TRPC3-Nox2 complex mediates doxorubicin-induced myocardial atrophy

Tsukasa Shimauchi,1,2,3 Takuro Numaga-Tomita,1,4 Tomoya Ito,1 Akiyuki Nishimura,1,4 Ryosuke Matsukane,1,2 Sayaka Oda,1,4 Sumio Hoka,3 Tomomi Ide,5 Norimichi Koitabashi,6 Koji Uchida,7 Hideki Sumimoto,8 Yasuo Mori,9 and Motohiro Nishida1,2,4,10

1Division of Cardiocirculatory Signaling, Okazaki Institute for Integrative Bioscience (National Institute for Physiological Sciences), National Institutes of Natural Sciences, Aichi, Japan. 2Department of Translational Pharmaceutical Sciences, Graduate School of Pharmaceutical Sciences, and 3Department of Anesthesiology and Critical Care Medicine, Graduate School of Medical Sciences, Kyushu University, Fukuoka, Japan. 4School of Life Science, The Graduate University for Advanced Studies (SOKENDAI), Department of Physiological Sciences, Aichi, Japan. 5Department of Cardiovascular Medicine, Graduate School of Medical Sciences, Kyushu University, Fukuoka, Japan. 6Department of Medicine and Biological Sciences, Graduate School of Medicine, Gunma University, Gunma, Japan. 7Department of Applied Biological Chemistry, Graduate School of Agricultural and Life Sciences, The University of Tokyo, Tokyo, Japan. 8Department of Biochemistry, Graduate School of Medical Sciences, Kyushu University, Fukuoka, Japan. 9Department of Synthetic Chemistry and Biological Chemistry, Graduate School of Engineering, Kyoto University, Kyoto, Japan. 10PRESTO, JST, 4-1-8 Honcho, Kawaguchi, Saitama, Japan.

Introduction

Cardiac plasticity is defined as structural remodeling of the heart in response to environmental demands and is an intrinsic compensatory mechanism to reduce hemodynamic workload (1, 2). Exercise, pregnancy, and postnatal growth all promote physiologic myocardial growth, whereas neurohumoral activation, hypertension, and myocardial injury can all cause pathological hypertrophy, which increases the risk of heart failure by reducing myocardial flexibility through myocardial stiffening. Cardiac atrophy, defined as an acquired reduction in the size and mass of the heart, is a complication of protracted bed rest, prolonged weightlessness during space travel, and mechanical unloading through the use of a ventricular assist device. The atrophied myocardium also exhibits reduced contractility due to decreased myocardial flexibility. Why myocardial flexibility is reduced during the processes of both cardiac atrophy and pathological hypertrophy is largely unknown.

Doxorubicin (DOX) is a highly effective anticancer agent used to treat a variety of hematologic and solid malignancies (3–6). However, its dose-dependent cardiotoxicity limits its clinical use (3–6). The cardiotoxic effects of DOX range from asymptomatic increases in left ventricular (LV) wall stress to reductions in ejection fraction, arrhythmias, and highly symptomatic congestive heart failure, which are all associated with high mortality (6, 7). DOX initially forces the heart to shrink, which leads to induction of myocardial atrophy.
apoptosis and interstitial fibrosis at later stages of LV dilated cardiomyopathy (8, 9). There is growing evidence that the DOX-induced cardiac apoptosis and fibrosis are mainly caused by excess production of ROS (10–13), which is initiated by NADPH oxidase (Nox) activation (13, 14).

Two Nox enzymes, Nox2 and Nox4, are predominantly expressed in mammalian hearts (15–17). Nox2 is activated by angiotensin II, endothelin-1, tumor necrosis factor–β, growth factors, and mechanical force, and Nox2-dependent ROS production reportedly mediates congestive heart failure in humans (18). Nox2 also participates in mechanochemical transduction in hearts through ROS production during LV diastolic filling (i.e., mechanical stretch of cardiomyocytes) (19, 20). Notably, Nox2-deficient mice reportedly exhibit resistance to DOX-induced cardiac atrophy (13), but how Nox2 contributes to DOX-induced myocardial atrophy remains obscure.

Transient receptor potential (TRP) channels, originally identified as a visual phototransduction mutant, have been attracting attention as multimodal cation channels regulated by several environmental factors, including mechanical stress, in vertebrates (21). In particular, two TRP canonical (TRPC) subfamily members, TRPC3 and TRPC6, reportedly participate in the development of pathological hypertrophy caused by neurohumoral factors (22, 23) and mechanical stress (24–26). Our recent studies using TRPC-KO mice revealed that TRPC3, but not TRPC6, functions as a positive regulator of ROS, leading to induction of mechanical stress–induced maladaptive fibrosis (15, 20, 27). TRPC3 interacts with Nox2 via TRPC3-specific C-terminal sites, thereby protecting Nox2 from proteasome-dependent degradation and amplifying Ca\(^{2+}\)-dependent Nox2 activation through TRPC3-mediated background Ca\(^{2+}\) entry (20). Nox2 also stabilizes TRPC3 proteins to enhance mechanical stress–induced background Ca\(^{2+}\) entry into cardiomyocytes. Whether the TRPC3-Nox2 axis participates in DOX-induced myocardial atrophy is unknown.

Here, we demonstrate that increased levels of a TRPC3-Nox2 complex correlate well with the severity of DOX-induced cardiac atrophy and that inhibition of TRPC3-Nox2 functional coupling suppresses DOX-induced myocardial atrophy in mice. By contrast, voluntary exercise, which induces physiological cardiac hypertrophy, reduced levels of the TRPC3-Nox2 complex. Considering that the TRPC3-Nox2 axis also mediates pressure overload–induced pathological cardiac remodeling, stable induction of the TRPC3-Nox2 complex is apparently key to the transition from adaptation to maladaptation in the heart.

**Results**

**Deletion of TRPC3 channels attenuates DOX-induced myocardial atrophy in mice.** DOX-induced congestive heart failure is characterized by a severe reduction of LV mass and systolic dysfunction, as well as bursts of ROS production in the heart (6). Treating mice with DOX caused severe reductions in heart weight and myocardial cell size as compared with vehicle-treated (sham) mice (Figure 1, A and B). DOX also significantly increased cardiac levels of TRPC3 (Figure 1C). We therefore examined the involvement of TRPC3 channels in DOX-induced heart failure using TRPC3-deficient (TRPC3–/–) mice and their WT (TRPC3+/+) littermates. We confirmed that the TRPC3 protein band detected in TRPC3+/+ hearts was not present in TRPC3–/– hearts (Figure 1C). DOX elicited significant reductions in heart weight and myocardial cell size in TRPC3+/+ mice, but these effects were almost completely suppressed in TRPC3–/– mice (Figure 1, D and E). Likewise, DOX-induced reductions of LV systolic and diastolic functions observed in TRPC3+/+ mice were canceled in TRPC3–/– mice (Figure 1, F and G and Tables 1 and 2). In addition, production of malondialdehyde (MDA), an index of oxidative stress, was significantly lower in DOX-treated TRPC3–/– than TRPC3+/+ hearts (Figure 1H). We confirmed that the TRPC6 mRNA expression level was never compensatively upregulated in TRPC3–/– hearts (Supplemental Figure 1; supplemental material available online with this article; https://doi.org/10.1172/jci.insight.93358DS1). These results clearly indicate that TRPC3 participates in DOX-induced myocardial atrophy, oxidative stress, and LV dysfunction in mice.

