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Cyclic ADP-Ribose and NAADP in Vascular Regulation and Diseases

Pin-Lan Li*, Yang Zhang, Justine M. Abais, Joseph K. Ritter, and Fan Zhang

Department of Pharmacology and Toxicology, Medical College of Virginia Campus, Virginia Commonwealth University, VA 23298, USA

Abstract

Cyclic ADP-ribose (cADPR) and nicotinic acid adenine dinucleotide phosphate (NAADP), two intracellular Ca^{2+} mobilizing second messengers, have been recognized as a fundamental signaling mechanism regulating a variety of cell or organ functions in different biological systems. Here we reviewed the literature regarding these ADP-ribosylcyclase products in vascular cells with a major focus on their production, physiological roles, and related underlying mechanisms mediating their actions. In particular, several hot topics in this area of research are comprehensively discussed, which may help understand some of the controversial evidence provided by different studies. For example, some new models are emerging for the agonist receptor coupling of CD38 or ADP-ribosylcyclase and for the formation of an acidic microenvironment to facilitate the production of NAADP in vascular cells. We also summarized the evidence regarding the NAADP-mediated two-phase Ca^{2+} release with a slow Ca^{2+} -induced Ca^{2+} release (CICR) and corresponding physiological relevance. The possibility of a permanent structural space between lysosomes and sarcoplasmic reticulum (SR), as well as the critical role of lysosome trafficking in phase 2 Ca^{2+} release in response to some agonists are also explored. With respect to the molecular targets of NAADP within cells, several possible candidates including SR ryanodine receptors (RyRs), lysosomal transient receptor potential-mucolipin 1 (TRP-ML1) and two pore channels (TPCs) are presented with supporting and opposing evidence. Finally, the possible role of NAADP-mediated regulation of lysosome function in autophagy and atherogenesis is discussed, which may indicate a new direction for further studies on the pathological roles of cADPR and NAADP in the vascular system.

Keywords

Calcium Mobilization; Lysosomal Channels; Signal Transduction; ADP-Ribose; Vasoconstriction; Autophagic Flux; Vesicle Trafficking; Intracellular Ca^{2+} Stores

INTRODUCTION

It is widely accepted that in vascular cells, which mainly include vascular endothelial cells (ECs) and vascular smooth muscle cells (VSMCs), Ca^{2+} signaling is one of the most

important signaling mechanisms leading to vascular cell activation, producing vascular tone and vasomotor responses to various agonists and stimuli (Berridge, 1994; Berridge, 1997; Himpens et al., 1995; Nelson et al., 1990). Over the last 30 years, numerous studies have shown that intracellular Ca^{2+} concentration ($[\text{Ca}^{2+}]_i$) in vascular cells, particularly in VSMCs, is determined by both the influx of extracellular Ca^{2+} and the mobilization of Ca^{2+} from intracellular stores. In the search for intracellular Ca^{2+} mobilizing second messengers, the discovery both of inositol 1,4,5-tris-phosphate (IP_3) in the early 1980s (Streb et al., 1983) and of cyclic ADP-ribose (cADPR) and nicotinic acid adenine dinucleotide phosphate (NAADP) in the late 1980s (Lee et al., 1989) made a paradigm shift in how we understand intracellular Ca^{2+} mobilization in vascular cells and its associated physiological and pathological relevance. In this review we briefly summarized the current progress of knowledge on cADPR and NAADP in vascular cells and provided an overview regarding the role of both second messengers in the regulation of vascular function and development of related vascular diseases.

PRODUCTION OF cADPR AND NAADP IN VASCULAR CELLS

cADPR was first discovered by Lee and associates in sea urchin eggs (Lee et al., 1989) and then detected in a variety of mammalian tissues or cells such as heart, liver, spleen, brain and red blood cells, lymphocytes, pituitary cells and cultured renal epithelial cells (Beers et al., 1995; Galione et al., 1991; Koshiyama et al., 1991; Lee and Aarhus, 1993; Takasawa et al., 1993). Basal concentrations of cADPR in cardiac muscle, liver and brain are estimated at 100–200 nM (Ladd et al., 1996). Homogenates prepared from dissected small bovine coronary arteries, cultured arterial ECs and smooth muscle cells produced cADPR and its metabolite ADPR, when incubated with NAD (Li et al., 2000a; Li et al., 2000b; Li et al., 1997; Li et al., 1998). Tissue cADPR concentration in the coronary smooth muscle is about 150 nM (Li et al., 1998). The cADPR extracted from the reaction mixture of NAD with cultured smooth muscle cells or arterial homogenates is capable of stimulating Ca^{2+} release *in vitro* using single cell Ca^{2+} fluorospectrometry. Recently, we detected cADPR in coronary arterial ECs, also in the nM range (Zhang et al., 2006b).

Similarly, homogenates or microsomes from VSMCs converted NADP^+ along with nicotinic acid into NAADP in a concentration-dependent manner at pH of 4.5, which had similar efficiency to that observed for cADPR production under pH 7.4, indicating that NAADP is an enzymatic product of NADP^+ in these vascular cells. In VSMCs from other vascular beds such as renal, cerebral and pulmonary vasculatures, NAADP was also detected with a range of 4–16 nM (Churamani et al., 2004; Kinnear et al., 2004). More recently, intracellular NAADP levels were also detected in ECs ($1.774\ 88 \pm 0.65$ pmol/mg protein), which could be produced by selective histamine 1 receptor (H1R) stimulation (Esposito et al., 2011). It is clear that both VSMCs and ECs are capable of producing NAADP as a specific second messenger to mobilize Ca^{2+} from intracellular stores.

Enzymatic Products of ADP-Ribosylcyclase

cADPR—cADPR can be synthesized from NAD via the action of ADP-ribosylcyclase. Once formed, cADPR can be further hydrolyzed by cADPR hydrolase to ADPR. Therefore, the cellular cADPR level is determined by the expression and activity of these enzymes.

Both ADP-ribosylcyclase and cADPR hydrolase are membrane-bound enzymes in a wide range of mammalian tissues including arterial smooth muscle (Franco et al., 1994; Zocchi et al., 1993). It has been reported that the human lymphocyte differentiate antigens CD38 and CD157 are highly homologous with *Aplysia* ADP-ribosylcyclase, which possesses multiple forms of enzymatic activity including NAD glycohydrolase, ADP-ribosylcyclase and cADPR hydrolase activity (Adebanjo et al., 2000; Franco et al., 1994; Zocchi et al., 1993). These CD proteins are considered to be a molecular switch in regulating the cellular levels of cADPR by balancing its synthesis and hydrolysis. In response to stimuli, this multi-functional enzyme can be aggregated and internalized into the cytoplasm where it can more efficiently produce or metabolize cADPR. By Western blot analysis and RT-PCR, we demonstrated that CD38 was detectable in coronary arterial smooth muscle. In these experiments, two immunoreactive bands with molecular sizes of 42 and 90 kDa were recognized by a monoclonal antibody against CD38 in coronary arterial homogenates and microsomes (Li et al., 1997). Removal of CD38 by immunoprecipitation significantly decreased the production and catabolism of cADPR in these arterial homogenates. In CD38^{-/-} mice, very low cADPR levels and no detectable ADP-ribosylcyclase activity were observed in lung, kidney, coronary arterial tissue, and renal afferent arterioles dissected from these mice (Boini et al., 2011; Deshpande et al., 2005; Li et al., 2000a; Tegatz et al., 2005a; Zhang et al., 2010). Interestingly, this CD38-associated enzyme in coronary VSMCs not only produces cADPR, but also metabolizes cADPR into ADPR through its bifunctional domain. Therefore, intracellular cADPR levels could be dynamically maintained by switching of different functional CD38 domains (Li et al., 1998). It is now well accepted that an enzymatic pathway responsible for the formation and metabolism of cADPR is present in vascular cells. cADPR is formed from NAD⁺ by ADP-ribosylcyclase and metabolized into ADPR by the hydrolase activity of the same enzyme, which could be associated with the multifunctional activity of CD38 (Li et al., 1997; Li et al., 2002; Li et al., 1998; Zhang et al., 2001).

NAADP—Although the metabolites of NADP⁺ was reported to induce Ca²⁺ release from sea urchin egg microsomes by Lee and his associates in 1987 (Clapper et al., 1987), this NADP⁺ metabolite was only identified as NAADP nearly a decade later (Chini et al., 1995; Lee and Aarhus, 1995). Since then, the enzymatic pathways for NAADP production and metabolism have been characterized. Similar to cADPR production, soluble protein *Aplysia* ADP-ribosylcyclase and its membrane-bound homologs, CD38 and CD157, have also been reported to be involved in the production of NAADP (Aarhus et al., 1995; Galione et al., 1993; Lee, 1997; Lee, 2005). These enzymes can exchange the terminal nicotinamide group of the NADP⁺ with nicotinic acid to produce NAADP through a base—exchange reaction, which has been shown in a variety of cells and tissues such as sea urchin eggs, pancreatic acinar cells, human T lymphocytes, rat brain, and smooth muscle cells (Ge et al., 2002; Ge et al., 2003; Lee and Aarhus, 2000; Li et al., 2001). In addition to membrane-bound CD38 and CD157, a cytosolic soluble ADP-ribosylcyclase isoform or CD38 are also interestingly found in VSMCs (Lee and Aarhus, 1991; Rusinko and Lee, 1989). Our previous studies demonstrated that cytosolic ADP-ribosylcyclase activity may be primarily derived from internalized CD38 in coronary arterial smooth muscle, which may be associated with lipid raft clustering and endocytosis in seconds (Jia et al., 2008). Other studies also demonstrated

that CD38 internalization is important in mediating cADPR production in cell cytosol (Chidambaram and Chang, 1999; Han et al., 2002; Zocchi et al., 1999). Therefore, based on the current understanding, conversion of NAAP⁺ to NAADP due to high levels of ADP-ribosylcyclase activity may be associated with the internalization of a membrane-bound enzyme, although there remains to be some unidentified pathways. This internalized cytosolic ADP-ribosylcyclase or CD38 is responsible for catalyzing the exchange of the nicotinamide group of NADP⁺ with nicotinic acid to produce NAADP under acidic conditions in VSMCs (Zhang et al., 2006a). Our recent studies in mouse coronary artery indeed demonstrated that CD38 and its cytosolic isoforms are responsible for NAADP production in response to death receptor activation, endothelin and oxidant stimulation (Xu et al., 2012b; Zhang et al., 2010).

