

Supplementary Materials for  
**An *RYR1* mutation associated with malignant hyperthermia is also associated with bleeding abnormalities**

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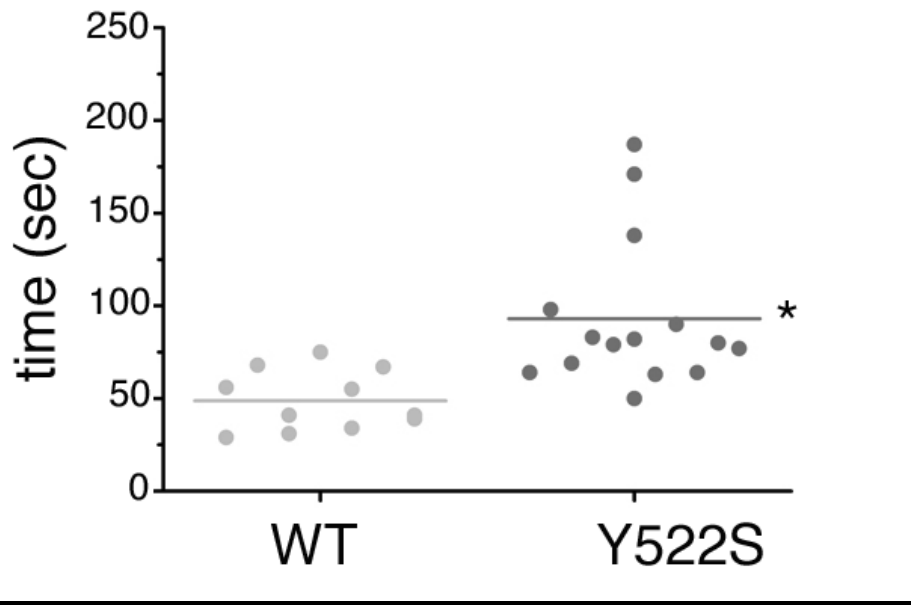
**The PDF file includes:**

- Fig. S1. *RYR1*<sub>Y522S</sub> female mice also exhibit prolonged bleeding times.
- Fig. S2. Heart rate and systolic blood pressure are similar in wild-type and *RYR1*<sub>Y522S</sub> mice.
- Fig. S3. Expression of *RYR2* and *RYR3* is not significantly different in the tail arteries of wild-type and *RYR1*<sub>Y522S</sub> mice.
- Fig. S4. Specificity of the anti-RyR1 antibodies used for immunohistochemistry.
- Fig. S5. Membrane potential measurements using the fluorescence potentiometric probe bis-oxonol.
- Table S1. Detailed analysis of full kinetic parameters of sparks in vascular smooth muscle cells from wild-type and *RYR1*<sub>Y522S</sub> mice.
- Legends for videos S1 to S5

**Other Supplementary Material for this manuscript includes the following:**  
(available at [www.sciencesignaling.org/cgi/content/full/9/435/ra68/DC1](http://www.sciencesignaling.org/cgi/content/full/9/435/ra68/DC1))

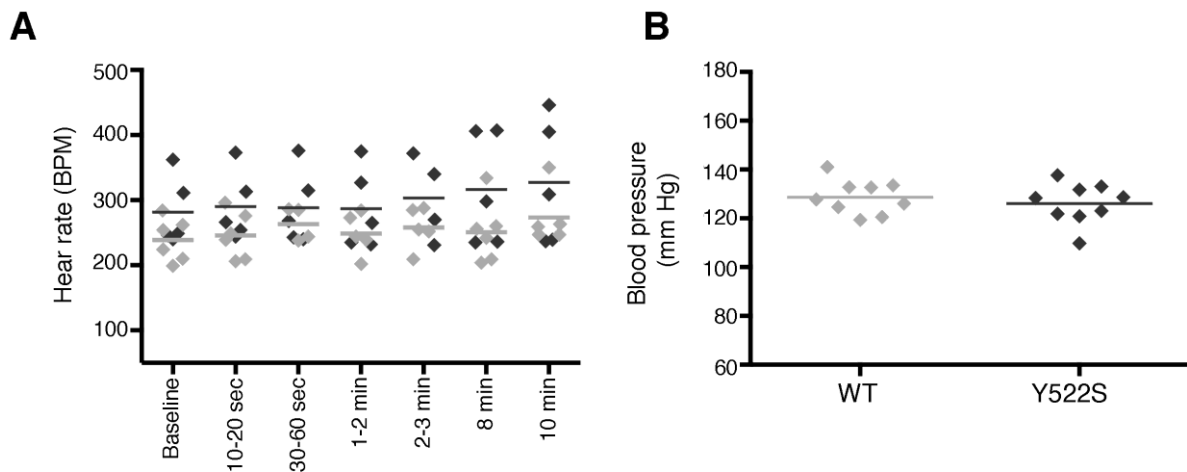
- Video S1 (.avi format). Sparks in smooth muscle cells from wild-type mice.
- Video S2 (.avi format). Sparks in smooth muscle cells from *RYR1*<sub>Y522S</sub> mice.
- Video S3 (.avi format). Sparks in smooth muscle cells from wild-type mice after treatment with 10  $\mu$ M ryanodine.
- Video S4 (.avi format). Sparks in smooth muscle cells from *RYR1*<sub>Y522S</sub> mice after treatment with 20  $\mu$ M dantrolene.

