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Reactive γ -Ketoaldehydes Promote Protein Misfolding and Preamyloid Oligomer Formation in Rapidly-Activated Atrial Cells

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Abstract

Rapid activation causes remodeling of atrial myocytes resembling that which occurs in experimental and human atrial fibrillation (AF). Using this cellular model, we previously observed transcriptional upregulation of proteins implicated in protein misfolding and amyloidosis. For organ-specific amyloidoses such as Alzheimer's disease, preamyloid oligomers (PAOs) are now recognized to be the primary cytotoxic species. In the setting of oxidative stress, highly-reactive lipid-derived mediators known as γ -ketoaldehydes (γ -KAs) have been identified that rapidly adduct proteins and cause PAO formation for amyloid β_{1-42} implicated in Alzheimer's. We hypothesized that rapid activation of atrial cells triggers oxidative stress with lipid peroxidation and formation of γ -KAs, which then rapidly crosslink proteins to generate PAOs. To investigate this hypothesis, rapidly-paced and control, spontaneously-beating atrial HL-1 cells were probed with a conformation-specific antibody recognizing PAOs. Rapid stimulation of atrial cells caused the generation of cytosolic PAOs along with a myocyte stress response (e.g., transcriptional upregulation of *Nppa* and *Hspa1a*), both of which were absent in control, unpaced cells. Rapid activation also caused the formation of superoxide and γ -KA adducts in atrial myocytes, while direct exposure of cells to γ -KAs resulted in PAO production. Increased cytosolic atrial natriuretic peptide (ANP), and the generation of ANP oligomers with exposure to γ -KAs and rapid atrial HL-1 cell stimulation, strongly suggest a role for ANP in PAO formation. Salicylamine (SA) is a small molecule scavenger of γ -KAs that can protect proteins from modification by these reactive compounds. PAO formation and transcriptional remodeling were inhibited when cells were stimulated in the presence of SA, but not with the antioxidant curcumin, which is incapable of scavenging γ -KAs. These results demonstrate that γ -KAs promote protein misfolding and PAO formation as a component of the atrial cell stress response to rapid activation, and they provide a potential mechanistic link between oxidative stress and atrial cell injury.

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CONFLICTS OF INTEREST

None declared.

Keywords

preamyloid oligomers; amyloidosis; atrial HL-1 cells; oxidant stress; atrial natriuretic peptide; γ -ketoaldehyde; levuglandin

1. INTRODUCTION

Atrial fibrillation (AF) is a progressive disorder, with remodeling during rapid atrial activation that increases arrhythmia susceptibility. There is abundant evidence linking oxidative stress to the pathogenesis and progression of AF [1, 2]. Risk factors that promote AF, as well as inflammation and activation of the renin-angiotensin-aldosterone system, cause oxidative damage in humans [1-3]. Oxidative injury is evident in the atria of patients with permanent AF [4], and superoxide production is increased in both experimental and human AF [5, 6]. While pharmacologic agents with antioxidant properties have shown promise in experimental AF models, clinical trials to date have largely been disappointing [7].

Previously, we showed that atrial cells rapidly stimulated in culture undergo remodeling similar to that observed in human AF [8]. Importantly, transcriptional profiling in paced cells exhibited substantial concordance with changes seen *in vivo* [9]. Unexpectedly, we observed conserved transcriptional upregulation for proteins implicated in amyloidosis, including numerous heat-shock proteins that serve as molecular chaperones to prevent protein misfolding and aggregation.

It is increasingly recognized that proteotoxicity such as amyloid plays an important role in disease pathogenesis, especially for aging-related degenerative disorders such as Alzheimer's disease [10, 11]. For systemic amyloidoses (e.g., with multiple myeloma), organ dysfunction is caused by the large quantity of amyloid fibrils that are deposited [12]. In contrast, for *organ-limited* amyloidoses such as Alzheimer's disease and type II diabetes, mature amyloid deposits have no correlation with the state of disease advancement. Rather, soluble protein aggregate intermediates are now recognized as the primary cytotoxic species that correlate with disease phenotype [10, 11]. Preamyloid oligomers (PAOs) cause cell injury and/or death by multiple mechanisms, including endoplasmic reticulum stress, increased cytoplasmic calcium concentration, mitochondrial injury with oxidative stress, reduced protein clearance, and probable cell membrane pore formation [13]. PAO complexes derived from different proteins possess a common structural epitope related to the peptide backbone that is irrespective of amino acid sequence, enabling the development of confirmation-specific antibodies. Importantly, a critical distinction between soluble protein oligomers and amyloid deposits is that the oligomers do not possess the structure required for binding of amyloid-detecting dyes such as Congo red, and hence they are not visible by standard amyloid staining methods. Recent studies have demonstrated the presence of PAOs in experimental and human heart failure [14, 15]. However, the role of these cytotoxic complexes in atrial pathophysiology has not been explored.

In the setting of oxidative injury and cyclooxygenase activation, arachidonic acid can undergo oxygenation and structural rearrangement to generate γ -ketoaldehyde compounds

given the trivial name of isolevuglandins or isoketals [16-18]. γ -Ketoaldehydes (γ -KAs) are the most reactive products of lipid peroxidation identified to date, and they rapidly adduct to lysine residues of proteins to form stable adducts and intermolecular crosslinks [19-21]. γ -KA adducts are increased in a number of pathologic conditions, including Alzheimer's disease, that are linked to oxidative injury and inflammation [22, 23]. Recent evidence demonstrates that these highly-reactive compounds can directly promote formation of PAOs derived from amyloid β_{1-42} , the highly fibrillogenic peptide involved in the development of Alzheimer's disease [24]. In addition, molecular scavengers have been discovered that rapidly and irreversibly react with and inactivate γ -KAs, thus preventing them from reacting with and damaging proteins [17, 25]. Salicylamine (SA) is a member of a family of phenolic amines that act as highly-effective γ -KA scavengers. Importantly, SA prevents the development of cognitive deficits in a mouse model of Alzheimer's disease [26].

Given the evidence implicating oxidative stress in the development of both Alzheimer's disease and AF, common pathophysiologic mechanisms may be operative for these seemingly disparate disorders. We hypothesized that rapid activation of atrial myocytes triggers a cellular stress response that includes oxidative injury and the generation of γ -KAs, to promote protein misfolding and PAO formation.

