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Lone Atrial Fibrillation: What is known and what is to come.

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Disclosures

None.

Abstract

Atrial fibrillation (AF) is the most prevalent sustained cardiac arrhythmia in adults, affecting >1% of general population. AF is commonly associated with structural heart disease and is a major cause of significant cardiovascular morbidity and mortality. AF sometimes develops in a subset of young patients (e.g. aged ≤ 60 years), with no evidence of associated cardiopulmonary or other comorbid disease (including hypertension), and has been referred to as 'lone AF'. The latter generally has a favorable prognosis; the prognostic and therapeutic implications of an accurate identification of patients with truly lone AF (that is, truly at low risk of complications), if any, would be of the utmost importance. The true prevalence of lone AF is unknown, varying between 1.6% and 30%, depending on the particular study population.

Nonetheless, novel risk factors for AF, including obesity, metabolic syndrome, sleep apnea, alcohol consumption, endurance sports, anger, hostility, subclinical atherosclerosis and others, have been increasingly recognized. Also, various underlying pathophysiologic mechanisms predisposing to AF, including increased atrial stretch, structural and electrophysiological alterations, autonomic imbalance, systemic inflammation, oxidative stress and genetic predisposition, have been proposed. The growing evidence of these diverse (and numerous) pathogenic mechanisms and factors related to AF inevitably raises the question of whether 'lone AF' does exist at all.

In this review article we summarize the current knowledge of the epidemiology, pathophysiology, clinical course and treatment of patients with so-called 'lone AF' and outline emerging insights into its pathogenesis and the potential therapeutic implications of a diagnosis of lone AF.

Review criteria

We searched MEDLINE (1950 through October 2010) and Google Scholar using the terms "lone atrial fibrillation", "lone atrial fibrillation epidemiology", "natural history of lone atrial fibrillation" and "lone atrial fibrillation what do we know" and manually reviewed the references in English language. Abstracts from international cardiovascular meetings were studied to identify unpublished data.

Message for the clinic

The patients diagnosed as so-called 'lone AF' comprise a heterogeneous group and should not *a priori* be classified as having a benign rhythm disorder. Detailed assessment of lone AF patients is warranted, both at presentation and regularly during follow-up. Emerging risk factors and pathophysiologic mechanisms predisposing to AF should also be considered and, clearly, there is a need for further research regarding stroke risk stratification and optimal antiarrhythmic treatment in these patients, including 'old' and new investigational antiarrhythmic drugs and catheter ablation of AF.

Introduction

Atrial fibrillation (AF) is the most prevalent sustained cardiac arrhythmia in adults, affecting >1% of general population. The incidence of AF is projected to rise over the next few decades and AF has therefore been referred to as a "non-contagious epidemic" [1-5].

AF is associated with significant cardiovascular morbidity and mortality, primarily due to ischemic stroke, congestive heart failure (CHF) and impaired quality of life, with the requirement for chronic use of medication [6-11]. Well-defined risk factors for AF include advanced age, hypertension, structural heart disease, CHF, diabetes mellitus and thyroid disease [12,13]. In turn, the risk of AF-related complications is strongly influenced by the nature and extent of underlying co-morbid disease and age [9,13,14].

AF sometimes develops in a subset of young patients (e.g. aged ≤ 60 years), with no evidence of associated cardiopulmonary or other comorbid disease (including hypertension), and has been referred to as 'lone AF', which generally has a favorable prognosis [15-23]. The term "lone" AF was introduced ~60 years ago and it is still a diagnosis of exclusion, which should be made only after careful assessment including detailed medical history, physical examination, laboratory testing with thyroid status, electrocardiogram, echocardiography and, possibly, chest x-ray and exercise testing [15,24].

Nonetheless, novel risk factors for AF, including obesity [25-28], metabolic syndrome [29,30], sleep apnea [31-35], increased alcohol consumption [36-41], endurance sport practice [42-46], increased pulse pressure [47], anger and hostility [48,49], subclinical atherosclerosis [50,51] and others, are increasingly recognized. Also, various underlying pathophysiologic mechanisms predisposing to AF, including increased atrial stretch, structural and electrophysiological alterations, autonomic imbalance, systemic inflammation, oxidative stress and genetic predisposition, have been proposed [52-56]. The growing evidence of these diverse (and numerous) pathogenic mechanisms and factors related to AF inevitably raises the question of whether lone AF does exist at all [57,58]. The prognostic and therapeutic implications of an

accurate identification of patients with truly lone AF (that is, truly at low risk of complications), if any, would be of the utmost importance [59].

