Chapter from the book *Cardiac Arrhythmias - New Considerations*
Downloaded from: http://www.intechopen.com/books/cardiac-arrhythmias-new-considerations

Interested in publishing with InTechOpen?
Contact us at book.department@intechopen.com
The Pathophysiological Implications of TRP Channels in Cardiac Arrhythmia

Ryuji Inoue, Duan Yubin, Hu Yaopeng and Jun Ichikawa

Department of Physiology, Fukuoka University School of Medicine
Japan

1. Introduction

The cardiac arrhythmia is a common cause of morbidity and mortality in the present world, especially in developed countries. The etiology of cardiac arrhythmia is quite broad involving both hereditary and secondary backgrounds. An increasing number of rare but lethal arrhythmogenic mutations have been identified by genome-wide association assays in genes associated with cardiac excitation, conduction and morphogenesis. Such well-characterized examples include causative mutations in voltage-dependent Na⁺, Ca²⁺ or K⁺ channel (Naᵥ, Caᵥ, Kᵥ) genes for the prolongation of QT interval which sometimes result in premature sudden death and may act as predisposing factors to drug-induced arrhythmia (Chen et al., 1998; Roden, 2004; Saenen & Vrints, 2008; Towbin & Vatta, 2001; Zareba et al., 2004). Furthermore, some of heart malformations (e.g. atrial septal defect) and idiopathic cardiomyopathy (e.g. dilated cardiomyopathy) with symptoms of arrhythmias have also proved to have some genetic linkage to impaired impulse conduction (Benson et al., 1999; Bonne, et al., 2001; Schott et al., 1998). However, much more prevalent are rhythm disturbances and conduction failures occurring in ‘remodeling’ hearts which progressively undergo structural and electrical changes in prolonged metabolically and mechanically stressed states, e.g. chronic heart failure and myocardial infarction (Nattel, 2007; Olson, 2005). The well-known consequences of cardiac remodeling are; altered expression and gating properties of ion channels shaping the action potential and generating pace-making activities; reduced electrical coupling which leads to aberrant conduction of excitation. These changes are thought to work as the ‘substrates’ increasing the risk of lethal cardiac arrhythmia (Janse, 2004; Nattel, 2007). It has also been known that abnormal handling of intracellular Ca²⁺ occurring in the remodeled heart may cause spontaneous membrane depolarizations triggering ectopic excitations (Dobrev, 2010; Janse, 2004; Nattel, 2007). However, the available knowledge is still limited largely to ion channels/transporters/pumps whose roles in the cardiac excitation-contraction cycle have relatively been well established.

The transient receptor potential (TRP) channels constitute a newly-emerging non-selective cation channel (NSCC) superfamily activated by a plethora of physico-chemical stimuli other than voltage change. Because of this unique activation profile as well as the ability to permeate Ca²⁺ (except for TRPM4/TRPM5), TRP channels have attracted great attention as promising candidate molecules elucidating a variety of biological functions and disorders.
associated with slow sustained Ca\textsuperscript{2+} influx initiated by neurohormonal factors, pheromones, mechanical (membrane stretch/bending, osmotic change, shear force etc.) and thermal (from cold through cool and warm to heat) stresses, noxious stimuli (acid, respiratory irritants and toxicants) and many gustatory and pungent/cooling agents (camphor, citral, capsaicin, eucalyptol, icilin, menthol, allicin, mustard oil, sweet, umami and bitter tastants etc.) (Holzer, 2011; Vay et al., 2011; Wu et al., 2010). In the cardiovascular system, recent investigations have revealed pathophysiological implications of TRP channels in vasospasm, hypertension, occlusive vascular diseases, cardiac hypertrophy, cardiomyopathy and cardiac arrhythmia (Dietrich et al., 2010; Inoue et al., 2009b; Watanabe et al., 2008). Especially, as will be described below, involvement of some TRP channels in abnormal intracellular Ca\textsuperscript{2+} handling and increased responses to mechanical and noxious stresses may make them particularly relevant to the pathogenesis of acquired cardiac arrhythmias tightly associated with cardiac hypertrophy and failure, myocardial infarction and atrial fibrillation (Nattel, 2007; Ter Keurs & Boyden, 2007).

This chapter will deal with the arrhythmogenic potential of several TRP isoforms identified in the cardiovascular system, with brief introduction to the general concepts of cardiac arrhythmia and its connection to cardiac diseases and with several examples in which the roles of TRP channels have been established or suggested.

2. Factors contributing to cardiac arrhythmia

Cardiac arrhythmia is the abnormality of cardiac rhythm, the highly coordinated and integrated electrophysiological behavior of multiple ion channels/transporters/exchangers residing in tens of billions of myocytes and non-myocytes consisting of the heart. Clinically, cardiac arrhythmia can be defined as any anomalous excitations out of normal sinus control, and is conventionally classified into bradyarrhythmia (<50 beats per minute) and tachyarrhythmia (>100). Faulty or abnormal excitation of sinus node and various extents of conduction blocks mainly explain the former. In contrast, ectopic excitations occurring outside the sinus node (triggered activity, ectopic automaticity) and perpetuation of spiral/scroll waves rotating around a central core/filament (reentry) are thought to contribute to the latter. Although the mechanism remains still incompletely understood, shortened refractoriness, slowed and anisotropic conduction facilitate the occurrence of reentry (Jalife, 2000). Thus, altered expression or activities of ion channels contributing to the upstroke (Na\textsubscript{v}) and repolarization of action potential (K\textsubscript{v}) as well as those determining the conduction velocity and cell-to-cell coupling [Na\textsubscript{v} and connexins (Cx), respectively] are involved in the reentrant mechanism.

Ectopic excitations result from premature depolarizations before the next normal excitation arrives. Early afterdepolarization (EAD) occurs before the repolarization of action potential completely terminates, thereby prolonging it and evoking premature action potentials due to re-opening of Na\textsubscript{v} and/or Ca\textsubscript{v} channels. This is a mechanistic background for the initiation of a specific form of polymorphic ventricular arrhythmias (Torsades de pointes) observed in both congenital and drug-induced Long QT syndromes; the function of ‘repolarization reserve’, i.e. K\textsubscript{v} channels, is compromised, or residual activity of Na\textsubscript{v} channel sustains because of its incomplete inactivation (Roberts & Gollob, 2010).