Video S5 (.avi format). Sparks in smooth muscle cells from RYR1<sub>Y522S</sub> mice after treatment with 250 nM xestospongin C.



**Supplementary Figure 1: RYR1<sup>Y522S</sup> female mice also exhibit prolonged bleeding times.**

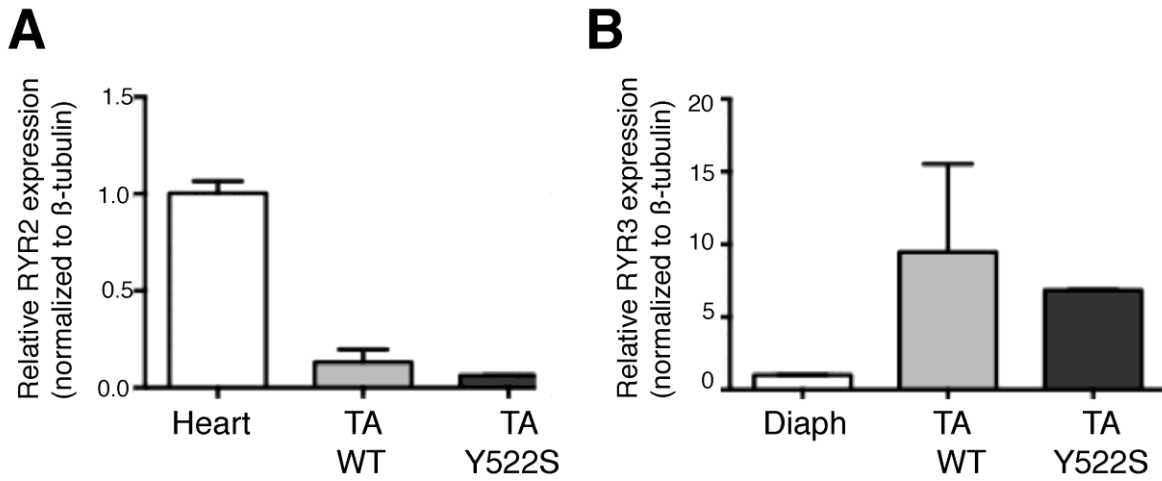
Each symbol represents the bleeding time (in sec) of a single wild type (light grey) and RYR1<sup>Y522S</sup> (dark grey) female mouse. \*P<0.005 Students *t*-test.



**Supplementary Figure 2: Heart rate and systolic blood pressure are similar in wild-type and RYR1<sup>Y522S</sup> mice.**