2. MATERIALS AND METHODS

2.1 Atrial HL-1 Cell Culture and Stimulation

Atrial HL-1 myocytes were grown as described previously [8]. Nearly confluent HL-1 cells were subjected to rapid stimulation for 6hr at 5Hz (18V, 4ms) using a C-Pace cell culture stimulator (Ion Optix Corp) in the absence and presence of treatment, with spontaneously-beating control cells cultured in parallel. Optimization of pacing conditions was performed as described in Supplementary Data, to ensure reproducible stimulation of atrial HL-1 cells at 5 Hz, a rate that causes electrophysiologic and transcriptional remodeling [8, 9].

2.2 Immunocytochemistry: Preamyloid Oligomers

Rapidly-stimulated and control spontaneously-beating cells were subjected to immunocytochemistry using a rabbit polyclonal antibody (A-11; 1:200, EMD Millipore) recognizing the conformational epitope common to all preamyloid oligomers [11, 27], as detailed in the Supplementary Data.

2.3 Image Acquisition

Cells were imaged using a 40X/1.3 Plan-Neofluar objective on a confocal microscope (LSM-510, Carl Zeiss). Fluorescence images were acquired at 0.5 μ m focus intervals with a confocal pinhole set to 1 Airy unit, thus optimizing contrast and resolution. The Alexa Fluor 488-labeled A-11 and TO-PRO-3 were excited at 488 nm and 633 nm, respectively, and detected with 505-550 nm band pass and 650 nm long pass filters, respectively. DIC images were acquired as a separate scan and registered with the corresponding fluorescence images.

2.4 Detection of Superoxide Production

Atrial HL-1 cells were subjected to rapid stimulation as described above, except that for the final 30min, Claycomb medium was replaced with 50mM Krebs-HEPES buffer (in mM: NaCl 145; KCl 4.86; NaH₂PO₄ 5.7; CaCl₂ 0.54; MgSO₄ 1.22; glucose 5.5; pH 7.4) containing 10μM dihydroethidium (DHE, Molecular Probes). Cells were washed with PBS and fixed with 3% PFA in PBS as usual. In some experiments, tempol (10μM, Enzo Life Sciences) was included in the buffer.

2.5 PAO Formation by γ -Ketoaldehydes

Atrial HL-1 myocytes were incubated with the synthetic γ -KA 15-E₂-isolevuglandin/isoketals [28] (E₂-IsoKs; 1μM) for 6hr in HBSS medium (Mediatech, Inc). Cells were fixed with 4% PFA in PBS for 10min and permeabilized with 0.1% Triton in PBS for 7min. Unreacted aldehyde groups were blocked with 50mM NH₄Cl, while non-specific antigen binding sites were blocked using Power block (BioGenex) for 1hr. Immunostaining was performed using the A-11 antibody as described above. After incubation with E₂-IsoKs, exclusion of Trypan blue was also used to determine cell viability.

2.6 Immunocytochemistry: γ -Ketoaldehyde Adducts

Rapidly-stimulated and control spontaneously-beating cells were subjected to immunocytochemistry using an anti- γ -KA-lysyl adduct single-chain antibody (D11 ScFv) as described previously [29] and in the Supplementary Data.

2.7 Immunocytochemistry: Atrial Natriuretic Peptide

ANP immunostaining of atrial HL-1 cells was performed using a primary rabbit polyclonal anti- α -ANP (1-28) antibody (1:200, Phoenix Pharmaceuticals, Inc.) and donkey anti-rabbit Alexa 488-conjugated secondary antibody (1:500, Molecular Probes), using the protocol described above for A-11. Immunostaining was visualized by using confocal microscopy.

2.8 Western Analysis

Cell lysate from both rapidly-stimulated and control atrial HL-1 cells was diluted with LDS sample buffer, denatured at 70°C for 10min, and electrophoresed onto a 4-12% gradient gel (Life Technologies). Proteins were then transferred onto a PVDF membrane (0.2μm pore size, BioRad) and exposed overnight to either rabbit A-11 polyclonal antibody (1:500) or rabbit anti- α -ANP (1-28) polyclonal antibody (1:500, Phoenix Pharmaceuticals, Inc.). After removal of primary antibodies, membranes were incubated with HRP-conjugated secondary antibody (1:5000, goat anti-rabbit, Jackson ImmunoResearch) for 1hr. Membranes were stripped (Restore stripping buffer; Thermo Scientific) for 15min at room temperature and reprobbed with anti- β -actin mouse monoclonal antibody (1:5000, Sigma) as a protein loading control. An enhanced chemiluminescent kit (Pierce ECL Western Blotting Substrate) was used for detection of protein bands.

2.9 α -ANP Peptide Oligomer Formation

Synthetic α -ANP peptide (1-28) (SLRRSSCFGGRMDRIGASGLGCNSFRY- disulfide bond [C7-C23]) was synthesized by RS Synthesis, LLC. To test for ANP oligomerization,

peptide (10 μ M) was prepared in PBS buffer (pH 7.4) and incubated at room temperature for 6 days (positive control), while a separate sample was incubated for 24hr with either synthetic E₂-IsoKs (2 molar equivalents) or vehicle (DMSO). After incubation, Western analysis was performed using the anti- α -ANP antibody (1:500) and goat anti-rabbit HRP-conjugated secondary antibody (1:5000, Jackson ImmunoResearch). E₂-IsoKs were prepared as described previously [28].

2.10 mRNA Extraction and Real-Time Quantitative RT(q)-PCR

mRNA was extracted from cells, analyzed, and subjected to cDNA synthesis, and expression of *Nppb* and *Hspa1a* was evaluated by q-PCR as described previously [9] and in the Supplementary Data.

2.11 Data Analysis

Following normalization for 18S, the average Ct (cycle threshold) value for control cells was subtracted from that for paced cells. Fold change was defined as the residual Ct value for paced cells, using the 2^{-Ct} method, with an unpaired t-test used to determine statistically significant changes. Nonparametric analysis with a Kruskal-Wallis test for the Ct values was used to evaluate for a drug effect for each gene. Pairwise comparisons for vehicle and either salicylamine or curcumin exposure were performed with a Wilcoxon rank sum test.