**Deletion of TRPC3 suppresses hypoxic stress–mediated upregulation of Nox2 and myocardial atrophy in DOX-treated mice.** Using hypoxyprobe, we found that DOX induced hypoxic stress in cardiac tissue (Figure 2A), which in turn led to increases in the abundance of hypoxia-inducible factor 1α (HIF1α) and heme oxygenase-1 (HO-1) in mouse hearts (Figure 2, B and C). HIF1α is known to enhance production of ROS by increasing expression of Nox2 (28), and DOX also increased the abundance of Nox2 (Figure 2, B–D). The upregulation of Nox2 expression correlated well with the reduction in heart weight caused by DOX treatment (Figure 2D). This DOX-induced cardiac atrophy and Nox2 upregulation were significantly diminished in TRPC3–/– mice, and the degree to which myocardial atrophy was suppressed by TRPC3 deletion correlated well with the Nox2 expression (Figure 2, E–G). On the other hand, TRPC3 deletion had no
impact on the increased abundance of HIF1α. These data suggest that TRPC3 participates in hypoxic stress–mediated upregulation of Nox2 during DOX treatment, which leads to ROS-dependent cardiac atrophy and LV dysfunction in mice.

**Formation of a TRPC3-Nox2 complex contributes to DOX-induced cardiomyocyte atrophy.** TRPC3 has two functions: activation of local Ca2+ influx–mediated Ca2+ signaling induced by physicochemical stimulation and activation of context-dependent ROS signaling through physical interaction with Nox2 (29, 30). We used neonatal rat cardiomyocytes (NRCMs) to determine whether functional coupling between TRPC3 and Nox2 is required for DOX-induced myocardial atrophy. DOX treatment decreased the size of NRCMs. TRPC3 knockdown suppressed this cardiomyocyte atrophy (Figure 3A) and increased 2',7'-dichlorofluorescein diacetate (DCFH2DA) fluorescence intensity (Figure 3B). Despite the high sensitivity of DCFH2DA to ROS, this probe is limited to use for measuring intracellular ROS (31). Therefore, we used a genetically encoded fluorescent sensor for hydrogen peroxide, Hyper, and confirmed that a DOX-induced increase in Hyper fluorescence intensity was suppressed by the treatment with siRNA for TRPC3 or Nox2 in H9c2 cardiac myoblasts (Supplemental Figure 2A). We previously reported that TRPC3 interacts with Nox2 via a TRPC3-specific C-terminal region — which protects Nox2 from proteasome-dependent degradation, leading to its stabilization — and that overexpression of a GFP-fused TRPC3-C terminal fragment peptide (C3-C-GFP) interfered with the interaction between Nox2 and TRPC3 without reducing TRPC3 channel activity (20) (Figure 3C).
us to test the effect of C3-C-GFP on DOX-induced cardiomyocyte atrophy. To avoid the overlap of GFP fluorescence from DCF fluorescence, ROS production was measured using H9c2 cells expressing HA-tagged TRPC3-C terminal fragment peptide (C3-C-HA) instead of C3-C-GFP. We observed that C3-C-GFP and C3-C-HA expression significantly reduced the ROS production, cell shrinkage, Nox2 upregulation, and increase of the interaction between Nox2 and TRPC3 (Figure 3, D–F and Supplemental Figure 2B). These results suggest that functional coupling between TRPC3 and Nox2 is essential for DOX-induced cell atrophy and ROS production in cardiac myocytes.

Cardiomyocyte-specific expression of C3-C-GFP suppresses DOX-induced cardiomyopathy. We also examined whether cardiomyocytes are cellular origins of DOX-induced cardiomyopathy in vivo. Using a cardiomyocyte-specific, cardiac troponin-T promoter–driven, C3-C-GFP–encoding adeno-associated virus, C3-C-GFP was selectively expressed in cardiomyocytes in C57BL/6J mice. Compared with control (GFP-expressing) mice, the C3-C-GFP–expressing mice showed significant suppression of the myocardial atrophy, LV dysfunction, and oxidative stress normally seen in DOX-treated mice (Figure 4, A–E, and Tables 3 and 4). We also confirmed that C3-C-GFP was widely expressed in nearly all cardiomyocytes of the LV myocardium (Figure 4F). C3-C-GFP expression significantly suppressed the increases of Nox2 and TRPC3 protein abundances in DOX-treated hearts (Figure 4G). These results suggest that cardiomyocytes are a cellular origin of the cardiac atrophy mediated via the TRPC3-Nox2 axis in DOX-treated mice.

Pharmacological inhibition of TRPC3-Nox2 interaction suppresses DOX-induced cardiomyopathy. The above results led us to consider that a compound that inhibits the TRPC3-Nox2 interaction could be an ideal

<table>
<thead>
<tr>
<th>Table 1. Echocardiographic parameters of TRPC3+/+ and TRPC3–/– mice</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
</tr>
<tr>
<td></td>
</tr>
<tr>
<td>Mortality</td>
</tr>
<tr>
<td>BW (g)</td>
</tr>
<tr>
<td>HR (bpm)</td>
</tr>
<tr>
<td>IVSd (mm)</td>
</tr>
<tr>
<td>LVPWd (mm)</td>
</tr>
<tr>
<td>LVIdd (mm)</td>
</tr>
<tr>
<td>LVIds (mm)</td>
</tr>
<tr>
<td>EF (%)</td>
</tr>
<tr>
<td>FS (%)</td>
</tr>
</tbody>
</table>

BW, body weight; HR, heart rate; IVSd, interventricular septum diastolic thickness; LVPWd, left ventricular posterior wall diastolic thickness; LVIdd, left ventricular internal diameter at end-diastole; LVIds, left ventricular internal diameter at end-systole; EF, ejection fraction; FS, fractional shortening. Data are shown as mean ± SEM. Significance was determined by one-way ANOVA followed by Tukey’s comparison test. aP < 0.05 vs. TRPC3+/+ Sham, TRPC3–/– Sham. bP < 0.05 vs. TRPC3+/+ Sham. cP < 0.01 vs. TRPC3+/+ Sham, TRPC3–/– Sham, TRPC3–/– DOX.

