Prospective Evaluation of Clinico-pathological Predictors of Post-operative Atrial Fibrillation: An Ancillary Study from the OPERA Trial

Running title: Corradi & Saffitz et al.; Predictors of post-operative atrial fibrillation

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Abstract:

Background - Post-operative atrial fibrillation (POAF) occurs in 30-50% of patients undergoing cardiac surgery and is associated with increased morbidity and mortality. Prospective identification of structural/molecular changes in atrial myocardium that correlate with myocardial injury and precede and predict risk of POAF may identify new molecular pathways and targets for prevention of this common morbid complication.

Methods - Right atrial appendage (RAA) samples were prospectively collected during cardiac surgery from 239 patients enrolled in the OPERA trial, fixed in 10% buffered formalin and embedded in paraffin for histology. We assessed general tissue morphology, cardiomyocyte diameters, myocytolysis (perinuclear myofibril loss), accumulation of perinuclear glycogen, interstitial fibrosis, and myocardial gap junction distribution. We also assayed NT-proBNP, hscTnT, CRP, and circulating oxidative stress biomarkers (F2-isoprostanes, F3-isoprostanes, isofurans) in plasma collected before, during, and 48h after surgery. POAF was defined as occurrence of post-cardiac surgery atrial fibrillation or flutter of at least 30 seconds duration confirmed by rhythm strip or 12-lead ECG. The follow-up period for all arrhythmias was from surgery until hospital discharge or post-operative day 10.

Results - 35.2% of patients experienced POAF. Compared to the non-POAF group, they were slightly older and more likely to have chronic obstructive pulmonary disease or heart failure. They also had a higher EuroSCORE and more often underwent valve surgery. No differences in left atrial size were observed between POAF and non-POAF patients. The extent of atrial interstitial fibrosis, cardiomyocyte myocytolysis, cardiomyocyte diameter, glycogen score or Cx43 distribution at the time of surgery was not significantly associated with incidence of POAF. None of these histopathologic abnormalities were correlated with levels of NT-proBNP, hs-cTnT, CRP or oxidative stress biomarkers.

Conclusions - In sinus rhythm patients undergoing cardiac surgery, histopathological changes in the RAA do not predict POAF. They also do not correlate with biomarkers of cardiac function, inflammation, and oxidative stress.

Key words: pathology; atrial fibrillation; remodeling; biomarker; postoperative complication arrhythmia; fibrosis; cardiomyocyte; right atrial appendage

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Nonstandard Abbreviations and Acronyms

AF: atrial fibrillation

BNP: brain natriuretic peptide

CRP: C-reactive protein

Cx43: connexin 43

ECG: electrocardiogram

EuroSCORE: European system for cardiac operative risk evaluation

n-3 PUFA: long-chain ω -3 polyunsaturated fatty acid

NT-proBNP: N-terminal pro brain natriuretic peptide

OPERA: omega-3 fatty acids for prevention of post-operative atrial fibrillation

PAS: periodic acid–Schiff

PCI: prior percutaneous coronary intervention

POAF: post-operative atrial fibrillation

RAA: right atrial appendage



Introduction

Post-operative atrial fibrillation (POAF) is the most common complication of cardiac surgery ranging from 30% to 50% of cases and usually occurring within 2 to 4 days after the surgical procedure (with a peak incidence on post-operative day 2).¹⁻⁴ POAF, in its turn, is associated with increased post-operative thromboembolic risk and stroke, hemodynamic compromise, ventricular dysrhythmias, as well as significantly higher both in-hospital and 6-month mortality.⁵ In addition to the type of surgery, the only well-established predictor for the development of POAF is advanced age.⁶⁻¹¹

Remarkably, despite decades of recognition of POAF and attempts to mitigate its development, the underlying pathophysiologic mechanisms for POAF remain unclear. One hypothesis holds that both initiation and maintenance of this arrhythmia depend on a structural substrate that arises as a result of the multiple risk factors listed above³. However, a major limitation in developing preventive strategies has been a lack of understanding of the specific preceding anatomic and molecular changes in the atrial myocardium at the time of cardiac surgery that might influence the risk of POAF.