Delayed afterdepolarization (DAD) is thought to reflect the generation of transient inward currents (I\textsubscript{i}) activated by diastolic Ca\textsuperscript{2+} release from the sarcoplasmic reticulum (SR) of
myocytes (Venetucci et al., 2008). If the magnitude of DAD exceeds the threshold of Na$_V$ channel activation, an extrasystolic discharge of action potentials occurs. Sustained discharges from one or more foci may propagate around to cause tachyarrhythmias (focal excitation). In ventricular myocytes from rabbit failing heart which shows spontaneous ventricular tachyarrhythmia, $I_h$ has been ascribed exclusively to enhanced forward-mode Na$^+$/Ca$^{2+}$ exchanger type 1 (NCX1) current (Pogwizd et al., 2001; Pogwizd & Bers, 2004). However, there is evidence that implicates Ca$^{2+}$-dependent Cl$^-$ and nonselective cationic conductances in the genesis of DAD (Han & Ferrier, 1992; Hill et al., 1988; Kass et al., 1978; Laflamme & Becker, 1996; Wu & Anderson, 2000). The mechanism underlying the diastolic Ca$^{2+}$ release is likely to reflect Ca$^{2+}$ overload into the sarcoplasmic reticulum (SR) caused by a net increase in Ca$^{2+}$ influx into the SR (increased Ca$^{2+}$ influx and SRECA2a Ca$^{2+}$-ATPase activity), or the dysfunction of ryanodine type 2 receptor (RyR2) which may, especially when intensively phosphorylated by protein kinase A (Wehrens et al., 2003) or Ca$^{2+}$/calmodulin-dependent kinase II (Ai et al, 2005), causes a diastolic Ca$^{2+}$ leak from the SR (Bers, 2006). Thus, under excessive sympathetic activities (hard exercise, mental stress, chronic heart diseases) or upon the application of drugs increasing the cAMP level (caffeine, catecholamines, phosphodiesterase inhibitors) or [Na$^+$]$_i$ (digitalis) in cardiomyocytes which increase Ca$^{2+}$ loading into the SR, the occurrence of DAD and resultant tachyarrhythmias may be greatly enhanced (Pogwizd, 2003; Sipido, 2007; Venetucci, 2008). In addition, in some pathological settings such as unphysiologically decreased extracellular K$^+$ level and excessive sympathetic activity, ectopic automaticity, particularly in the Purkinje fiber, can also be abnormally increased to cause tachyarrhythmias (Osadchii, 2010).

Clinical significance of mechanical loading in inducing arrhythmia has been well recognized (Dean & Lab, 1989). It is widely accepted that both acute and chronic stretch of myocardial tissue significantly affects its electrophysiological properties to increase the propensity for arrhythmia (Janse, 2003; Ravens, 2003). Experimentally, a transient diastolic stretch of isolated left ventricle was shown to induce ventricular arrhythmia which was inhibited by micromolar Gd$^{3+}$ (Hansen et al. 1991; Stancy et al, 1992). Atrial fibrillation elicited by an acute increase in intra-atrial pressure in isolated rabbit hearts was also found effectively blocked by MSCC blockers, Gd$^{3+}$ and a Tarantula toxin GsMTx-4 (Franz and Bode, 2003; Suchyna, 2000). At the single cell level, Kamkin et al. (2000a) demonstrated that direct stretch of a ventricular myocyte could cause membrane depolarization and prolongation of action potential resembling EAD, which, at extensive stretch, led to an extrasystolic depolarization. This stretch-induced depolarization was attributed to the activation of Gd$^{3+}$-sensitive mechanosensitive NSCCs (MSCCs) (Hamill & Martinac, 2001; Inoue et al., 2009b). The stretch sensitivity of MSCCs was significantly enhanced in hypertrophied cardiomyocytes with increased susceptibility to stretch-induced arrhythmia (Kamkin et al., 2000a). Considering that the heart is continuously subjected to hemodynamic stresses and deformation due to contraction, mechanical stresses may play a significant role in the causality and severity of arrhythmia, particularly under pressure- and volume-overloaded conditions.

3. Connection of cardiac diseases to cardiac remodeling and arrhythmia

The cardiac remodeling can be defined as the restructuring and reactivation of differentiated cardiac tissues comprised of myocytic and non-myocytic populations, and is initiated and
progressed by a complex interplay of genetic, environmental and aging factors (Cohn, et al., 2000; Fedak, et al., 2005; Swynghedauw, 1999). Chronic heart failure, myocardial infarction and atrial fibrillation (AF) are three common pathological states accompanying the remodeling process with arrhythmic changes; in which excessively activated sympathetic nervous and renin-angiotensin-aldosterone systems, increased generation of inflammatory cytokines and reactive oxygen/nitrogen species, and sustained mechanical stresses likely play active roles. The remodeling process is initially ‘adaptive’ to compensate the impaired pumping function, but gradually becomes ‘maladaptive’ with disturbances in rhythm formation and conduction as common clinical complications.

The key consequences of cardiac remodeling associated with the appearance of arrhythmia include both structural and electrical alterations. Enhanced collagen synthesis promotes the fibrotic replacement of damaged myocardial tissues thereby increasing the electrical heterogeneity. Altered expression and activities of ion channels, transporters and exchangers also bring about significant changes in the shape of action potential and its conduction properties as well as induce the susceptibility to premature excitations (Nattel, et al., 2007; Janse, 2004).

In chronic heart failure, the major electrophysiological changes are pronounced prolongation of action potential which often accompanies EAD-like membrane oscillations. These changes likely occur through decreased expression or activity of transient outward current (I_{to}; K_{v4.3}) and voltage-dependent K+ channels forming the repolarization phase of action potential (I_{kr}, I_{ks}) (Bers, 2006; Janse, 2004; Nattel, et al., 2007; Ravens, 2010). The reduced inward rectifying K+ current (I_{ki}) may also destabilize the membrane (i.e. increases the diastolic membrane resistance) and thereby enhance extrasystolic depolarizing responses (DAD) (Nattel, et al., 2007; Pogwitz et al, 2001). Furthermore, upregulation of hyperpolarization-activated current (I_{f}) and its mRNA (HCN2/4) (Cerbai, 1994; Fernandez-Velasco et al., 2003) and reduced expression of gap junction channel (Cx43; Dupont et al, 2001) have been reported to contribute to abnormally enhanced automaticity and impaired conduction, respectively. However, more notable changes observed in failing heart are abnormalities in intracellular Ca^{2+} handling (Bers, 2006; Janse, 2004). Despite a reduction in the SR Ca^{2+} content due to decreased expression of SERCA2a, the propensity for triggered activity based on DAD is enhanced. This likely reflects the other two major changes in Ca^{2+} handling, i.e. (1) the increased Ca^{2+} sensitivity of RyR2 under intensive phosphorylation by protein kinase A (however controversial now; Wehrens et al, 2003) or Ca^{2+}/calmodulin kinase II (Ai et al, 2005) which causes diastolic Ca^{2+} leak from the SR, and (2) upregulation of Na^{+}/Ca^{2+} exchanger (NCX1) protein which can carry depolarizing inward currents in the diastole. It has been suggested that increased β-adrenergic drive, which is prominent in chronic heart failure, may greatly facilitate the occurrence of DAD by increasing the Ca^{2+} content in the SR above the threshold of spontaneous Ca^{2+} release (Bers, 2006; Pogwizd, 2003).