More recently, CD38 has also been found to hydrolyze NAADP to ADP-ribose 2'-phosphate. This activity of CD38 in the degradation of NAADP was greatly increased at acidic pH, which is determined by acidic residues at the active site of this enzyme. X-ray crystallography of the CD38 complex or purified ADP-ribosylcyclase with substrates demonstrated that acidic residues at the active sites of both enzymes determine NAADP synthesis or hydrolysis and that these residues are Glu-146 and Asp-155. Changing Glu-146 or Asp-155 by site-directed mutagenesis could eliminate their strong pH dependence (Graeff et al., 2006). However, in myometrial cells, NAADP could not be produced by either CD38 or a base-exchange reaction (Soares et al., 2007), but by some unidentified pathway. It seems that different cell types use different enzymatic pathways to produce NAADP.

Although NAADP production has been reported in various tissues and cells, it remains a mystery how an acidic reaction of ADP-ribosylcyclase or its human homologue CD38 produces NAADP either at the cell membrane or in the cytosol, where the pH is around 7.4. Furthermore, it is unknown why some agonists or stimuli can selectively elicit such acidic reaction of ADP-ribosylcyclase. Based on our current studies on redox signaling and lipid raft-associated transmembrane signaling mechanisms (Jia et al., 2008; Xu et al., 2012a; Xu et al., 2012b), we believe that an acid microenvironment may be generated locally in response to some agonists or stimuli in vascular cells. For example, we found that both ET-1 and FasL preferably stimulate NAADP production in VSMCs (Zhang et al., 2010; Zhang et al., 2006a), which may be associated with their ability to generate a local acidic microenvironment at the cell membrane, facilitating a base exchange reaction via ADP-ribosylcyclase or CD38. It is assumed that lipid raft clustering may be involved in the formation of this local acidic environment if agonist receptors are linked to lipid rafts. As shown in our previous studies and by others, CD38 or ADP-ribosylcyclase activity may be activated through membrane lipid raft clustering, (Deaglio et al., 2007; Jia et al., 2008; Munoz et al., 2003; Zilber et al., 2005). Upon agonist stimulation, membrane raft clustering is usually induced by lysosome fusion to the cell membrane, which translocates not only critical molecules for raft clustering such as acid sphingomyelinase (ASM) (only present in lysosomes), but also lysosomal vacuolar H⁺-ATPase. The latter provides a local acidic environment which maintains the activity of translocated ASM to amplify the production of ceramide and form lipid raft platforms (Xu et al., 2012a). Such membrane microenvironment of acidic pH generated and maintained by vacuolar H⁺-ATPase may be critical for ADP-ribosylcyclase or CD38 to produce NAADP within lipid raft platforms.

The formation of this acidic microenvironment may be an important step for the enzymatic activity of CD38, which is actively regulated by agonists or other stimuli outside the cells. Since such acidic environment is formed through rapid lysosome trafficking and fusion and subsequent lipid rafts clustering, the substrates in CD38-mediated enzymatic response such as nicotinic acid or other related substrates may be transported and maintained enough concentrations in the local area of lipid raft platforms. It has been well known that lysosome membrane has numerous transporter systems including nucleosides and nucleotides (Pisoni and Thoene, 1991). However, so far there is no study done to elucidate how these transporters work to provide nicotinic acid for NAADP production, which will be another interesting topic for future studies.

Activation of ADP-Ribosylcyclase in Vascular Cells

VSMCs—Many studies have been done in VSMCs to test whether ADP-ribosylcyclase can be activated in response to different agonists or stimuli (Bai et al., 2005; Evans et al., 2005; Zhang and Li, 2006). In this regard, we have reported that incubation of coronary VSMCs with oxotremorine, a specific M₁ mAChR agonist, produced time- and concentration-dependent activation of ADP-ribosylcyclase, which was blocked by both ADP-ribosylcyclase inhibitor nicotinamide and by specific M₁ mAChR antagonist, pirenzepine (PIR). The activation of ADP-ribosylcyclase occurred rapidly within the first minute of oxotremorine incubation with coronary arterial smooth muscle cells (CASMCs) (Ge et al., 2003). In pulmonary circulation, activation of ADP-ribosylcyclase has been reported as a primary trigger of hypoxia-induced contraction of small arteries or arterioles (Dipp and Evans, 2001; Dipp et al., 2001). In addition, Ang II has been demonstrated to activate ADP-ribosylcyclase in neonatal rat cardiac myocytes (Higashida et al., 2000), and ET-1 may increase cADPR production via activation of ADP-ribosylcyclase in rat mesenteric small arteries (Giuliumian et al., 2000) or shark anterior mesenteric arteries (Fellner and Parker, 2004). However, recent studies have demonstrated that ET-1 may also stimulate the production of NAADP (Kinnear et al., 2004; Lee, 2005; Zhang et al., 2006a). More recently, we reported that death receptor ligand FasL markedly increased NAADP production in CASMCs from wild-type mice (CD38^{+/+}), but not in CASMCs from CD38 knockout (CD38^{-/-}) mice (Zhang, et al., 2010). Other vascular agonists that increase ADP-ribosylcyclase activity and thereby enhance cADPR production include β -adrenergic agonists (Boittin et al., 2003) and urocortin (Sanz et al., 2003).

On the other hand, NO donor sodium nitroprusside (SNP) was reported to decrease ADP-ribosylcyclase activity, inhibiting the production of cADPR in VSMCs, either from coronary arteries (Yu et al., 2000) or from airway (White et al., 2002). However, these results are not in concordance with the findings of previous studies in nonvascular cells, where NO increases the production of cADPR in nonvascular cells such as rat parotid acinar cells (Looms et al., 2001) and urchin eggs (Willmott et al., 1996). It remains unknown why NO decreases ADP-ribosylcyclase activity in VSMCs, but increases it other cells.

ECs—It has been reported that in non-vascular tissues CD38 or ADP-ribosylcyclase may be a link from bradykinin receptors to its effector enzymes such as NOS (Deshpande et al., 2003; Higashida et al., 1996; Higashida et al., 2001). Given that bradykinin is a typical

stimulator of endothelium-dependent vasodilator, it is possible that cADPR production via CD38-ADP-ribosylcyclase may be a critical mechanism regulating endothelial Ca^{2+} and corresponding function. By direct measurement of cADPR production in ECs, we indeed demonstrated that bradykinin induced cADPR production and thereby evoked a Ca^{2+} release from RyR-sensitive stores, which was accompanied by an increase in NO production (Yu et al., 2000; Zhang et al., 2005; Zhang et al., 2006b). Interestingly, a recent study reported that in rat ECs, NAADP production may be activated by endothelium-dependent vasodilator acetylcholine (Brailoiu et al., 2010c). In addition, human ECs were also shown to produce NAADP in response to the specific histamine H1 receptor agonist, 2-[(3-Trifluoromethyl)phenyl] histamine dimaleate (TMPH), which mediates H1R-induced Ca^{2+} release from acidic organelles and the endoplasmic reticulum (Naylor et al., 2009). It has been suggested that NAADP production and subsequent Ca^{2+} release may underlie histamine-mediated endothelial activation and the exocytosis of VWF (Esposito et al., 2011).

Receptor-Effector Coupling—As discussed above, many agonists can activate or inhibit ADP-ribosylcyclase to produce cADPR and NAADP, which mediate Ca^{2+} release from intracellular stores. It has been shown that these agonists exert action through their receptors in different cells, such as muscarinic acetylcholine receptors in NG108–15 neuronal cells (Higashida et al., 2007), Ang II or β -adrenergic receptors in ventricular myocytes (Gul et al., 2008), and M1 muscarinic receptors, ET-1 receptors and Fas in VSMCs (Ge et al., 2003; Zhang et al., 2010; Zhang et al., 2006a). However, little is known regarding the link between activation of these agonist receptors and ADP-ribosylcyclase. The major mammalian analogues of ADP-ribosylcyclase, namely CD38 and CD157, are pleiotropic ectoenzymes, which act independently as both receptors and enzymes. However, it still remains unknown how CD38 or CD157 is able to produce intracellular signaling molecules. Although there is evidence that internalization and aggregation of CD38 are importantly involved in signaling of this cADPR-producing enzyme (Franco et al., 1994; Howard et al., 1993; Koguma et al., 1994; Lacapere et al., 2003; Lee et al., 1993; Zocchi et al., 1995; Zocchi et al., 1999), how CD38 internalization and aggregation is linked to agonist receptors is still an unresolved issue.

In some studies, various receptors have been proposed to be directly coupled with ADP-ribosylcyclase, where their activation may result in increased production of cADPR (Higashida et al., 1999; Higashida et al., 1997). With respect to bradykinin, there is evidence that B2 receptors are directly coupled to the membrane-bound form of ADP-ribosylcyclase in human airway smooth muscle cells (Deshpande et al., 2003) and NG108–15 neuronal cells (Higashida et al., 1996), suggesting that bradykinin could stimulate cADPR production via this direct coupling mechanism. G-proteins may contribute to this linking of agonist receptor to ADP-ribosylcyclase since it is well known that G-proteins play an important role in linking cell surface receptors to intracellular second messengers. In this context, G-proteins may link Ach receptors to ADP-ribosylcyclase and therefore Ach stimulates the production of cADPR through G-protein activation. This view has been supported by several reports from vascular and non-vascular tissues or cells (Higashida et al., 1999; White et al., 2003; Wu et al., 1997). Given the ectoenzyme nature of CD38, how such G-protein-

mediated receptor-effector coupling occurs is still speculative and further studies are needed to address the direct structural or functional connection between G-proteins and any enzyme determining intracellular cADPR levels.