3. RESULTS

3.1 Formation of Preamyloid Oligomers in Rapidly-paced Atrial HL-1 Cells

To test the hypothesis that rapid activation of atrial cells triggers protein misfolding and generation of PAOs, both rapidly-paced and control, spontaneously-beating atrial HL-1 cells were probed with a conformation-specific anti-oligomer antibody, A-11, recognizing the structural epitope common to all PAOs. As shown in Figure 1, rapid pacing caused the appearance of diffuse cytoplasmic immunoreactive protein (indicated by the green color in **panel B**) that was absent in control cells (**panel A**). Immunostaining was also absent when the primary antibody was omitted or IgG was substituted (Figure 1C and 1D), demonstrating the specificity of immunolabeling using A-11. Western blotting was also performed using A-11 for control and paced cells [30]. As shown in Figure 1E, rapid stimulation led to the appearance of bands at 34 and 95 kDa, representing different-sized PAOs (n=3 separate experiments), not present in lysate for control cells. Collectively these data indicate that rapid stimulation of atrial cells causes PAO formation. Given that these protein complexes are cytotoxic, these findings identify PAOs as potential candidates to promote myocyte injury during atrial cell remodeling.

3.2 Generation of Oxidative Stress with Rapid Activation

Reactive intermediates generated during oxidative stress can modify, cross-link, and aggregate proteins. Given that superoxide (O₂⁻) production is increased in animal models and human AF, we hypothesized that this reactive oxygen species (ROS) would also accumulate in rapidly-stimulated atrial HL-1 cells. In the presence of DHE (a non-fluorescent probe that becomes highly fluorescent after reacting with O₂⁻), spontaneously

beating atrial HL-1 cells generated minimal $O_2^{\cdot -}$ under control conditions (Figure 2A); however, with rapid activation, $O_2^{\cdot -}$ production is markedly enhanced (Figure 2B). In the presence of the small molecule cell membrane-permeable superoxide dismutase mimetic tempol, $O_2^{\cdot -}$ levels were essentially reduced to baseline (Figure 2C), demonstrating the specificity of the DHE-generated fluorescent signal. These findings confirm that, as *in vivo*, rapid activation of atrial HL-1 cells leads to the production of ROS.

3.3 Role of γ -Ketoaldehydes in PAO Formation

As described above, γ -KAs are highly-reactive products of oxidative stress that can crosslink proteins to initiate their aggregation. To investigate whether these compounds could play a role in generating PAOs in our experimental preparation, atrial HL-1 cells were incubated with a physiologic concentration of a synthetic E-series γ -KA (E₂-IsoKs; 1 μ M) for 6 hr, followed by immunostaining with A-11. As illustrated in Figure 3A and 3B, exposure to E₂-IsoKs led to the production of cytosolic PAOs similar to that observed with rapid pacing. Furthermore, immunostaining using an anti- γ -KA lysyl adduct single chain antibody (D11 ScFv) demonstrated the presence of abundant adducts in rapidly-stimulated but not control cells (Figure 3C and 3D). Taken together, these results are highly suggestive that γ -KAs can serve as a mechanistic link between oxidative stress and PAO formation in our experimental preparation.

3.4 Atrial Natriuretic Peptide and PAO Formation

Rapid activation of atrial cells causes atrial granule release, leading to elevated concentrations of natriuretic peptides such as ANP in plasma. ANP is known to be fibrillogenic (i.e., to form amyloid fibrils), and this hormone is primarily responsible for the development of aging-related cardiac amyloidosis in the atrium, a condition known as isolated atrial amyloidosis [31, 32]. We have previously demonstrated upregulation of *Nppa*, which encodes ANP, in rapidly-stimulated atrial HL-1 cells, with release of natriuretic peptides into the culture medium [9]. Therefore, immunostaining for ANP was performed on control and rapidly-paced atrial HL-1 cells. Figure 4B demonstrates increased intracellular production of ANP following rapid activation, with a diffuse cytosolic distribution resembling PAO formation. To test the hypothesis that ANP could contribute to PAO formation under these circumstances, purified ANP peptide was incubated with E₂-IsoKs, followed by Western blotting using an anti- α ANP antibody. As shown in Figure 4C, incubation of the peptide with E₂-IsoKs led to oligomer formation (lane 2), detected as bands of increased molecular size (predominantly at ~10 kDa), compared to monomeric ANP (3kDa; lane 3). Analogous to amyloid β_{1-42} , Figure 4C also demonstrates that ANP can spontaneously generate PAOs during incubation at room temperature for 6 days (lane 1), illustrating the fibrillogenic nature of this peptide. Finally, Western analysis of cell lysate from control and paced cells was performed for ANP. As shown in Figure 4D, the blot demonstrates the appearance of bands with rapid stimulation having nearly the identical size as the new bands that appeared on the A-11 Western blot (32 and 95 kDa; Figure 1E) and the predominant oligomer-related band (10 kDa) formed during the *in vitro* experiments with purified ANP shown in **panel C**. These data link both γ -ketoaldehydes and rapid pacing to the production of ANP oligomers, and they provide strong support for the

hypothesis that ANP-derived oligomers are a significant component of the PAOs that form with rapid stimulation of atrial HL-1 cells.

3.5 Effect of the γ -Ketoaldehyde Scavenger Salicylamine

To confirm an essential role for γ -KAs in atrio-myocyte PAO formation, cells were rapidly paced in the absence and presence of the γ -KA scavenger salicylamine. As illustrated in Figure 5A and B, pacing-induced PAO formation was essentially eliminated when salicylamine was present in the culture medium.

We have previously shown that rapid activation of atrial cells causes a myocyte stress response, a component of which includes transcriptional upregulation of *Nppb* and *Hspa1a* (encoding heat shock protein 70; Figure 6, left bars) [9]. However, in the presence of salicylamine, this response was blunted (Figure 6, middle bars; see Supplementary Data), providing additional evidence to support a cytoprotective response when γ -KAs are scavenged.

