-Keuls test if $P < 0.05$.

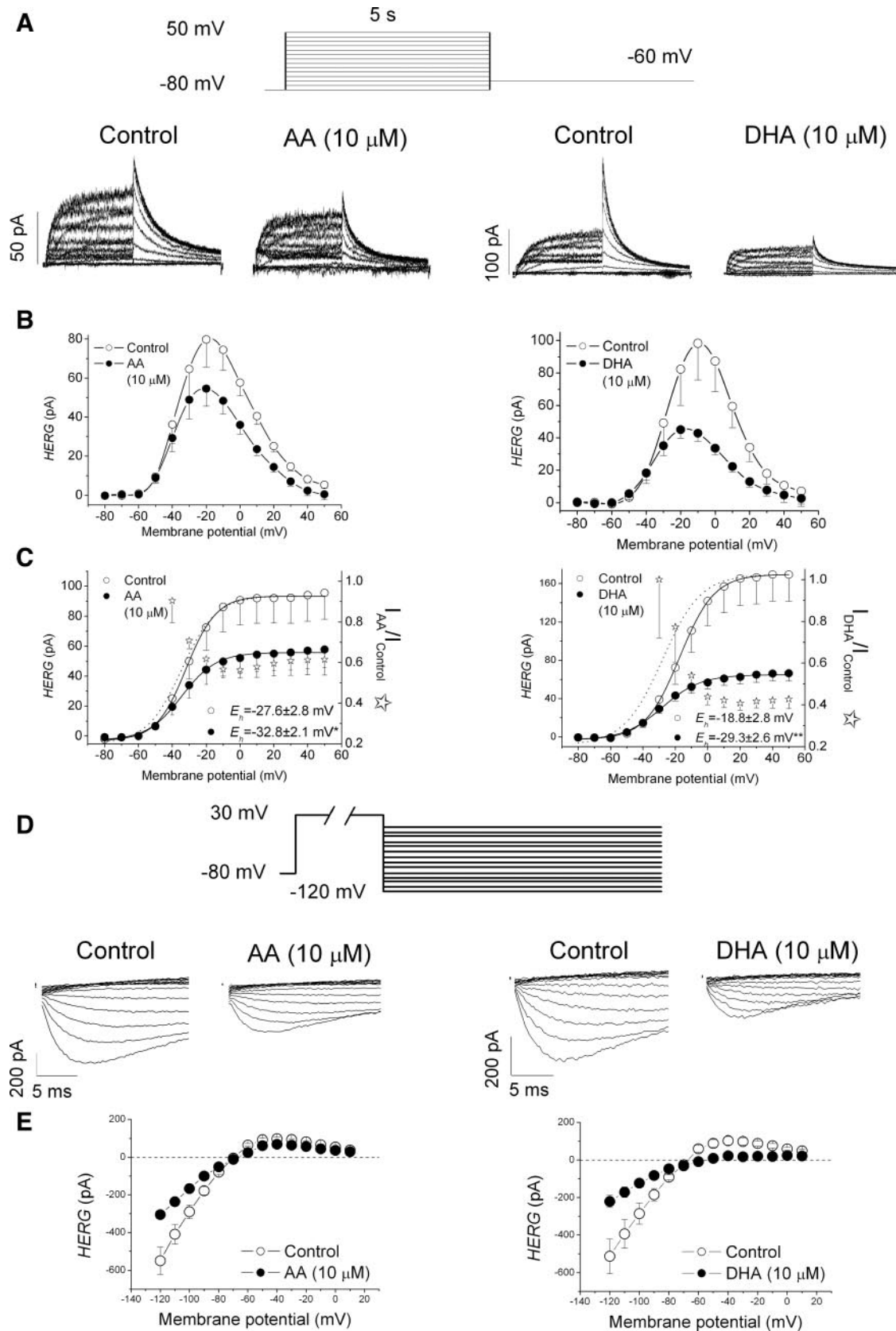
RESULTS

Free fatty acids are generally toxic to cells and are kept at low micromolar concentrations in plasma. These levels can vary greatly depending on the hormonal, metabolic, and nutritional state of the individual. Around 99.9% of free fatty acids are carried in plasma bound to albumin. The low plasma free fatty acid concentrations are maintained by the competition between the affinity of the albumin-binding sites and cell membrane phospholipids for free fatty acids. The range of free AA and DHA in human plasma is 5.3–13.1 μM and <2.8 μM , respectively (6). The choice of the concentration of DHA (10 μM) is based on the reported EC_{50} for the effects of PUFAs on ion channels that ranges between 1 and 30 μM (14, 43). This concentration was also used to show antiarrhythmic properties of this fatty acid in single cell (16–18). Following the same reasoning and for better comparisons between ω -3 and ω -6 PUFAs effects on *HERG* channels, the same AA concentration was used.