Prior pathological studies have typically been retrospective (with tissue collected after onset of atrial fibrillation, AF), and have combined tissues from patients who had already experienced AF for variable intervals (e.g. new onset, paroxysmal, and persistent). These studies have also relied on what were likely non-representative samples obtained at convenience rather than pre-specified sampling by protocol. These limitations have made it impossible to identify and define rigorously the critical tissue substrates of AF, and to distinguish changes in tissue structure that increase the risk of AF from those caused by AF itself.

Therefore, we set out to assess structural and molecular changes in atrial myocardium and circulating biomarkers collected prospectively in a large number of cardiac surgery patients to determine if any might be independently associated with the subsequent development of POAF.

Methods

The data that support the findings of this study are available from the corresponding author upon reasonable request.

OPERA patient population and ω-3-PUFA supplementation

The Omega-3 Fatty Acids for Prevention of Post-Operative Atrial Fibrillation (OPERA) study was a prospective, double-blind, placebo-controlled, multinational, randomized clinical trial that tested the hypothesis that perioperative long-chain ω -3 polyunsaturated fatty acid (n-3 PUFA)

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supplementation reduces the occurrence of POAF after cardiac surgery.^{4, 12} The 1,516 patients enrolled in this trial were in sinus rhythm at the time of surgery. POAF was defined as "any documented AF or atrial flutter following cardiac surgery of at least 30 seconds duration and documented by rhythm strip or 12-lead electrocardiogram (ECG). If only a shorter duration ECG is available, then the diagnosis of AF/atrial flutter is based on the arrhythmia being present at onset or termination". The time period of follow-up for all arrhythmia endpoints was from the time of cardiac surgery until hospital discharge or post-op day 10, whichever occurred first.⁴ The protocol specified collection of blood samples before, during, and after surgery, and biopsy of the right atrial appendage (RAA) at the time of surgery.

As a pre-specified substudy, RAA samples were prospectively collected during cardiac surgery from 292 OPERA patients. Out of these, 239 cases had RAA biopsies suitable for a reliable histopathological analysis and were the object of the present study. The baseline characteristics of this subsample was similar to the overall OPERA trial population (Supplementary Table 1). All histological and biochemical analyses were performed centrally by skilled personnel unaware of patient characteristics, study treatment or outcomes.

This study was approved by the IRB of all participating institutions and conducted according to international standards of Good Clinical Practice (FDA Title 21 part 312, International Conference on Harmonization guidelines). All patients provided written informed consent.

Additional details are reported in the Data Supplement.

Histopathologic and morphometric methods

After excision, tissue specimens were fixed in 10% buffered formalin solution for at least 24 hours and then embedded in paraffin tissue blocks. From each block, 4 μ m-thick histological

sections were cut for histopathological analyses aimed at exploring the most common changes observed in the setting of AF structural remodeling. RAA samples were initially observed in hematoxylin-eosin stained sections to evaluate general tissue morphology (Figures 1A-1B). To quantitate cardiomyocyte hypertrophy, the transverse diameters of 100 longitudinally oriented cardiomyocytes were measured (through the nucleus region) in PAS-stained sections using Cell^F software (Olympus Soft Imaging Solutions, version 2.6) (Figures 1C-1D)¹³. The presence of severe perinuclear myofibril loss in cardiomyocytes (so-called "myocytolysis") was also determined in the same PAS-stained sections¹³. The percentage of cardiomyocytes showing myocytolysis was assessed at a magnification of x400. Accumulation of glycogen in the perinuclear area in cells showing myocytolysis is an indicator of a metabolic shift from the use of fatty acids to glucose. The presence of glycogen granules was evaluated at a magnification of x400 and scored semi-qualitatively into three categories: *none* (absent), *mild-moderate*, *severe*. Examples are shown in Figure 2.

The percentage of myocardium occupied by interstitial fibrosis was assessed at a magnification of x400 in Van Gieson-stained sections (Figure 3). To quantify fibrosis, a light microscope equipped with an ocular reticule containing 42 sampling points was used. The fraction of points overlying interstitial fibrosis was accurately counted and multiplied by 100.¹⁴

Connexin 43 distribution

A primary antibody against the major myocardial gap junction protein connexin 43 (Cx43; SIGMA Aldrich, polyclonal, code C6219, dilution 1:400, overnight at 4°C) was used to characterize the distribution of gap junctions in cardiomyocytes.