<table>
<thead>
<tr>
<th>Table 2. Hemodynamic parameters of TRPC3+/+ and TRPC3–/– mice measured by left ventricular catheter</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
</tr>
<tr>
<td>HR (bpm)</td>
</tr>
<tr>
<td>LVEESP (mmHg)</td>
</tr>
<tr>
<td>LVEDP (mmHg)</td>
</tr>
<tr>
<td>LVPd/dt max (mmHg/s)</td>
</tr>
<tr>
<td>LVPd/dt min (mmHg/s)</td>
</tr>
<tr>
<td>τ (msec)</td>
</tr>
</tbody>
</table>

HR, heart rate; LVEESP, left ventricular end-systolic pressure; LVEDP, left ventricular end-diastolic pressure; LVPd/dt min, left ventricular maximal rate of pressure development; LVPd/dt max, left ventricular maximal rate of decay of pressure; τ, monoexponential time constant of relaxation. Data are shown as mean ± SEM. Significance was determined by one-way ANOVA followed by Tukey’s comparison test. aP < 0.01 vs. TRPC3+/+ Sham, TRPC3–/– Sham. bP < 0.05 vs. TRPC3+/+ Sham, TRPC3–/– Sham, TRPC3+/+ DOX.
drug for the treatment of heart failure. We therefore assessed the effect on the TRPC3-Nox2 interaction of six compounds previously reported to selectively inhibit TRPC3 channel activity (Supplemental Figure 3) (32–34). Among these, pyrazole-3 (Pyr3) most effectively suppressed the DOX-induced increase in Nox2 in NRCMs (Figure 5A). Pyr3 also suppressed DOX-induced cell shrinkage, Nox2 upregulation, and ROS production in either NRCMs or H9c2 cells (Figure 5, B–D, and Supplemental Figure 4). Prolonged treatment of NRCMs with Pyr3 reduced the basal abundance of Nox2 in a concentration-dependent manner (Figure 5E) without affecting TRPC3 and Nox2 mRNA levels (Figure 5G), and Pyr3 inhibited the interaction of TRPC3 with Nox2 (Figure 5F). Pyr3 thus acted as a potent inhibitor that suppressed the TRPC3-Nox2 interaction and DOX-induced toxicity in cardiomyocytes. We further examined whether Pyr3 directly suppresses the interaction between Nox2 and TRPC3 using cell-free system. Treatment of cell lysates with Pyr3 reduced the complex formation between Nox2 and TRPC3 in a concentration-dependent manner, while Pyr3 never affected the interaction between C3-C-GFP and Nox2 (Figure 5, H and I). We previously reported that Pyr3 specifically binds to extracellular N-terminal region of TRPC3 (25); these results suggest that Pyr3 inhibits the TRPC3-Nox2 complex formation by allosterically changing conformation of TRPC3 to reduce the binding affinity to Nox2, rather than directly disturbing the interaction between TRPC3 and Nox2.
To test whether Pyr3 could suppress DOX-induced cardiac atrophy in vivo, osmotic pumps were used to i.p. infuse Pyr3 in C57BL/6J mice for 3 days prior to DOX treatment. The infusion of Pyr3 significantly attenuated DOX-induced decreases in LV systolic and diastolic function (Figure 5, J–L, and Tables 5 and 6), as well as the increase in MDA production and myocardial atrophy (Figure 5M and Supplemental Figure 5). These results suggest Pyr3 attenuates the DOX-induced myocardial atrophy in mice by destabilizing Nox2 through disruption of the TRPC3-Nox2 interaction.

The mechanism by which DOX induces cardiotoxicity is known to differ from its antitumor actions (35). It opens the possibility of establishing a strategy for reducing or preventing DOX-induced cardiotoxicity without interfering with its antitumor effects. We therefore examined the effect of Pyr3 on DOX-induced cytotoxicity in tumor cells. A549 cells, a cell line derived from human lung adenocarcinoma, were treated with Pyr3 before DOX treatment. DOX significantly suppressed A549 cell proliferation, and Pyr3 did not inhibit the DOX-induced antitumor effect (Supplemental Figure 6). Instead, Pyr3 selectively suppressed cardiotoxicity without diminishing the antitumor effect of DOX.

Hypoxia mediates the DOX-induced upregulation of the TRPC3-Nox2 complex in cardiomyocytes. Because hypoxic stress underlies the upregulation of TRPC3 and Nox2 induced by DOX (36), we also investigated the contribution made by TRPC3 channels to ROS production under hypoxia. The level of TRPC3 mRNA expression in DOX-treated hearts was 2-fold higher than that in sham hearts (Figure 6A). Similarly, levels of TRPC3 mRNA in NRCMs were increased 3-fold by hypoxic stress (Figure 6A). The level of HO-1 expression was increased under 1% hypoxia and was not affected by Pyr3 (Figure 6B). TRPC3 inhibition appears not to diminish hypoxic stress–induced gene expression. However, Nox2 expression was actually increased by hypoxia, and TRPC3 inhibition diminished the
hypoxia-induced Nox2 upregulation (Figure 6C). Consistent with this, siRNA-mediated silencing of TRPC3 and Nox2 significantly suppressed hypoxia-induced ROS production determined by dihydroethidium (DHE) intensity (Figure 6D). Treating NRCMs with Pyr3 or diphenyleneiodonium (DPI), a Nox inhibitor, also reduced hypoxia-induced ROS production (Supplemental Figure 7). These results suggest that hypoxia-mediated formation of a TRPC3-Nox2 complex underlies DOX-induced ROS production in rat cardiomyocytes.

Physiological role of TRPC3 in LV diastolic function in mouse hearts. Finally, we used a pressure-volume catheter to assess whether TRPC3 is involved in determining physiological LV contractility and flexibility. As the basal Nox2 protein abundance was slightly but significantly reduced in TRPC3–/– mouse hearts (Figure 2F), the physiological role of TRPC3 was also expected to mediate mechanical stress–induced Nox2-dependent ROS signaling by stabilizing Nox2 protein in heart. Basal LV systolic function and heart rate were similar in TRPC3+/+ and TRPC3–/– mice (Tables 1 and 2). The slope of the end-systolic pressure volume relationship (Ees), an index of myocardial contractility, was similar in TRPC3–/– and TRPC3+/+ hearts. By contrast, the slope of the end-diastolic pressure volume relationship β coefficient (EDPVRβ), an index of ventricular stiffness, was significantly lower in TRPC3–/– than TRPC3+/+ hearts (Figure 7, A–C). Moreover, under acute volume loading, TRPC3–/– mice exhibited significantly larger LV stroke volumes than TRPC3+/+ mice (Figure 7D). The ability of the heart to change its force of contraction in response to changes in venous return (i.e., the Frank-Starling mechanism) was also significantly greater in Pyr3-treated mice (Figure 7E). These results indicate that TRPC3 positively regulates LV stiffness, and inhibition or reduced expression of TRPC3 enhances myocardial flexibility in mice.