Myocardial infarction is initiated by a sudden cessation of blood supply to heart tissues (i.e. myocardial ischemia), most frequently by thrombotic obstruction of coronary arteries. The time course of ischemic changes in the heart is variable and complex, and can mechanistically be distinguished between early and late acute phases, and subsequent postinfarction period vulnerable to structural and electrical remodeling (Clements-Jewery et al., 2005; Janse & Wit, 1989; Nattel et al., 2007). In acute phases, rapid depletion of intracellular ATP and accumulation of intracellular ADP, extracellular K+ and lactate
occur because of anaerobic glycolysis, and loss of intracellular K\textsuperscript{+} and intracellular acidosis follows. The extracellular accumulation of K\textsuperscript{+} leads to the depolarization of myocyte membrane that attenuates the amplitude and upstroke velocity of action potential (facilitated Na\textsubscript{v} inactivation) and the shortening of action potential duration (due to enhanced K\textsubscript{v} activities) or refractoriness. Simultaneously, intercellular accumulation and release of many biochemical substances occurs including catecholamines, ATP, lysophosphatidylcholine, cytokines (e.g., TNF\textalpha), reactive oxygen species (ROS), and platelet-activating factors (Clements-Jewery et al., 2005; de Jong & Dekker, 2010). All these possess arrhythmogenic potential to induce ventricular premature excitations (DAD, EAD) and reentry. In postinfarction period, down-regulation of K\textsuperscript{+} channels (I\textsubscript{to}, I\textsubscript{Ks}, I\textsubscript{Kr}, I\textsubscript{K1}) occurs in border-zones adjacent to infarct areas which impairs the repolarization of action potential leading to EAD, and altered intracellular Ca\textsuperscript{2+} handling facilitates spontaneous subcellular Ca\textsuperscript{2+} release events that can trigger arrhythmic episodes (Nattel, et al., 2007).

Atrial fibrillation (AF) is a common supraventricular tachyarrhythmia with rapid and highly irregular firing, being closely associated with aging and cardiovascular diseases such as heart failure, myocardial infarction, valvular diseases and hypertension (Nattel et al, 2007; Ravens, 2010). AF per se is thought to serve as an arrhythmogenic remodeling process, because the occurrence of AF itself progressively aggravates electrophysiological features and facilitates fibrotic changes of atrial tissues in favor of more frequent and sustained occurrences. This clinical feature is described “AF begets AF” and experimentally confirmed by the finding that rapid atrial pacing causes significant shortening of atrial refractoriness and persistent AF (Wijffels, et al. 1995). The major arrhythmogenic changes found for AF are marked shortening of action potential duration which reduces the refractoriness and increases the susceptibility to reentry (Nattel et al, 2007; Ravens, 2010). In AF, this change is combined with abnormal Ca\textsuperscript{2+} handling with increased Ca\textsuperscript{2+} release from the SR and augmented NCX1 expression which facilitates ectopic premature depolarizations (i.e. DAD) that serve as a trigger to initiate the reentry (Dobrev, 2010). Both reduced/increased or unchanged expression of gap-junction channels have been reported, but their increased regional heterogeneity may contribute to various extents of conduction failure and reentry (Nattel, et al., 2007).

4. TRP channels and cardiac arrhythmia

The above considerations strongly suggest that, although alterations in the genesis and conduction properties of action potential are undoubtedly of central importance, other factors, e.g. Ca\textsuperscript{2+} overload, increased stretch sensitivity and noxious stimuli generated in ischemia/reperfusion, may also play a vital role in arrhythmogenicity. It is also possible that structural remodeling (necrotic/apoptotic and fibrotic changes) may contribute to the electrical heterogeneity of the myocardium which may act as the pro-arrhythmic substrates for altered conduction and reentry. As described above, Ca\textsuperscript{2+-} and stretch-sensitivities, nociception as well as association to remodeling are the hallmark features of TRP channel members (Inoue et al., 2009b; Nishida & Kurose, 2008; Watanabe et al., 2008; Wu et al, 2010; Vay et al., 2011). Therefore, in the following, we would like to discuss about the pathophysiological relevance of several TRP isoforms particularly in these aspects of acquired arrhythmogenicity (see Table 1).
<table>
<thead>
<tr>
<th>Intrinsic activator/modulator</th>
<th>possible arrhythmogenic mechanisms</th>
<th>chemical agonists</th>
<th>inhibitors</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Cardiomyocyte</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>TRPC1 store depletion, stretch?, GPCR stimulation (TRPC1/TRPC4 or C5), Ca(^{2+})/CaM</td>
<td>stretch-induced EAD/DAD in hypertrophy?</td>
<td>Gd(^{3+}), SKF, GsMTX-4?</td>
<td></td>
</tr>
<tr>
<td>TRPC3 const-act., store depletion, GPCR stimulation, BDNF, DAG, oxidative stress (TRPC3/C4), Ca(^{2+})/CaM, PKC, PKG, Src</td>
<td>oxidative stress? and other mediators (ATP, UTP) during ischemia/reperfusion</td>
<td>Gd(^{3+}), La(^{3+}), SKF, 2-APB, flufenamate, Pyr-3</td>
<td></td>
</tr>
<tr>
<td>TRPC5 const-act., store depletion, GPCR stimulation, LPC, S-nitrosylation, thioredoxin, oxidative stress</td>
<td>possible relation to ischemia/oxidative stress in failing heart</td>
<td>Gd(^{3+}), SKF, GsMTX-4?</td>
<td></td>
</tr>
<tr>
<td>TRPC6 store depletion, stretch?, GPCR stimulation, GF, DAG, 20-HETE, PiP(_3)/PiP(_2), Ca(^{2+})/CaM, CamKII, PKC, PKG, Fyn</td>
<td>stretch-inuced EAD/DAD in hypertrophy?</td>
<td>flufenamate</td>
<td>Gd(^{3+}), La(^{3+}) (high), SKF, 2-APB, BTP-2, ML-7, ML-9, propol, halothane, chloroform</td>
</tr>
<tr>
<td>TRPM2 ADPR, cADPR, NAADP, AMP, [Ca(^{2+})](_i), pH</td>
<td>oxidative stress in ischemia/reperfusion?</td>
<td>flufenamate, ACA, 2-APB</td>
<td></td>
</tr>
<tr>
<td>TRPM4 [Ca(^{2+})](_i), GPCR stimulation, PiP(_2), MgATP, ATP, AMP, voltage, PKC, CaM, spermine,</td>
<td>familial conduction block (PFHBI, ICCD), stretch-induced EAD/DAD?, QT-elongation in hypertrophy, DAD in AF?</td>
<td>decavanadate, BTP-2</td>
<td>Gd(^{3+}), flufenamate, clotrimazol, 9-PA</td>
</tr>
<tr>
<td>TRPC6 see above</td>
<td>anti-fibrogenesis</td>
<td>see above</td>
<td>see above</td>
</tr>
<tr>
<td>TRPM7</td>
<td>fibrogenesis in AF</td>
<td>2-APB (high)</td>
<td>Gd(^{3+}), SKF, 2-APB (low), LOE908,</td>
</tr>
</tbody>
</table>