Rather than focusing on this classical receptor-effector coupling model of transmembrane signaling, we recently developed a new model for temporospatial activation of ADP-ribosylcyclase in VSMCs in response to different agonists or stimuli. In this model, selected agonists or stimuli will concurrently activate membrane NADPH oxidase to convert NAD(P)H to NAD(P)⁺ and O₂⁻. This membrane NADPH oxidase-derived O₂⁻ is released out of VSMCs and nearby activates ADP-ribosylcyclase (CD38) through its rapid internalization and dimerization. Activated CD38 uses increased NAD⁺ as the substrate to produce cADPR (or NADP⁺ to produce NAADP). The coupling of agonist receptors to ADP-ribosylcyclase is via membrane NADPH oxidase rather than G-proteins. This cross talk between NADPH oxidase and ADP-ribosylcyclase may explain why some agonists link to ADP-ribosylcyclase, but others do not (Fig. 1). There are several lines of evidence supporting this working model for agonist receptor-ADP-ribosylcyclase coupling. First, NADPH oxidase can be activated by many agonists such as Ang-II, Oxo, ET-1, and FasL (Bao et al., 2010; Zhang et al., 2008; Zhang et al., 2007). NADPH oxidase is also reported to produce and release O₂⁻ outside VSMCs (Zhang et al., 2006b). Importantly, one of the intracellular products from NADPH oxidase, NAD⁺ or NADP⁺ can provide substrate for ADP-ribosylcyclase. Second, it has been reported that ADP-ribosylcyclase or CD38 dimerizes in response to increases in intracellular oxidants, which enhances the catalytic activity of ADP-ribosylcyclase (Chidambaram et al., 1998; Guida et al., 1995). Several studies have demonstrated that cysteine residues in CD38 or ADP-ribosylcyclase determine the enzyme function as ADP-ribosylcyclase or cADPR hydrolase (Tohgo et al., 1994). The oxidation of cysteine molecules may lead to the formation of one or several disulfide bonds, which induces dimerization of the enzyme protein. In other studies, the activity of ADP-ribosylcyclase was found to be inhibited by disulfide bond reducing reagents (Berruet et al., 1998; Galione et al., 1993). Lastly, some agonists such as oxtremorine, ET-1 and FasL have been reported to stimulate membrane lipid raft clustering and internalization of CD38 (Gambara et al., 2008; Jia et al., 2008; Trubiani et al., 2004). Internalized CD38 or ADP-ribosylcyclase can form dimers to produce maximal activity. As mentioned above, lipid raft clustering of NADPH oxidase and CD38 with or without vacuolar H⁺-ATPase may determine the production of cADPR or NAADP since vacuolar H⁺-ATPase is important for the generation and maintenance of local acidic microenvironment (Xu et al., 2012a). If the local pH is around 4 in those raft platforms, NAADP may be produced; otherwise, cADPR may be synthesized and released.

Ca²⁺ MOBILIZING ACTION OF cADPR AND NAADP IN VASCULAR CELLS

Independent of IP₃, cADPR stimulates Ca²⁺ release from intracellular Ca²⁺ stores when given directly in VSMCs or in response to different agonists or stimuli. Kannan et al., reported that cADPR induces SR Ca²⁺ release in β -escin-permeabilized smooth muscle cells freshly isolated from porcine coronary arteries. In α -toxin permeabilized cells, we found that cADPR produces SR Ca²⁺ release in both cultured and freshly dissociated bovine coronary and rat renal VSMCs (Li et al., 2000a; Yu et al., 1996). This cADPR-induced Ca²⁺

release from the SR can be completely blocked by cADPR antagonist, 8-Br-cADPR, but not by IP₃R blockers. It is concluded that cADPR mobilizes intracellular Ca²⁺ through a mechanism independent of IP₃ in VSMCs. Recently, we also determined whether bradykinin-induced vasodilator response is directly linked to cADPR-mediated Ca²⁺ release from the endoplasmic reticulum (ER) in bovine coronary arterial ECs (Zhang et al., 2006b). Using a newly developed fluorescence imaging system to simultaneously measure Ca²⁺ transient and NO production in the intact arterial endothelium, we showed that bradykinin produced a rapid and transient increase in [Ca²⁺]_i that was accompanied by enhanced NO production (Zhang et al., 2006b). However, bradykinin-induced Ca²⁺ release and NO production were significantly attenuated by pretreatment of the arteries with cADPR-RyRs signaling inhibitors such as nicotinamide, 8-Br-cADPR or ryanodine. This supports the view that bradykinin-induced Ca²⁺ increase in arterial ECs is through cADPR-mediated Ca²⁺ release from the ER and via RyR activation.

NAADP has been proposed to act as a ubiquitous Ca²⁺ messenger, and this nucleotide is one of the most potent intracellular Ca²⁺ mobilizing molecules (Evans et al., 2005; Galione, 2006; Lee, 2005; Yamasaki, et al., 2005a). The Ca²⁺ mobilizing action of NAADP is even stronger than that induced by commonly known Ca²⁺ mobilizing second messengers IP₃ and cADPR (Clapper et al., 1987; Lee and Aarhus, 1995). Two working models have been proposed to interpret the different actions of NAADP in mobilizing intracellular Ca²⁺ (Yamasaki et al., 2005a). In the first model, the ER or SR that expresses IP₃R and RyRs is responsible for NAADP-induced Ca²⁺ release, where NAADP may interact either directly with RyRs or via a separate protein that may indirectly activate RyRs (Dammermann and Guse, 2005; Gerasimenko et al., 2003b). This model may work in several cell types such as T-lymphocytes (Dammermann and Guse, 2005), cardiac cells (Mojzisova et al., 2001) and skeletal muscle (Hohenegger et al., 2002). In these cells, the target for the actions of NAADP is the RyR on the ER or SR (Dammermann and Guse, 2005). The second model relates to a two-pool mechanism, which is based on the assumption that an NAADP-sensitive Ca²⁺ store exists in arterial myocytes, which is possibly a thapsigargin-insensitive acidic store (Churchill et al., 2002). This NAADP-sensitive Ca²⁺ store is responsible for a localized signal, which triggers CICR to cause global Ca²⁺ increases through IP₃R and RyRs on the SR (Cancela et al., 1999; Cancela et al., 2000; Churchill and Galione, 2000; Churchill and Galione, 2001). Considering a wide variety of cellular processes regulated by changes in intracellular Ca²⁺ concentrations including fertilization to cell death, it is possible that this temporospatial Ca²⁺ signaling pattern related to NAADP importantly contributes to the regulation of different cell functions, either through its action as second messenger or via its activity to synchronize the actions of other second messengers (Albrieux et al., 1998; Brailoiu et al., 2005; Brailoiu et al., 2001; Chameau et al., 2001; Johnson and Mislser, 2002; Kinnear et al., 2004; Masgrau et al., 2003; Yamasaki et al., 2004; Yamasaki et al., 2005b).

Over the last two decades, numerous studies have been conducted to explore the possible molecular mechanisms mediating the action of cADPR and NAADP to release Ca²⁺ from the intracellular stores. It is clear that several molecules centered on RyRs may be the targets of cADPR to mobilize Ca²⁺ from the SR in VSMCs. For NAADP, in addition to targeting RyRs on the SR, lysosomal channels are reported to be involved in a lysosomal burst of

Ca²⁺ that triggers CICR, leading to global Ca²⁺ mobilization. Below is an overview of these intracellular targets for the action of cADPR and NAADP.

Intracellular Targets of cADPR Action

RyR in SR—There is considerable electrophysiological evidence showing that the RyR/Ca²⁺ release channels reconstituted into a planar lipid bilayer are activated by cADPR in a variety of tissues or cells. In coronary arterial smooth muscle, a calcium channel with 245 pS conductance is present on the SR membrane and cADPR was found to increase the NP_O of these RyR/Ca²⁺ release channels in a concentration-dependent manner (Li et al., 2001). In the presence of ryanodine (50 μM), cADPR-induced activation of these channels was completely abolished. These results provided direct evidence that cADPR activates RyRs and therefore may serve as an endogenous activator or modulator of RyRs in these VSMCs. However, this view has been challenged by studies using other tissues such as neurons, myocardium and other smooth muscle, where cADPR was found to release Ca²⁺ independently of RyR (Kannan et al., 1996; Lahouratate et al., 1997; Sitsapesan et al., 1994). It seems that there exists tissue specific effects of cADPR on RyRs, which may be associated with the intermediate proteins or accessory proteins that regulate RyR activity.

It has been reported that cADPR participates in KCl, CaCl₂, Bay K 8644 (Ca²⁺ channel activator) and caffeine-induced Ca²⁺ release response in coronary and renal VSMCs, suggesting that cADPR contributes to Ca²⁺-induced Ca²⁺ release (CICR) (Teggatz et al., 2005b). In these smooth muscle cells, high extracellular Ca²⁺ (5 mM CaCl₂) and agonist Ach produced 1–1.5 Hz oscillations, which were blocked by CICR inhibitor tetracaine and cADPR antagonist 8-Br-cADPR (Li et al., 2000a). Kannan et al., have also reported that cADPR increased Ach-induced Ca²⁺ oscillations which was blocked by cADPR antagonist 8-amino-cADPR in porcine tracheal smooth muscle cells (Kannan et al., 1997). Taken together, these results demonstrated that cADPR is necessary for CICR and intracellular Ca²⁺ oscillation and that RyRs are the mechanistic link between cADPR and CICR or Ca²⁺ oscillations.

There are also two mechanistic models proposed to elucidate the role of endogenous cADPR in mediating vascular Ca²⁺ mobilization through RyRs. First, cADPR acts as a mediator to activate RyRs. In this regard, various agonists or stimuli activate ADP-ribosylcyclase to produce cADPR, which induces Ca²⁺ release from the SR by its direct action on the RyRs. Second, cADPR serves as a modulator of CICR or RyR reactivity. In this way, cytosolic cADPR sensitizes the RyRs, enhancing CICR activated by agonists or Ca²⁺ influx.

FKBP 12.6 Proteins—FKBP12.6 is a ubiquitous 12.6-kDa cytosolic protein that binds to one RyR monomer and its activity is inhibited by the immunosuppressant drug FK506 and rapamycin. In nonvascular cells, Ca²⁺ release from the SR is inhibited when FKBP12.6 is bound to the RyR, and dissociation of FKBP 12.6 from the RyR releases Ca²⁺. This 12.6 kDa protein is also expressed in coronary arterial smooth muscle (Tang et al., 2002). Blockade, dissociation or removal of FKBP12.6 protein from the RyR substantially abolished cADPR-induced activation of RyR/Ca²⁺ release channels on lipid bilayer membrane. Ligand binding experiments have demonstrated that cADPR can directly bind to

FKBP12.6 in islet microsomes (Tang et al., 2002). Using confocal fluorescence imaging, we have demonstrated that FKBP12.6 colocalizes with RyRs in renal arterial myocytes (Teggatz et al., 2005b). Ca^{2+} influx by CaCl_2 significantly decreased this colocalization, and 8-Br-cADPR reversed CaCl_2 effects suggesting that cADPR is involved in the dissociation of FKBP 12.6 protein from RyRs under this condition. These results indicate that cADPR exerts its action by dissociating FKBP12.6, resulting in Ca^{2+} release from the SR in VSMCs. Recently, studies from other groups also demonstrated such contribution of FKBP to the action of cADPR in the mobilization of Ca^{2+} (Morita et al., 2006; Noguchi et al., 1997; Wang et al., 2004). However, some studies could not obtain similar results (Bradley et al., 2003; Copello et al., 2001; Zhang et al., 2009b). Therefore, further investigations are needed to solve such controversy by demonstrating the possible tissue specific action of this intracellular regulatory mechanism.