Detailed descriptions of the standard immunofluorescence methods have previously been published.^{15, 16} Immunopositive signals were detected using a Cy3-conjugated goat anti-rabbit

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(H+L) (Jackson Immunoresearch, Cambridgeshire, UK; catalog number 111-165-144; dilution 1:400) and the localization of Cx43 was qualitatively evaluated through a confocal microscope (Zeiss 510, Belgium) at an original magnification of x40. No attempt was made to quantify immunoreactive signals. Rather, we optimized the staining protocol to identify the lowest antibody concentration that shows a strong signal at intercalated disks in normal myocardial samples. We then assessed changes in the amount and/or distribution of signal in the tissue samples. As shown in Figure 4, Cx43 signal was categorized as a) *control-like*, when strong signal occurred in a normal junctional distribution; b) *depressed*, when the amount of signal was reduced; c)*heterogeneous*, if both a) and b) conditions coexisted; or d) *absent*. The extent to which Cx43 immunoreactive signal was redistributed to the lateral borders of cardiomycytes was categorized as either *minimal/absent* or *present*.

Statistical analyses

Multivariate linear and logistic regression analyses were employed to assess the relation of circulating markers [brain batriuretic peptide (BNP), troponin, C-reactive protein (CRP), plasma and urine F2-isoprostanes, F3-isoprostanes, isofurans] with the atrial histopathological endpoints. Detailed description of the performed analyses is described in the *Supplement Material*.

Supplementary Methods

Methods regarding assay of circulating cardiac markers and oxidative stress biomarkers are detailed in the *Supplement Material*.

Results

Characteristics of patients undergoing atrial biopsy

RAA tissue samples were obtained from 239 patients, of whom 141 were American participants and 98 Italian patients. Characteristics of the overall population and those in the current histological sub-study were similar and are summarized in Supplementary Table 1.

The mean age of study patients was 64.8±12.6 years; 72.4% were male, and 53/239 patients (21.2%) had heart failure, 8% had experienced previous AF, 17.2% had prior percutaneous coronary interventions (PCI), and 4.3% had undergone previous cardiac surgery. The median logistic EuroSCORE (European System for Cardiac Operative Risk Evaluation) value was 3.8 (interquartile range [IQR], 1.9-8.4). Valve surgery was performed in 58.8% of Heart Associated patients; see also Table 1.

Eighty-four patients (35.2%) developed POAF. Compared to patients who did not develop POAF, these patients were older (67.2 vs. 63.5, p=0.03) and more likely to have chronic obstructive pulmonary disease (22.6% vs 12.3%, p=0.04). Further, they presented more often with heart failure (29.3% vs 18.2%, p=0.04), experienced more prior AF episodes (17.9% vs 2.6%, p<0.001), and had a higher EuroSCORE (4.8 vs 3.1, p=0.04). They were also likely to have undergone valve surgery (69.1% vs 53.3%, p=0.02), compared to those in the non-POAF group. Left atrial size (41.5 vs 42.2, p=0.56) and left ventricular ejection fraction (58.4 vs 57.3, p=0.47) were comparable between the two groups (Table 1).

Histopathological, morphometric and Connexin 43 analyses

None of the histological variables (amount of interstitial fibrosis, cardiomyocyte myocytolysis, cardiomyocyte diameter, glycogen score, Cx43 distribution analysis) significantly differed between POAF and non-POAF patients (Figures 1-4 and Table 2). In a multivariable logistic

regression model adjusted for all statistically significant clinical variables and including all histological features, the following factors independently associated with incident POAF: age (OR: 1.05 95% CI: 1.01-1.09, p=0.021), history of prior AF (OR: 11.6, 95% CI: 2.26-59.0, p=0.003), and congestive heart failure (OR=0.37, 95% CI:0.16-0.85, P=0.018). None of the histological features (e.g. fibrosis, myocytolysis, cardiomyocyte diameter, glycogen accumulation, Cx43 patterns) reached statistical significance. A non-significant trend was seen between presence of Cx43 lateralization at the time of cardiac surgery and lower future risk of POAF (OR=0.460, 95% CI=0.191-1.104, P=0.084).