In this review article we summarize the current knowledge of the epidemiology, pathophysiology, clinical course and treatment of patients with so-called 'lone AF' and outline emerging insights into its pathogenesis and the potential therapeutic implications of making a diagnosis of 'lone AF'.

Epidemiology of lone AF

The overall prevalence of AF increases with age: <1% of individuals younger than 60 and $\sim10\%$ of those >80 years old have AF [3]. The true prevalence of lone AF is unknown, varying between 1.6% and 30%, depending on the patients' age and criteria used in the study [16-18,20,60].

The lifetime risk for development of AF, in subjects aged above 40, is 1:4 both for males and females [61]. Nevertheless, males have a higher incidence of AF in all age groups. Indeed, AF develops in males at 1.5-fold greater rate than the rate in females, when adjusted for other risk factors [12,62]. There is a male preponderance in lone AF, with a male-to-female ratio of 3-4:1 [16,17,20]. However, male predominance is less evident among confirmed familial AF probands or affected relatives, when compared with nonfamilial (sporadic) lone AF patients [63]. To explain the increased frequency of sporadic lone AF in males, the possibility of an unrecognized X-linked recessive AF in males with negative family history and apparently sporadic AF (and mothers and sisters as the healthy carriers) has been raised [63]. Of note, familial and sporadic lone AF share similar phenotypes and are clinically indistinguishable [63].

A genetic contribution to AF was recognized nearly 70 years ago, when Wolff documented an autosomal dominant transmission of lone AF in one family [64]. Since then, the familial aggregation of lone AF has been increasingly reported and certain genetic mutations (often confined to the particular family, and not found outside the family) have been described [56,65-

73]. The heritability of AF in the general population has been documented in several populationbased studies [74-77]. For example, in the Framingham Study, parental AF increased the risk for offspring AF with an Odds Ratio of 3.2 (95%CI: 1.7-5,9, p<0.01), when the sample was limited to those age <75 years old without antecedent cardiovascular morbidity [74].

In the past decade, an increasing number of genetic loci and causal mutations of genes (primarily those encoding components of potassium, sodium and L-type calcium ion channels) have been identified in lone AF patients, and significant overlap of AF with congenital long QT, short QT and Brugada syndrome, and with dilated cardiomyopathy has been observed (Table 1) [56,65-83]. However, large-scale sequencing in lone AF cohorts suggests that these channel mutations are not a major cause of AF [56,83].

Nevertheless, patients at comparable risk for AF often do not exhibit uniform susceptibility to the arrhythmia, which could possibly be attributed to a genetically determined "AF diathesis" [56,83,84]. The introduction of genome-wide association studies, which compare non-familial AF cases to age-related and gender-matched controls, has enabled some insight into the genetics of acquired AF, and several common gene or single nucleotide polymorphisms including those encoding the renin-angiotensin-aldosterone system (RAAS), calcium handling, neurohumoral and lipoprotein pathways, gap junction proteins, ion channels, interleukins, signaling molecules and mediators of other molecular pathways have been examined [56,83,84]. However, common polymorphisms identified thus far perhaps explain only <10% of the inherited contribution to AF [83-86].

To date, genetic studies have revealed diverse mechanisms of susceptibility to AF (Table 1). Single-gene disorders provide an insight into the electrophysiological mechanisms which predispose to AF, but it is very likely that vulnerability to AF primarily arises from the influence of multiple genes which, either alone or in combination, alter the atrial structural and functional properties. The recognition of those genes and further insight into their effect on atrial electrophysiology could facilitate the implementation of a pharmacogenomic treatment approach for treatment of AF in the future [56].

Pathophysiology and pathogenesis of lone AF

The occurrence of AF requires a trigger and appropriate atrial substrate to sustain reentry of electrical impulses. Various triggers can create propagating wavelets, which may lead to reentrant circuits and AF in the presence of functionally and structurally altered atrial myocardium (Figure 1) [87,88]. The electrophysiological hallmarks of an atrial pro-fibrillatory state are a reduced atrial effective refractory period (ERP), attenuated ERP rate adaptation and reduced impulse conduction velocity [87].