Intracellular Targets of NAADP Action

RyRs in SR—Depending on the tissues or cells studied, RyRs were reported to be a possible target for the action of NAADP in mobilizing Ca^{2+} from intracellular stores. There is evidence that reconstituted RyR1 and RyR2 channels from cardiac cells (Mojzisova et al., 2001) and skeletal muscle (Hohenegger et al., 2002) are sensitive to NAADP to increase Ca^{2+} channel activity. In some other tissues or cell preparations, in particular T-lymphocytes, NAADP was found to target RyRs localized to the ER (Dammermann and Guse, 2005; Gerasimenko et al., 2003a; Gerasimenko et al., 2003b; Langhorst et al., 2004). However, other studies agreed more with the findings that in sea urchin eggs, there is an acidic compartment related to lysosomes (Churchill et al., 2002). It has been proposed that an NAADP-sensitive acidic Ca^{2+} store exists in some type of cells including arterial myocytes, which is a thapsigargin-insensitive acidic store mainly shown in lysosomes or lysoendosomal compartments. NAADP is demonstrated to first activate Ca^{2+} bursts as a triggering mechanism and then lead to global Ca^{2+} mobilization through IP_3 R and RyRs in the SR, a so-called two-pool mechanism. It is assumed that the NAADP-sensitive Ca^{2+} store or acidic Ca^{2+} store is responsible for a localized signal, where latter triggers CICR to cause global Ca^{2+} increases through RyRs or IP_3 R on the SR (Kinnear et al., 2004; Kinnear et al., 2008; Zhang et al., 2010). In VSMCs, this two-pool mechanism has been demonstrated to function in response to different agonists such as ET-1 and FasL or by delivery of NAADP into the cells (Zhang et al., 2010; Zhang et al., 2006a). Lipid bilayer reconstitution and Ca^{2+} imaging in intact CASMCs did not confirm that NAADP directly activates RyRs on the SR from VSMCs (Zhang et al., 2009a; Zhang and Li, 2007; Zhang et al., 2006a). However, RyR is an important SR receptor mediating CICR upon NAADP stimulation, and therefore blockade of RyRs still abolishes global Ca^{2+} increases, which may be of physiological significance in many biological processes.

Lysosome (Lyso)-SR Junction—Although the two-pool mechanism is attractive in its interpretation of a two phase Ca^{2+} release induced by NAADP, some issues remain to be addressed. For example, NAADP-induced CICR has a long delayed second phase Ca^{2+} release (seconds to minutes) (Evans and Cannell, 1997; Zhang et al., 2010; Zhu et al., 2010a), which is very different from the classical CICR reported previously (Fleischer and Inui, 1989; Franco et al., 1994; Galione 1993; Galione et al., 1991; Hirst et al., 1994). It

remains unknown why such a long delay occurs and what physiological relevance this delayed second phase Ca^{2+} in CICR may have. Given the principle of economic design in the biological system, the significance of such unconventional CICR through lysosomal triggering is imperative to define.

The Lyso-SR junction between lysosomal clusters and a subpopulation of SR was proposed to form a trigger zone, where Ca^{2+} released from lysosomes activates a global Ca^{2+} response via CICR in pulmonary VSMCs and some other cells (Kinnear et al., 2004; Zhang et al., 2010; Zhang et al., 2006a). However, the structural and functional characteristics of this Lyso-SR junction have yet to be completely revealed. In collaboration with Dr. van Breemen, who has extensive experience in characterizing plasma membrane-SR (PM-SR) and mitochondria-SR (Mito-SR) junctions in smooth muscle cells under resting and contracting condition (Dai et al., 2005; Poburko et al., 2004), we performed electronic microscopy and found that there are Lyso-SR junctions around 30–80 nm in VSMCs, which are relatively large compared to PM-SR and Mito-SR junctions. However, these Lyso-SR junctions are rather heterogeneous within arterial myocytes (unpublished data). Although it seems that some Lyso-SR junctions are present within VSMCs, there may not be a permanent structural space between lysosomes and SRs. Given the great mobility of lysosomes within cells, it is possible that Lyso-SR junctions depend upon lysosome trafficking and aggregation toward the SR as we hypothesized in CASMCs (Zhang et al., 2009a). It has been proposed that the global Ca^{2+} release following small Ca^{2+} bursts from lysosomes may be associated with lysosomal trafficking to the SR. Although small amounts of Ca^{2+} released from lysosomes may not be enough to activate global Ca^{2+} release from the SR, it may be enough to drive lysosome movement or aggregation. When these clustered or aggregated lysosomes work together, global Ca^{2+} release from the SR is activated (Zhang et al., 2009a). This interaction of lysosome and SR was also proposed later by Zhu et al., in pulmonary VSMCs (Zhu et al., 2010a). In some preliminary studies, we detected colocalization of lysosomal marker (GFP-Lamp1) with RyR3 or SR tracker (red fluorescent labeled) suggesting that there is lysosomal trafficking and aggregation toward the SR in coronary VSMCs when they were stimulated by death factor, FasL (Xu et al., 2011). In another study, we indeed demonstrated that NAADP stimulates lysosome trafficking through its Ca^{2+} mobilizing action (Zhang et al., 2011).

TRP-ML Channels in Lysosomes—The transient receptor potential-mucolipin (TRP-ML) subfamily of TRP channels consists of three mammalian members (TRP-ML1–3), which are relatively small proteins consisting of <600 amino acid residues with an expected molecular mass of approximately 56–65 kDa. TRP-ML1 is widely expressed and mainly resides in late endosomes/lysosomes (Bach 2005; Laplante et al., 2002; Laplante et al., 2004). It has been demonstrated that the protein encoded by the TRP-ML1 gene, MCOLN1, has six predicted transmembrane domains, a putative channel pore, and is predominantly expressed in endosomes or lysosomes. Due to cleavage and other modifications, TRP-ML1 may be detected in different sizes from 36–75 kDa in lysosome membranes of native cells (Kiselyov et al., 2005; Yamaguchi et al., 2011). Within TRP-ML1, there is a TRP channel-homologous region located within amino acids 331–521 and an internal Ca^{2+} and Na^{+} channel pore region between amino acids 496 and 521. Mutations of MCOLN1 are

implicated in the pathogenesis of a neurological disease, namely, mucopolisaccharidosis Type IV (MLIV). This disease is a lysosomal storage disorder that is characterized by severe neurologic and ophthalmologic abnormalities (Bach 2001; Kiselyov et al., 2005; Laplante et al., 2004; Raychowdhury et al., 2004; Slaugenhaupt, 2002; Sun et al., 2000). Compared to TRP-ML1, the functions of TRP-ML2 and TRP-ML3 are as extensively characterized until recently (Nilius et al., 2007).

In the search for the molecular target of NAADP action in lysosomes from native cells or tissues, reconstitution of liver lysosome preparations to characterize the possible Ca^{2+} channels was first done in our laboratory using lipid bilayer. Liver lysosomes were used since lysosome preparations were well established with using large volumes of tissues. Our work, published in 2007, demonstrated a 174 pS of a reconstituted Ca^{2+} channel with all the biophysical and pharmacological features of TRP-ML in liver lysosomes (Zhang and Li, 2007). We further characterized this TRP-ML in bovine coronary arterial muscle cells using lysosome preparations by constitution in lipid bilayer. It was demonstrated that the reconstituted lysosomal channel is also a voltage-dependent Ca^{2+} channel with a conductance of 145 pS in arterial muscle preparation (Zhang et al., 2009a). At the same time, Xu and his associates developed an elegant approach using patch clamp techniques and recorded Fe^{2+} and Ca^{2+} channel activity directly from lysosome membrane in different cell types with TRP-ML transgenes, discovering that TRP-ML1 is an inwardly rectifying, proton-impermeable, Ca^{2+} and $\text{Fe}^{2+}/\text{Mn}^{2+}$ dually permeable cation channel (Dong et al., 2008; Dong et al., 2010; Dong et al., 2009). More recently, the same group used a genetically encoded Ca^{2+} indicator (GCaMP3) attached directly to TRP-ML1 to directly measure Ca^{2+} release from lysosomes via TRP-ML1 (Martelli et al., 2012), which further confirm the nature of TRP-ML1 as a lysosomal Ca^{2+} release channel.

NAADP was found to activate TRP-ML channels reconstituted from liver and coronary arterial muscle lysosome preparations in a concentration-dependent manner, which was featured by a self-desensitization at high concentrations of NAADP. In particular, when the bilayer preparations containing coronary arterial muscle lysosomal channels were pretreated with a subthreshold concentration of NAADP, the activity of these channels in response to higher concentrations of NAADP was substantially attenuated, suggesting that these lysosomal channels can be desensitized. This self-desensitization property of lysosomal ion channels has also been shown in the actions of NAADP as a Ca^{2+} releasing second messenger in other cells such as sea urchin egg fractions or intact egg cells (Aarhus et al., 1996; Bach, 2005; Genazzani et al., 1996). Although the mechanism mediating this self-desensitization is not yet clear, it is assumed that the NAADP receptor allosteric site transformation between high- and low- binding affinity to NAADP may play an important role, which is very similar to the regulatory machinery observed in well-studied IP_3 receptors (Hirata et al., 1990). However, more studies will be needed to further elucidate the underlying mechanism responsible for such lysosomal channel desensitization and to address the possible physiological significance of this phenomenon, in particular, in the regulation of NAADP-mediated Ca^{2+} signaling and related function in the vasculature.

Pharmacologically, these reconstituted lysosomal channels in both liver and coronary arterial lysosomes were blocked by commonly used antagonists of TRP-ML1 channels such

as dihydropyridine derivatives nifedipine and verapamil, sodium channel antagonist amiloride and an NAADP receptor antagonist PPADS (Yusufi et al., 2002; Zhang and Li, 2007). Furthermore, silencing the expression of TRP-ML1 gene in CASMCs with its specific siRNA substantially attenuated reconstituted lysosomal NAADP-sensitive Ca^{2+} release channel activity. Similarly, immunoprecipitation of TRP-ML1 from lysosome preparations of CASMCs with a specific antibody that was raised against its 101–150th amino acids, an epitope antigen sequence located on the lysosomal lumen-oriented TRP-ML1 loop between segment one and two, almost completely removed the channel activity and related response to NAADP. The use of an anti-TRP-ML1 antibody that was raised against a peptide mapping at the C terminus of TRP-ML1, a channel pore forming region, also attenuated NAADP-induced activation of reconstituted lysosomal Ca^{2+} channels. It is known that TRP-ML1 is a protein with full length of 580 amino acid containing six transmembrane segments and cytoplasm-resided C- and N- termini. The 101–150 antigen epitope is located on the lysosomal lumen-oriented loop between segment 1 and 2. The TRP cation channel domain of amino acids 331 to 521 spans transmembrane segments 3 to 6 (Cheng et al., 2010; Sun et al., 2000). All the results presented above that were obtained by gene silencing, deprivation of TRP-ML1 protein or interference of its channel pore formation strongly suggest that the activity of this reconstituted lysosomal NAADP-sensitive Ca^{2+} release channel represents a function of TRP-ML1 in CASMCs (Zhang et al., 2009a).