**cardiac fibroblast**

**respiratory sensory neuron**

<table>
<thead>
<tr>
<th>Intrinsic activator/modulator</th>
<th>possible arrhythmogenic mechanisms</th>
<th>chemical agonists</th>
<th>inhibitors</th>
</tr>
</thead>
<tbody>
<tr>
<td>TRPA1 cold (&lt;17°C), [Ca(^{2+})](_i), alkalosis, stretch, GPCR, PiP(_2), PKA.</td>
<td>autonomic imbalance</td>
<td>environmental irritants (acrolein), allicin, allyl isothiocyanate, citral, NSAIDs, eugenol, cinnamaldehyde, 2-APB, GsMTX-4</td>
<td>HC-030031</td>
</tr>
</tbody>
</table>
| TRPV1 heat (>43°C) · acidity (pH<5.9) · 12-HETE, 12-LOX metabolites, anandamide, chemokines · GPCR (e.g. bradykinin, ATP, PAR2) · NGF, GDNF, PKA, PKC, PI3K, Ca\(^{2+}\)/CaM, CaMK II, src, calcineurin, Ras/MAPK, SCF | autonomic imbalance | environmental irritants, capsaicin, 2-APB, camphor, allicin, citral, resi
iniferatoxin, gingerol, eugenol, EtOH | capasazeine, ABT-102, AMG-517, GRC-6211, MK-2295, SB-366791, SB-705498, |

Table 1. Arrhythmogenic TRP channels
Abbreviations; CaM, calmodulin; GPCR, G-protein-coupled receptor (Gq-coupled); const-act, constitutively active; BDNF, brain-derived neurotrophic factor; DAG, diacylglycerol; PKC, protein kinase C; PKG, protein kinase G; LPC, lysophosphatidylcholine; S1P, sphingosin 1-phosphate; EGF, epidermal growth factor; PGE2, prostaglandin E2; 20-HETE, 20-hydroxyeicosatetraenoic acid; PIP3, phosphatidylinositol 3,4,5-trisphosphate; PKA, protein kinase A; 12-HETE, 12-hydroxy-5,8,10,14-eicosatetraenoic acid; LOX, lipooxygenase; PAR2, Proteinase-activated receptor 2; NGF, nerve growth factor; GDNF, glial cell-line derived neurotrophic factor; MAPK, mitogen-activated protein kinase; SCF, stem cell factor; SKF, SK&F96365: BTP-2, 4-methyl-4-[3,5-bis(trifluoromethyl)-1H-pyrazol-1-yl]-1,2,3-thiadiazole-5-carboxanilide; Pyr-2, Ethyl-1-(4-(2,3,3-trichloroacrylamide)phenyl)-5-(trifluoromethyl)-1H-pyrazole-4-carboxylate; 2-APB, 2-aminoethoxydiphenyl borate; 9-PA, 9-phenoanthrol.

4.1 Stretch-induced arrhythmia in pathological settings
As described above, acute stretch of cardiac muscle can induce arrhythmic responses in both atrium and ventricle, which are inhibited by widely used MSCC blockers Gd³⁺ and GsMTx-4. This has led to the speculation that activation of nonselective cationic MSCCs causes premature depolarizations. There are at least three candidates for these MSCCs in TRPs expressed in the heart, i.e. TRPC1, TRPC6 and TRPM4.

In both heterologous and native overexpression systems, TRPC1 and TRPC6 have been found to activate in response to mechanical stimuli (Maroto et al., 2005; Spassova et al., 2006). Albeit controversial (e.g. Inoue et al., 2009a), the so-called MSCC-selective peptide blocker GsMTx-4 has been found to inhibit the mechanical activation of both TRPC1- and TRPC6-mediated MSCC activities (Maroto et al., 2005; Spassova et al., 2006). TRPC1 and TRPC6 are ubiquitously expressed in whole myocardial tissues (Huang et al., 2009; Ward et al., 2008), and GsMTx-4 suppresses stretch-induced force development and concomitant [Ca²⁺]i increase in mouse left ventricular trabecular muscle (Ward et al., 2008) and pressure-induced atrial fibrillation (Franz and Bode, 2003). Moreover, the expression of TRPC1 and TRPC6 is greatly increased in hypertrophied heart under prolonged pressure overload (Kuwahara, et al; 2006; Ohba, et al., 2007), where the susceptibility to mechanically-induced arrhythmia is also enhanced (Kamkin et al., 2000a). All these observations favor the view that TRPC1 and TRPC6 contribute to stretch-induced arrhythmias as MSCCs in some pathological settings. However, there is controversy over the mechanosensitivity of these two TRPC channels. In knock-out mice deficient in TRPC1 or TRPC6 expression, pressure-induced vasoconstriction (myogenic response), which is believed to reflect the depolarizing effects of MSCC activation, was found intact (Dietrich et al., 2005; Gottlieb et al., 2007). Although this fact could not totally negate the roles of these TRP channels in the heart, it is essential to test whether mechanical activation of myocardial MSCCs and mechanical arrhythmogenicity are indeed impaired in these knock-out mice or vice versa in TRPC1- or TRPC6-overexpressing mice. Alternatively, it may also deserve to test the possible involvement of a recently identified GsMTx-4-sensitive MSCC, Piezo1/Piezo2, in stretch-induced arrhythmia in the heart (Bae et al., 2011).

In cerebral arterial myocytes, TRPM4 has been proposed as a MSCC responsible for myogenic response (Earley et al., 2004). However, again, the results from TRPM4-deficient mice do not support this role (for more detail, see below) (Mathar et al., 2010). Nevertheless,
it is noteworthy that TRPM4 can be secondarily activated by mechanical stretch through stretch-induced Ca\(^{2+}\) release from ryanodine-sensitive stores in arterial myocytes (Morita et al., 2007), since similar axial stretch-induced Ca\(^{2+}\) release from the SR has been demonstrated in cardiomyocytes (Iribe, et al., 2009). Considering that heart wall is periodically distended by diastolic filling pressure, this mechanosensitive mechanism may have considerable pathophysiological significance for the genesis of DAD in remodeled heart in which the expression of TRPM4 and Ca\(^{2+}\) loading into SR are prominently enhanced (see below).