DOX-induced cardiomyopathy is reportedly attenuated by exercise (37), and exercise causes physiological hypertrophic growth of the heart (1). We therefore examined the degree to which the TRPC3-Nox2 complex mediates the three major types of cardiac plasticity: atrophy, physiological hypertrophy, and pathological hypertrophy. Voluntarily exercising mice showed a marked (10-fold) increase in daily walking distance compared with nonexercising mice (Figure 7F). Voluntary exercise for 4 weeks also caused a significant increase in heart weight with preservation of cardiac function, suggesting the development of physiological hypertrophy (Figure 7G and Table 7). Under acute volume loading, voluntarily exercising mice showed a significantly greater increase in LV stroke volume than nonexercised mice (Figure 7H). Voluntary exercise decreased the cardiac protein abundances of TRPC3 and Nox2 (Figure 7I), correlating well with the magnitude of physiological
hypertrophy in mice (Figure 7J). Voluntary exercise had no impact on the expression levels of HIF1α and HO-1 proteins (Figure 7I), or on TRPC3 and Nox2 mRNAs in heart (Supplemental Figure 8), indicating that the downregulation of Nox2 is mediated by a posttranslational mechanism. These results strongly suggest that formation of a TRPC3-Nox2 complex negatively regulates LV compliance and flexibility, which increases the risk of pathological cardiac remodeling induced by volume load.

Discussion

Although cardiac plasticity is an intrinsic compensatory mechanism to reduce hemodynamic workload, pathological atrophy or hypertrophy actually confers risk for chronic heart failure. By contrast, physiological hypertrophy, caused by hemodynamic load during postnatal growth, exercise, and pregnancy, is also a major cardiac plasticity but one in which cardiac contractility is maintained. We first demonstrated that inhibition of TRPC3 prevents DOX-induced myocardial atrophy by suppressing Nox2-mediated amplification of ROS signaling in cardiomyocytes. We also found that voluntary exercise causes hypertrophy with preserved cardiac contractility by enhancing LV compliance and flexibility through destabilization of Nox2 (Figure 7, G–J, and Table 7). Taken together with our earlier observations made in a pressure-overloaded mouse heart model (20, 27), our present findings, to our knowledge, provide a new concept that a TRPC3-Nox2 complex functions as a common major risk factor for chronic heart failure, driving pathological cardiac remodeling. Furthermore, they suggest that, by suppressing Nox2-mediated ROS production in cardiomyocytes, TRPC3 inhibition may be an effective strategy for reducing the risk of DOX-induced heart failure (Figure 7K).

Cardiac atrophy is mainly caused by sarcopenia associated with aging, cachexia, and malnutrition (2). Although it is not known whether these three major causes are involved in DOX-induced cardiac atrophy, our findings suggest the possible contribution of the TRPC3-Nox2 complex to the induction of cardiac atrophy. Future studies using animal models of sarcopenia or cancer cachexia will be required to establish the pathological role of the TRPC3-Nox2 complex formation in the cardiac atrophy seen with those ailments.

### Table 3. Echocardiographic parameters in GFP- and C3-C-GFP–expressing mice

<table>
<thead>
<tr>
<th></th>
<th>GFP Sham (n = 6)</th>
<th>C3-C-GFP Sham (n = 6)</th>
<th>GFP DOX (n = 6)</th>
<th>C3-C-GFP DOX (n = 6)</th>
</tr>
</thead>
<tbody>
<tr>
<td>BW (g)</td>
<td>20.5 ± 0.4</td>
<td>21.0 ± 1.0</td>
<td>17.3 ± 1.8**</td>
<td>19.8 ± 2.1</td>
</tr>
<tr>
<td>HR (bpm)</td>
<td>549 ± 21</td>
<td>547 ± 25</td>
<td>524 ± 24</td>
<td>528 ± 28</td>
</tr>
<tr>
<td>IVSd (mm)</td>
<td>0.7 ± 0.0</td>
<td>0.8 ± 0.0</td>
<td>0.7 ± 0.0</td>
<td>0.7 ± 0.0</td>
</tr>
<tr>
<td>LVPWd (mm)</td>
<td>0.8 ± 0.0</td>
<td>0.9 ± 0.0</td>
<td>0.6 ± 0.0**</td>
<td>0.7 ± 0.0</td>
</tr>
<tr>
<td>LVIDd (mm)</td>
<td>2.9 ± 0.1</td>
<td>2.8 ± 0.1</td>
<td>2.8 ± 0.1</td>
<td>2.9 ± 0.1</td>
</tr>
<tr>
<td>LVIDs (mm)</td>
<td>1.3 ± 0.1</td>
<td>1.4 ± 0.1</td>
<td>1.7 ± 0.1**</td>
<td>1.4 ± 0.1</td>
</tr>
<tr>
<td>EF (%)</td>
<td>89.5 ± 1.0</td>
<td>88.2 ± 1.0</td>
<td>76.7 ± 1.5**</td>
<td>86.0 ± 1.0</td>
</tr>
<tr>
<td>FS (%)</td>
<td>53.2 ± 1.5</td>
<td>51.8 ± 2.0</td>
<td>39.7 ± 1.3**</td>
<td>49.2 ± 1.3</td>
</tr>
</tbody>
</table>

Data are shown as mean ± SEM. Significance was determined by one-way ANOVA followed by Tukey’s comparison test. **P < 0.05 vs. GFP Sham, C3-C-GFP Sham. *P < 0.05 vs. GFP Sham, C3-C-GFP Sham, C3-C-GFP DOX.

### Table 4. Hemodynamic parameters in GFP- and C3-C-GFP–expressing mice measured by left ventricular catheter

<table>
<thead>
<tr>
<th></th>
<th>GFP Sham (n = 5)</th>
<th>C3-C-GFP sham (n = 5)</th>
<th>GFP DOX (n = 6)</th>
<th>C3-C-GFP DOX (n = 5)</th>
</tr>
</thead>
<tbody>
<tr>
<td>HR (bpm)</td>
<td>458 ± 22</td>
<td>467 ± 20</td>
<td>419 ± 33</td>
<td>409 ± 23</td>
</tr>
<tr>
<td>LVESP (mmHg)</td>
<td>113 ± 5</td>
<td>126 ± 10</td>
<td>106 ± 7</td>
<td>126 ± 7</td>
</tr>
<tr>
<td>LVEDP (mmHg)</td>
<td>3.6 ± 0.5</td>
<td>5.9 ± 1.1</td>
<td>12.7 ± 1.0**</td>
<td>3.5 ± 1.2</td>
</tr>
<tr>
<td>LVdP/dt max (mmHg/s)</td>
<td>11,410 ± 1,053</td>
<td>11,930 ± 467</td>
<td>6,823 ± 778**</td>
<td>13,250 ± 860</td>
</tr>
<tr>
<td>LVdP/dt min (mmHg/s)</td>
<td>7,802 ± 899</td>
<td>8,203 ± 1,202</td>
<td>4,030 ± 610**</td>
<td>5,941 ± 859</td>
</tr>
<tr>
<td>τ (msec)</td>
<td>9.4 ± 0.8</td>
<td>11.7 ± 1.1</td>
<td>23.5 ± 5.1**</td>
<td>11.0 ± 0.4</td>
</tr>
</tbody>
</table>