More recently, TRP-ML channel activity was reconstituted by using lysosomal preparation from wild-type (TRP-ML1^{+/+}) human fibroblasts, but not from TRP-ML1^{-/-} cells. Reconstituted TRP-ML1 channels in wild-type cells were stimulated by NAADP (0.01–1.0 μM) in a concentration-dependent manner. However, when a TRP-ML transgene was expressed in TRP-ML1^{-/-} cells, the channel activity can be observed and NAADP enhanced this channel activity. In intact cell experiments, microscopic Ca^{2+} imaging showed that NAADP significantly increased intracellular [Ca^{2+}] in TRP-ML1^{+/+} cells, but had no effect in TRP-ML1^{-/-} cells. If a TRP-ML1 transgene was expressed in TRP-ML1^{-/-} cells, the Ca^{2+} response to NAADP was restored to the level comparable to TRP-ML1^{+/+} cells. This further support the view that NAADP increases lysosomal TRP-ML1 channel activity to release Ca^{2+} and TRP-ML channels in lysosomes may regulate local compartmental Ca^{2+} which may be important for the control of lysosome function such as trafficking or intracellular signaling (Zhang et al., 2011).

Since the report of TRP-ML1 as a NAADP-sensitive lysosomal Ca^{2+} channel, some studies doubt that TRP-ML1 serves as a target of NAADP because different ligand binding experiments did not demonstrate TRP-ML1 binding to NAADP agonistic or antagonistic probes (Lin-Moshier et al., 2012; Pryor et al., 2006; Walseth et al., 2012). Given the multiple sizes of TRP-ML1 channel from 36–75 kDa in the lysosome membrane of native cells and its possible cleavage after function (Kiselyov et al., 2005; Yamaguchi et al., 2011), the results with negative binding of intracellular cytosol, particles or cell membrane with overexpressed TRP-ML1 gene may not rule out the role of TRP-ML1 as a target of NAADP or NAADP-sensitive lysosomal channels. It should be noted that in our original report (Zhang and Li, 2007), the action of NAADP through TRP-ML1 channels does not

necessarily indicate that it must use this channel as its receptor. Since there are evidence that mucolipin-1 is able to oligomerize and/or form complexes with other proteins (Manzoni et al., 2004; Miedel et al., 2006) and that this heteromeric formation to constitute cation-permeable pores is common in TRP channels (Clapham, 2003), it is possible that stimulation of some regulatory protein or some common accessory NAADP binding protein may facilitate NAADP-associated Ca^{2+} release from lysosomes (Zhang and Li, 2007; Zhang et al., 2011). Most recently, Dr. Guse further proposed NAADP binding proteins as a unifying hypothesis for the action of NAADP to activate multiple channels in different Ca^{2+} stores (Guse, 2012).

Interestingly, a recent study reported that TRP-ML1 and two-pool channels (TPCs) are present in the same complex, yet function as two independent organellar ion channels (Yamaguchi et al., 2011). Although TRP-ML1 channels were not confirmed to be the target for NAADP, it seems that such conclusion is only for TRP-ML1 channels in the plasma membrane rather than lysosomes, because the channel activity in those studies were only recorded in the plasma membrane of cells with overly expressed TRP-ML1 gene. However, these studies using patch clamp recordings of calcium activated ionic currents in acinar cells indeed showed that membrane channel activity is extensively correlated with both local (short-lived) and global (longer-lived) calcium increases. In these experiments, the lack of difference in Ca^{2+} -activated Cl^- current oscillation upon NAADP stimulation between wild-type and knockout cells may not be specific enough to define the role of TRP-ML1, because this Ca^{2+} -activated Cl^- current oscillation depends upon global Ca^{2+} increase, which may be influenced by NAADP action on cell depolarization (Brailoiu et al., 2009b; Moccia et al., 2006a; Moccia et al., 2006b). The cell depolarization will activate Ca^{2+} -activated Cl^- current. Furthermore, when cells are clamped at certain membrane potentials, NAADP may amplify the globalization of Ca^{2+} signals (Cancela et al., 2002). Therefore, direct recording of lysosomal TRP-ML1 currents or more direct measurements of Ca^{2+} release from lysosomes, rather than global Ca^{2+} levels are needed for a solid conclusion to specify the action of TRP-ML1 channels as an NAADP-sensitive lysosome responder.

Another concern relates to the use of cell lines or transgenic cells in many studies for testing the target of NAADP action (Kiselyov et al., 2005; Yamaguchi et al., 2011), where TRP-ML1 was very overly expressed and spread to other compartments of cells in addition to lysosomes. It has been reported that although the majority of TRP-ML1 is expressed in intracellular compartments when a transgene was introduced into cell lines, some of the overexpressed TRP-ML1 can be targeted to the plasma membrane, while under a moderate overexpression condition TRP-ML1 variants were not found at the plasma membrane. This is because saturation of the protein trafficking pathway by marked overexpression forces expression of significant amounts of TRP-ML1 on the plasma membrane. Such plasma membrane mistargeting provides opportunity for studies on TRP-ML1 channel properties using the whole cell patch clamp configuration of the cell plasma membrane (Kiselyov et al., 2005; Soyombo et al., 2006). Given that TRP-ML1 on the plasma membrane may not occur in native organs or cells, whether the results obtained from the plasma membrane regarding TRP-ML1 channels indicate its channel property in lysosomes may be questionable. In some studies, the results that TRP-ML1 failed to act as an NAADP sensitive channel were obtained by manipulating TRP-ML1 expression or modifying TRP-

ML integrability in SKBR3 cell lines (Yamaguchi et al., 2011). However, as mentioned above, overexpressed TRP-ML1 by these approaches may not reflect the real circumstances of TRP-ML1 in the native cells in terms of location and configuration. In addition, recent studies (Lin-Moshier et al., 2012; Walseth et al., 2012) have demonstrated that there may be some specific NAADP binding proteins, which may function as a cofactor to activate a Ca^{2+} channel such as TRP-ML1 or TPCs in response to NAADP. It is possible that the overexpressed TRP-ML1 has differences from the native channel protein in terms of the access of the NAADP binding proteins.

TPC Channels in Lysosome-Like Acidic Organelles—In 2009, Zhu, Evans, Galione and their associates in collaboration with other groups published a work indicating that TPCs may be an NAADP-targeted Ca^{2+} channel in lysosome-like acidic organelles mediating lysosomal Ca^{2+} bursts and consequent global Ca^{2+} release in HEK293 cells transfected with human TPC2 channels (Calcraft et al., 2009). Since then, TPC1 and TPC2 were reported to serve as major NAADP-sensitive Ca^{2+} -permeable channels in different cells with overly expressed transgenes of these TPCs (Brailoiu et al. 2009a; Brailoiu et al., 2010a; Brailoiu et al., 2010b; Bright et al., 2005; Bund and Lee, 2003; Churchill and Galione 2001; Yamaguchi et al., 2011; Zinchuk et al., 2007). With some positive results in ligand binding experiments (Calcraft et al., 2009), these investigators proposed that like IP_3 receptors in the SR, TPCs in lysosome-like acidic organelles serve as an NAADP receptor with Ca^{2+} release channel activity, whereby NAADP binds and elicits Ca^{2+} release from these acidic organelles. In addition, blockade of endogenous NAADP responses by siRNA, gene knockout and/or use of dominant negative constructs also demonstrated that TPCs may mediate NAADP-induced Ca^{2+} response (Brailoiu et al., 2009a; Calcraft et al., 2009; Pereira et al., 2011; Rybalchenko et al., 2012). Some pharmacological and functional characteristics of this NAADP-TPCs working model (including binding and activation) have been described in several reviews by the investigators who first reported the role of TPCs as NAADP-sensitive channels (Galione, 2011; Galione et al., 2009; Zhu et al., 2010b), and the readers are directed to these reviews for details.

Unfortunately, this NAADP-TPC working model for NAADP-mediated Ca^{2+} response together with its hypothetical basis, namely, the two-pool Ca^{2+} release mechanism, has not yet widely been accepted as a general pathway of Ca^{2+} mobilization, at least in the area of vascular biology. Several lines of evidence have challenged this NAADP-TPC working model. First, photoaffinity binding analysis using more potent and specific probes has recently shown that NAADP did not bind to TPCs in different cells. Walseth and colleagues developed a photoaffinity probe for the NAADP receptor, 5- N_3 -NAADP (Lin-Moshier et al., 2012; Walseth et al., 2012) and confirmed that its binding proteins are 30, 40, and 45 kD in sea urchin egg homogenates and 22- and 23-kD doublet proteins in mammalian SKBR3 cells, HEK 293 cells, and mouse pancreas, which were much smaller than TPC channel proteins. In particular, such binding patterns were not changed by overexpression or knocking out of TPC genes in different cells (Lin-Moshier et al., 2012). These results suggest that TPCs may not be NAADP binding proteins and challenge the view of TPCs as an NAADP receptor. The NAADP-TPC working model for NAADP-mediated Ca^{2+} response requires further evidence for an NAADP-binding domain of TPCs.

Second, previous studies have reported that TPCs can be detected in the plasma membrane (Ishibashi et al., 2000) and that TPCs probably assemble as dimers in cell membrane through differential interactions between transmembrane regions (Churamani et al., 2012). These results suggest that TPCs may be a channel in the plasma membrane, which is able to mediate the action of NAADP to regulate Ca^{2+} influx and membrane depolarization (Moccia et al., 2006a; Moccia et al., 2004). Although the localization of TPCs in the cell plasma membrane may be due to overexpression of TPC genes during experiments, a majority of evidence supporting TPCs as an NAADP receptor or target is also from experiments using TPC transgenes in different cell lines (Calcraft et al., 2009; Zong et al., 2009). Until more evidence shows the similar role of TPCs in mediating NAADP mobilization of Ca^{2+} from lysosome-like acidic organelles or vesicles of native cells or cells isolated from animal tissues or organs without exogenously introduced TPC genes, a cautious conclusion should be made regarding the role of TPCs in the action of NAADP intracellular Ca^{2+} mobilization and its related physiological relevance. In this regard, we recently demonstrated that TPC2 was ubiquitously expressed in various compartment of arterial smooth muscle cells isolated from bovine hearts, which was colocalized with plasma membrane marker caveolin-1, endoplasmic reticulum marker protein-disulfide isomerase, lysosome marker LAMP-1, as well as mitochondrial marker Mito-tracker. While mitochondrial expression of TPCs was much higher compared to other organelles (unpublished data), it is interesting to note that the SR and lysosomes had almost equal abundance of TPC proteins suggesting that NAADP may also act on the SR TPCs to release Ca^{2+} . Lysosomal Ca^{2+} release is perhaps coinciding event with SR Ca^{2+} release in VSMCs where NAADP may simultaneously act on the TPCs of lysosomes and SR to mediate different cellular regulation. However, more studies are certainly needed to confirm this hypothesis.