We also explored the univariate relationships between baseline characteristics of the population and the histological features of the right atrium at the time of cardiac surgery. Patients with angina pectoris had a higher fibrosis percentage (p=0.034; Supplementary Table 2); men had more myocytolysis than women (p=0.022); and patients with diabetes mellitus showed more perinuclear myofibril loss and moderate accumulation of glycogen (p=0.022, p=0.003 respectively). Cx43 junctional signal (Figure 4) was depressed at intercalated disks in patients with COPD (p=0.006) and with higher EuroSCORE index (p=0.043). Interestingly, patients randomized to the n-3 PUFA treatment had less frequent lateralization of Cx43, compared with the placebo group (p=0.035) (data not shown). These exploratory analyses were not corrected for multiple comparisons and should be interpreted with caution.

Since the administration of n-3 PUFA correlated with reduced Cx43 lateralization, we determined if it also impacted POAF development in the subgroup in which Cx43 lateralization was present. However, compared with placebo, n-3 PUFA treatment did not significantly alter the POAF risk in the 202 patients with Cx43 present (p=0.656)

Histopathological features and biomarker concentrations

In multivariable-adjusted analyses, biomarker levels of cardiac strain, inflammation, and oxidative stress were generally not significantly associated with histopathological features of the right atrium at the time of cardiac surgery (Tables 3 and 4). The most notable association was an inverse association between N-terminal pro brain natriuretic peptide (NT-proBNP) levels and those of myocytolysis (p=0.001): mean NT-proBNP levels were about twice as high in the lowest tertile of myocytolysis, compared to the highest tertile.

Discussion

POAF and its currently-known predisposing factors

POAF commonly complicates cardiac surgery, and represents a major contributor to postoperative morbidity and increased length of stay and other healthcare utilization.¹ Its incidence has increased significantly over the few past decades, likely because of the aging of the patient populations undergoing cardiac surgery. Despite substantial investigation, the pathologic and pathophysiologic variables that can reliably predict development of POAF and might provide mechanistic insight into its pathogenesis remain elusive.¹⁷

POAF onset peaks on the second post-operative day, which suggests a key role for acute surgery-induced factors.¹⁷ Inflammation and oxidative stress related to atrial trauma during surgery, systemic inflammation, and stress related to cardiopulmonary bypass may all contribute to supraventricular arrhythmogenesis. Atrial enlargement due to intravenous fluid administration post-operatively may also predict POAF. All these early changes have the potential of creating atrial conduction disturbances and enhancing triggers for POAF.

In addition to the above-mentioned factors directly related to surgery, various features reflecting chronic cardiac remodeling might increase POAF risk by creating anatomical substrates that promote arrhythmia maintenance. Aging itself correlates strongly with POAF presumably due, at least in part, to age-related collagen deposition within atrial myocardium which may favor fragmentation and reentrant arrhythmogenesis. Associated co-morbidities, such as systolic dysfunction, hypertension, COPD, and obesity, may also increase POAF risk.

Major strengths of the present investigation

The present investigation – an ancillary study to the large, placebo-controlled, double-blind randomized OPERA clinical trial – is the most comprehensive prospective evaluation to-date of how key structural and molecular characteristics of the right atrial myocardium at the time of surgery relate to the subsequent risk of POAF. Prior studies of myocardial histopathology and AF have nearly always been cross-sectional, i.e., with myocardial sampling occurring after AF is already present; and utilized convenience samples rather than a more general, prospectively ascertained population of patients undergoing cardiac surgery. Major strengths of the present study include (a) analysis of the largest number of prospectively collected (prior to AF onset) atrial tissue samples; (b) analysis of a defined, well characterized multi-center patient population in sinus rhythm at the time of cardiac surgery (previously, it has been impossible to distinguish changes in tissue structure that increased risk of AF from those associated to persistent AF); and (c) the broad inclusion criteria of OPERA and its multi-center, multi-country design which maximized generalizability of our findings to the real-world population of patients, while minimizing potential bias from patient selection. Other strengths include participation by two experienced cardiac pathologists who were members of the OPERA Biological Studies Committee and have expertise in morphological/immuno-histochemical analysis of atrial

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myocardium and anatomic substrates of cardiac arrhythmias, ensuring a coordinated and rigorous approach; and a generalizable patient population with overall characteristics and incidence of POAF resembling those reported in numerous previous large patient samples.