Voltage-dependent block of *HERG* channels induced by AA and DHA. Figure 1A shows *HERG* currents obtained after applying 5-s pulses from a holding potential of -80 to $+50$ mV in 10-mV steps in the absence and in the presence of 10 μM AA or DHA. Deactivating tail currents were measured at -60 mV. Figure 1B shows the *I-V* relationships obtained by plotting the amplitude of the *HERG* current measured at the end of the 5-s pulses vs. membrane potential in the absence and in the presence of AA (Fig. 1B, left) or DHA (Fig. 1B, right). Under control conditions, the *I-V* exhibits the characteristic bell shape that increases from -50 mV to -10 mV (-12.4 ± 1.8 mV, $n = 19$) and, due to the fast C-type inactivation of *HERG* channels, it decreased with further depolarizations (38, 39). Thus steady-state drug-induced block was measured at the end of 5-s pulses to -10 mV. Blockade induced by AA ($37.7 \pm 2.4\%$, $n = 10$) was similar to that produced by DHA ($50.2 \pm 8.1\%$, $n = 7$; $P > 0.05$). However, block induced by DHA was higher than that produced by AA measured at the maximum peak of the tail current recorded at -60 mV after applying a 5-s depolarizing test pulse to 0 mV ($47.4 \pm 4.2\%$, $n = 10$, vs. $58.3 \pm 2.7\%$, $n = 7$, in the presence of AA and DHA, respectively; $P < 0.05$). AA and DHA shifted the midpoint of activation curves without modifying the slope factors, being the voltage shift induced by DHA higher than that produced by AA (-11.2 ± 1.1 mV, $n = 7$ vs. -5.1 ± 1.8 mV, $n = 10$; $P < 0.05$) (Fig. 1C). These results, 1) the higher degree of block observed in the tail currents, and 2) the negative shift of the activation curve, suggest an open-channel block mechanism.

To determine the fully activated *I-V* relationships (Fig. 1E), a double pulse protocol was applied (Fig. 1D). A 1-s step to $+30$ mV to activate *HERG* channels and a test pulse to potentials from -120 to $+10$ mV in 10-mV steps were

Fig. 1. Effects of arachidonic acid (AA) and docosahexanoic acid (DHA; 10 μM) on human *ether-a-go-go*-related gene (*HERG*) currents. A: current records obtained upon depolarization from a holding potential of -80 up to $+50$ mV in 10-mV steps and upon repolarization to 60 mV. Current records obtained in the absence and in the presence of AA (left) and DHA (right). B: current-voltage (*I-V*) relationships (5-s isochronal) of *HERG* channels obtained in the absence and in the presence of each fatty acid. C: activation curves of *HERG* channels obtained under control conditions and in the presence of AA or DHA. Dotted lines reflect the normalized activation curves obtained in the presence of polyunsaturated fatty acids (PUFAs) and matching control values. Open stars represent the relative current obtained at different membrane potentials. Each point represents the means \pm SE of 7–10 experiments. $E_{1/2}$, half-maximal voltages. * $P < 0.05$ and ** $P < 0.01$ vs. control. D: records obtained during the application of the pulse protocol shown in the top of the figure in the absence and in the presence of AA (left) or DHA (right). E: fully activated *I-V* relationships of *HERG* channels obtained in the absence and in the presence of AA or DHA.



applied. The pulse to +30 mV was positive enough to induce full conductance of *HERG* channels but also inactivated a large number of channels (inward rectification from potentials positive to -40 mV). Block induced by 10 μ M AA and DHA was not voltage-dependent and when measured at -120 mV averaged $41.8 \pm 7.1\%$ and $56.2 \pm 3.5\%$ ($n = 4$; $P > 0.05$), respectively.

AA metabolism occurs via three principal pathways: cyclooxygenase, lipoxygenase, and epoxygenase catalysis. To test whether the AA effects were due to its direct interaction with *HERG* channels or to the actions of some of its metabolites, the electrophysiological effects of its nonmetabolizable analog ETYA were analyzed (Fig. 2). ETYA (10 μ M) inhibited *HERG* current by $37.4 \pm 3.7\%$ and $42.5 \pm 5.8\%$, measured at the end of 5-s pulses to -10 mV and at the maximum peak tail currents, respectively ($n = 7$; $P > 0.05$), similar to the inhibition produced by AA at the same concentration. Moreover, as it can be observed in Fig. 2, *B-D*, the kinetics of block induced by ETYA was similar to that observed in the presence of AA. Therefore, these results suggest that the AA effects observed on *HERG* channels are not due to AA metabolites.

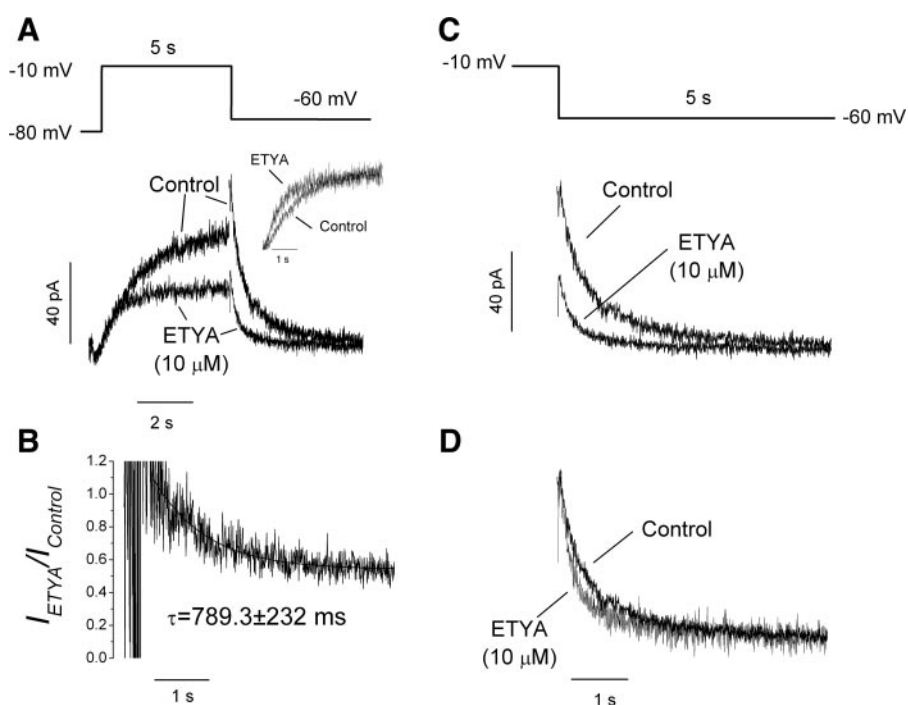
Time-dependent block of *HERG* channels induced by AA and DHA. AA and DHA accelerated the time course of the maximum outward current and decreased its amplitude (Fig. 3A). The initial increase of the current is likely due to the negative voltage shift of the activation curve induced by both PUFAs. In this figure, the control traces exponentially rise during depolarization, whereas in the presence of both PUFAs, the current remains constant after ~ 1 s. Thus the relative current ($I_{\text{Fatty-Acid}}/I_{\text{Control}}$) vs. time of depolarization was plotted (Fig. 3A). These ratio traces are a composite of the block, which seems to reach a maximum within the first second, which could represent a mixture of tonic- and time-dependent block. The relative current decreased in the presence of both PUFAs with a similar time constant ($\tau = 549.6 \pm 117.9$ ms, $n = 8$ vs. 731.6 ± 109.6