Finally, in a recent study large efforts were made to distinguish the trigger event of TPCs mediated by NAADP from its amplification via ER Ca^{2+} stores by targeting TPCs on the cell membrane (Brailoiu et al., 2010b). This is because resolving triggering events of TPCs in lysosomes is relatively difficult. It is true that many studies aiming to define the role of TPCs in the Ca^{2+} response to NAADP used relatively indirect methods of measuring Ca^{2+} release from lysosome-like acidic organelles (Brailoiu et al., 2010c; Calcraft et al., 2009; Morgan and Galione, 2007; Yamasaki et al., 2004) by testing bafilomycin-sensitive or -dependent Ca^{2+} release. Although bafilomycin is considered to be a selective vacuolar H^{+} -ATPase inhibitor, its wide spectrum for inhibition of ATPase (Bowman et al., 1988) has also been reported. In addition, the findings that the vacuolar H^{+} -ATPase is also expressed in the plasma membrane and other cellular compartments or organelles (Forgac, 2007; Gluck, 1992; Rojas, et al., 2006; Tapper and Sundler, 1995) further raise a concern about the reliability of bafilomycin-sensitive lysosomal Ca^{2+} release as a parameter to define the role of TPCs in lysosomes or acidic organelles. In many studies, global Ca^{2+} transient response and plasma membrane Ca^{2+} -sensitive Cl^{-} channel or K^{+} channel activity were used to study the role of TPC as a lysosomal Ca^{2+} release channel to trigger intracellular global Ca^{2+} increase (Arredouani et al., 2010; Calcraft et al., 2009; Zong et al., 2009). Findings from these studies that measure global Ca^{2+} response may not differentiate the triggering action of TPCs in lysosomes since no lysosomal Ca^{2+} burst could be detected in those experiments. Therefore, it is imperative to develop useful measurements that can be used to directly

monitor lysosomal Ca^{2+} release such as Ca^{2+} indicator (GCaMP3) attached directly to lysosome proteins, as was done for TRP-ML1 (Shen et al., 2012), and localization of Ca^{2+} release signals with lysosome markers (Zhang et al. 2010). Such direct measurements may provide reliable evidence for the Ca^{2+} channel nature of TPC as a trigger of NAADP-induced global Ca^{2+} release. Unfortunately, a most recent study by direct patch clamp recording of ion channels in endosome/lysosomes demonstrated that TPCs are not activated by NAADP and that TPC currents are absent in pancreatic β -cell lines that exhibit NAADP-induced lysosomal Ca^{2+} release. In addition, both TPC1 and TPC2 are not required for NAADP-or glucose-induced Ca^{2+} responses in pancreatic islets. It is concluded that TPC proteins are phosphoinositide-activated sodium-selective ion channels in endosomes and lysosomes (Wang et al., 2012).

Based on the current knowledge, NAADP can be produced in response to ET-1, Ang II, norepinephrine, FasL, hypoxia and other agonists in VSMCs (Thai and Arendshorst, 2009; Zhang et al., 2006a). NAADP may induce lysosomal Ca^{2+} release through TRP-ML channels or TPCs either by direct binding or through a cytosolic binding protein. Such local Ca^{2+} bursts promote lysosome trafficking toward the SR, where aggregated lysosomes further release Ca^{2+} to activate RyRs or IP_3 R to produce large Ca^{2+} release from the SR, thereby increasing global Ca^{2+} concentrations within cells (Fig. 2).

FUNCTIONAL RELEVANCE OF cADPR OR NAADP-MEDIATED SIGNALING

Vascular Regulation

Vascular Tone—Vascular smooth muscle (VSM) usually operates in a contracted state, which is referred to as vascular tone. It has been proposed that $[\text{Ca}^{2+}]_i$ importantly contributes to the production of this “resting” vascular tone. Under resting conditions, $[\text{Ca}^{2+}]_i$ in VSMCs is dependent upon Ca^{2+} influx, spontaneous brief releasing bursts of Ca^{2+} from the SR into the cytoplasm and CICR (Berridge, 1997). cADPR participates in the control of resting Ca^{2+} levels in smooth muscle cells through RyRs and CICR. Therefore, cADPR plays an important role in forming basic vascular tone. In isolated, perfused and pressurized small coronary arteries, basic vascular tone or spontaneous tension can be developed during a 1.5-hour equilibration period. Under this condition, SR Ca^{2+} -ATPase inhibitor thapsigargin decreased the arterial diameter, and CICR blocker tetracaine and cADPR antagonist 8-Br-cADPR slightly dilated these arteries, suggesting that $[\text{Ca}^{2+}]_i$ associated with the cADPR-RyR signaling pathway and CICR is one determinant of basic vascular tone (Boini et al., 2011; Deshpande et al., 2005; Li et al., 2000a; Li et al., 1997; Li et al., 2002; Li et al., 1998; Tegatz et al., 2005b; Zhang et al., 2001; Zhang et al., 2010).

Little is known so far whether NAADP-mediated two-phase Ca^{2+} release is implicated in the control of basic vascular tone. Recently, we reported that a death receptor agonist, FasL, can induce a two-phase Ca^{2+} release and enhance U46619-induced vasoconstriction in mouse coronary arteries. Such enhanced vasoconstrictor response was consistent with the time frame of the FasL-induced two-phase Ca^{2+} response (Zhang et al., 2010). However, FasL itself did not alter vascular tension. It seems that the slow development of FasL-induced slow development of vascular tension is different from that of the instant vasoconstriction provoked by some classical vasoactive agonists (e.g., norepinephrine, Ang

II, or ATP), which is dependent on PLC-mediated direct Ca^{2+} release from the SR. Thus, FasL-induced two-phase Ca^{2+} release may be an important mechanism maintaining vascular tone where the NAADP signaling pathway in VSM is critical for the development of sustained vascular tone. In some other experiments using perfused small renal artery, we indeed found that blockade of the CD38-NAADP pathway by nicotinamide or PPADS had no effect on activation of arterial contraction in response to phenylephrine at the beginning of tension increase. However, it significantly attenuated maintenance of vascular tension when it reached a plateau (unpublished data). This suggests that NAADP-induced two-phase or multiple phase Ca^{2+} release is an important source of intracellular Ca^{2+} to regulate the related molecular mechanisms for the maintenance of vascular tone such as protein kinase C and myosin light-chain kinase (Gao et al., 2001; Rasmussen et al., 1987).

Another important mechanism regulating vascular tone is the production of endothelium-derived relaxing factors (EDRFs), where blood vessels, particularly arteries, can produce relaxation in response to blood flow, shear stress and circulatory vasodilators, which may importantly contribute to the control of vascular tone under physiological or pathological conditions (Edwards et al., 2010; Qi et al., 2011; Vanhoutte et al., 2009). It is well known that Ca^{2+} activation of ECs is critically implicated in the production of EDRFs such as nitric oxide (NO), epoxyeicosatrienoic acids (EETs) and prostacyclins (Yi et al., 2002; Zhang et al., 2005; Zhang et al., 2004). Upon stimulation with different factors such as bradykinin, thrombin, histamine, bradykinin and oxidants, $[\text{Ca}^{2+}]_i$ could increase 5–10 fold compared with the basal level. Increased Ca^{2+} stimulates the binding of $\text{Ca}^{2+}/\text{CaM}$ to endothelial NO synthase (eNOS), resulting in rapid conversion of L-arginine into L-citrulline, producing NO (Freichel et al., 2001; Putney, 1999; Tirupathi et al., 2002). Interestingly, ADP-ribosylcyclase gives rise to bradykinin signal transduction from receptors to its effector enzymes (Deshpande et al., 2003; Higashida et al., 1996; Higashida et al., 2001), suggesting that cADPR/RyR signaling may be present in ECs to modulate endothelial function by regulating EDRF production. In this regard, we indeed demonstrated that inhibition of cADPR production or antagonism of its action significantly attenuated bradykinin-induced concentration-dependant coronary arterial vasodilation, endothelial Ca^{2+} release from RyR-sensitive stores and increases in NO production. Measurement of endothelial ADP-ribosylcyclase activity and intracellular cADPR concentrations confirmed that bradykinin induced cADPR production via enhanced ADP-ribosylcyclase activity (Zhang et al., 2006b). It is concluded that such bradykinin-induced intracellular Ca^{2+} increase and NO response are not mainly associated with IP_3 signaling, but with cADPR levels in coronary ECs cells that participates in endothelium-dependent vasodilation. More recently, NAADP acetoxyethyl ester (NAADP-AM), a cell-permeant NAADP analog, was also demonstrated to increase cytosolic Ca^{2+} concentration in aortic ECs. This increase in intracellular Ca^{2+} and those evoked by acetylcholine were accompanied by hyperpolarization of ECs and NO production. Correspondingly, NAADP-AM was found to dilate aortic rings in an endothelium- and NO-dependent manner. In anesthetized rats, intravenous administration of NAADP-AM markedly decreased mean arterial pressure. Taken together, this study suggest that NAADP may regulate endothelial function by participating in the control of vascular tone and arterial blood pressure (Brailoiu et al., 2010c).