Data are shown as mean ± SEM. Significance was determined by one-way ANOVA followed by Tukey’s comparison test. **P < 0.01 vs. GFP Sham, C3-C-GFP Sham, C3-C-GFP DOX. *P < 0.05 vs. GFP Sham, C3-C-GFP Sham, C3-C-GFP DOX.
Anthracyclines, including DOX, are inexpensive, highly effective antimutu drugs used to treat breast cancer, lung cancer, lymphoma, and leukemia. However, the cardiotoxicity caused by anthracyclines limits their utility. In the present study, we revealed the molecular mechanism underlying the initiation of DOX-induced cardiomyopathy and suggest that inhibiting the TRPC3-Nox2 complex could be an effective strategy to prevent/reverse DOX-induced chronic heart failure. In this study, we demonstrated that Pyr3, a compound that can inhibit TRPC3 channel activity (25), potently prevented DOX-induced myocardial atrophy and LV dysfunction in mice (Figure 5, Tables 5 and 6, Supplemental Figure 5). Pyr3 was also found to suppress DOX-induced ROS production by destabilizing Nox2 protein through disrupting the TRPC3-Nox2 complex formation in NRCMs (Figure 5). As Pyr3 reportedly inhibits several other channels, such as CRAC channels and TREK and TASK2 potassium channels (24, 38), it cannot be simply assumed that the effects of Pyr3 are mediated by disrupting the TRPC3-Nox2 interaction. However, our results clearly show that pharmacological effects by Pyr3 are quite similar to those by knockdown or genetic deletion of TRPC3 in cardiomyocytes. Pyr3 had no impact on upstream factors regulating the TRPC3-Nox2 complex formation, such as upregulation of HIF1α protein and basal mRNA levels of TRPC3 and Nox2. We also revealed that Pyr3 directly acted on the complex formation between Nox2 and full-length TRPC3, but not the C-terminal fragment of TRPC3, in a cell-free system (Figure 5, H and I), suggesting the involvement of allosteric conformational changes in TRPC3 for disruption of the TRPC3-Nox2 complex. These findings strongly support the idea that Pyr3 prevents DOX-induced Nox2 upregulation by disrupting the TRPC3-Nox2 complex formation. Other approaches that reportedly prevent or mitigate DOX-induced dilated cardiomyopathy include exercise (37); treatment with cilostazol (39), an inhibitor of phosphodiesterase type 3 (PDE3), antioxidants (10), or iron chelators (40); and expression of Gαq-inhibitory peptide in cardiomyocytes (41). We previously observed that inhibition of PDE3 (42) or Gαq-mediated phosphatidylinositol turnover (22) attenuates TRPC3/6 channel activity, and we observed here that exercise reduces TRPC3 expression in mouse hearts (Figure 7I). These findings are consistent with our new finding that TRPC3 in cardiomyocytes is a critical mediator of DOX-induced dilated cardiomyopathy.

Although the pathological significance of the TRPC3-Nox2 complex in DOX-induced myocardial atrophy in mice is evident, the physiological role of TRPC3 remains elusive. Redox/cysteine modification of proteins that regulate Ca2+ handling has been reported to affect cardiac contractility (43), including Nox2-dependent ROS signaling (X-ROS signaling) in the diastolic filling heart (19). Actually, TRPC3-deficient mice had less protein abundances of TRPC3 and Nox2, as well as less basal MDA production in NRCMs with or without Pyr3 pretreatment (36 h, n = 3). Pyr3 had no impact on upstream factors regulating the TRPC3-Nox2 complex formation, such as upregulation of HIF1α protein and basal mRNA levels of TRPC3 and Nox2. We also revealed that Pyr3 directly acted on the complex formation between Nox2 and full-length TRPC3, but not the C-terminal fragment of TRPC3, in a cell-free system (Figure 5, H and I), suggesting the involvement of allosteric conformational changes in TRPC3 for disruption of the TRPC3-Nox2 complex. These findings strongly support the idea that Pyr3 prevents DOX-induced Nox2 upregulation by disrupting the TRPC3-Nox2 complex formation. Other approaches that reportedly prevent or mitigate DOX-induced dilated cardiomyopathy include exercise (37); treatment with cilostazol (39), an inhibitor of phosphodiesterase type 3 (PDE3), antioxidants (10), or iron chelators (40); and expression of Gαq-inhibitory peptide in cardiomyocytes (41). We previously observed that inhibition of PDE3 (42) or Gαq-mediated phosphatidylinositol turnover (22) attenuates TRPC3/6 channel activity, and we observed here that exercise reduces TRPC3 expression in mouse hearts (Figure 7I). These findings are consistent with our new finding that TRPC3 in cardiomyocytes is a critical mediator of DOX-induced dilated cardiomyopathy.

Table 5. Echocardiographic parameters of mice with Pyr3 administration.

<table>
<thead>
<tr>
<th>Veh sham (n = 6)</th>
<th>Pyr3 sham (n = 6)</th>
<th>Veh DOX (n = 9)</th>
<th>Pyr3 DOX (n = 9)</th>
</tr>
</thead>
<tbody>
<tr>
<td>BW (g)</td>
<td>23.9 ± 0.5</td>
<td>23.5 ± 0.5</td>
<td>20.1 ± 0.5</td>
</tr>
<tr>
<td>HR (bpm)</td>
<td>471 ± 10</td>
<td>429 ± 14</td>
<td>425 ± 8</td>
</tr>
<tr>
<td>LVIDd (mm)</td>
<td>0.8 ± 0.0</td>
<td>0.8 ± 0.0</td>
<td>0.7 ± 0.0</td>
</tr>
<tr>
<td>LVPWd (mm)</td>
<td>1.0 ± 0.0</td>
<td>1.0 ± 0.1</td>
<td>0.7 ± 0.0</td>
</tr>
<tr>
<td>LVDD (mm)</td>
<td>2.6 ± 0.1</td>
<td>2.6 ± 0.0</td>
<td>2.6 ± 0.1</td>
</tr>
<tr>
<td>LVDs (mm)</td>
<td>1.1 ± 0.1</td>
<td>1.1 ± 0.1</td>
<td>1.6 ± 0.1</td>
</tr>
<tr>
<td>EF (%)</td>
<td>90.5 ± 1.7</td>
<td>91.8 ± 1.1</td>
<td>75.9 ± 2.2</td>
</tr>
<tr>
<td>FS (%)</td>
<td>56.8 ± 2.7</td>
<td>58.0 ± 2.3</td>
<td>39.4 ± 2.0</td>
</tr>
</tbody>
</table>

Data are shown as mean ± SEM. Significance was determined by one-way ANOVA followed by Tukey’s comparison test. *P < 0.05 vs. Veh Sham, Pyr3 Sham. **P < 0.01 vs. Veh Sham, Pyr3 Sham, Pyr3 DOX. ‡P < 0.05 vs. Veh Sham, Pyr3 Sham, Pyr3 DOX.
content in the heart (Figure 1, C and H, and Figure 2F). This implies that the TRPC3-Nox2 complex formation underlies TRPC3-mediated negative regulation of LV compliance in both physiological and pathological conditions. This idea is supported by the fact that voluntary exercise, an effective treatment to prevent the heart from DOX-induced cardiotoxicity, decreased protein abundances of Nox2 and TRPC3 (Figure 7I). As the phenotypic change of mouse heart induced by voluntary exercise seems consistent with those in TRPC3-deficient mice, we suggest that TRPC3 is a prerequisite for the basal expression of Nox2 because it prevents constitutive degradation in cardiac cells. Although Nox2-mediated ROS production induced by physiological stretching of cardiomyocytes (i.e., X-ROS signaling) is reported to participate in enhancement of cardiac contractility, Nox2 deletion has no impact on cardiac compliance compared with the impact by TRPC3 inhibition (13). This indicates that Nox2-derived ROS signaling has two roles: one is to promote cardiac contractility through oxidative modification of Ca^{2+}-handling proteins, and the other is to suppress cardiac compliance through unknown mechanism(s). Our results suggest that the context-dependent ROS signaling mediated by the TRPC3-Nox2 complex plays a key role in cardiac stiffness. Future identification of a new TRPC3-mediated ROS target protein will elucidate its underlying mechanism.