It has been shown that repeated cyclic stretch induces hypertrophic responses of cardiac myocytes, which are significantly attenuated by antagonists for a PLC-linked G-protein coupled receptor (GPCR), angiotensin type 1 (AT\(_1\)) receptor (Komuro and Yazaki, 1993). This is explained by direct mechanical activation of unoccupied AT\(_1\) receptor as demonstrated by the substituted cysteine accessibility mapping technique (Yasuda et al., 2008). Similar mechanical activation also appears to occur for many other PLC-linked GPCRs including endothelin ET\(_A\), vasopressin V\(_{1A}\) and muscarinic M\(_5\) receptors (Mederos y Schnitzler et al., 2008). TRPC6 and its homologue TRPC3 are activated by stimulation of PLC-coupled GPCRs via generation of diacylglycerol (DAG) (Hofmann, et al., 1999) and Ca\(^{2+}\) influx through activated TRPC3 and TRPC6 channels has been shown to be essential for hypertrophic responses of caridomyocyte via calcineurin/NFAT pathway (Kuwahara et al., 2006; Bush et al., 2006; Onohara et al., 2006). In failing hearts where pressure overload is sustained, the release of catecholamines from sympathetic nerves and adrenal gland is increased (see above), and the production of angiotensin II (AngII) and expression of AT\(_1\) receptor and angiotensin converting enzyme are greatly enhanced (Goette, A et al., 2000; Ihara M et al, 2000; Kaprielian RR et al., 1997). Moreover, the mechanosensitivity of TRPC6 channel is remarkably enhanced by the simultaneous stimulation of PLC-linked GPCRs via concerted actions of two lipid messengers DAG and 20-HETE (Inoue et al., 2009a). Thus, taken together, it is highly conceivable that sustained mechanical loads promote the hypertrophic remodeling of cardiomyocytes through synergistic interplay between excessive activation of GPCR-PLC pathways and enhanced receptor/mechanical activation of TRPC6 (and TRPC3) channels. This would in turn exacerbate Ca\(^{2+}\) overload into the myocyte SR to increase the propensity for pro-arrhythmic depolarizations. Consistent with this scenario, cardiac-specific TRPC6 transgenic mice have been found to exhibit much increased susceptibility to mechanical stress with increased incidence of sudden death accompanied by severe macroscopic and histological signs of cardiomyopathy (Kuwahara et al., 2006). The same study also found that expression of TRPC6 was several-fold upregulated in human failing heart. Obviously, detailed electrophysiological analyses are required to corroborate whether the above changes could indeed induce the arrhythmogenicity.

### 4.2 Ca\(^{2+}\)-dependent arrhythmia

Abnormal Ca\(^{2+}\) handling or cycling in cardiac tissues is the main cause for arrhythmia (Ter Keurs & Boyden, 2007). As briefly introduced above, diastolic Ca\(^{2+}\) release (or leak) from the SR is a key event to initiate arrhythmogenic premature depolarizations DAD. Presently, the mechanism whereby DAD is generated by the Ca\(^{2+}\) release can be accounted for by the activation of three ionic conductances which can generate I\(_{\text{ti}}\) near the resting membrane potential; an electrogenic forward-mode NCX1 current which allows the influx of three Na\(^{+}\) ions in exchange of one Ca\(^{2+}\) efflux after the repolarization has completed; and Ca\(^{2+}\)-activated Cl\(^{-}\) and nonselective cation channel (CAN) currents. Although the most accepted hypothesis suggests that I\(_{\text{ti}}\) is carried exclusively by enhanced forward-mode NCX1 inward
current in failing ventricle (Pogwizd, et al. 2001; Pogwizd, 2003; Wu and Anderson, 1999), there is also experimental evidence suggesting the involvement of Ca\(^{2+}\)-activated Cl\(^-\) current (Han et al., 1996; Laflamme et al., 1996) and CAN (Hill et al., 1988; Kass et al., 1978; Wu & Anderson, 2000) in the genesis of I\(_{\text{ti}}\).

CAN constitutes a large heterogenous family of cation channels with varying unitary conductances, Ca\(^{2+}\)-sensitivities, voltage dependence and regulations, and distributes broadly in neurons, smooth muscle, heart, exocrine and endocrine glands, and other epithelial tissues (Siemen, 1993). The most frequently recorded CAN from cardiac tissues, first identified in a cultured rat neonatal ventricular myocyte (Colquhoun, 1981), is a 20-40pS channel with almost equal selectivity over monovalent cations and poor permeability to divalent cations (Guinamard et al., 2006). Albeit rather great variations, these cardiac CANs are activated by the micro- to mili-molar range of Ca\(^{2+}\) and voltage-dependent. The molecular identification that best fits this type of CAN is now thought to be TRPM4 protein (Launay et al., 2002). Expressed TRPM4 channel shows a ~25pS conductance (in near-physiological ionic milieu), is monovalent cation-selective, and, in cell-free conditions, undergoes Ca\(^{2+}\)-dependent activation and depolarization-dependent enhancement of open probability only in their unphysiologically high ranges. However, it has been found that the Ca\(^{2+}\)- and voltage-sensitivities of TRPM4 rapidly shift to their much higher ranges immediately after the excision of patch membrane (‘desensitization’ or ‘rundown’), and this is substantially prevented by pretreatment with MgATP, protein kinase C activators, maneuvers replenishing phosphatidylinositol 4,5-bisphosphate and increased temperature (Nilius, 2006). These findings suggest that, in vivo, TRPM4 may be more effectively regulated by physiological ranges of [Ca\(^{2+}\)]\(_i\) and membrane potential. Expressed TRPM4 channel also undergoes negative regulation of free adenosine nucleotide phosphates (efficacy; ADP > ATP = AMP), and is subject to spermine block, so that severe ischemic conditions might affect this channel activity (Nilius et al., 2004).

In human, cardiac expression of TRPM4 is ubiquitous in the order of abundance; Purkinje fibers, septum, atrium, left and right ventricles (Kruse et al., 2009). Guinamard et al (2002) found that expression of TRPM4-like CAN, which was virtually absent at the time of cell isolation, was progressively enhanced in rat dedifferentiating ventricular myocytes with the time of culture. Studies from the same group also found that purinergic receptor activation by ATPyS, PKC activators and DAG, all of which act as interventions to facilitate cardiac remodeling, enhanced the occurrence of TRPM4-like channel in cultured ventricular myocytes. This led to the postulation that remodeling of ventricular myocytes in response to excessive hypertrophic signals may enhance the activity of TRPM4 thereby increasing arrhythmogenic propensity mostly likely by increasing DAD. Consistent with this, in the hypertrophic heart of spontaneously hypertensive rat, the elongation of Q-T interval in electrocardiogram has been reported together with increased expression of TRPM4 protein and density of CAN (Guinamard et al., 2006). Recent reports indicate that TRPM4 is more abundantly expressed in atrium and sinoatrial node (Demion et al, 2007; Guinamard, 2004; Kruse et al., 2009), where DAD-based tachyarrhythmias have been well established (Dobrev, 2010; Guinamard 2006; Nattel 2007; Raves, 2010). However, TRPM4-deficient mice showed no obvious abnormalities in heart rate, cardiac output, ejection fraction and cardiac contractility at basal conditions questioning a vital role of TRPM4 in physiological regulation of cardiac functions (Mathar et al., 2009). Rather unexpectedly, the genetic deletion of TRPM4 caused postnatal development of high blood pressure due to increased sympathetic catecholamine secretion. These seemingly paradoxical observations might point to the pathological contribution of TRPM4 to
arrhythmogenicity in diseased remodeling heart rather than to normal rhythm formation, in which the expression and activity of TRPM4 are enhanced.