The condition of the atrial substrate is strongly influenced by the underlying disease process and ageing. Furthermore, AF itself initiates the complex processes of atrial electrical, contractile and structural remodeling, which further contributes to the AF persistence and progression (that is, "AF begets AF") (Table 1) [52,89]. Such atrial electrical and contractile remodeling occurs rapidly (within hours and days) and is reversible, whilst structural changes occur after weeks or months of persistent AF and are usually irreversible [52]. Thus, the early restoration of sinus rhythm (SR) should, at least theoretically, prevent further progression of AF. Unfortunately, a large body of evidence suggests that "SR does not beget SR" and that, in fact, vulnerability to AF and the risk of AF perpetuation and stability often extend beyond the time course of electrical remodeling reversal, which led to the hypothesis that "a second factor" (that is, some structural changes) is responsible for AF recurrence [90,91].

In the setting of lone AF, whereby structural disease is absent by definition, AF may be viewed as a primary electrophysiological disorder with ectopic focal discharges from the pulmonary veins and posterior wall of the left atrium playing an important role in both the initiation and maintenance of AF [88,92,93]. However, atrial fibrosis and loss of muscle mass have been observed in lone AF patients, along with isolated atrial myocardial perfusion abnormalities and coronary flow reserve impairment indicative of microvascular dysfunction [94,95]. For example, in a recent study of 25 patients with paroxysmal lone AF, bi-atrial abnormalities remote from arrhythmia (and thus very unlikely to result from AF-induced remodeling) were documented, including conduction abnormalities, structural changes and sinus node dysfunction [96]. More

recently, one study of 58 lone AF patients also documented distinct abnormalities in atrial electrophysiological properties [97].

After the initial description of isolated atrial structural changes consistent with myocarditis in lone AF patients [94], many studies have documented a relationship between circulating markers of inflammation (including C-reactive protein and interleukins) and AF. Indeed, inflammation plays a significant role in the initiation and perpetuation of AF and the AF-related prothrombotic state [53,98-100]. Inflammatory biomarkers are associated with future development, recurrence and burden of AF and the likelihood of successful cardioversion [101-105]. However, inflammation may be associated not with AF per se, but rather with underlying cardiovascular disease [106]. Whether the initiation of AF activates direct inflammatory reactions or the presence of pre-existing inflammation promotes AF is not completely clarified, and both mechanisms may be operating. For example, rapid atrial activation accompanied with calcium accumulation may result in calcium overload and apoptotic loss of atrial myocytes, with the consequent low-grade inflammatory response as a part of structural remodeling [107]. Alternatively, in patients with triggering atrial foci, systemic inflammation with increased circulating C-reactive protein may predispose to AF via the classic complement pathway activation and atrial tissue damage, or by binding to phosphocholine which may contribute to membrane dysfunction by the alterations in sodium and calcium handling [98].

Other potential biomarkers of a predisposition to apparently lone AF have also been described. Atrial natriuretic peptide (ANP) and brain natriuretic peptide (BNP) are sensitive biomarkers of cardiac contractile dysfunction, and both peptides are elevated in AF patients with underlying structural heart disease [108]. In lone AF patients, a discordant pattern of elevated N-terminalpro-BNP and normal N-terminal-pro-ANP levels can be observed, but the precise mechanism of ANP and BNP uncoupling is not clarified [109]. The question of whether the increased BNP levels are the result of AF or simply a sign of subclinical cardiac alterations which predispose to AF still remains unanswered.

Apelin is an endogenous peptide hormone, which is normally involved in the counter-regulation of the angiotensin and vasopressin systems. A significant reduction of mean apelin levels has

been found in patients with lone AF, indicating certain alterations of the cardiac humoral axis [110].

Variations in autonomic nervous system tone are also involved in AF. Sympathetic stimulation often initiates AF in patients with structural heart disease, whilst lone AF patients more frequently have paroxysmal AF in the setting of increased vagal tone [87,111]. Although both sympathetic and parasympathetic components play a role in AF, the cholinergic component appears to be important for spontaneous initiation of AF; in the experimental setting, for example, electrical stimulation of the left atrial ganglionic plexi or the autonomic nerve endings with retrograde activation of the ganglia induces spontaneous firing from pulmonary veins followed by AF [112].