The mechanism by which exercise downregulates TRPC3 and Nox2 is also obscure. As a DOX-induced increase in TRPC3 mRNA level was associated with upregulation of HIF1α, we first thought that exercise treatment downregulates TRPC3 and Nox2 mRNAs by suppressing hypoxic stress and/or HIF1α upregulation. However, exercise had no impact on basal HIF1α abundance, which was comparable with that in sedentary mice (Figure 7I). Exercise treatment never decreased the expression levels of TRPC3 and Nox2 mRNAs (Supplemental Figure 8), but it significantly reduced TRPC3 and Nox2 protein abundances (Figure 7I).
result strongly suggests the involvement of posttranslational modulation or modification of TRPC3 or Nox2 proteins in the TRPC3-Nox2 stable complex formation. Further studies for identification of some intrinsic molecule(s) that can disrupt the interaction between TRPC3 and Nox2 from exercised heart will be required.

All Ca2+ transporters present on the myocardial plasma membrane are generally thought to participate in positive inotropic and/or chronotropic responses to physiochemical stimuli. Although several reports, including one from our group, have demonstrated the pathological effects of TRPC channels in vivo in rodent hearts (44), TRPC channels are believed to be hardly expressed in normal hearts and to be uninvolved in normal cardiac physiology, as TRPC KO had no impact on cardiac contractility or heart rate in mice. In the context of our in vitro finding that TRPC3 mediates mechanical stress–induced background Ca2+ entry in cardiomyocytes (20), our present observation using TRPC3-deficient mice suggests TRPC3 contributes to the negative regulation of myocardial compliance and flexibility in response to volume load. We anticipate that these findings will provide the basis for a new strategy to maintain the cardiac performance while driving cardiac plasticity.

**Methods**

*Mice, chemicals, and reagents.* The 129Sv mice with homozygous deletion of the gene-encoding TRPC3 were provided by the National Institute of Environmental Health Sciences. Genotyping was performed using PCR the primers TRPC3-A (5′-GAATCCACCTGCTTACAACCATGTG-3′) and TRPC3-B (5′-GGTGGAGGTAACACACAGCTAAGCC-3′). The PCR was performed using Phusion High-Fidelity DNA polymerase (Thermo Scientific). C57BL/6J mice were purchased from Japan SLC Inc. Mice were maintained in a specific-pathogen–free area under a 12-hour/12-hour light/dark cycle. Male 7-week-old mice were administered DOX (15 mg/kg, Wako) i.p. and sacrificed 2 weeks later. Osmotic pumps (ALZET) for sustained administration of Pyr3 (30 μg/kg/day) were implanted i.p. 3 days before DOX administration. The TRPC3/6 blocker, GSK-417651A, was purchased from Focus Biomolecules. Other TRPC3/6 inhibitors were synthesized or made to order as described previously (25, 32–34). Cell lines (H9c2 and A549) were purchased from American Type Culture Collection (ATCC).

*Voluntary exercise experiment in mice.* Male 8-week-old mice (C57BL/6J) were exposed to voluntary exercise in a cage with a running wheel. The mice were allowed to use the voluntary running wheel for 4 weeks. Non-exercised (sedentary) mice were kept in identical cages with an immovable wheel. Spontaneous activity was recorded every minute and walking distances were summed up for 24 hours using Actimo-100 (Shinfactory).

*Transthoracic echocardiography and cardiac catheterization.* Echocardiography and cardiac catheterization were performed as described previously (15, 20). Echocardiography was performed using a Nemio-XG echocardiograph (Toshiba) with a 14-MHz transducer after anesthetizing the mice using a mixture of domitor (Zenoaq), midazolam (Sando), and butorphanol (Meiji Seika Pharma). LV hemodynamic parameters were assessed after DOX treatment in the anesthetized mice using a micronanometer catheter (Millar I.4F, SPR 671, Millar Instruments).

---

**Table 6. Hemodynamic parameters of mice with Pyr3 administration measured by left ventricular catheter**

<table>
<thead>
<tr>
<th></th>
<th>Veh Sham (n = 6)</th>
<th>Pyr3 Sham (n = 6)</th>
<th>Veh DOX (n = 7)</th>
<th>Pyr3 DOX (n = 7)</th>
</tr>
</thead>
<tbody>
<tr>
<td>HR (bpm)</td>
<td>392 ± 30</td>
<td>415 ± 16</td>
<td>352 ± 21</td>
<td>342 ± 8</td>
</tr>
<tr>
<td>LVESP (mmHg)</td>
<td>100 ± 6</td>
<td>116 ± 6</td>
<td>100 ± 8</td>
<td>115 ± 5</td>
</tr>
<tr>
<td>LVEDP (mmHg)</td>
<td>1.5 ± 0.4</td>
<td>1.3 ± 0.3</td>
<td>10.4 ± 0.7^a</td>
<td>2.4 ± 1.8</td>
</tr>
<tr>
<td>LVdP/dt_max (mmHg/s)</td>
<td>10,866 ± 1,441</td>
<td>12,467 ± 811</td>
<td>5,221 ± 535^a</td>
<td>9,643 ± 849</td>
</tr>
<tr>
<td>LVdP/dt_min (mmHg/s)</td>
<td>5,606 ± 599</td>
<td>5,823 ± 263</td>
<td>3,810 ± 449^a</td>
<td>5,510 ± 343</td>
</tr>
<tr>
<td>τ (msec)</td>
<td>10.9 ± 1.7</td>
<td>11.1 ± 1.0</td>
<td>23.4 ± 3.4^a</td>
<td>13.9 ± 1.4</td>
</tr>
</tbody>
</table>

Data are shown as mean ± SEM. Significance was determined by one-way ANOVA followed by Tukey’s comparison test. ^P < 0.01 vs. Veh Sham, Pyr3 Sham, Pyr3 DOX. *P < 0.05 vs. Veh Sham, Pyr3 Sham, Pyr3 DOX.