A recent genetic analysis of human families with lethal conduction failure has found a gain-of-function mutation (c.19G→A in exon 1 or p.E7K) in the N-terminal domain of TRPM4 channel (PFHBI; progressive familial heart block type I) (Kruse et al., 2009). More recently, other TRPM4 mutations with very similar biological impacts (p.R164W, p.A432T, p.G844D) have also been assigned to autosomal dominant isolated cardiac conduction block (ICCD) of Lebanese and French families (Liu et al., 2010). Biochemical and immunocytochemical data suggested that these mutations increase the cell surface expression of TRPM4 protein due to deregulated SUMOylation/deSUMOylation process which results in impaired endocytotic protein degradation. The accompanying electrophysiological changes for these mutations are consistent with the biochemical/immunocytochemical data; the density of whole-cell CAN current was increased with little noticeable changes in macroscopic Ca\(^{2+}\) sensitivity and voltage dependence or in unitary conductance and open probability at single channel level. These results suggest that the number of functional channels were increased without altered biophysical properties. The precise pathogenic mechanism for these mutations to cause conduction block remains unclear. However, RT-PCR analysis indicated the highest expression of TRPM4 in Purkinje fiber, and detailed immunohistochemical examination showed strongly TRPM4-positive subendocardial bundles of Purkinje fibers branching and penetrating toward subepithelial layer (Kruse et al., 2009; Liu et al., 2010). These morphological findings, combined with a broadened QRS complex in ECG in both PFHBI and ICCD patients, led the authors to speculate that elevated TRPM4 expression may increase the membrane leak conductance thereby disabling action potential propagation along the Purkinje fibers. There are several previous studies showing that Ca\(^{2+}\) overload causes membrane oscillations in Purkinje fibers due to increased spontaneous Ca\(^{2+}\) release when Na\(^+\) extrusion via Na\(^+\)/K\(^+\)-ATPase was pharmacologically inhibited (Kass et al., 1978; Kass & Tsien, 1982; Lederer & Tsien, 1976). Considering that TRPM4 may also act as a Na\(^+\) entry pathway in response to GPCR stimulation (Launay, et al., 2002), the resultant increase in [Na\(^+\)] might then elevate [Ca\(^{2+}\)], via the reversed mode operation of NCX1 in Purkinje fiber cells (Bers, 2006; Pogwizd, 2003). This would not only facilitate Ca\(^{2+}\) overload and DAD generation due to increased TRPM4 channel activation, but could also induce cell death leading to cardiac injury/remodeling if the elevation of [Ca\(^{2+}\)] would persist. Fibrotic replacement observed in the His-Purkinje fiber system of a PFHBI patient (Kruse et al., 2009) may support the latter possibility. Thus, the depolarizing and Na\(^+\)-permeating properties of TRPM4 channel could bring about both acute (induction of DAD) and chronic (conduction block due to remodeling) pro-arrhythmic effects.

4.3 Arrhythmia associated with ischemia and oxidative stress

In the acute phases of myocardial infarction, many arrhythmogenic substances are released (see above). ATP and UTP are amongst these, being released from cardiomyocytes during ischemia and may promote the occurrence of ventricular tachyarrhythmias (Dutta et al., 2004; Kuzmin et al., 1998). In vitro, ATP can induce DAD-based arrhythmic depolarizations in single cardiomyocytes when combined with Ca\(^{2+}\) increasing agents such as catecholamines (Song & Belardinelli, 1994). In voltage-clamped adult ventricular myocytes, ATP/UTP can activate a sustained inward current via P2Y2 receptor and enhance the opening of 14 and 23pS single channel activities in a PLC-dependent manner. TRPC3 and TRPC7 proteins from rat ventricular myocytes immunoprecipitated, and the macroscopic current induced by
ATP was suppressed by intracellular application of anti-TRPC3 antibody from the patch pipette (Alvarez et al., 2008). These findings are collectively interpreted as indicating that ATP/UTP may cause arrhythmia by activating TRPC3/TRPC7 heterotetrameric Ca\(^{2+}\) and Na\(^{+}\) entry channels whereby to facilitate intracellular Ca\(^{2+}\) overloading and trigger electrical activities. Consistent with this scenario, the same researchers’ group has recently shown that flash photolysis of caged ATP can evoke one or even a train of extrasystolic contractions. Further, intraperitoneal injection of creatine, which is capable of buffering a sudden ATP/UTP release and clinically exhibits anti-arrhythmic effects, markedly reduced ventricular tachyarrhythmias and early death events in a rat coronary-ligature myocardial infarction model. These ameliorating effects of creatine were defective in an inactive creatine analogue β-guanidinopropionate (Vassort et al., 2010). Cell surface expression of TRPC3 was found tightly controlled by vesicular trafficking, cytoskeletal actin dynamics and interaction with caveolar or subsarcolemmal proteins and molecules (Groschner K, et al., 2005). And PLC stimulation by AngII in cardiomyocytes was reported to facilitate the recruitment of TRPC3 complexed with NCX1 to cell membrane, resulting in the reverse mode Ca\(^{2+}\) entry through NCX1 (Eder et al., 2007). Thus, although the precise connection between TRPC3 (and possibly TRPC7) channel activation by ATP and arrhythmogenicity is unclear, the above mechanism may have particular pathophysiological significance in Ca\(^{2+}\)-dependent arrhythmogenesis during postinfaction remodeling period or sudden ischemic insult of hypertrophied heart, where enhanced expression of TRPC3 and sympathetic nerve activity play pivotal roles. Seemingly in line with these results, a recent independent study has reported that cardiomyocytes from TRPC3-overexpressing mice show an increased susceptibility to apoptotic death due to Ca\(^{2+}\) overload when subjected to hypoxia/reoxygenation (Shan et al., 2007).