Ultrastructural alterations and cellular electrophysiology in lone AF

A decreased duration of action potential (AP), depressed AP plateau and attenuated response of AP duration to depolarization rate are the principal electrophysiological findings at the atrial cellular level [113,114]. These alterations are mediated through changes in sodium, potassium and calcium channels, which are not likely to be the principal cause of AF, but rather a nonspecific response to variety of stresses such as an altered oxidative state in the setting of increased metabolic demand, elevated sympathetic tone and increased levels of circulating cytokines. The redox state and metabolic reserve are also age-dependent, and regulation of channels is largely genetically determined [87].

Connexins are protein subunits of gap junctions - the clusters of transmembrane channels that link adjacent cells and mediate cell-to-cell electrical coupling and communication. Decreased numbers and altered spatial arrangement of connexin40 in AF have been documented, both in the atrial myocardium and thoracic veins [115-117]. These findings correlate with facilitated trigger activity in the pulmonary veins and discontinuous and slowed conduction due to intercellular uncoupling in the atrial myocardium [118,119]. In addition, connexin40 gene mutations or polymorphisms underlie an inherited predisposition to AF [82].

Atrial ultrastructural changes in AF patients are identified at the level of cardiomyocytes and extracellular matrix. Whilst fibrosis and myocyte degeneration could be attributed to the underlying structural disease, AF itself induces de-differentiation of cardiomyocytes and regression toward a more immature developmental stage with depleted contractile apparatus and fetal actin and myosin forms [55,87]. Apoptotic cell death is also more pronounced in the fibrillating atria [55]. An increased accumulation of various proteins in the extracellular matrix, including collagen, fibronectin-1 and fibrillin-1, has also been documented, along with the upregulation of matrix metalloproteinases, transforming growth factor B1 and AT-1 angiotensin II receptor [52,55,87]. These changes may significantly alter the atrial micro-architecture (which is already complex under normal conditions) and increase vulnerability to AF in the setting of an acute stretch, which promptly activates the immediate gene program for cell hypertrophy, augments the synthesis of angiotensin II, influences the stretch-activated and other ion channels, receptors and enzymes with cytoskeletal connections, thus creating a fertile electrophysiological and anatomical environment for initiation and propagation of AF [87]. Reduced calcium systolic release and myolysis underlie contractile remodeling, which could trigger thrombus formation and atrial dilatation [120].

Risk factors for lone AF

Apart from the traditional AF risk factors, there are numerous conditions which have been associated with AF in apparently healthy individuals.

Independent of other cardiovascular risk factors, obesity confers an increased risk of new-onset AF (3%-8% with each unit of increase in body mass index), which has been attributed to diastolic dysfunction due to myocardial thickening, elevated plasma volume and increased neurohormonal activation resulting in left atrial dilatation [25-27]. Lone AF patients have also been reported to be taller and leaner than other AF patients [28].

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Obstructive sleep apnea may cause various hemodynamic alterations and autonomic imbalance predisposing to AF, and, conversely, a paroxysm of AF may result in central sleep apnea caused by an acute decrease in the left ventricular function accompanied by an increase in pulmonary wedge pressure and consequent stimulation of pulmonary vagal receptors [31-35].

A "holiday heart syndrome" of paroxysmal AF after occasional excessive alcohol intake was described over 30 years ago. Alcohol has a direct toxic effect on cardiomyocytes, causes a hyperadrenergic state with impaired vagal tone and may increase the intra-atrial conduction time. However, moderate alcohol intake (2-3 drinks/week) was not related to the increased risk of AF in the Cardiovascular Health Study [36-41]. Other stimulants including caffeine and nicotine have also been associated with AF [57].

In contrast to mild regular exercise, endurance sport practice is associated with a higher prevalence of AF, and >1500 lifetime hours of practice seems to be the threshold for the association. Enlargement of the cardiac chambers, an increase in left ventricular mass and left atrial dilatation, along with increased vagal tone resulting in bradycardia and shortening of the atrial ERP, transient hypovolaemia and changes in electrolytes due to sweating have been proposed to explain the vulnerability to AF in sportsmen. There is also anecdotal evidence of AF development after the use of anabolic steroids, and changes in the autonomic function and the baro-reflex could be the underlying mechanism [42-46,57].

Various social characteristics, including the type of personality, anger and hostility and acute life stress have been shown to increase the vulnerability to AF [48,49]. Drug-induced AF should be considered when inotropic agents (dopamine), cholinergics (acetylcholine), adenosine, broncodilatators (especially sympathicomimetic inhalants), corticosteroids, cytostatics, central nervous system drugs (anticholinergics, dopamine agonists, antidepressants, antipsychotics, anaestetics) and others are administered [121].