**Table 7. Echocardiographic parameters of mice with voluntary exercise**

<table>
<thead>
<tr>
<th></th>
<th>Sedentary (n = 6)</th>
<th>Exercise (n = 7)</th>
</tr>
</thead>
<tbody>
<tr>
<td>BW (g)</td>
<td>26.0 ± 0.3</td>
<td>25.3 ± 0.3</td>
</tr>
<tr>
<td>HR (bpm)</td>
<td>512 ± 20</td>
<td>481 ± 12</td>
</tr>
<tr>
<td>IVSd (mm)</td>
<td>0.8 ± 0.0</td>
<td>0.9 ± 0.0^a</td>
</tr>
<tr>
<td>LVPWd (mm)</td>
<td>0.8 ± 0.0</td>
<td>1.0 ± 0.0^a</td>
</tr>
<tr>
<td>LVIDd (mm)</td>
<td>2.7 ± 0.1</td>
<td>2.6 ± 0.1</td>
</tr>
<tr>
<td>LVIDs (mm)</td>
<td>1.4 ± 0.1</td>
<td>1.2 ± 0.1</td>
</tr>
<tr>
<td>EF (%)</td>
<td>85.2 ± 2.0</td>
<td>90.3 ± 1.5</td>
</tr>
<tr>
<td>FS (%)</td>
<td>48.7 ± 2.6</td>
<td>56.0 ± 2.2</td>
</tr>
</tbody>
</table>

Data are shown as mean ± SEM. Significance was determined by unpaired t test. ^P < 0.01 vs. Sedentary.
Figure 7. Roles of TRPC3-Nox2 complex in myocardial flexibility and plasticity. (A) Representative LV pressure-volume loops after inferior vena cava occlusion in TRPC3+/+ and TRPC3−/− mice. (B and C) Results of LV systolic indexes (end-systolic elastance; Ees) (B) and diastolic indexes (end-diastolic pressure-volume relationship β; EDPVRβ) (C) (n = 5). (D) Cardiac performance under volume overload reflecting the Frank-Starling law in TRPC3+/+ and TRPC3−/−.
**Voluntary exercise can induce physiological hypertrophy under hemodynamic load by reducing the TRPC3-Nox2 complex.**

Increases in the abundance of the TRPC3-Nox2 complex underlie pathological hypertrophy and atrophy. Voluntary exercise can induce physiological hypertrophy under hemodynamic load by reducing the TRPC3-Nox2 complex.

**In vivo pressure-volume analysis.** Pressure-volume analysis was performed according to established protocols (45, 46). Pyr3 (100 μg/kg/day) was administered using an osmotic minipump for 7 days before the analysis. Each mouse was anesthetized (induction: 3%-4% isoflurane, maintenance: 1%-2% isoflurane), intubated, and mechanically ventilated (MiniVent, model 845, Harvard Apparatus) with 100% O2. The animal was then placed on a temperature-controlled heating platform to maintain body temperature. After accessing the internal jugular vein using a 30-gauge needle connected to polyethylene tubing, 12.5% albumin in normal saline was infused at 300 μl/hour after an initial 50-μl bolus to counteract the peripheral vasodilation and hypotension caused by the anesthetia (46). The diaphragm and ribs were then carefully dissected using an electrosurgery device (Bovie Medical), the LV apex was punctured with a 27-gauge needle, and a catheter (Millar 1.4F, SPR 839, Millar Instruments) was introduced through the puncture. Mice were allowed to stabilize for 5 minutes, after which transient occlusion of inferior vena cava was performed to calculate Ees for evaluation of load-independent systolic function, and EDPVRβ was determined to evaluate end-diastolic stiffness (45, 46). Ees was calculated as ESP = Ees × ESV + Vo, and EDPVRβ was calculated as EDP = α × exp β × EDV, where ESP represents end-systolic pressure, ESV represents end-systolic volume, Vo represents calculated pressure when LV volume equals 0, α represents the stiffness and scaling coefficient, and EDV represents end-diastolic volume. To investigate the cardiac responsiveness to volume overload, 12.5% albumin was transfused at 2 ml/hour. Hypertonic saline (10% NaCl, 10 μl) was injected to calculate parallel conductance.

**AAV-mediated expression of TRPC3 C-terminal fragment in mouse hearts.** The Nox2-interacting fragment from the TRPC3 C-terminus (C3-C fragment) was PCR amplified and cloned into pEGFP-N1 vector. The EGFP-fused C3-C fragment (C3-C-GFP) cassette under the control of cardiac troponin T promoter was subcloned into pZac2.1 (Penn Vector). Viral vectors encoding EGFP or C3-C-GFP were generated as described previously (47). The AAV vectors (1 × 10^{10} genomic copies) were i.p. injected into male 6-day-old C57BL/6J mice. Expression of C3-C-GFP in the LV myocardium was verified using a fluorescence microscope (BZ-X710, Keyence).

**Morphological analysis.** Hearts were removed from mice, washed in PBS to remove the blood, and fixed by 10% neutral buffered formalin. To assess (cross-sectional area) CSA in the cardiomyocytes, sections were stained with wheat germ agglutinin (WGA) conjugated with Alexa Fluor 488 (Invitrogen) and observed using a fluorescence microscope. Five LV regions were selected at random for each specimen, and the average values were calculated using ImageJ software (NIH).

**Assessment of oxidative stress in vivo.** Total cardiac MDA was assessed in mice using a Bioxytech MDA-586 kit (OxisResearch) according to the manufacturer’s instructions, as described previously (15, 20). Briefly, frozen mouse heart samples were homogenized in potassium phosphate extraction (KPE) buffer (pH 7.5) containing 100 mM potassium phosphate, 5 mM EDTA, 0.1% Triton X-100, 0.6% sultosalicilic acid, and 5 mM butylated hydroxytoluene (OxisResearch). The resultant lysates were centrifuged at 20,400 g for 10 minutes at 4°C. Samples of the supernatant were then allowed to react with N-methyl-2-phenylindole for 1 hour at an acidic pH at 45°C, after which the samples were centrifuged at 20,400 g for 10 minutes at 4°C and read at 586 nm using a Spectra Max i3 (Molecular Devices).

**Real-time PCR.** Total RNA was isolated from frozen mouse heart samples using an RNeasy Fibrous Tissue Mini Kit (Qiagen) or from cardiomyocytes using an RNeasy Mini Kit (Qiagen) according to the manufacturer’s instructions. Quantitative real-time PCR was performed using an ABI PRISM 7500 Real-Time PCR System (Applied Biosystems) with OneStep RT-PCR Kit (Qiagen) or Power SYBR green PCR master mix (Applied Biosystems) according to the manufacturer’s instructions. The information of Taqman probes and primer sets were as previously described (20).

**Western blotting and IP.** Mouse hearts were homogenized in RIPA buffer containing 0.1% SDS, 0.5% sodium deoxycholate, 1% NP-40, 150 mM NaCl, 50 mM Tris-HCl (pH 7.4), and protease inhibitor cocktail (Nacalai). Samples (10 μg) were then fractionated by SDS-PAGE and transferred onto PVDF membranes (Millipore). The objective proteins were detected using the indicated antibodies. After incubation
with the secondary antibody, the bands were visualized using Western Lightning Plus ECL (PerkinElmer). Images were captured with an ImageQuant LAS 4000 (GE Healthcare Life Science) and quantified using ImageQuant TCL software (GE Healthcare Life Sciences). Primary antibodies against Nox2 (1:1,000, sc-130543, Santa Cruz Biotechnology Inc.), αβ-tubulin (1:2,000, 2143, Cell Signaling Technology), GAPDH (1:2,000, sc-25778, Santa Cruz Biotechnology Inc.), HIF1α (1:1,000, NB100-105, Novus Biologicals), and HO-1 (1:1,000, ADI-SPA-895, Enzo) were used.