Vasomotor Response—As discussed above, acetylcholine (ACh) M_1 -type receptor (mAChR) agonist, oxotremorine was demonstrated to markedly enhance the activity of ADP-ribosylcyclase and to increase the production of cADPR in cultured CASMCs, which was blocked by M_1 mAChR blocker pirenzepine and by ADP-ribosylcyclase inhibitor nicotinamide (Ge et al., 2003). It seems that ADP-ribosylcyclase is directly coupled to M_1 mAChRs through G-proteins. In isolated, perfused and pressurized small coronary arteries, vasoconstriction induced by ACh or oxotremorine was also attenuated by the inhibition of ADP-ribosylcyclase and blockade of cADPR action. These results confirmed that cADPR is linked to M_1 mAChRs and mediates the vasoconstrictor response through activation of this subtype of mAChRs in CASMCs (Ge et al., 2003). This oxotremorine-induced vasoconstriction is also directly associated with cADPR-mediated Ca^{2+} release from the SR in these cells because oxotremorine-induced Ca^{2+} release with Ca^{2+} -free extracellular solution was significantly attenuated by inhibition of cADPR production by nicotinamide and by blockade of cADPR action by 8-Br-cADPR (Ge et al., 2003). In isolated and perfused small coronary septal arteries from CD38^{-/-} mice, oxotremorine produced much smaller vasoconstrictor response than in the same arteries from wild-type mice, and oxotremorine-induced intracellular Ca^{2+} increase was significantly lowered in freshly isolated septal arterial VSMCs from CD38^{-/-} mice than in VSMCs isolated from wild-type mice (Teggatz et al., 2005a). Taken together, these results provide direct evidence that endogenous cADPR contributes to oxotremorine-induced Ca^{2+} mobilization in VSMCs and that cADPR serves as a second messenger to activate M_1 receptors and mediate the vasoconstrictor response (Prakash et al., 1998). In renal circulation, Arendshorst and his associates demonstrated that basal ADP-ribosylcyclase activity is important in the control of renal blood flow since its inhibition resulted in increased renal blood flow in anesthetized rats (Thai et al., 2007). They also showed that inhibitors of ADP-ribosylcyclase attenuated renal vascular responses to Ang II and norepinephrine by approximately 60% when injected into the renal artery (Thai and Arendshorst 2008; Thai et al., 2007). In these experiments, it was also found that more pronounced renal ADP-ribosylcyclase inhibition produced local vasodilation without altering arterial pressure, indicating that the basal levels of ADP-ribosylcyclase activity contribute to resting renal vascular resistance. This role of ADP-ribosylcyclase activity in mediating the vasoconstrictor response was further confirmed by using CD38^{-/-} mice. It was shown that acute renal vasoconstrictor responses to Ang II and norepinephrine were reduced by approximately 50% in CD38^{-/-} mice compared with wild-type mice. All these studies suggest that CD38/ADP-ribosylcyclase is importantly involved in mediating renal vasoconstrictor responses to stimulation of G-protein coupled receptors in mice and rats (Thai and Arendshorst, 2008; Thai et al., 2007).

ET-1 is a potent endothelium-derived vasoconstrictor peptide that increases intracellular Ca^{2+} via activation of ET_A and/or ET_B receptors in different vascular beds. Recent studies have shown that ET-1-induced Ca^{2+} response is associated with cADPR/RyR signaling. In rat mesenteric arteries, Giulumian et al., reported that ET-1-induced Ca^{2+} increase and vasoconstriction were significantly attenuated by ADP-ribosylcyclase inhibitor nicotinamide and RyR Ca^{2+} release channel inhibitor dantrolene (Giulumian et al., 2000). In isolated pulmonary arteries, membrane-permeant cADPR antagonist 8-Br-cADPR was demonstrated to block sustained hypoxic pulmonary vasoconstriction (Dipp and Evans, 2001). In rat

peritubular VSMCs, Barone et al., observed that both ET_A - and ET_B -mediated Ca^{2+} signaling were completely abolished by cADPR antagonist 8-NH₂-cADPR (Barone et al., 2002). In porcine airway smooth muscle, White et al., showed that ET-1-induced Ca^{2+} response was inhibited by cADPR antagonist 8-Br-cADPR (White et al., 2003). Although there was direct detection of cADPR production in response to ET-1 in some vascular beds in these studies, these functional data have suggested that cADPR/RyR Ca^{2+} signaling may importantly participate in ET-1-induced vasoconstrictor response.

In addition to cADPR, NAADP-mediated Ca^{2+} signaling was also found to regulate agonist-induced vasoconstrictor response. In this regard, Evans and associates first reported that intracellular dialysis of NAADP induced spatially restricted “bursts” of Ca^{2+} release that initiated a global Ca^{2+} wave and contraction in pulmonary artery smooth muscle cells. Depletion of SR Ca^{2+} stores with thapsigargin and inhibition of RyRs with ryanodine both blocked the global Ca^{2+} waves by NAADP (Boittin et al. 2002). They suggest that NAADP may act in concert with cADPR to promote hypoxic pulmonary vasoconstriction (Evans 2010). Consistent with a previous study by Kinnear et al., in coronary arteries, we also found that ET-1 induced NAADP production, which mobilized intracellular Ca^{2+} in a manner dependent of normal lysosome function. ET-1-induced maximal coronary arterial constriction was substantially blocked by lysosome function inhibitor bafilomycin A1 and NAADP antagonist PPADS. It is obvious that a lysosome-mediated Ca^{2+} regulatory mechanism via NAADP contributes to ET-1-induced Ca^{2+} mobilization in CASMCs and consequent vasoconstriction of coronary arteries (Zhang et al., 2006a). More recently, we further demonstrated that FasL also increased NAADP production, but FasL itself did not produce vasoconstriction in coronary arterial preparation. However, FasL significantly enhanced IP₃-producing agonist U46619-induced coronary arterial contraction, suggesting that NAADP may also sensitize arterial contraction when its production is increased (Zhang et al., 2010). Despite these reports, more studies are needed to differentiate the role of intracellular Ca^{2+} stores or signaling pathways in the mediation or modulation of vasomotor response. In particular, definition of the temporospatial action of NAADP, cADPR and IP₃ in VSMCs is imperative.

Vascular Diseases

Although cADPR and NAADP have been extensively investigated as Ca^{2+} signaling second messengers with their related molecular signaling mechanisms, the pathological role of both signaling molecules is still under studied. Based on acute experiments in cells or isolated vessels, cADPR or NAADP are indicated to be involved in the development of hypertension and pulmonary hypertension (Evans et al., 2005). However, so far there is no direct evidence showing that both ADP-ribosylcyclase-derived second messengers are implicated in any vascular diseases. More recently, our laboratory is working on the potential contribution of NAADP-mediated regulation of lysosome function in the development of atherosclerosis. The general hypothesis is that NAADP regulates lysosome function, which plays an essential role in the control of lysosome trafficking or fusion to autophagosomes (APs), and regulates autophagy via its effect on autophagic flux (Fasano et al., 2012; Ryter et al., 2010; Seedorf et al., 1995; Xu et al., 2011).

Given the different cell types in the artery wall, the role of autophagy in the development of atherosclerosis is complex due to its action on different vascular cell functions. It has been recently assumed that autophagy may have both protective and detrimental roles during atherosclerosis, depending upon the status of autophagy or stages of atherosclerosis (Bampton et al., 2005; Martinet and De Meyer, 2009). Since autophagy is important in the degradation of damaged materials, it is possible that autophagy in the arterial wall helps clean up damaged components and recover cells from the damage upon atherosclerotic stimuli. In addition, autophagy activation interferes with cell apoptosis due to engulfment of defective or damaged mitochondria by APs, which limits the release of proapoptotic proteins (Gutierrez et al., 2004; Kim et al., 2008; Zhu et al., 2007). This autophagic process may protect arterial cells from atherogenic injury. However, if acute or persistent oxidative stress occurs during atherosclerosis, lysosomes may be damaged to release hydrolases, engage as part of oxidative stress, and enhance cellular damages (Jia et al. 2006; Martinet and De Meyer 2008; Martinet and De Meyer 2009; Xu et al., 2010). Enhanced or reduced autophagy plays different roles in the development of atherosclerosis depending on the different cells involved. In macrophages, autophagy increases cholesterol transport out of these cells, which may prevent lipid droplet formation reducing foam cell formation. Similarly, enhanced autophagic death of macrophages also possibly attenuates foam cell formation, reducing atherosclerotic injury. However, excessive activation of autophagy in ECs may lead to damage of the endothelium enhancing atherogenic injury. In arterial SMCs, enhanced autophagy induces their modulation to a differentiated, quiescent, and contractile phenotype, decreasing cell proliferation and preventing fibrosis. Nevertheless, excessive autophagy in arterial SMCs may result in their death, increasing the instability of atherosclerotic plaques (Jia et al., 2007; Jia et al., 2006; Schrijvers et al., 2007; Verheye et al. 2007; Xu et al., 2010). Although the role of augmented autophagy in atherosclerosis has been extensively studied, there is no evidence that a defective or reduced autophagy is involved in the pathogenesis of atherosclerosis.

In some of our preliminary studies, we demonstrated that defective autophagy is also importantly involved in atherogenesis, which is associated with molecular dysregulation of lysosome function. As shown in Figure 3, proatherogenic stimuli such as 7-keto-cholesterol (7-keto) or ox-LDL activates autophagy in arterial SMCs, leading to the formation of APs. Under normal condition, lysosome trafficking and fusion to APs are controlled by CD38-ADP-ribosylcyclase-mediated regulation, particularly by NAADP production, which leads to the formation of autophagolysosomes (APLs) and subsequent breakdown of the autophagic vesicles within the cells. This regulated autophagic process via NAADP signaling pathway protects SMCs from atherosclerotic injury upon atherogenic stimulations. When the controlling mechanism of lysosome function is insufficient, such as impaired CD38-ADP-ribosylcyclase activity or reduced NAADP production, the formation of APLs and breakdown of autophagic vesicles also become impaired, which activates cell dedifferentiation, proliferation and growth, thereby stimulating production of extracellular matrix and ultimately inducing or accelerating atherosclerosis. Indeed, CD38^{-/-} mice developed atherosclerosis when they were exposed to the atherogenic diet (Xu et al., 2011). Under such condition, dysregulation of lysosome function due to the lack of CD38 product, mainly NAADP, caused deficient autophagy by failed formation of APLs and impaired

degradation of autophagic content, which resulted in arterial smooth muscle remodeling by enhanced cell proliferation and extracellular matrix production.