To investigate the specificity of the effects of SA, similar experiments were conducted using the antioxidant/anti-inflammatory compound curcumin [33]. This drug prevents soluble oligomer aggregation and fibril formation for amyloid β_{1-42} *in vitro* and *in vivo* [34]. However, unlike salicylamine, curcumin lacks an amine group, so that it cannot scavenge γ -KAs. In the presence of curcumin, rapid stimulation of atrial HL-1 cells resulted in robust generation of PAOs (Figure 5C and 5D) similar to that observed under drug-free conditions, as well as activation of stress response genes (Figure 6, right bars). Collectively, these data establish a critical role for γ -KAs in PAO formation in rapidly-activated atrial myocytes.

4. DISCUSSION

The basic molecular processes that promote atrial remodeling and disease progression during AF are not well understood. In the present study, we have identified a novel pathologic component of the myocyte stress response that occurs during rapid activation of atrial cells, namely the generation of protein oligomers. Our findings also indicate that a fundamental molecular mechanism driving the formation of these protein complexes is the production of highly-reactive γ -ketoaldehydes by ROS (Figure 7). The capacity of PAOs to cause injury of cells (including cardiomyocytes) is well documented [13]. Thus, these data support a growing body of evidence that “proteotoxicity” can occur in the heart and contribute to cardiac dysfunction in humans [2, 35].

A wide variety of physiologic and pathophysiologic factors can promote protein misfolding, including alterations in cellular temperature, mechanical strain, and oxidative stress [35]. In response, these stimuli trigger the production of molecular chaperones designed to protect newly-synthesized or damaged proteins from aggregation. Among these chaperones, heat shock proteins are particularly abundant in cardiac tissue, and we have previously shown that they are among the most highly-upregulated genes during rapid activation of atrial HL-1 cells [9]. Expression of heat shock proteins is increased in the atria of both animal models and patients with AF [36-38], while pharmacologic upregulation appears to be protective [38]. These data imply a prominent role for protein misfolding *in vivo*, and thus

proteotoxicity, as a novel potential mechanism of myocyte injury that could increase AF susceptibility (Figure 7). Because PAOs are not visible by standard amyloid staining methods, the identification of these cytotoxic protein oligomers has become possible only recently with the development of appropriate antibodies, such as A-11. Moreover, as demonstrated in the brain, PAOs can cause organ dysfunction in the absence of demonstrable cellular pathology [39].

For degenerative disorders related to aging, factors such as impaired protein homeostatic mechanisms and acquired mutations can contribute to protein misfolding [35]. Senile amyloidosis can develop exclusively in the atria, known as isolated atrial amyloidosis [31, 40, 41]. The incidence of this atrial-specific disorder increases with age, exceeding 90% in the ninth decade of life. In several studies, isolated atrial amyloidosis has been linked to an increased risk of AF [31, 32, 41]. Given that PAOs are intermediate complexes in the formation of amyloid fibrils (Figure 7), it is logical that they should be detectable in human atria, which we have recently demonstrated [42]. Moreover, amyloid fibrils in atrial amyloidosis are primarily composed of ANP, and our data demonstrate that ANP is a likely component of PAOs in rapidly-stimulated atrial HL-1 cells. Based on the organ-specific nature of isolated atrial amyloidosis, it is likely that as for the brain in Alzheimer's disease, PAOs play a role in atrial pathophysiology.

Increasingly, toxic protein oligomers have been recognized to play a role in experimental and human cardiac disease, in particular cardiomyopathy [35]. α B-Crystallin (CryAB) is a small heat shock protein in the heart that binds the intermediate filament desmin. Mutations in CryAB cause a cardiomyopathy characterized by PAO formation that are toxic to proteasomes and implicated in myocardial dysfunction [15, 35]. Recent data indicate that PAOs can be detected in ventricular tissue of patients with systolic and/or diastolic dysfunction in the setting of dilated, hypertensive, and hypertrophic cardiomyopathy [14, 43, 44]. Given our data that rapid activation induces protein misfolding in atrial myocytes as well, we hypothesize that mechanical strain is a stimulus for PAO formation in cardiomyocytes.

Oxidative stress has been linked to both Alzheimer's disease and AF, supporting the concept that common pathophysiologic mechanisms may be present for these diseases. Upstream therapy with drugs having antioxidant properties (e.g., statins, omega (n)-3 polyunsaturated fatty acids, and RAAS inhibitors) have not proven beneficial for primary or secondary prevention of AF [7, 45]. An important omission in nearly all of the studies is that the ability of the intervention to reduce ROS was not measured. Thus, one potential explanation is insufficient potency of existing drugs. Indeed, recent studies have shown that therapeutically-used doses of both vitamin E [46, 47] and fish oil [48] are ineffective to reduce *in vivo* measures of oxidative stress. This limitation of existing antioxidants has led to the concept of identifying molecular scavengers that specifically target critical mediators of atrial oxidant injury. Reactive aldehyde products of fatty acid oxidation have recently been identified as strong candidates for this pivotal role, given their capacity to react extremely rapidly with lysyl residues and covalently modify, crosslink, and aggregate proteins, as well as DNA [17, 21, 49-51]. Additionally, γ -KA adducts have been identified in affected regions of the brain in patients with Alzheimer's disease [52]. γ -KAs have been

shown to directly promote the formation of soluble oligomers derived from amyloid β_{1-42} [24], while inactivation of these reactive γ -KAs with the small molecule scavenger SA prevented development of cognitive impairment in a mouse model of Alzheimer's disease [26]. A recent study has identified a required role for γ -KAs in T cell-mediated hypertension in mice, and SA (referred to as 2-hydroxybenzylamine or 2-HOBA) was effective to lower blood pressure [53]. Our data, demonstrating an inhibitory effect of salicylamine, support a similar mechanism for γ -KAs in PAO formation in rapidly-paced atrial HL-1 cells.