Oxidative stress causes tissue damage in a variety of pathological states including aging, cancer, neurodegenerative disorders, autoimmune diseases, atherosclerosis and ischemia/reperfusion injury of myocardium (Chandra et al., 2000; Langley et al., 2004; Misra et al., 2009). Many reactive oxygen spices (ROS; O\(_2^-\), OH, H\(_2\)O\(_2\), NO, ·ONOO-) are generated as the result of ischemia and reoxygenation and may confer the susceptibility to arrhythmia in direct and indirect ways (via modifying autonomic nerve activity (Danson et al., 2006; Misra et al., 2009). Several TRP members including TRPM2, TRPM7, TRPC5 and a TRPC3/TRPC4 heterooligomer are known to be activated by oxidative stresses (Miller & Zhang, 2011; Naylor et al., 2011; Poteser et al., 2006). TRPM2 is the first reported ROS-sensitive Ca\(^{2+}\)-permeable nonselective cation channel activated by H\(_2\)O\(_2\), ADP-ribose, TNFα, and β-amyloid peptide, and is implicated in both physiological functions and pathophysiology including oxidant (H\(_2\)O\(_2\))-induced cell damage/death in pancreatic β-cells and neurons. TRPM2 is positively regulated by intracellular Ca\(^{2+}\), and its cell damaging action appears to be mediated at least in part by sustained [Ca\(^{2+}\)]\(_i\) elevation (Takahashi et al., 2011). In myocardial infarction, reperfusion injury and congestive heart failure, elevated plasma levels of TNFα have been reported (Sack et al., 2000). TNFα can activate both cytotoxic and cytoprotective signaling pathways in many cell types including mitochondrial ROS production through the recruitment of caspase-8 into the death-inducing signaling complex (Kroemer et al., 2007; Zhu et al., 2006). Further, altered redox state changes the activity of cardiac ion channels and transporters regulating Ca\(^{2+}\) dynamics in the cardiomyocyte, i.e. L-type Ca\(_v\), RyR2, NCX1 and SERCA2a (Zima & Blatter, 2006). In keeping with these observations, activation of TRPM2 by oxidative stress (H\(_2\)O\(_2\)) has been found to cause apoptotic/necrotic changes in single cardiomyocytes with ultrastructural
changes characteristic of cardiac ischemia/reperfusion injury (Yang et al, 2006). Although whether these changes are pro-arrhythmic remains to be explored, it is possible that sustained [Ca$^{2+}$], elevation through activated TRPM2 channel would disturb basal Ca$^{2+}$ homeostasis underlying rhythmic excitations in cardiomyocytes thereby inducing Ca$^{2+}$-dependent arrhythmogenicity. The arrhythmogenic effect of TRPM2 may not necessarily be restricted to direct cell-damaging actions via sustained [Ca$^{2+}$], elevation. It could occur indirectly through the production of inflammatory cytokines from ambient immune cells, as was demonstrated in an animal model of inflammatory bowel syndrome (Yamamoto et al., 2008).

In addition, expression of TRPC4 and TRPC5 was found to increase 2-3-fold in cultured rat cardiomyocytes subjected to downregulation of SERCA by siRNA technology which mimics the remodeling process of diseased heart (Seth et al., 2004). TRPC5 is known to be activated by many noxious stimuli such as H$_2$O$_2$ and LPC (Flemming et al., 2006; Naylor et al., 2011), and thus could act as a pro-arrhythmic mediator in ischemic remodeling heart. This possibility will be an intriguing subject of future investigation, together with other TRP isoforms implicated in the pathogenesis of cardiomyopathy (TRPV2, TRPC7; Iwata et al., 2003; Satoh et al., 2007).

### 4.4 Arrhythmia related to fibrosis

Cardiac fibroblasts occupy about 70% of whole cardiac cell populations, and, once placed in pathological states such as myocardial injury, oxidative stress and excessive mechanical stretch, start to proliferate and differentiate into the active phenotype 'myofibroblast'. Activated myofibroblasts secrete extracellular matrix proteins (e.g. collagen), matrix metalloproteinases, cytokines and growth factors thereby promoting the pathological restructuring of diseased heart with fibrotic replacement of damaged myocardial tissue (Souders, et al., 2009). Cardiac fibroblasts have a shallow resting membrane potential (-30 - -10mV) and are non-excitable themselves, but show mechanically-induced depolarizations well synchronized with spontaneous contractions of myocardium (Kamkin et al, 2000b; Kamkin et al, 2003). These electrophysiologic properties allow them, by coupling electrically to myocytes as a leaky capacitor, to modify the electrical properties of myocyte action potential and its propagation (Yue et al., 2011). In contrast, fibrotic tissues generated by myofibroblasts act as a physical barrier insulating the spread of electrical currents between the bundles of cardiomyocytes. Thus, disturbances in the myocyte-fibroblast coupling and alterations in myocardial architecture due to spatially inhomogenous fibrosis can lead to cardiac arrhythmias associated with reentry or conduction block.

It has been known that fibrotic remodeling is a fundamental process underlying the perpetuation of AF. Atrial fibrosis is dependent on Ca$^{2+}$, but what source of Ca$^{2+}$ is involved therein had been elusive. A recent study of Du et al. (2010) demonstrated that upregulation of TRPM7, a constitutively active Ca$^{2+}$/Mg$^{2+}$ entry channel (Wu et al, 2010), and resultant Ca$^{2+}$ influx was crucial for the progression of atrial fibrosis in AF patients. This was supported by the following evidence; (1) human atrial fibroblasts express TRPM7 protein abundantly, and show basal Ca$^{2+}$ influx and spontaneously active Ca$^{2+}$-permeable inward currents with the fingerprint features of heterologously expressed TRPM7 channels [potentiation by low pH and inhibition/potentiation by low/high concentrations of 2-aminoethoxydiphenylly boroate (2-APB) respectively: Li et al., 2006, 2007]; (2) TRPM7-like current and accompanying Ca$^{2+}$ influx were strikingly increased in parallel with the upregulation of TRPM7 expression in differentiated fibroblasts from AF patients; (3) all
these changes were largely eliminated by knockdown of TRPM7 expression with small hairpin RNA; and (4) in vitro promotion of myofibroblastic differentiation by TGF-β1, a major stimulator of atrial fibrosis, was correlated with the upregulation of TRPM7. Although in this study a direct linkage between Ca$^{2+}$ influx through TRPM7 channel, the fibrogenesis and arrhythmogenesis in AF have not been explored, the above results may provide a new therapeutic target against the progression of AF, by disclosing an essential role of TRPM7-associated Ca$^{2+}$ influx in fibroblast proliferation and differentiation.

In rat neonatal cardiac fibroblasts, endothelin-1 (ET-1) or AngII can induce myofibroblastic differentiation and collagen synthesis via the $G_{12/13}$ signaling. ET-1 (via $E_{T_A}$ receptor) and AngII can also selectively enhance the expression of TRPC6, thereby increasing basal Ca$^{2+}$ influx in the fibroblasts. Importantly, increased TRPC6 expression is causally correlated with anti-fibrotic effects via calcineurin/NFAT pathway (Nishida et al., 2007). The upregulation of TRPC6 likely occurs through the $G_{12/13}$- (Rac/NOX)-ROS-JNK signaling, which is reminiscent of TRPC6-mediated abnormal proliferation of pulmonary artery smooth muscle cells induced by platelet-derived growth factor, a pathogenic model of pulmonary artery hypertension (Yu et al., 2003). The concentrations of ET-1 or AngII required to upregulate TRPC6 are significantly higher than those for the differentiation of fibroblasts (Nishida et al., 2007). Thus, in an intriguing contrast to the pro-fibrotic role of TRPM7, the anti-fibrotic effects of enhanced TRPC6 channel activity may serve as a negative feedback mechanism to limit excessive fibrogenesis via ET-1/AngII signaling during cardiac remodeling. It remains to be determined whether this mechanism works beneficially in human AF patients or other fibrosis-associated arrhythmias.