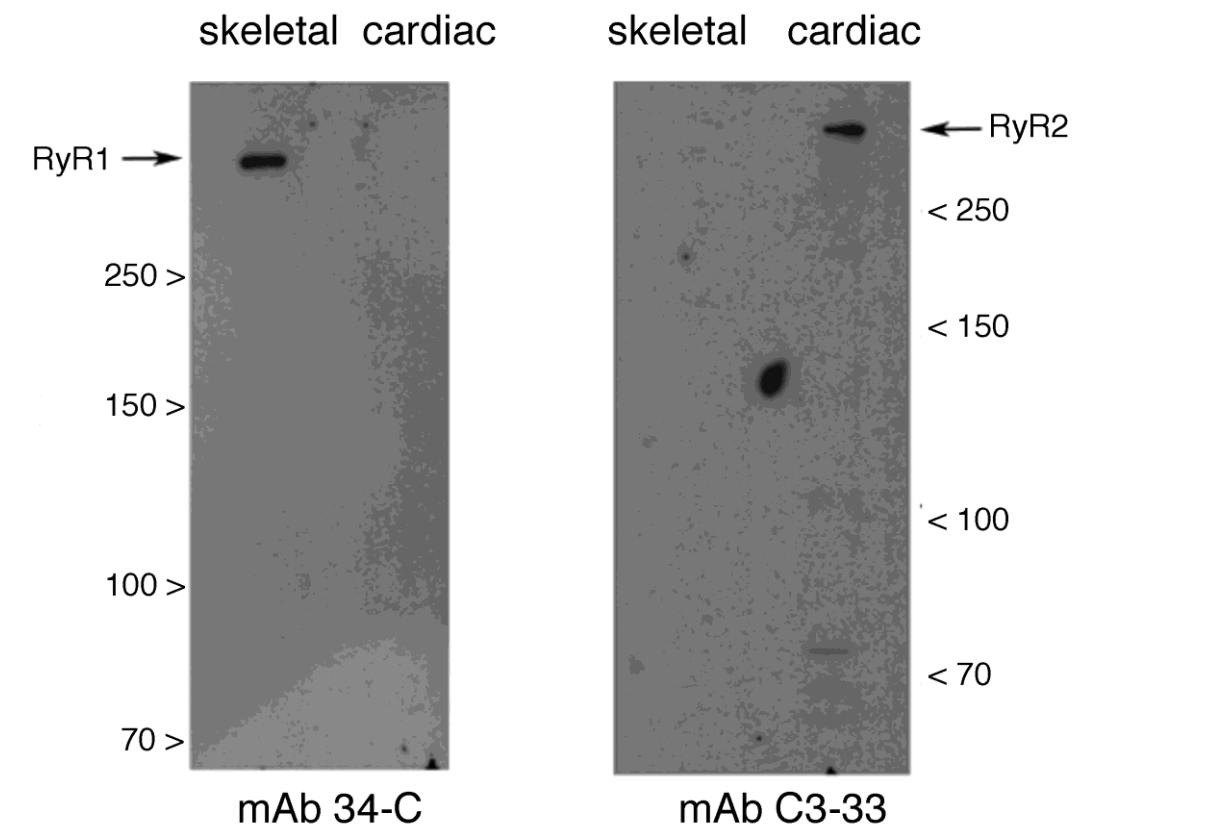
**A.** Heart rates were obtained while performing Echo Doppler in lightly anesthetized animals at different time points after incision of the caudal artery. Each symbol represents a WT mouse (light grey) and RYR1<sup>Y522S</sup> mouse (dark grey). There was a trend for

higher heart rates in the RYR1<sub>Y522S</sub> mice that did not reach statistical significance. **B.** Systolic blood pressure in non-anesthetized animals did not differ between WT (light grey) and RYR1<sub>Y522S</sub> (dark grey) mice.