POAF occurrence and the levels of circulating cardiac/oxidative stress biomarkers are not correlated with the structural remodeling of the RAA

Despite the above-mentioned strengths, this investigation did not demonstrate any consistent statistically significant relationships between structural remodeling of the RAA (e.g., interstitial fibrosis, percentage of myocytolytic cardiomyocytes, muscle-cell hypertrophy, Cx43 distribution) and development of POAF. Similarly, levels of circulating cardiac biomarkers (e.g., NT-proBNP, high-sensitivity cardiac troponin T, high sensitivity C-reactive protein)¹⁸ and oxidative stress biomarkers (e.g., F2-isoprostanes, F3-isoprostanes, isofurans)¹⁹ did not correlate with the presence of these histopathologic abnormalities. One exception is the counterintuitive highly significant relation (Table 3, p=0.003) between plasma F2-isoprostanes and fibrosis. This is, most likely, a chance finding attributable to the multiplicity of comparisons.

To our knowledge, the present study represents the largest prospective investigation todate of these crucial relationships in cardiac surgery patients. Yet, while the sample size of nearly 250 patients is impressive for a prospective collection, it may have been insufficient to convincingly detect more subtle relationships between structural remodeling of the RAA, biomarkers of cardiac stress and inflammation, and risk of POAF. The findings relating to Cx43 lateralization, randomized n-3 PUFA treatment, and POAF; to NT-proBNP and myocytolysis; and to other associations that did not achieve statistical significance, provide an important foundation for further hypothesis generation and subsequent prospective studies to elucidate the crucial unanswered questions on atrial structural remodelling and risk of POAF.

Review of the literature

With these issues in mind, we reviewed 49 clinical and experimental studies focusing on the relationship between RAA structural remodeling and, broadly speaking, AF (either postoperative or non-surgery associated; Table 5 and Table 6, respectively). This included 19 studies on POAF and 30 (5 in animal models) on non-POAF supraventricular arrhythmia. Studies on POAF assessed the predictive value of histological alterations for incident AF, while in the other 30 studies associations between ongoing AF and histological alterations were assessed. Specific experimental designs differed, and relatively few studies comprehensively investigated multiple aspects of structural remodeling of the RAA including fibrosis, myocytolysis, cardiomyocyte hypertrophy, and connexins. The results varied considerably regarding specific histopathological features that correlated with AF. None of these previous studies were double-blind, placebocontrolled, multinational, and randomized in nature as the present OPERA sub-study was and in which these biases were minimized. These methodological differences may be largely responsible for the higher frequency of significant relationships between histological features and risk of POAF seen in previous studies. Another concern involves use of automatic measurements, used in many prior studies, which may incorrectly classify and overestimate the extent of histological changes. In this regard, 19 previous studies (39%) used automatic morphometry methods, 8 (16%) used semi-automatic measurements, and 22 (49%) involved non-automatic (manual) histopathological evaluations. Only 13 of 19 investigations on POAF (68%) and 6 of 30 studies on non-POAF (20%) employed multivariate data analysis. Finally, the sample size of the various populations under investigation differed substantially between studies (96±55, range 24-259 in POAF manuscripts; 72±59, range 8-245 in non-POAF papers).

Limitations of the study

One potential limitation was the RAA sampling site, selected for both ethical and practical reasons in this large trial. While the RAA would be expected to undergo remodeling in response to AF, which involves the entire atrial myocardium, it may not exhibit the full range of pathogenic changes in a pre-AF stage as it may be more protected than the left atrium from the effects of many AF-associated conditions. Only a minority of OPERA patients had isolated right heart dysfunction. In addition, AF usually arises in the pulmonary veins and then progressively affects the left atrium, especially its posterior wall.²⁰ Thus, the RAA may not be an ideal biopsy site to identify structural and molecular determinants of POAF in patients with mainly left-sided cardiac disease.

Conclusions

In conclusion, in this large, prospective clinico-pathological analysis of patients in sinus rhythm undergoing cardiac surgery, histopathological features of the RAA at the time of surgery did not significantly correlate with the subsequent development of POAF, and baseline biomarkers of cardiac inflammation and oxidative stress also generally did not associate with the RAA histopathological features. These novel findings provide an important foundation for further hypotheses and prospective studies to elucidate the crucial unanswered questions on atrial structural remodeling and POAF risk.