ms, $n = 10$, in the presence of AA and DHA, respectively, $P < 0.05$).

Time dependency of block was also apparent in the tail currents, as shown in Fig. 3B, which shows superimposed tail currents recorded under control conditions and in the presence of AA or DHA. The time course of deactivation of *HERG* channels was fitted to a biexponential function (τ_{fast} and τ_{slow}). AA accelerated τ_{fast} (from 377.7 ± 30.8 ms to 243.8 ± 45.6 ms, $n = 8$; $P < 0.05$) and τ_{slow} (from 1923.7 ± 257.9 ms to 1599.9 ± 198.1 ms, $n = 8$; $P < 0.05$) of deactivation without modifying the ratio of amplitudes [$A_{\text{fast}}/(A_{\text{fast}} + A_{\text{slow}})$] (0.31 ± 0.09 vs. 0.34 ± 0.06 , $n = 8$, $P < 0.05$). DHA accelerated τ_{fast} (from 351.6 ± 41.2 to 222.3 ± 24.9 ms, $n = 9$; $P < 0.05$) without modifying τ_{slow} (1713.7 ± 264.0 vs. 1345.7 ± 85.7 ms, $n = 9$; $P > 0.05$) or the ratio of amplitudes (0.51 ± 0.05 vs. 0.48 ± 0.04 , $n = 9$, $P < 0.05$).

To analyze the effects of AA or DHA on the inactivation kinetics, the three-pulse protocol shown in the top of Fig. 4 was applied. Holding potential was maintained at -80 mV, and after a 2-s step to +40 mV that fully activates *HERG* channels, a 20-ms pulse to -120 mV that promotes the recovery from the fast inactivation of *HERG* channels was applied. Then, 20-ms test pulses to membrane potentials between 0 and +50 mV were applied. The degree of block induced by DHA when measured at the maximum peak current recorded during the application of the test pulse to 0 mV was higher than that induced by AA. Neither DHA nor AA modified the time constant of inactivation (τ_{Inac}) at membrane potentials between 0 and +40 mV. However, at membrane potentials positive to +40 mV, AA and DHA increased the τ_{Inac} , which can be explained either by a fatty acid-induced stabilization of the open state of *HERG* channels or a destabilization of the inactivated state. Figure 4B shows the degree of block induced by AA and DHA at the end of 5-s pulses to 0 mV (End 5-s) and at the maximum peak current recorded during the application

Fig. 2. Effects of 5,6,11,14-eicosatetrayenoic acid (ETYA; 10 μ M) on *HERG* currents. *A*: current records obtained upon depolarization from a holding potential of -80 mV up to -10 mV and upon repolarization to -60 mV in the absence and in the presence of ETYA. *Inset*, first 5 s of the current traces shown in *A*. *B*: relative current ($I_{\text{ETYA}}/I_{\text{Control}}$) vs. time of depolarization. Current exponentially decreased during depolarization. *C*: tail currents recorded upon repolarization to -60 mV after a 5-s pulse to -10 mV in the absence and in the presence of ETYA. *D*: tail current traces shown in *C* normalized to match control values.