To analyze expression of endogenous Nox2 in vitro and in vivo, the total membrane fraction was isolated as described previously (30). To investigate the abundance of TRPC3, endogenous TRPC3 was concentrated by IP. Briefly, TRPC3 proteins were immunoprecipitated from mouse heart lysates (500 μg total protein), cell lysates from Pyr3-treated NRCM, or cell lysates from DOX-treated H9c2 cells using anti-TRPC3 antiserum (1.5 μg) obtained from guinea pig (20) and protein G sepharose beads (GE Healthcare). After incubation for 24 hours, the immune complexes were washed 3 times with lysis buffer, suspended in SDS sample buffer containing 0.1 M DTT, and incubated for 1 hour at room temperature. TRPC3 was detected with rabbit anti-TRPC3 antibody (1:1,000, ACC-016, Alomone). For the IP with Pyr3 in a cell-free system, HEK293 cells expressing Flag-tagged Nox2 and either TRPC3-EGFP or C3-C-GFP were lysed in 1% Triton X-100, 140 mM NaCl, 1 mM EDTA, 20 mM Tris-HCl (pH 7.4), and protease inhibitor cocktail (Nacalai). Equal amounts of cell lysates were aliquoted to 5 tubes and incubated with Flag M2 agarose for 12 hours to immunoprecipitate Flag-Nox2 in the presence of Pyr3 at the indicated concentration. Immune complexes were washed 3 times with the lysis buffer and eluted with 0.1 mg/ml Flag peptide (Sigma-Aldrich) in lysis buffer.

Detection of hypoxia using a hypoxyprobe kit. Hypoxia in mouse hearts was detected using a hypoxyprobe kit (Hypoxyprobe HPI Inc.) according to the manufacturer’s instructions. Hypox probe solution (60 mg/kg) was i.p. injected 60 minutes prior to sacrifice. Frozen tissue was cut into 4-μm-thick sections and incubated first in 1% BSA in PBS for 1 hour at 4°C and then with anti-hypoxyprobe antibody (1:50 dilution) in 1% BSA in PBS overnight at 4°C. After washing, the sections were incubated with secondary antibody for 1 hour at room temperature. Sections were then observed using a confocal laser microscopy, and fluorescence intensity was analyzed using ImageJ (NIH) software.

Isolation of cardiomyocytes from neonatal rat. Cardiomyocytes were isolated from neonatal rats as described previously (48). For TRPC3 and Nox2 knockdown, the cells were transfected with siRNAs (20 nM; ref. 20) for 72 hours using Lipofectamine RNAiMAX (Thermo Fisher scientific).

Measurements of cell size and ROS production in rat cardiac cells. NRCMs and H9c2 cardiac myoblasts were visualized by Alexa Fluor 488 Phalloidin (Thermo Fisher Scientific), and cell surface area was observed using a fluorescence microscope and analyzed in at least 50 cells in each experiment using ImageJ software. Production of ROS in NRCMs and H9c2 cardiac myoblasts was measured using DCF-H$_2$DA as described previously (15). Cells were loaded with DCFH$_2$DA (10 μM) for 10 minutes at 37°C. The DCF fluorescence excited by illumination at 488 nm was observed at room temperature using a confocal laser microscope. For the detection of hydrogen peroxide, H9c2 cells were transfected with cDNA encoding Hyper (Evrogen) by electroporation (2 pulses of 1,200 mV for 10 milliseconds; Neon Transfection System, Invitrogen). After 24-hour recovery, cells were plated on glass-base dishes, cultured for another 24 hours, and treated with DOX for 12 hours. Hyper was excited at 488 nm, and emissions were collected at 536 nm. Images of single cell fluorescence were collected (40 cells per experiment) and analyzed by Metafluor (Molecular Device). DHE staining of siRNA-transfected NRCMs was performed by the incubation with DHE (2 μM) for 1 hour. Cells (2 × 10$^5$ cells/35 mm dish) were fixed in 2% paraformaldehyde for 10 minutes to stop the reaction. To avoid the possibility of cell population dependency, the averaged DHE fluorescence intensity was introduced by averaging the DHE florescence intensity of 40–50 cells in each experiment.

Tumor cell viability. Tumor cell viability was analyzed using an MTT assay kit (Sigma Aldrich). A549 cells were treated with DOX (10 μM) with or without Pyr3 (3 μM) for 24 hours. Cells were further incubated in MTT-containing (50 μg/ml) medium for 6 hours and then lysed with DMSO. The absorbance at 570 nm was measured using Spectra Max i3.

Statistics. All results are presented as the mean ± SEM from at least 3 independent experiments. Statistical comparisons were made using unpaired t tests for two-group comparisons or using one-way ANOVA with Tukey’s post hoc test or two-way ANOVA with Bonferroni’s post hoc test for comparisons among 3 or more groups.
**Study approval.** All protocols using animals were approved by the ethics committees at the National Institutes of Natural Sciences or the Animal Care and Use Committee and were performed according to the institutional guidelines concerning the care and handling of experimental animals.

**Author contributions**

MN supervised and conceived the project; TS designed experiments, interpreted data, and prepared the manuscript; TS, TNT, T. Ide, AN, and RM performed experiments; SO, SH, T. Ito, NK, KU, HS, and YM contributed reagents/analytic tools; and MN edited the manuscript.

**Acknowledgments**

TRPC3−/− mice were provided from Lutz Birnbaumer of the NIEHS, Research Triangle Park, North Carolina, USA. This research was funded by grants from the Japan Science and Technology Agency; Precursory Research for Embryonic Science and Technology Program (JPMJPR1336 to MN); Grants-in-Aid for Scientific Research (25293018 and 16776376 to MN); Scientific Research on Innovative Areas (Research in a Proposed Research Area ‘Oxygen Biology’ [14430291 to MN] and ‘Living in Space’ [16H01656 to TNT]); and Platform for Drug Discovery, Informatics, and Structural Life Science (to MN) from the Ministry of Education, Culture, Sports, Science, and Technology of Japan.

Address correspondence to: Motohiro Nishida, Division of Cardiocirculatory Signaling, Okazaki Institute for Integrative Bioscience (National Institute for Physiological Sciences), National Institutes of Natural Sciences, Higashiyama 5-1, Myodaiji-cho, Okazaki, Aichi 444-8787, Japan. Phone: 81.564.59.5560; Email: nishida@nips.ac.jp.

22. Onohara N, et al. TRPC3 and TRPC6 are essential for angiotensin II-induced cardiac hypertrophy. *EMBO J.*