Finally, what seems to be apparently 'lone AF' may occur in the setting of subclinical atherosclerosis (determined by carotid intima-media thickness and/or mitral annular calcification

[MAC]). In the Framingham Study, for example, MAC was an independent predictor for incident AF (hazard ratio 1.6, 95%CI: 1.1-2.2) [50,51].

Clinical course of lone AF

In some studies, lone AF patients may have a similar risk of thromboembolism, CHF and mortality as the general population [17,21,22]. However, other studies suggest otherwise. For example, in the Paris Prospective Study I, lone AF was associated with increased mortality in middle aged working men [16,23]. Also, ageing and development of underlying heart disease strongly influence the long-term prognosis of (what was originally) lone AF [14,21]. Therefore, regular follow-up with evaluation of cardiovascular risk factors is necessary in patients diagnosed as 'lone AF'. This is supported by the observation that as many as 44% of apparently lone AF patients may actually have occult arterial hypertension [122].

Patients originally diagnosed with lone AF may follow divergent clinical courses based on the left atrial volume. In contrast to those with normal atria, patients with increased left atrial volume (either at diagnosis or during the follow-up) subsequently experienced adverse cardiovascular events including stroke, which occurred at left atrial volumes >32mL/m² [123].

It has also been suggested that paroxysmal lone AF carries a better prognosis than chronic lone AF, regarding thromboembolism and mortality [20]. The vast majority of lone AF patients present with paroxysmal arrhythmia and have a relatively low rate of progression to permanent AF during the follow-up, with an estimated risk of progression of approximately 30% over 30 years, being significantly influenced by ageing and development of comorbidities [21]. Clinical AF types are summarized in Figure 2.

Management of lone AF

The principal aims of AF treatment are to reduce symptoms and to prevent AF-related complications, using antithrombotic therapy, ventricular rate control and treatment of concomitant cardiovascular risk factors and/or comorbidities. Symptom relief may additionally require rhythm control including cardioversion(s), long-term use of antiarrhythmic drugs and/or catheter-ablation (Figure 3) [124].

Stroke and thromboembolism is a severe complication of AF. More recently, greater efforts have been directed towards the identification of "truly low risk" AF patients who are at low risk of stroke and, therefore, do not need any antithrombotic therapy; all other patients with ≥ 1 stroke risk factors merit oral anticoagulation therapy (Figure 3) [59]. This approach is acknowledged in the latest 2010 European Society of Cardiology Guidelines for AF [124].

The cumulative risk of stroke is very low in true 'lone AF' patients (1.3% over 15 years) and no therapy (or aspirin) is advised [21,124]. Of note, the thromboprophylactic effect of aspirin in lone AF patients is questionable. In the Japan AF Stroke trial, aspirin150-200mg daily was worse than no therapy (the annual incidence of primary outcomes was 3.1% vs. 2.4%, respectively), and the use of aspirin was associated with non-significant increased risk of major bleeds (1.6% with aspirin vs. 0.4% in the control group) [125]. Meta-analysis of 7 trials comparing aspirin with placebo in AF patients showed a non-significant 19%-reduction in stroke incidence (95%CI -1% to 35%), which could easily be attributed to an effect of aspirin on atherosclerosis and atherothrombosis [126].

Dedicated re-assessment of thromboembolic risk factors during regular follow-ups is mandatory in lone AF patients, since the stroke risk increases with ageing or development of comorbidities (including hypertension). However, various atrial ultrastructural changes, contractile dysfunction and endothelial perturbation have been documented in apparently lone AF [101-105,108-110,120,127]. With growing insights into these alterations, all of which could influence the stroke risk, and increasing number of potentially useful biomarkers, further refinement of thromboembolic risk assessment should be expected.

Moreover, new oral anticoagulants are being developed for stroke prevention in AF. Those are the oral direct thrombin inhibitors (dabigatran etexilate, AZD0837) and oral factor Xa inhibitors (rivaroxaban, apixaban, endoxaban, betrixaban, YM150 and others). In contrast to the oral vitamin-K-antagonists (VKA), such as warfarin, these new agents have a rapid onset of action, a low propensity for food and drug interactions and a predictable anticoagulant effect after fixeddose administration and coagulation monitoring is unnecessary. The efficacy and safety of dabigatran compared with warfarin for the prevention of stroke and systemic embolism in AF patients was investigated in the Randomized Evaluation of Long-term anticoagulant therapy with dabigatran etexilate (RE-LY) trial [128]. The Apixaban VERsus acetylsalicylic acid to pRevent strOkES (AVERROES) study was stopped early because of clearly evident superiority of apixaban 5mg bid over aspirin in patients intolerant of or unsuitable for VKA, with similar rates of major bleeding with aspirin and the oral anticoagulant, and aspirin was less well tolerated [124].