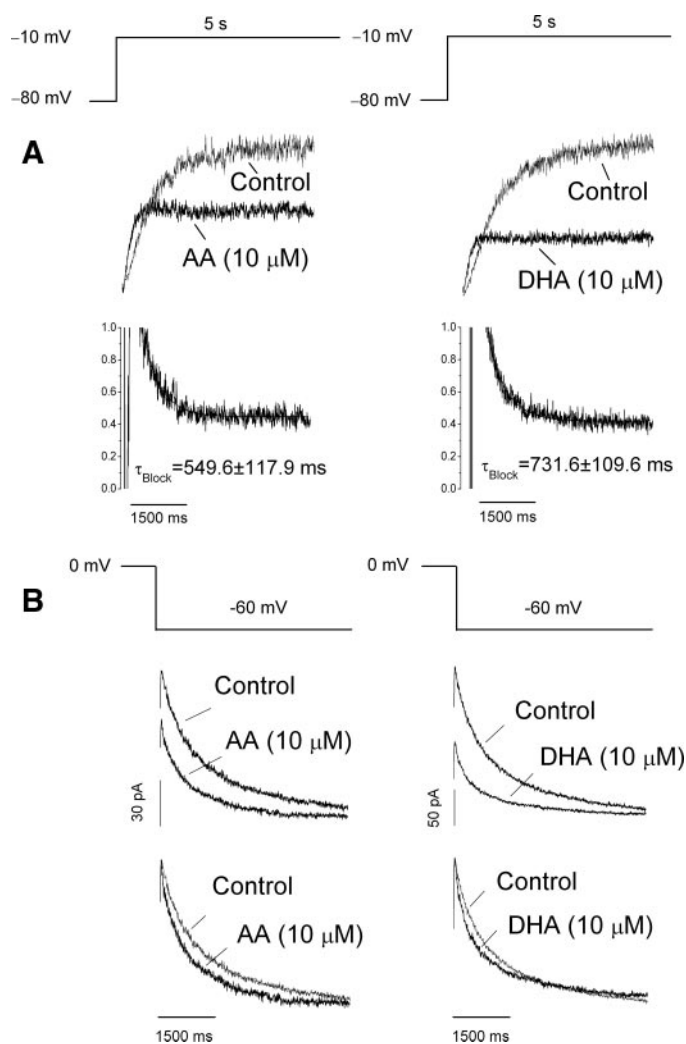


Fig. 3. Time-dependent block induced by AA and DHA (10 μM). *A*, top: current traces obtained after depolarizing the cell membrane from -80 to -10 mV in the absence and in the presence of AA or DHA. *Bottom*, plot of the relative current ($I_{\text{Fatty Acid}}/I_{\text{Control}}$) vs. time of depolarization. Current exponentially decreased during depolarization. *B*, top: tail currents recorded upon repolarization to -60 mV after a 5-s pulse to 0 mV in the absence and in the presence of AA or DHA. *Bottom*, tail current traces normalized to match control values.

of the test pulse to 0 mV (Maximum). DHA induced a higher block ($P < 0.05$) when measured at the maximum peak current than at the end of 5-s depolarizing pulses, thus suggesting an open channel block. However, block of *HERG* channels induced by AA was similar under both experimental conditions.

Effects of AA and DHA on *HERG* inactivation availability. To analyze the effects of AA and DHA on the inactivation availability, a three-pulse voltage clamp protocol was used (Fig. 5) (38, 39). Figure 5A shows the effects of 10 μM AA and DHA on the voltage dependence of *HERG* channels ("inactivation availability"). The dynamic nature of *HERG* currents precludes direct measurements of "steady-state" inactivation, but the data shown approximate a voltage dependence of the distribution of channels between open and inactivated states. The 1-s step to $+30$ mV fully activates and inactivates the channels. The second step to the test potential (between -120 and $+10$ mV) allows some fraction of the channels to recover

from inactivation. The instantaneous current after the third step to $+30$ mV allows measurements of the fraction of channels that have recovered from inactivation during the preceding test step. Figure 5B shows a plot of the current measured at $+30$ mV as a function of voltage. *HERG* current falls off at negative potentials because the channels also begin to deactivate during the 20-ms pulse. We corrected the current magnitude for the amount of deactivation at each voltage by fitting the deactivating tail currents with an exponential function and then back-extrapolated this fit to the beginning of the hyperpolarizing pulse, which permitted us to estimate the fraction of channels deactivating during the 20-ms pulse and then increase the outward current accordingly (38). Figure 5C shows the Boltzmann fit of the corrected data in the absence and in the presence of the two PUFAs. AA and DHA shifted the apparent voltage dependence of channel availability toward more positive potentials by $+12.0 \pm 3.9$ mV ($n = 4$, $P < 0.05$) and $+15.8 \pm 4.3$ mV ($n = 4$, $P < 0.05$), respectively, without modifying the slope factor.

State- and use-dependent effects of AA and DHA on *HERG* channels. To further analyze the interaction between AA or DHA and *HERG* channels, we applied the voltage protocol shown in the top of Fig. 6. From a holding potential of -80 mV, *HERG* channels were rapidly activated by a 5-ms step to $+180$ mV, followed by a 200-ms step to 0 mV, before returning to -60 mV. With this protocol, *HERG* current amplitude at 0 mV, under control conditions, was nearly constant. In the presence of AA or DHA, the amplitude of *HERG* current exponentially declined to a steady-state level as additional block developed during the step to 0 mV. The time course of AA- and DHA-induced *HERG* current decay was determined by dividing the current trace obtained in the presence of the fatty acid and the control trace. This current was fitted as a monoexponential decay to calculate the time constant of block of the open state of *HERG* channels (τ_B) (27.9 ± 5.6 ms, $n = 3$, and 39.2 ± 5.7 ms, $n = 5$, for AA and DHA). This pulse protocol also allowed us to measure the recovery kinetics by fitting the hook of the tail current at -60 mV (Fig. 6A). AA or DHA, at 10 μM , did not modify the recovery process (6.4 ± 1.3 vs. 6.5 ± 1.0 ms; $n = 4$, $P > 0.05$ in the absence and in the presence of AA and 5.9 ± 0.6 vs. 4.9 ± 0.5 ms; $n = 5$, $P > 0.05$ in the absence and in the presence of DHA). This voltage pulse protocol was applied at 1.0 Hz and, as can be observed, AA and DHA decreased in a use-dependent manner the amplitude of *HERG* current. This pattern is consistent with a preferential block of the open state of the channel. In the presence of DHA, the maximum peak current was higher than under control conditions, which can be due to the slower τ_{Inac} at positive potentials ($+180$ mV).

AA and DHA *HERG* current inhibition during cardiac action potential. Because of the complex gating of *HERG*, it is difficult to predict the effects of AA and DHA on ionic currents during a cardiac action potential (AP), where the membrane voltage is continually changing with time. To assess this issue, the effects of AA and DHA were analyzed using the AP clamp technique as shown in Fig. 7. The frequency of stimulation used was 1 Hz. Under control conditions, *HERG* current rapidly inactivates during the upstroke. As the action potential repolarizes, *HERG* channels rapidly recover from inactivation to reopen and slowly deactivate, increasing their occupancy in the open state. The current deactivates with the final phase of

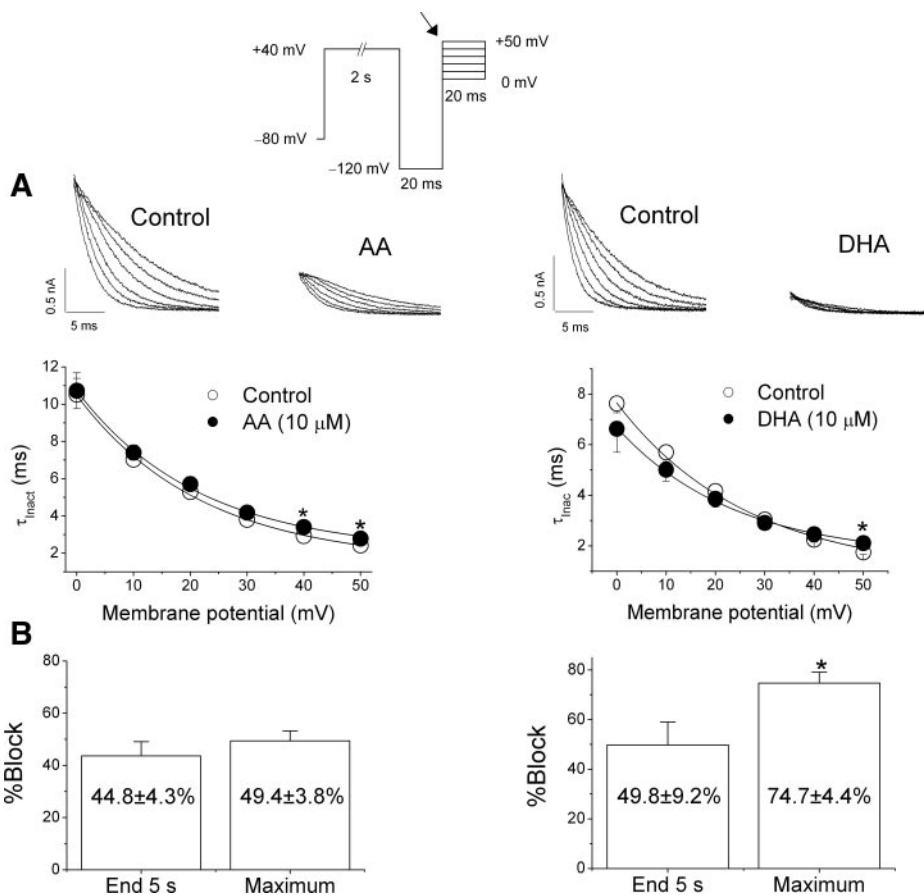


Fig. 4. Effects of 10 μ M AA or DHA on *HERG* inactivation kinetics. *A*, top: current records of test pulses obtained in the absence and in the presence of each fatty acid. Bottom, plots of the time constant of inactivation at different membrane potentials in the absence and in the presence of each agent. *B*: graph showing the degree of block induced by AA or DHA measured at the end of 5-s pulses to 0 mV (End 5-s) or at the maximum peak current recorded at a test pulse to +40 mV after a test pulse to -120 mV during 2-s and followed by a 20-ms pulse to -120 mV (Maximum) indicated by an arrow in the pulse protocol. Each point represents the means \pm SE of 7–9 experiments. * $P < 0.05$ vs. control conditions.

repolarization. AA (10 μ M) decreased the maximum peak current by $43.3 \pm 1.6\%$ ($n = 3$). DHA increased *HERG* current during the upstroke and decreased it at the end of the AP to an extent similar to that of AA ($48.1 \pm 8.3\%$, $n = 3$, $P > 0.05$).

DISCUSSION

In the present study, the effects of AA and DHA on *HERG* channels have been studied. We found that, at relevant plasma concentrations, both PUFAs block *HERG* channels to a similar extent. Moreover, the effects of AA on *HERG* channels appear to be due to this PUFA and not to its metabolites because its actions are mimicked by the nonmetabolizable analog ETYA.

Effects of AA and DHA on *HERG* channels. AA and DHA block *HERG* channels in a voltage-, time-, and state-dependent manner, which is consistent with an open channel block mechanism. In fact, block induced by both PUFAs steeply increased in the range of membrane potentials that coincides with the range of *HERG* channel activation, suggesting that their binding may derive a significant fraction of its voltage sensitivity through coupling to channel gating. Unfortunately, at depolarized voltages the open and inactivated conformations of *HERG* channels are in rapid equilibrium, making it difficult to unequivocally identify the state(s) with which these two PUFAs interact.

Whereas AA induced a similar inhibition of the *HERG* current when measured at the end of depolarizing pulses to -10 mV and at the maximum tail currents, DHA inhibited this current to a higher extent when measured at the maximum tail current than at the end of depolarizing steps to -10 mV.

During depolarization, *HERG* channels inactivate faster than they activate and thus the amplitude of the current is reduced. On repolarization, closed channels transit through the open state, resulting in tail currents with higher amplitude (38, 39). In agreement with these results, block induced by AA when measured at the maximum peak current of a test pulse to 0 mV applied after a hyperpolarizing pulse to -120 mV (that promotes the I \rightarrow O transition) was similar to that observed at the same voltage at the end of a 5-s depolarizing pulse. However, DHA-induced block, when measured at the maximum peak current of a test pulse to 0 mV applied after a hyperpolarizing pulse to -120 mV, was higher than that observed at the same voltage at the end of a 5-s depolarizing pulse. Block induced by AA and DHA was also time dependent, being evident after a prepulse to +180 mV, suggesting a rapid drug binding to activated *HERG* channels, as previously described for cocaine and bupivacaine-type local anesthetics (11, 44). This time dependency was also evident in the deactivation process of *HERG* channels that was accelerated in the presence of both AA and DHA. However, AA and DHA did not modify the onset kinetics of the inactivation process or the recovery process. The faster deactivation induced by both PUFAs, together with their lack of effect on the recovery kinetics, suggests that a very fast dissociation rate constant from *HERG* channels is consistent with an open channel block mechanism, as has been proposed for propafenone (2). Another piece of evidence of an open channel interaction between both PUFAs and *HERG* channels is the use-dependent inhibition of the current. Taken together, all these results suggest that both AA

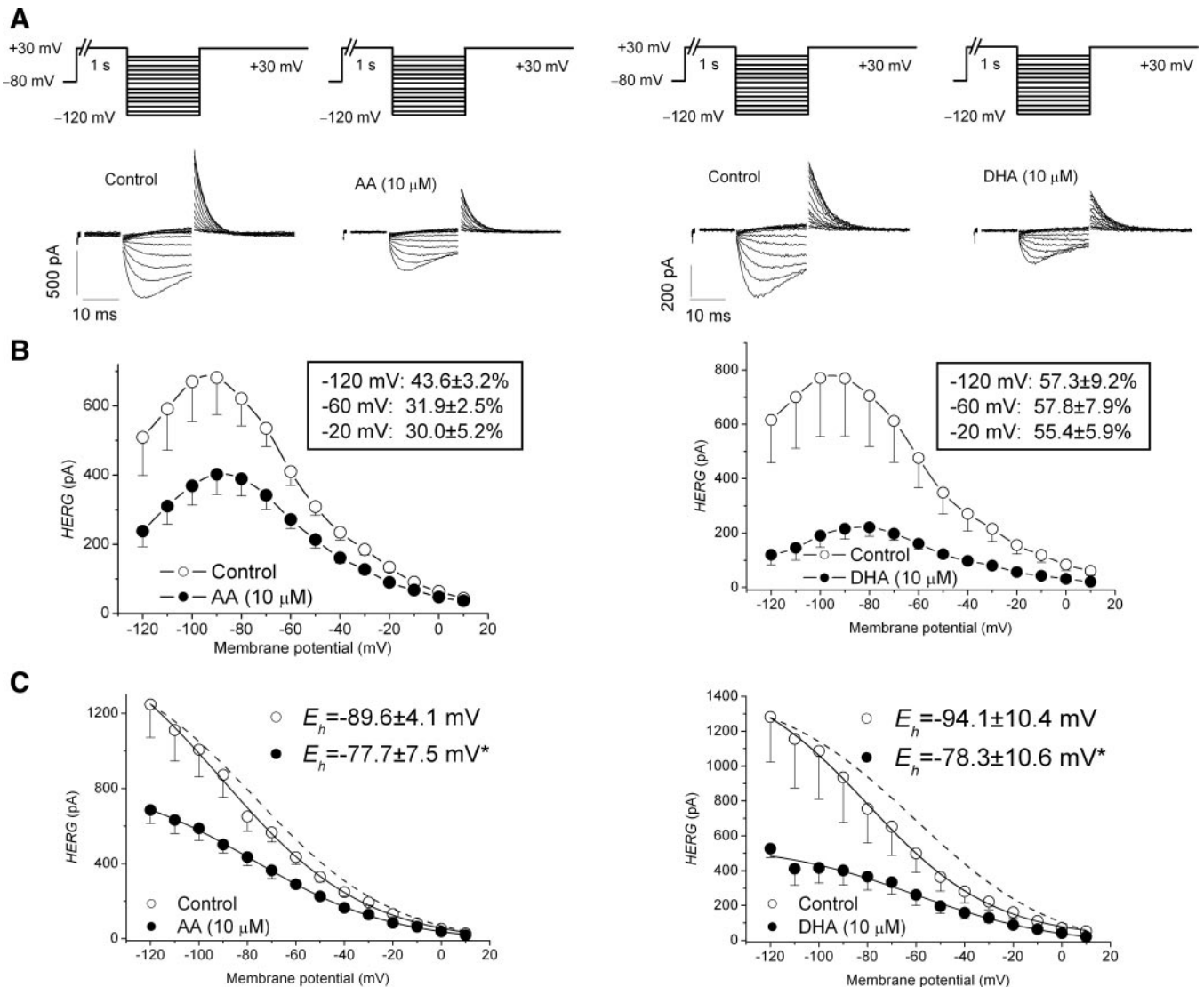


Fig. 5. Apparent voltage dependence of channel availability. The pulse protocol used to obtain each data point is shown at *top*. *A*: original traces obtained after applying such pulse protocol in the absence and in the presence of 10 μ M AA and DHA. *B*: peak current observed immediately after stepping to +30 mV was then plotted vs. test potential in the absence and in the presence of 10 μ M of AA (*left*) and DHA (*right*). Currents decreased at negative potentials due to deactivation during the 20-ms pulse. *C*: corrected data for deactivation (see RESULTS) together with the Boltzmann fit. Each point represents the mean \pm SE of 4 experiments. The dashed lines represent the normalized fits to matching control. * $P < 0.05$.

and DHA preferentially bind to the open state of *HERG* channels, and that DHA exhibits a higher affinity for this state of the channel.