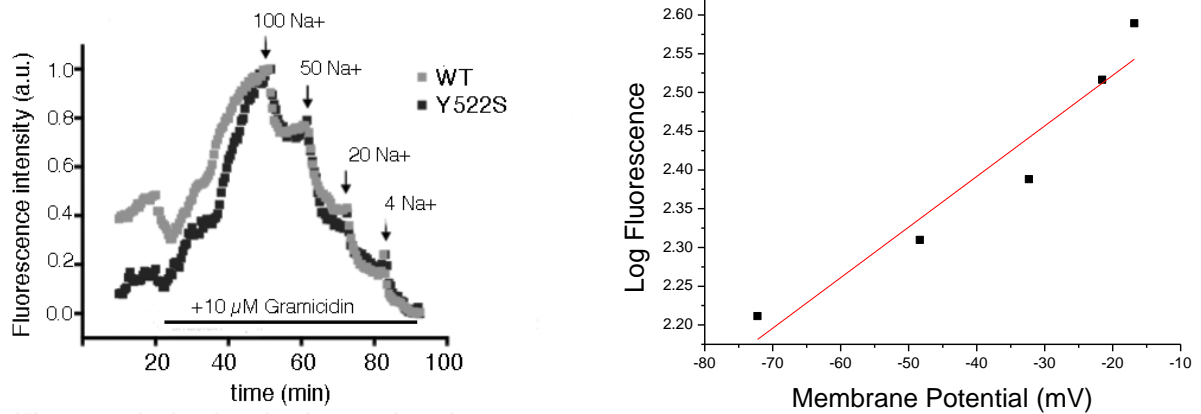


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**Supplementary Figure 3: Expression of *RYR2* and *RYR3* is not significantly different in the tail arteries of wild-type and *RYR1*<sub>Y522S</sub> mice.** Expression of *RYR2* and *RYR3* were determined by qPCR and was normalized to  $\beta$ -tubulin. Results are the mean  $\pm$  SEM of 3 replicates on tail arteries isolated from 4 different mice. Results are expressed as relative expression compared to *RYR2* expression in mouse heart (A) and *RYR3* expression in mouse diaphragm (B).



**Supplementary Figure 4: Specificity of the anti-RyR1 antibodies used for immunohistochemistry.** Thirty micrograms of sarcoplasmic reticulum proteins from skeletal muscle (left lane) or heart (right lane) were separated on a 6% SDS PAG and blotted onto nitrocellulose. Blots were blocked with 1% blocking buffer (Roche) in TBS for 60 min and probed with 1:5000 dilution monoclonal anti-RyR1 specific mAb34-C (left panel) or anti-RyR2 mAb C3-33 (right panel) for 60 min in TBST, rinsed 3 times with TBST, incubated with peroxidase-conjugate anti-mouse IgG (1:200,000 in TBST), rinsed 3 times and developed using the enhanced chemiluminescence kit from Thermo Scientific. As shown, the anti-RyR1 mAb does not cross react with cardiac RyR2 under the experimental conditions used in our laboratory. The blot on the right shows a positive control for cardiac RyR2. Representative of 2 independent experiments.



**Supplementary Figure 5. Membrane potential measurements using the fluorescence potentiometric probe bis-oxonol. Left:** Representative traces of the changes in fluorescence obtained in an arterial smooth muscle cell from WT mice (light symbols and heterozygous RYR1<sub>Y522S</sub> mice (dark symbols) expressed as Fluorescence Intensity arbitrary units (AU). The arrows indicated addition of Gramicidin or the indicated [Na<sup>+</sup>] in mM. **Right:** Values derived from the formula  $E_m = 60 \text{ Log}([Na^+]_o + [K^+]_o) / ([Na^+]_i + [K^+]_i)$  with the best linear fitting regression ( $R^2=0.90$ ;  $Y = 0.0065X + 2.6525$ ) of the log of the fluorescence obtained at each [Na<sup>+</sup>].

**Supplementary Table 1. Detailed analysis of full kinetic parameters of sparks in vascular smooth muscle cells from wild-type and RYR1<sub>Y522S</sub> mice.**

Spark Morphology							
Amplitude (DF/F <sub>0</sub> )	FWHM ( $\mu$ m)	FDHM (ms)	Full Wid ( $\mu$ m)	Full Duration (ms)	TtP (ms)	Tau (ms)	Freq (Sparks/image)
<b>WT</b>							
0.83±0.01 (N=877)	1.72±0.04	22.09±0.75	4.38±0.10	69.93±2.05	20.72±0.4	128.56±33.6	6.96±0.49
<b>RyR1<sub>Y522S</sub></b>							
0.79±0.0* (N=1388)	1.56±0.0*	19.52±0.6*	4.04±0.0*	59.32±1.3*	19.76±0.4	72.81±9.13	10.68±0.55*

- Values are expressed as mean  $\pm$  SEM. \* = P < 0.05 Student's *t*-test. Experiments were performed on 28 cells isolated from 6 WT and 6 RYR1<sub>Y522S</sub> mice respectively.
- †N indicates the number of individual sparks that were analyzed.



**Supplementary Video 1:** Sparks in smooth muscle cells from WT mice.

**Supplementary Video 2:** Sparks in smooth muscle cells from RYR1<sub>Y522S</sub> mice.

**Supplementary Video 3:** Sparks in smooth muscle cells from wild-type mice after treatment with 10  $\mu$ M ryanodine.

**Supplementary Video 4:** Sparks in smooth muscle cells from RYR1<sub>Y522S</sub> mice after treatment with 20  $\mu$ M dantrolene.

**Supplementary Video 5:** Sparks in smooth muscle cells from RYR1<sub>Y522S</sub> mice after treatment with 250 nM xestospongine C.