Rhythm control may be particularly successful in patients with paroxysmal lone AF, primarily due to a relatively low propensity to AF progression, and various strategies could be implemented including no therapy, a "pill-in-the-pocket" strategy or long-term use of antiarrhythmic drugs (Figure 3). In general, available antiarrhythmic drugs may not be particularly effective in acute cardioversion of AF, and are only moderately successful in long-term rhythm control. Should ß-blockers be ineffective, flecainide, propafenone, sotalol or dronedarone could be tried. Although amiodarone has a superior efficacy compared with other antiarrhythmic drugs, it should be reserved as an alternative because of its potential toxicity with long-term use [124,129]. Catheter ablation may be applied in symptomatic lone AF patients in whom antiarrhythmic drugs were not effective or at patient's preference [124]. Ablation strategies clearly offer better rhythm control during the short-term follow-up, but late AF recurrences are not uncommon [130-132]. Classic surgical 'maze procedure' and pulmonary vein isolation are highly effective in rhythm control, but novel procedures including thoracoscopic access and video assistance are promising [124,133,134].

Most of the currently available antiarrhythmic drugs exert their anti-AF effects via inhibition of cardiac ion channels, at the expense of an increased risk of life-threatening ventricular

arrhythmias, worsening of left ventricular systolic dysfunction or organotoxicity [135]. Until recently, large clinical trials evaluating the ability of antiarrhythmic drugs to improve all-cause mortality have yielded neutral outcomes at best [120,135]. However, there appears to be significant reduction in both cardiovascular morbidity and mortality in AF patients treated with dronedarone, which has encouraged an essential change of focus of rhythm control management [129,136]. Nevertheless, there was no significant reduction in the all-cause mortality in AF patients treated with dronedarone [136]. In the prevention of recurrent AF, dronedarone is better than placebo [137] and less successful, but with significantly fewer adverse effects, compared with amiodarone [129]. However, the use of dronedarone is not safe in patients with significantly reduced left ventricular systolic function (ejection fraction \leq 35%), in whom dronedarone has been shown to increase mortality, predominantly due to cardiovascular causes (most notably CHF) [138].

Novel pharmacological investigational approaches to rhythm control in AF include atrialselective multiple ion channel blockers which are expected to reduce the risk of ventricular arrhythmias (vernakalant, ranolazine, AVE0118, GsMTx4, tertiapin-Q, NIP-141, NIP-142, etc), improved derivatives of existing drugs (AZD1305, AVE1231, SSR149744C, ATI2042), gap junction modifiers (rotigaptide, AAP10, GAP-134, etc), and upstream therapy, which targets arrhythmogenic structural remodeling in the atria and/or factors that promote such remodeling including hypertension, heart failure and inflammation (angiotensin-converting enzyme inhibitors [ACEIs], angiotensin receptor blockers [ARBs], statins and omega-3 polyunsaturated fatty acids [PUFAs]) [120,124,135]. Vernakalant hydrochloride is a relatively atrial-selective, early-activating K⁺ and frequency-dependent Na⁺ channel blocker with a half-life of 2 to 3 hours. In a randomized clinical trial, vernakalant demonstrated rapid conversion of short-duration AF compared with placebo and was well tolerated [139].

Upstream therapy could prevent the development of AF (primary prevention) or recurrence and progression to permanent AF (secondary prevention). At present, upstream therapies are not recommended for primary prevention of AF in patients without cardiovascular disease [124]. However, the use of ACEIs and ARBs may be considered in patients with recurrent AF who are taking antiarrhythmic drug therapy. There is good experimental evidence of antifibrillatory and

antifibrotic effects of these drugs in AF patients, but the results of clinical studies are conflicting [124,140,141]. In animal models of AF, statins attenuate atrial electrical and structural remodeling and reduced inducibility of AF. However, clinical evidence in support of using statins in the secondary AF prevention, except for post-operative AF, is insufficient [124, 142,143].