Conclusion

Our results identify preamyloid oligomers as a novel component of myocyte stress remodeling in response to rapid atrial cell activation, and they identify γ -KAs as a critical molecular component of this pathophysiologic process. Future studies *in vivo* are needed to determine whether these ROS-generated mediators could represent a novel therapeutic target in the treatment of AF. Nonetheless, PAO formation by γ -KAs provides a potential mechanistic link between oxidative stress, atrial cell injury, and AF susceptibility.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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ABBREVIATIONS

AF	atrial fibrillation
PAO	preamyloid oligomer
γ-KAs	γ -ketoaldehydes
ANP	atrial natriuretic peptide
SA	salicylamine
RAAS	renin-angiotensin-aldosterone system
PFA	paraformaldehyde
PBS	phosphate buffered saline
BSA	bovine serum albumin

q-PCR	real-time quantitative RT-PCR
Ct	cycle threshold
CryAB	α B-crystallin
DHE	dihydroethidium
DAB	3,3-diaminobenzidine
ROS	reactive oxygen species
E₂-IsoKs	E ₂ -isolevuglandin/isoketals

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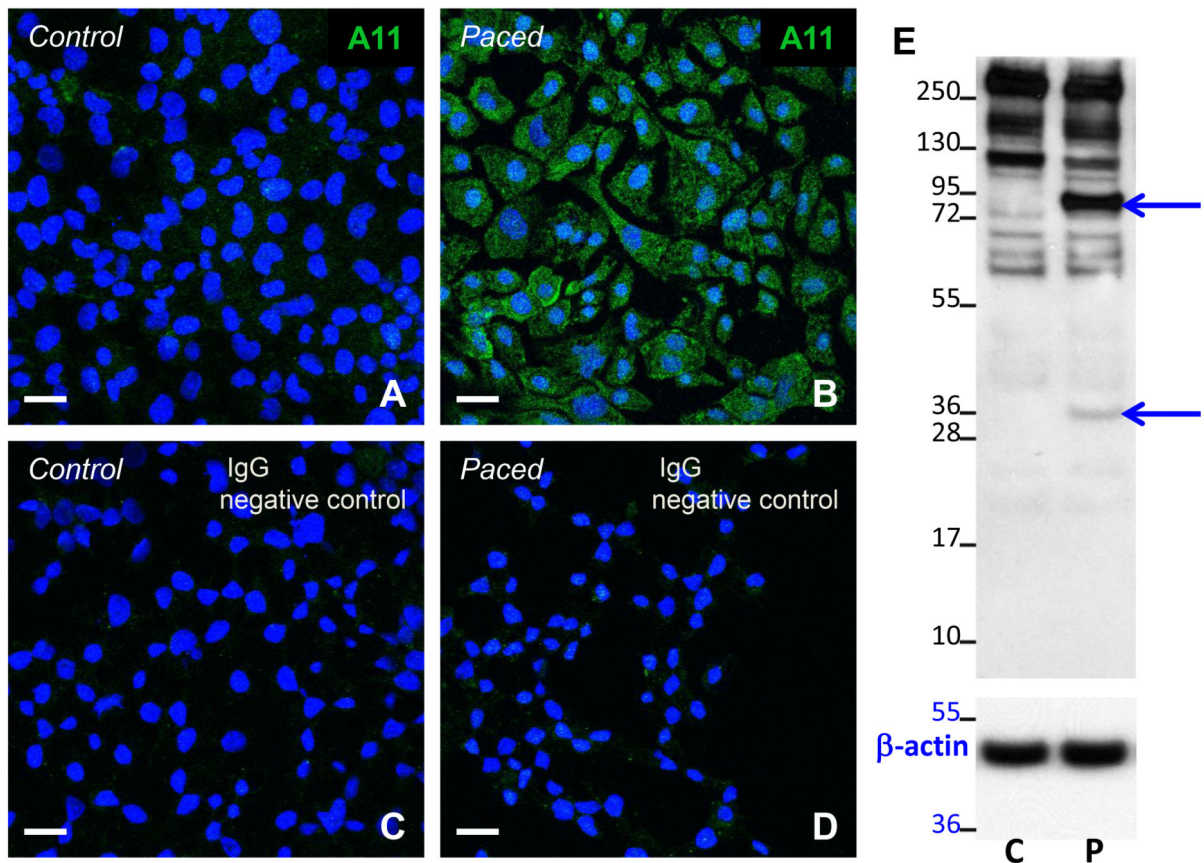


Figure 1. PAO formation following rapid stimulation of atrial HL-1 cells

Exposure of rapidly-paced cells to the anti-oligomer antibody A-11 revealed diffuse cytosolic immunostaining (**B**) that was absent in control, unpaced cells (**A**). Immunolabeling was not detected when IgG was substituted for A-11 (**C**, **D**). Scale bars = 20 microns. (**E**) Using A-11, Western analysis of cell lysate (20 μ g) from control (**C**) and rapidly-stimulated (**P**) atrial HL-1 cells demonstrates the development of bands at 34 and 95 kDa (arrows) with pacing. To control for protein loading, the blot was stripped and reprobed with an anti- β actin antibody (lower panel).

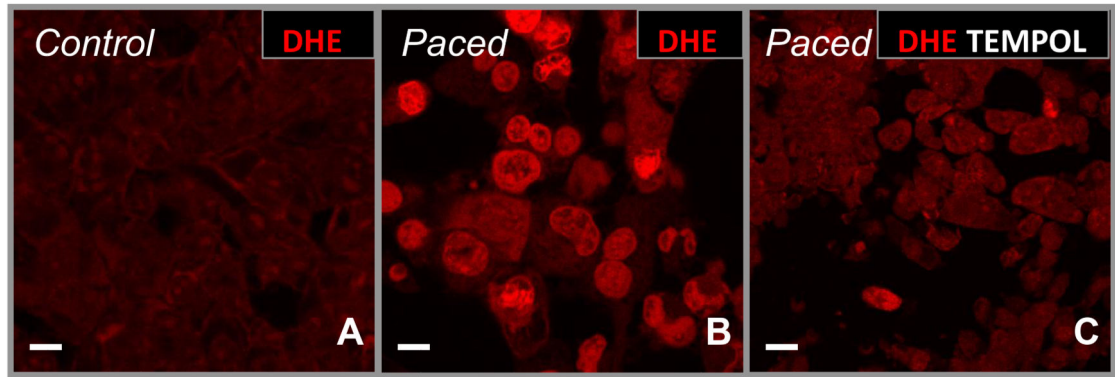


Figure 2. Production of superoxide ($O_2^{\cdot-}$) with rapid stimulation

After 6hr of rapid stimulation in culture, $O_2^{\cdot-}$ production by atrial HL-1 cells was detected by dihydroethidium (DHE; $10\mu M$; **B**, indicated by the bright red color) that is not present in control, spontaneously-beating cells (**A**). Superoxide generation was essentially abolished under these conditions by co-incubation of cells with the cell-permeable superoxide dismutase mimetic tempol (**C**). Scale bars = 10 microns.