4.5 Arrhythmia related to autonomic imbalance

The cardiac rhythm is under tight control of the autonomic nervous system. For instance, baroreceptors monitor blood pressure fluctuations and transmit the information to the brainstem vasomotor center via parasympathetic afferents, which then modulates the cardiac pumping force and rate via sympathetic efferents. The renin-angiotensin-aldosterone system is another well-established system centrally controlled via sympathetic nerves (via $\beta_2$-adrenoceptor). It fulfills a long-term control of body fluid and electrolyte balance, thereby effectively regulating the cardiac output (Guyenet, 2006). Any disturbances in these centrally-mediated autonomic regulations could therefore become the substrates for cardiac arrhythmias (Danson, et al., 2006; Janse 2004; Nattel, et al., 2007). Accumulating evidence however suggests that nociceptive reflexes via respiratory sensory neurons (e.g. C-fibers in nose and lung) have also some pathophysiological impact on cardiac functions through centrally-mediated autonomic mechanisms, as exemplified by air pollutant-induced changes in ‘repolarization’ parameters of ECG (Henneberger et al., 2005). The sensory nerve endings in respiratory airways contain two TRP members sensing noxious stimuli, i.e. TRPV1 and TRPA1 which likely participate in airway chemosensation and inflammation (Bessac & Jordt, 2008). It has thus been postulated that these TRP isoforms may be involved in nociceptive signaling in the respiratory system (Vay et al., 2011).

Two recent studies have revealed intriguing associations between respiratory sensory TRP channels and arrhythmogenicity. In one experimental model, activation of respiratory sensory neurons by inhalation of concentrated ambient particles significantly affected the cardiac rhythm with decreased heart rate, and shortened QT interval and P wave duration. Abrogation of these cardiac effects by a selective TRPV1 antagonist capsazepine suggested that TRPV1-mediated autonomic reflexes play a central role therein (Ghelfi et al., 2008).
another study employing a gaseous pollutant (Hazari et al., 2011), one-day exposure of rats to diesel exhaust gas resulted in increased heart rate with pro-arrhythmic ECG changes of prolonged action potential and shortened repolarization. Notably, these rats showed much heightened sensitivity to aconitine challenge that can induce moderate to lethal ventricular tachyarrhythmias dose-dependently. This pro-arrhythmic change was prevented by pretreatment with a TRPA1 antagonist HC-030031 or by sympathetic blockade with guanethidine, suggesting the involvement of increased sympathetic drive. In addition, partial involvement of TRPV1 in moderate aconitine-induced tachyarrhythmias was also suggested by pharmacological inhibition with a selective TRPV1 antagonist SB-366791. Although detailed mechanisms underlying remain unclear, these results clearly point to the pro-arrhythmic risk of environmentally-induced autonomic imbalance in which sensory chemosensing channels TRPA1 and TRPV1 may play a pivotal role.

5. Conclusions and therapeutic implications

The evidence presented above has disclosed that the unique activation profile (e.g. stretch- and Ca\(^{2+}\)-sensitivities, neurohormonal activation, sensitivity to noxious stimuli) and permeability to Ca\(^{2+}\) and Na\(^{+}\) (TRPM4/TRPM5 are however virtually Ca\(^{2+}\)-impermeable) render several TRP channels contribute to the acquired arrhythmogenesis during cardiac remodeling and other pathological processes. The examples so far available implicates, at least, TRPC1 and TRPC6 (and possibly their homologues) in stretch-induced arrhythmias in both acute and chronic manners, TRPM4 in Ca\(^{2+}\)-dependent arrhythmia and familial conduction block, TRPC3 and TRPM2 in ischemia-induced arrhythmia, TRPA1 and TRPV1 in autonomic imbalance-induced arrhythmia, and TRPM7 and TRPC6 in fibrosis-related arrhythmogenicity, especially in AF, as pro-fibrotic and anti-fibrotic factors respectively. Although the mechanism of the actions of each TRP isoform appears variable in different pathophysiological settings, these new lines of evidence have certainly put forward our essential understanding about the pathogenesis of commonly observed arrhythmias associated with cardiac diseases accompanying structural and electrical remodeling. Simultaneously it may open an avenue toward exploiting an entirely new generation of anti-arrhythmic drugs for increased mechanosensitivity, abnormal Ca\(^{2+}\) handling, oxidative stress, or nociception.

When considering the actual strategy for developing such drugs, however, there are at least two not easily tractable problems. The serious lack of structure-based chemical knowledge about the selectivity and efficacy for most TRP members is the first obvious one, although substantial progress is going on for analgesic therapy based on TRPV1 pharmacology (Szallasi et al., 2007). Virtually all drugs so far shown to be effective for cardiovascular TRP isoforms appear to be non-specific or have only narrow ranges of concentrations for their relatively selective actions, as exemplified by a pyrazole compound Pyr-3 for TRPC3 which reportedly inhibits experimentally-induced cardiac hypertrophy (Kiyonaka et al., 2009). The second type of difficulty for exploiting TRP-selective drugs is that connecting one TRP isoform to one end-function is almost always irrelevant oversimplification. This is because signaling pathways linked to TRP's appear to form a complex interwoven network with high degrees of divergence and convergence and with numerous feedback and forward regulations. In addition to this, activation of TRP channels depends intimately on local lipid dynamics in the cell membrane, which dramatically changes by lipid composition, voltage, temperature, membrane stretch, and enzyme-assisted catalysis (Hardie, 2007; Inoue et al.,
The Pathophysiological Implications of TRP Channels in Cardiac Arrhythmia

2009b). In the current absence of good knowledge about lipid physiology/pathophysiology for membrane proteins, these disadvantages would make it difficult to pharmacologically manipulate the gating of TRP channels. In this context, recent observations that activation of the cGMP-PKG signaling by nitric oxide, atrial natriuretic peptides and their structural and functional mimetics attenuate hypertrophic cardiac remodeling partly via inhibition of TRP channels (Inoue et al., 2010; Tsai & Kass, 2009) may provide a promising alternative maneuver to ameliorate the acquired arrhythmogenicity. Activation of this pathway is also known to improve myocardial damage after myocardial infarction (Burley et al., 2007; Garcia-Dorado et al., 2009). Obviously, there is much to be learned about the pathogenic roles and therapeutic potential of cardiovascular TRP channels for cardiac arrhythmias.

6. Acknowledgment

Part of this work is supported by Grants-in-aid for Scientific Research on Innovative Areas (No.22136008) and Scientific Research (C) (No. 21590246), and a grant from Seizon Kagaku Institute to R.I.

7. References


The Pathophysiological Implications of TRP Channels in Cardiac Arrhythmia


The most intimate mechanisms of cardiac arrhythmias are still quite unknown to scientists. Genetic studies on ionic alterations, the electrocardiographic features of cardiac rhythm and an arsenal of diagnostic tests have done more in the last five years than in all the history of cardiology. Similarly, therapy to prevent or cure such diseases is growing rapidly day by day. In this book the reader will be able to see with brighter light some of these intimate mechanisms of production, as well as cutting-edge therapies to date. Genetic studies, electrophysiological and electrocardiographyc features, ion channel alterations, heart diseases still unknown, and even the relationship between the psychic sphere and the heart have been exposed in this book. It deserves to be read!

How to reference
In order to correctly reference this scholarly work, feel free to copy and paste the following: