Hesperetin preferentially inhibits slow-inactivating currents of an LQT3 syndrome Na⁺ channel mutation.

Julio Alvarez-Collazo¹, Alejandro López-Requena¹, Loipa Galán², Ariel Talavera³, Julio L. Alvarez² and Karel Talavera¹

¹ Laboratory of Ion Channel Research, Department of Cellular and Molecular Medicine, KU Leuven; VIB Center for Brain & Disease Research, Leuven, Belgium. ² Laboratory of Electrophysiology. Institute of Cardiology and Cardiovascular Surgery, Havana, Cuba. ³ Molecular Recognition Unit, Department of Structural Biology, VIB, Brussels, Belgium.

INTRODUCTION | The citrus flavanone hesperetin (HSP) has been proposed for the treatment of several human pathologies, but its cardiovascular actions remain unexplored. Here we studied the effects of HSP on the human cardiac voltage-gated Na⁺ channel (hNa_v1.5) and compared it to its effects on a recombinant hNa_v1.5 channel baring a mutation (R1623Q) associated with lethal ventricular arrhythmias in the Long QT syndrome type 3 (LQT3). METHODS | Whole-cell patchclamp experiments were used to record Na⁺-currents (I_{Na}) in HEK293T cells expressing hNa_v1.5 wild type (WT) or mutant channels. RESULTS | HSP blocked the $hNa_v 1.5$ channels in voltage-dependent manner with an IC₅₀ \approx 100 μ M. Its inhibition was decreased by disruption of the F1760 residue. HSP preferentially accelerated the inactivation phase of I_{Na} and decreased the Na⁺ net influx into the cell. The effects of HSP on the inactivation phase and voltage-dependent

inhibition of I_{Na} were more marked in the LQT3 mutant. CONCLUSIONS | HSP could be used as a template to develop drugs against cardiac arrhythmias in LQT3.



Figure 1 | HSP preferentially inhibits the later phase of I_{Na} in HEK293T cells expressing the WT or LQT3 mutant channels. Time course of the I_{Na} from WT (a) or LQT3 (b) hNa_v1.5 expressing HEK293T cells in control condition and in presence of HSP 100 µM. Representative current traces in control (blue) and upon the application of HSP (orange) are shown in the insets and correspond with the coloured data points. Concentration-effect curves for the action of HSP on WT (c) or LQT3 (d) $hNa_v 1.5$ currents. The dots represent the mean \pm s.e.m. inhibition percentage during the application of different concentrations of HSP (n = 9). The estimated IC₅₀ are shown on the insets.

Figure 2 | **HSP decreases the cell Na⁺ load.** Effects of HSP 100 µM (orange) on the inactivation time course of I_{Na} in the WT (a) and R1623Q (b) channels. c) Concentration-dependent effects of HSP on the fast (τ_{fast}) and slow (τ_{slow}) inactivation time constants of I_{Na}. * P < 0.05 compared to its own control; n = 12, two-way ANOVA with Tukey's post hoc test. d) HSP 100 μ M reduced the amount of transported Na⁺ during the I_{Na}. Current traces were integrated and data expressed as charge normalized to membrane capacitance. * P < 0.05 with respect to its own control. § P < 0.05 with respect to WT; n = 32, two-way ANOVA with a Tukey's post hoc test.



Figure 3 | Effects of HSP on voltage-dependent kinetics of I_{Na} in the WT and the R1623Q mutant. Inactivation and activation curves for WT (a) and R1623Q (b) in control (blue) and in the presence of HSP 100 μ M (orange). The dots represent the mean ± s.e.m. of n = 13 normalized peak I_{Na} amplitude (for the inactivation curve) and channel conductance (for the activation curve) values. HSP 100 µM significantly shifted the inactivation curve compared to control in both the WT and R1623Q mutant channels (P < 0.05, paired t-test). Consequently, the Na⁺ window current is markedly decreased by HSP in both the WT (c) and the R1623Q mutant channels (d).

Figure 4 | HSP interacts with the hNa_v1.5 channel local anaesthetic (LA) binding site. a) Time course of the I_{Na} in a HEK293T cells expressing the hNa_v1.5 channel with the mutation F1760A. Representative current traces in control (blue) and upon the application of HSP 1 mM (orange) are shown in the insets and correspond with the coloured data points. b) Bar graph comparing the mean ± s.e.m. maximal inhibition of peak I_{Na} by HSP in WT and F1760A mutated channel (n = 9, * P < 0.05, two-sample t-test). c) and d) Molecular docking experiments in a model of the $hNa_v 1.5$ channel. HSP (yellow) was bound at the $hNa_v 1.5$ LA binding site (blue).