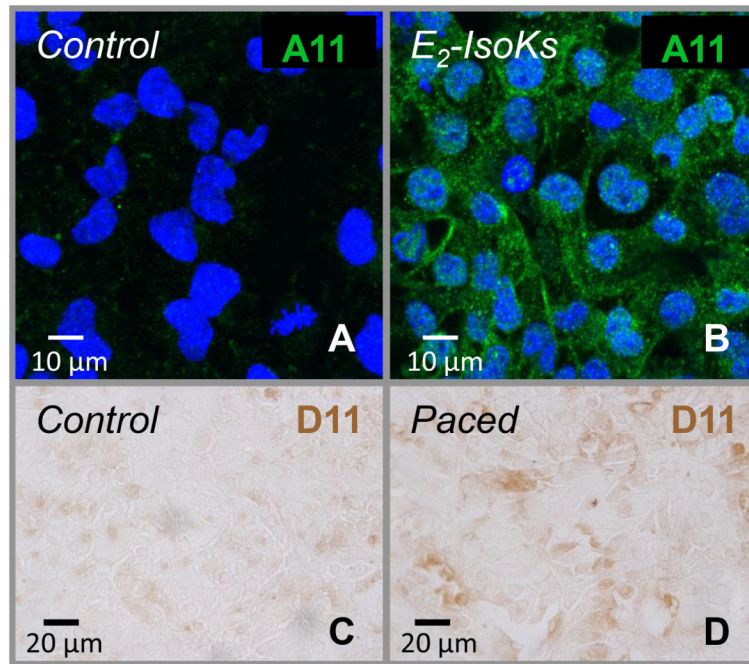


Figure 3. Evidence for a role of γ -ketoaldehydes in PAO formation for paced atrial HL-1 cells After incubation with synthetic isolevugland in/isoketals (E_2 -IsoKs; 1 μ M) for 6hr, immunofluorescent labeling with A-11 demonstrated abundant PAO formation in unstimulated atrial HL-1 cells (**top row**). Immunoreactivity using the anti- γ -KA adduct antibody (D11 ScFv) is shown in control and paced HL-1 cells (**bottom row**). Positivity (indicated by brown-colored DAB substrate) is evident in the paced cells.

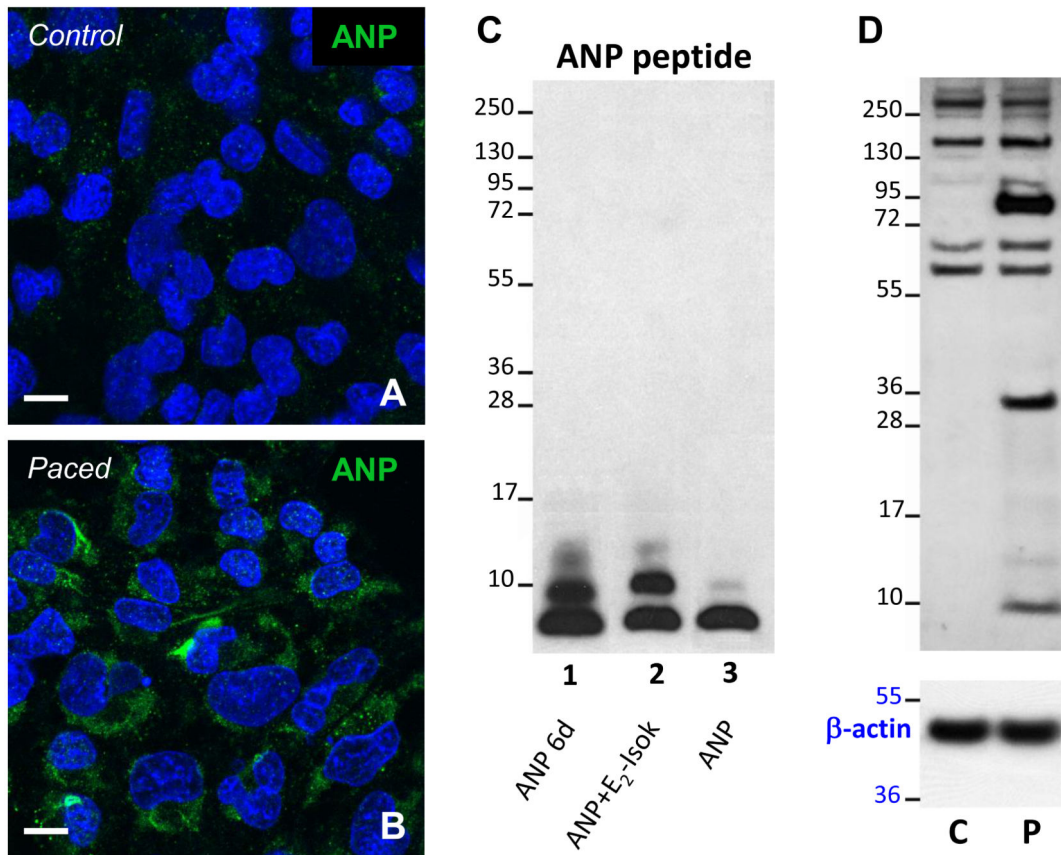


Figure 4. Increased production of atrial natriuretic peptide (ANP) and ANP oligomers in rapidly-paced atrial HL-1 cells

Immunofluorescent labeling with an anti- α -ANP antibody demonstrated a significant increase of ANP production in atrial HL-1 cells with rapid stimulation (**B**) compared to unpaced (control) cells (**A**). Scale bars = 10 microns. (**C**) Using the same antibody, Western analysis was performed on samples of purified ANP peptide (10 μ M) following incubation (at room temperature) for 6 days (lane 1), incubation for 24hr (lane 3), and incubation for 24hr in the presence of E₂-IsoKs (lane 2). The development of ANP oligomers in lanes 1 and 2 is indicated by the appearance of higher molecular weight species, predominantly at ~10 kDa. (**D**) Using the anti- α -ANP antibody, Western analysis of cell lysate from control (**C**) and rapidly-stimulated (**P**) atrial HL-1 cells indicates the development of bands with pacing that are nearly identical in size (32 and 95 kDa) to the pacing-induced bands shown in Figure 1E, as well as a 10 kDa band similar to the oligomer-related band in **panel C**.

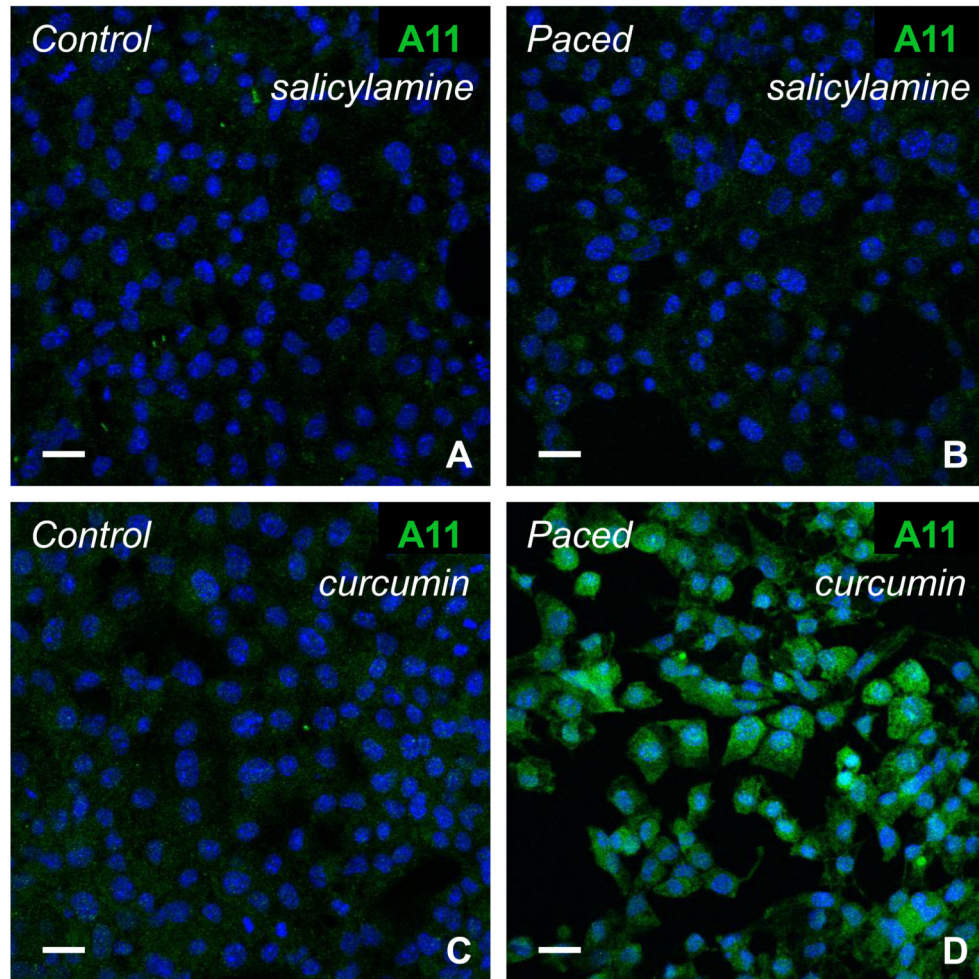


Figure 5. Salicylamine inhibited cytosolic PAO formation in rapidly-paced atrial HL-1 cells, while curcumin did not

Immunostaining with the A-11 antibody was minimal for control (A) and rapidly-paced (B) cells in presence of the γ -ketoaldehyde scavenger salicylamine (100 μ M). However, abundant PAO formation was detected for paced cells when cultured with the antioxidant curcumin (20 μ M; C, D). Scale bars = 20 microns.

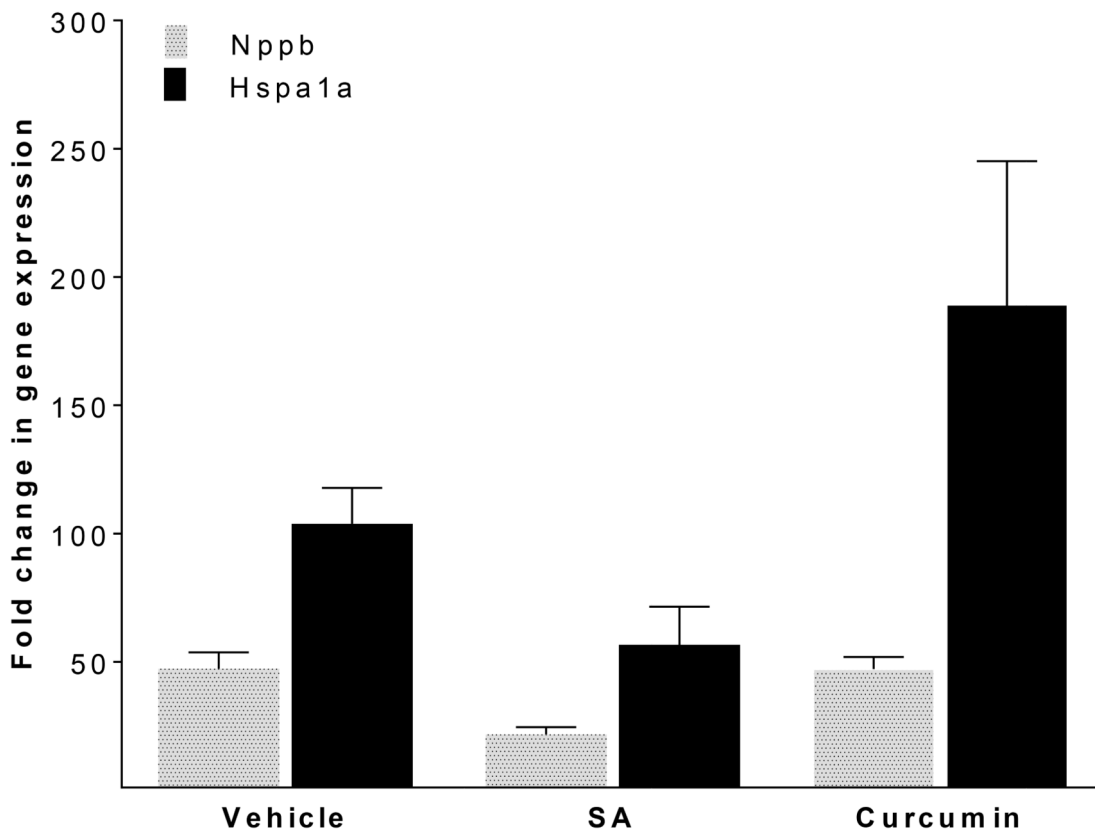


Figure 6. Effect of salicylamine (SA) and curcumin on transcriptional upregulation of *Nppb* and *Hspa1a* with rapid pacing

Real-time quantitative RT-PCR results are expressed as mean fold change for *Nppb* and *Hspa1a* for atrial HL-1 cells subjected to rapid stimulation for 6hr (n = 6; fold change expressed as the residual value calculated from paced and control cells Ct values). In the absence of SA or curcumin, mRNA expression was up-regulated for *Nppb* (Ct_{control} 18.56±0.91 vs. Ct_{paced} 13.67±0.68) and *Hspa1a* (Ct_{control} 21.00±0.22 vs. Ct_{paced} 15.19±0.31) with rapid-pacing (p=0.002 and p<0.001, respectively; see Supplementary Data). For *Nppb*, the group effect for drug was significant (p=0.047), while the group effect for *Hspa1a* was not (p=0.097). However, pairwise comparisons (performed with a Wilcoxon rank sum test) did not demonstrate significant changes for *Nppb* or *Hspa1a* with either salicylamine or curcumin.

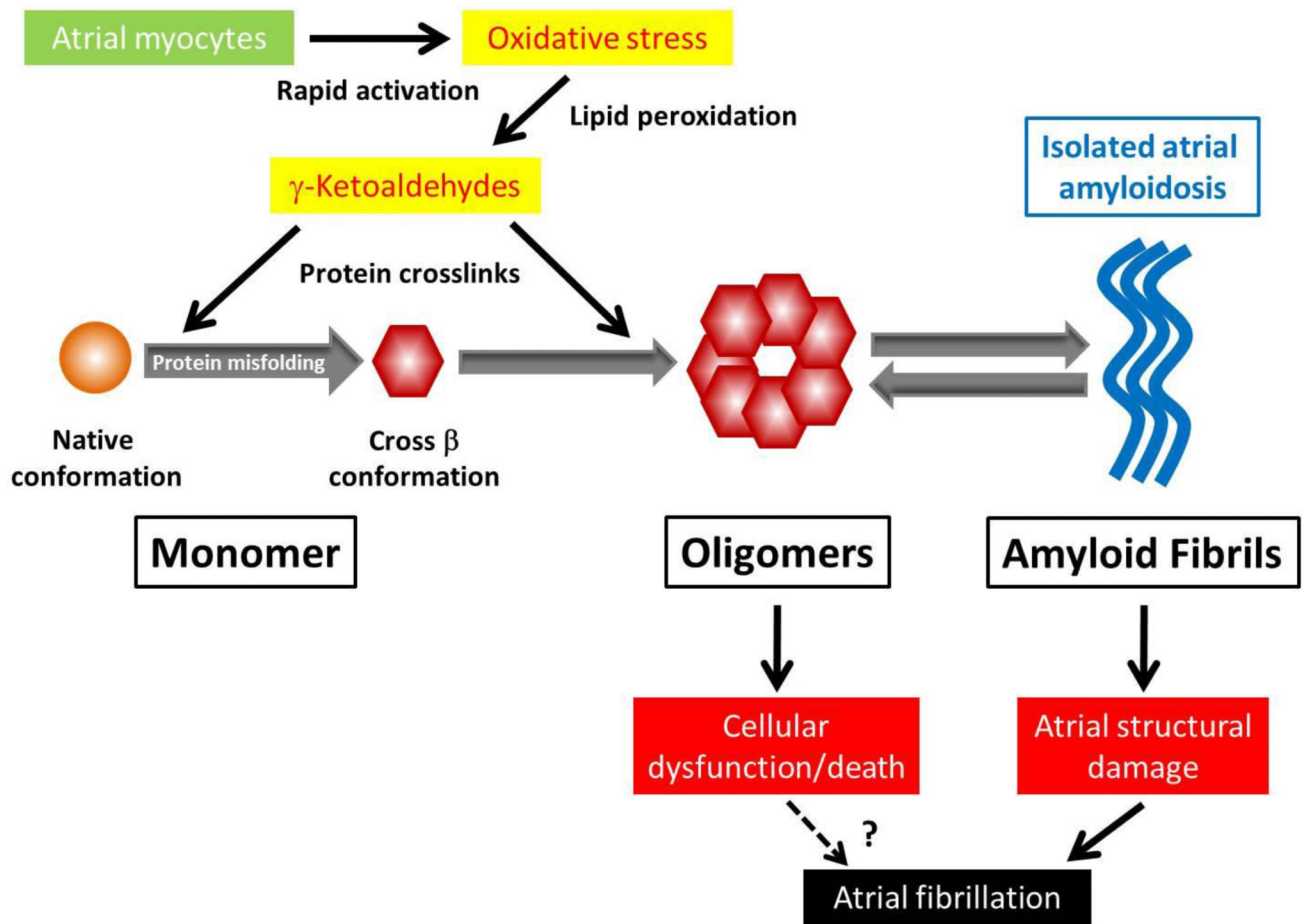


Figure 7. Hypothesis linking rapidly-activated atrial cells, PAO formation, and atrial pathology/arrhythmogenesis, compared with the known process of isolated atrial amyloidosis

Rapid activation of atrial cells causes oxidative stress, with superoxide generation. As a consequence of lipid peroxidation, highly-reactive γ -ketoaldehydes are formed that rapidly crosslink cellular proteins. This initiates protein misfolding and oligomerization in atrial cells, with ANP a principal component of PAOs. Protein oligomers can directly cause myocyte injury/death, or they can coalesce with aging to form amyloid fibril deposition in atrial myocardium. This process, known as isolated atrial amyloidosis, is associated with infiltrative structural damage in the atrium and an increased risk of atrial fibrillation.