We also observed that AA and DHA produced a positive shift in the inactivation curve. This could be explained either by 1) stabilization of the open state of *HERG* channels or 2) destabilizing the inactivation process; i.e., without modifying the onset but accelerating the offset of inactivation. In both cases, the shift of the inactivation curve would be the result of the interaction between PUFAs and a closed state of *HERG* channels (tonic block). This tonic block is likely to influence the apparent steady-state inactivation and perhaps the activation process because both PUFAs accelerate the deactivation process. All of these results suggest that AA and DHA preferentially block the open state of *HERG* channels, but also that they interact with a closed state, thus producing changes in channel gating. Finally, the similar degree of AA-induced

inhibition of the current at the end of 5-s depolarizing pulses (when most channels are inactivated) and at the maximum tail current or at the maximum peak current after applying a -120 mV step cannot permit us to rule out an interaction between AA and the inactivated state of *HERG* channels. Moreover, it has been shown that AA regulates the inactivation process in other K^+ channels in such a way that introduces rapid voltage dependent inactivation into noninactivating K_v channels (30). The authors explain these results under the framework that AA closes K_v channels by inducing conformational alterations in the selectivity filter region (30) and propose that K_v channel inactivation is lipid dependent and that this process has a high affinity, comparable to that of K_{ATP} channels for phosphoinositides (3). Oliver et al. (30) propose that AA inserts into the cell membrane from either side, interacts with the channel protein, and, allosterically, induces a fast closure of the open K_v channel pore through

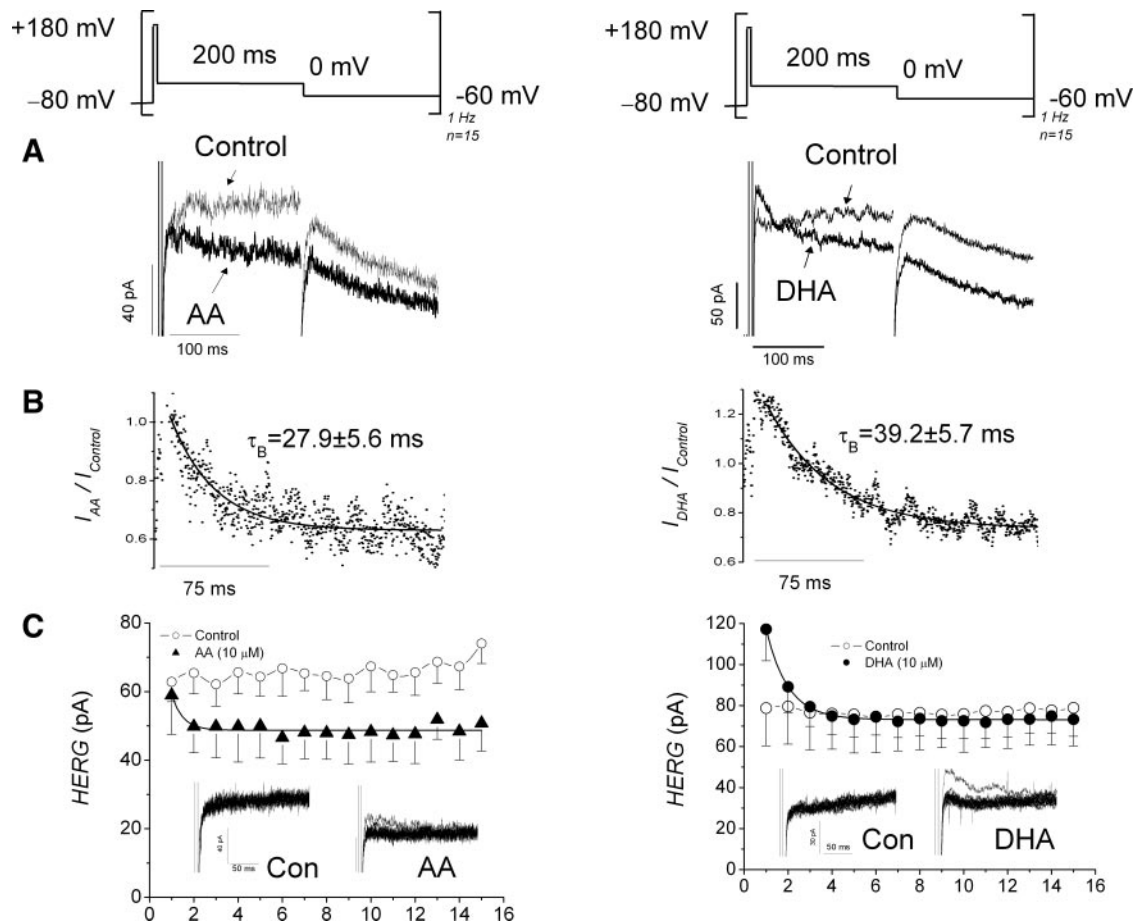


Fig. 6. Time-dependent block of AA and DHA (10 μ M) after a fast activation of *HERG* channels, using the pulse protocol shown at the top. *A*: first record obtained in the absence and in the presence of AA or DHA. *B*: relative current ($I_{FattyAcid}/I_{Control}$) during the step to 0 mV together with its monoexponential fit. *C*: use-dependent block of *HERG* channels induced by AA and DHA obtained after applying the pulse protocol shown at top at 1.0 Hz. Inset, records obtained in the absence and in the presence of AA or DHA.

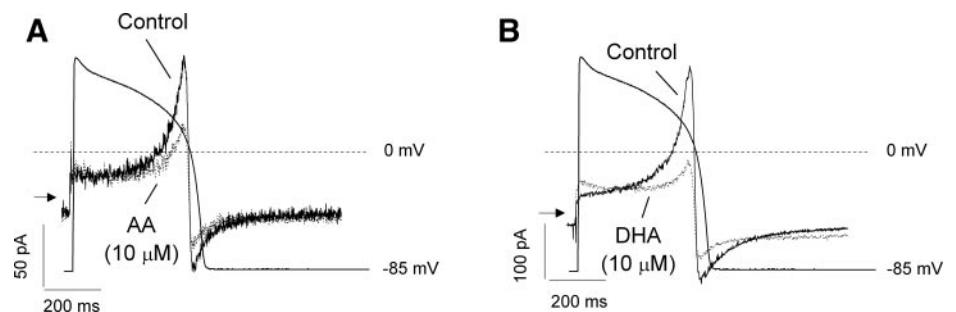
conformational modifications in the selectivity filter. Further studies are required to discern the possible role of AA on the inactivation of *HERG* channels.

Clinical implications of this study. Many studies suggest that ω -3 fatty acids have beneficial effects on human health. In contrast, a diet enriched in ω -6 fatty acids provokes atherosclerosis, carcinogenesis, and heart disease (20). In the present study we have demonstrated that AA and DHA block *HERG* channels. The plasma levels of these two PUFAs vary greatly depending on the hormonal, metabolic, and nutritional state of the individual. Around 99.9% of fatty acids are carried in plasma bound to albumin and the range of free AA and DHA

in human plasma is 5.3–13.1 μ M and <2.8 μ M, respectively (6). The EC_{50} values calculated from the blockade produced by both PUFAs at 10 μ M are well within their physiological plasma range. Therefore, we can conclude that both PUFAs block *HERG* channels at concentrations within their physiological plasma levels.

It has been described that AA is able to modulate several ion currents (28). In fact, AA decreases L-type I_{Ca} ($I_{Ca,L}$), T-type I_{Ca} ($I_{Ca,T}$), I_{to} , I_{Na} , and $Kv1.5$ currents (4, 14, 40, 41) and activates I_{K1} (25). AA plasma levels are known to be increased during ischemia and reperfusion at the intracellular and extracellular levels (19), which, in the light of previous and present

Fig. 7. Effect of 10 μ M AA (*A*) and DHA (*B*) on *HERG* currents evoked by an action potential waveform from a guinea pig ventricular myocyte. The frequency of stimulation was 1 Hz. Arrow indicates 0 current level.



results, could represent a “natural” protective mechanism to prevent arrhythmias under these pathological conditions. On the other hand, it has been shown that DHA exhibits potential cardioprotective effects that have been associated with their antiarrhythmic effects (21). DHA’s antiarrhythmic effects have been attributed to its actions on the cardiac ion channels responsible for the onset and maintaining of the cardiac action potential, mostly on its inhibitory effects on Na^+ and $I_{\text{Ca,L}}$ channels (22, 42), because DHA has inhibitory effects on I_{Na} , but not on $I_{\text{Ca,L}}$, that are higher than those of AA (41). DHA inhibitory effects on I_{Na} and I_{Ca} theoretically should shorten the cardiac action potential duration. Moreover, DHA enhances I_{Ks} , which would further shorten the cardiac action potential duration (9). However, experiments performed in rat ventricular myocytes demonstrate that DHA slightly prolongs the rat cardiac action potential (4, 26). In the present study we have demonstrated that DHA blocks *HERG* channels. Most drugs that selectively block *HERG* channels prolong the cardiac action potential (34). However, interactions of nonselective *HERG* channel blockers with other cardiac ion channels may mitigate or exacerbate the prolongation of action potential duration (27, 31, 33). Because DHA inhibits I_{Na} and I_{Ca} and enhances I_{Ks} (effects that would shorten the cardiac action potential), but also inhibits I_{to} , I_{Kur} , and *HERG* (which should produce a lengthening of the action potential duration), the result should be a modest effect on the time of repolarization. Besides these effects on the action potential duration, it should be stated that its inhibitory effects on Na^+ , Ca^{2+} , and several K^+ channels (Kv4 and *HERG* channels) would result in a lengthening of the refractory period and a decrease of cardiac excitability, thus contributing to its antiarrhythmic effects.

To our knowledge, this is the first demonstration that *HERG* channels are modulated by AA and DHA. The ω -3 antiarrhythmic effects have been attributed to their availability to modulate cardiac ion channels involved in the genesis and maintenance of the cardiac action potential (21). Our results suggest that AA and DHA block *HERG* channels mainly by binding to the open state of the channel. These PUFAs’ actions on *HERG* channels have to be taken into account to explain the antiarrhythmic effects of AA during ischemia and those previously reported for DHA in subjects consuming a diet rich in fish and fish oils.

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