CONCLUDING REMARKS

Since the discovery of cADPR and NAADP as Ca^{2+} signaling second messengers, a large body of evidence shows that they are importantly involved in the regulation of intracellular Ca^{2+} concentrations in vascular cells including ECs and VSMCs (Bai et al., 2005; Evans, 2010; Zhang and Li, 2006). In ECs, cADPR mediates agonist (such as bradykinin)-induced Ca^{2+} mobilization from the ER, resulting in production of NO or other EDRFs, participating in the EDVD response. In VSMCs, cADPR serves as a second messenger to stimulate Ca^{2+} release from the SR via RyRs and is involved in the regulation of CICR and consequent Ca^{2+} waves, producing global Ca^{2+} increase within these cells. cADPR activates RyRs by binding to FKBP12.6 and results in dissociation of this accessory protein from RyRs, whereby Ca^{2+} release from the SR is enhanced. There is considerable evidence showing that this cADPR-mediated Ca^{2+} signaling plays a critical role in Ca^{2+} release and vasoconstrictor response to different agonists, oxidative stress, cell membrane depolarization, and Ca^{2+} influx. Similarly, NAADP has been found to play similar roles in the regulation of intracellular Ca^{2+} mobilization, vascular tone and vasomotor responses (Zhang et al., 2010; Zhang et al., 2006a). It has been demonstrated that NAADP is a more potent Ca^{2+} mobilizer compared with other Ca^{2+} signaling second messengers. It first produces small Ca^{2+} bursts via lysosomes and then leads to a global Ca^{2+} increases in VSMCs due to CICR-mediated large Ca^{2+} release from the SR, known as a two-phase Ca^{2+} release. Functionally, NAADP-mediated Ca^{2+} release has been reported to also participate in the control of vascular tone and vasomotor response to different agonists, which may occur in a temporospatial way in concert with other pathways such as cADPR and IP_3 . In addition, NAADP-mediated Ca^{2+} regulation in vascular cells may be relevant to vascular function such as EDVD or cell apoptosis. Although many studies assume that cADPR and NAADP-mediated Ca^{2+} signaling may be implicated in the pathogenesis of some vascular diseases such as hypertension, pulmonary hypertension and atherosclerosis, there is not much direct evidence that both Ca^{2+} second messengers indeed contribute to the development of these diseases. This is an area necessary to be studied and addressed in the future.

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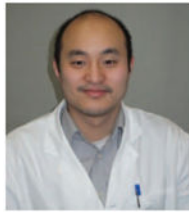
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Biographies



Pin-Lan Li is a Professor and Vice Chair in the Department of Pharmacology and Toxicology of the Virginia Commonwealth University. She was trained as M.D. in Yi-Chang Medical College and Tongji Medical University, China in 1973–1979 and then received her Ph.D. from the Heidelberg University, Germany in 1992. She joined the Medical College of Wisconsin as an Assistant Professor in 1994 and then promoted to full professor in 2004. She moved to the Virginia Commonwealth University as a Professor with tenure in 2005. Her research is mainly focused on the vascular signaling mechanisms

including cADPR, NAADP, ceramide, NO and redox-mediated signal transduction and their relevance to vascular diseases such as vascular inflammation, atherosclerosis and hypertension.



Yang Zhang is an Assistant Professor in the Department of Pharmacology & Toxicology, Virginia Commonwealth University (VCU). He obtained his Ph.D. in Pharmacology from the Medical College of Wisconsin in 2006. After two years postdoctoral training at the Department of Molecular Biology, University of Duisburg-Essen, Germany, he was promoted to a junior professor at the same department. He joined VCU as an Assistant Professor in 2011. His research interests are in studying the roles of ADP ribosylcyclase/CD38, NADPH oxidase, and acid sphingomyelinase in the control of vascular autophagy and inflammasome activity as well as their pathogenic roles in atherosclerosis, pulmonary fibrosis and chronic obstructive pulmonary disease.

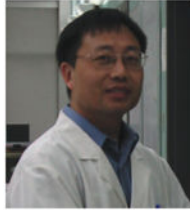


Justine M. Abais is a Ph.D. student in the Department of Pharmacology and Toxicology at the Virginia Commonwealth University (VCU), located in Richmond, Virginia. Justine received her Bachelor's degree in Chemistry from VCU in 2009. Her current research interest is in understanding the molecular mechanisms of hyperhomocysteinemia-associated vascular and renal degenerative diseases such as atherosclerosis and end-stage renal disease. She is currently a recipient of Ruth L. Kirschstein National Research Service Award for individual predoctoral fellows from the National Institute of Aging.



Joseph K. Ritter is an Associate Professor of Pharmacology and Toxicology at the Virginia Commonwealth University School of Medicine. He received his Ph.D. degree at the University of Utah in 1987 and was a postdoctoral fellow at the National Institute of Child

Health and Human Development in Bethesda, Maryland. His research interests are in the role of metabolism as a modulator of pharmacologic and toxic effects of xenobiotic and endogenous substances. Recent research projects in his laboratory focus on the functional relevance of endocannabinoids and associated signaling mechanisms to hypertension and kidney diseases.



Fan Zhang is an Assistant Professor of Pharmacology at the Virginia Commonwealth University (VCU). He obtained his Ph.D. in Pharmacology at the Tongji Medical College of Huazhong University of Science and Technology, China in 2002. After postdoctoral training in the Department of Pharmacology, VCU, he was promoted to Assistant Professor in 2009 at the same department. His research focus is on the vascular regulation by ADP ribosylcyclase/CD38 and its products, cADPR and NAADP. Recent research projects in his laboratory deal with lysosomal TRP-ML1 channels, regulation of lysosomal lipid metabolism in macrophages, and their roles in coronary arterial atherosclerosis.

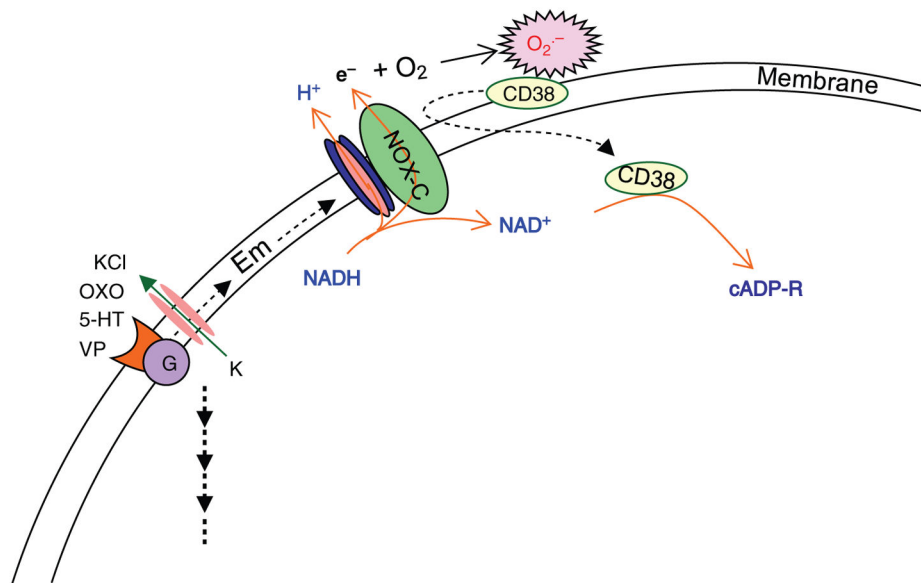


Figure 1. Cross talk between NADPH oxidase and CD38/ADP-ribosylcyclase in cell plasma membrane
 NOX-C: NADPH oxidase complex. VP: vasopressin. 5-HT: serotonin. Em: membrane potential.

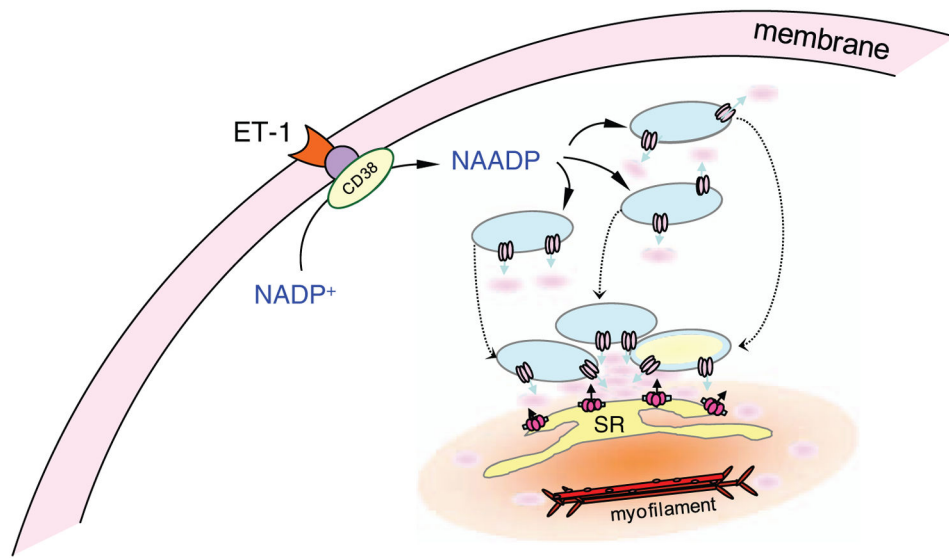


Figure 2. NAADP-induced lysosomal Ca^{2+} release, lysosome trafficking and CICR in VSMCs
SR: sarcoplasmic reticulum. ET-1: endothelin-1.

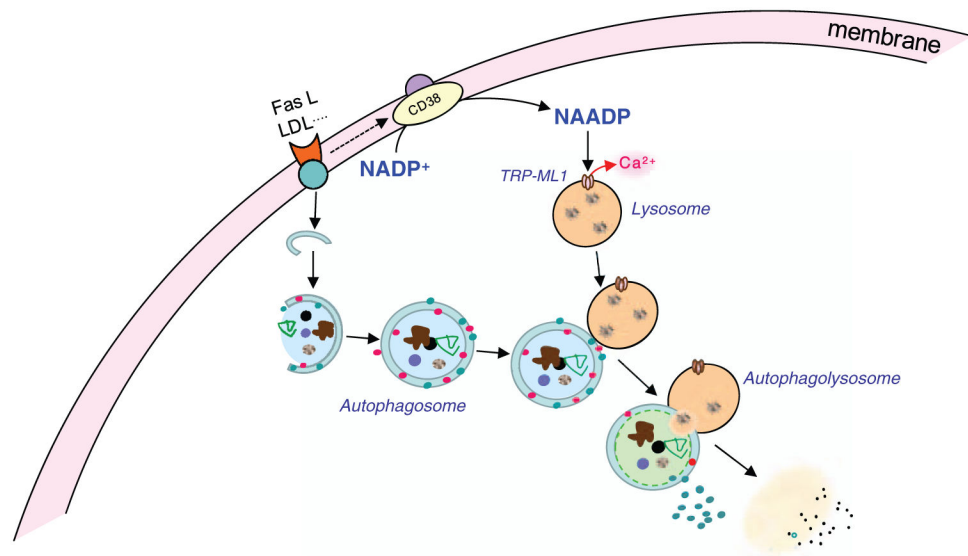


Figure 3. Lysosome trafficking and fusion to autophagosomes controlled by NAADP via CD38-ADP-ribosylcyclase

FasL: Fas ligand. LDL: Low density lipoprotein.