Conclusions

Accumulating insights into the pathogenesis of *apparently* lone AF suggest that the entity comprises heterogeneous subsets of AF patients who should not *a priori* be classified as having a benign rhythm disorder. Growing evidence of numerous common variations within the genome and atrial ultrastructural alterations that predispose to AF makes the classification of AF as secondary to structural disease or as a 'lone' arrhythmia increasingly meaningless. Detailed assessment of lone AF patients is warranted, both at presentation and regularly during the follow-up. Emerging risk factors and pathophysiological mechanisms predisposing to AF should also be considered and, clearly, there is a need for further research regarding stroke risk stratification and proper antiarrhythmic drug treatment in so-called lone AF patients.

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Table 1. The genetics of atrial fibrillation.

	Chromosome	Gene	Ion channel/ current/other	Electrophysiological effects of the mutation	Mechanism of AF	Associated phenotype/comment
nented, but rare causes of AF	11p15.5 21q22.1 17q23-24 X chrom. 7q36.1	KCNQ1 ^[67] KCNE2 ^[70] KCNJ2 ^[78] KCNE5 ^[79] KCNH2 ^[56]	I _{Ks} I _{Ks} [KCNQ1-KCNE2] I _{K1} I _{Ks} K _v 11.1	accelerated repolarization of atrial cardiomyocytes, shortening of AP duration reduction of ERP	multiple reentry wavelets	KCNQ1-causative gene for Long QT syndrome type 1 KCNE2, KCNJ2 and KCNE5 - normal QT interval KCNH2 and KCNQ1 – associated with Long QT type 2 and Short QT syndrome (~30% of cases)
	12p13	KCNA5 ^[71]	K _v 1.5/I _{Kur}	delayed atrial AP repolarization, prolonged both AP duration and atrial ERP hyperpolarization,	early afterdepolarizations initiate atrial polymorphic tachycardia "torsade de pointes" which degenerates to AE	- SCN5A-associated with Long QT type 3, Brugada
	12p13.1	CACNA1C, CACNB2 ^[81]	ICaL	prolonged AP duration enhanced heterogenity of repolarization	multiple reentry wavelets	syndrome and Sick sinus syndrome Associated both with Brugada syndrome and Short OT syndrome
cur	Gen. loci ^[65-69]	crici (b2		ropoliti induci		
her mutations (candidate genes) ^[56,68,82,83] Do	10q22-24	unknown	?	?	?	Overlaps with locus for DCM
	6q14-16	unknown	?	?	?	Overlaps with locus for DCM
	10p11-q21	unknown	?	?	?	Unknown
	5p11	unknown	?	?	?	Associated with prolonged P-wave
	Chrom. 3	SCN5A	Na _v 1.5	cell hyperexcitability enhanced automaticity of cardiomyocytes	electrical reentry with "mother- rotor" and spiral reentry waves	Not associated with Long QT syndrome
	5p13	NUP155	Nuclear pore protein	?	?	Associated with Emery-Dreifuss muscular dystrophy syndromes
	Chrom. 1	GJA5	Connexin40	heterogenity of conduction velocity enhanced vulnerability to electrical reentry	multiple reentry wavelets	<i>Connexins</i> : transmembrane proteins that form gap- junctions (intercellular pores) which serve as low- resistance conduction pathways between adjacent cells
	1p36-p35	NPPA	ANP	shortening of atrial AP, reduction of ERP	multiple reentry wavelets (in vitro)	As a mediator of inflammation, ANP may cause atrial fibrosis
	4q25	PITX2	RAAS (?)	?	?	Suppresses pacemaker cells from forming outside the sinus node during cardiogenesis
	16q22	ZFHX3	?	?	?	Associated with both ischemic and cardioembolic strokes
Otl	1q21	KCNN3	?	?	?	-

AF=atrial fibrillation, AP=action potential, ERP=effective refractory period, DCM=dilated cardiomyopathy, ANP=atrial natriuretic peptide, RAAS=rennin-angiotensin-aldosterone system.





PV=pulmonary veins, ERP=effective refractory period, CRP=C-reactive protein.





rial fibrillation.

Modified from Camm AJ, et al. [124].

AF=at





AF=atrial fibrillation, ACEI=angiotensin-converting-enzyme inhibitor, ARB=angitensinreceptor blocker, PUFA=polyunsaturated fatty acid, TE=thromboembolism.

Modified from Camm AJ, et al. [124].