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Disclosures: none.

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Variables		Total	Non-POAF	POAF	P *	
N (%)		239	155 (64.8%)	84 (35.2%)		
Age		64.8 ± 12.6	63.5 ± 13.3	67.2 ± 10.7	0.028	
g	Female	66 (27.6%)	41 (26.5%)	25 (29.8%)	0.585	
Sex	Male	173 (72.4%)	114 (73.6%)	59 (70.2%)		
	Placebo	122 (51.0%)	80 (51.6%)	42 (50.0%)	0.912	
Treatment	n-3 PUFA	117 (49.0%)	75 (48.4%)	42 (50.0%)	0.812	
Current smoking		25 (10.6%)	17 (11.1%)	8 (9.6%)	0.726	
COPD		38 (15.9%)	19 (12.3%)	19 (22.6%)	0.037	
Diabetes Mellitus		72 (30.1%)	45 (29.0%)	27 (32.1%)	0.617	
Hypertension		196 (82.0%)	123 (79.4%)	73 (86.9%)	0.147	
Chronic renal failur	·e	11 (4.6%)	8 (5.2%)	3 (3.6%)	0.751	
Prior myocardial in	farction	57/236 (24.2%)	38/154 (24.7%)	19/82 (23.2%)	0.797	
Angina pectoris		51/208 (24.5%)	30/134 (22.4%)	21/74 (28.4%)	0.336	
Heart Failure		53 (22.3%)	28 (18.2%)	25 (29.8%)	0.040	
Previous PCI		41 (17.2%)	25 (16.1%)	16 (19.1%)	0.568	
Previous AF		19 (8.0%)	4 (2.6%)	15 (17.9%)	<0001 tion.	
Prior cardiac surgery ⁺		10 (4.3%)	6 (3.9%)	4 (4.8%)	0.744	
Coronary bypass		5 (2.1%)	5 (3.2%)	0	1.000	
Valve surgery		3 (1.3%)	1 (0.7%)	2 (2.4%)	0.283	
Other cardiac surgery	7	3 (1.3%)	1 (0.7%)	2 (2.4%)	0.283	
LA size (mm)	UNIC	41.8 ± 7.2	41.5 ± 7.1	42.2 ± 7.5	0.562	
LVEF(%)		58.0 ± 10.4	58.4 ± 11.0	57.3 ± 9.3	0.469	
B-Blockers		113 (53.6%)	70 (51.1%)	43 (58.1%)	0.330	
Statins		113 (53.6%)	71 (51.8%)	42 (56.8%)	0.493	
ACE or ARB		116 (55.0%)	74 (54.0%)	42 (56.8%)	0.702	
Diuretics		58 (27.5%)	40 (29.2%)	18 (24.3%)	0.449	
ASA		128 (60.7%)	81 (59.1%)	47 (63.5%)	0.533	
Antiarrhythmics		9 (4.3%)	3 (2.2%)	6 (8.1%)	0.069	
EuroSCODE	Add-scale	5.1 ± 3.0	4.8 ± 3.1	5.7 ±2.7	0.041	
EUFOSCORE	Log-scale	3.8 [1.9 - 8.4]	3.1 [1.6 – 7.9]	4.8 [2.3 – 8.6]	0.040	
Value Sungan	No	98/238 (41.2%)	72/154 (46.8%)	26/84 (30.9%)	0.019	
varve Surgery	Yes	140/238 (58.8%)	82/154 (53.3%)	58/84 (69.1%)	0.018	
Pump time (hours)		1.64 ± 0.69	$1.59 \pm 0.71 \qquad 1.74 \pm .64$		0.105	
Cross time (hours)		1.20 ± 0.54	$1.16 \pm 0.57 \qquad 1.29 \pm 0.48$		0.072	
Cardioplegia		232 (97.5%)	148 (96.1%) 84 (100.0%)		0.093	
Blood transfusion		98 (41.2%)	63 (40.9%)	35 (41.7%)	0.910	

Table 1. Baseline characteristics prior to cardiac surgery in relation to Post-Operative Atrial

 Fibrillation.

* P-value for Chi² Test (discrete variables), ANOVA (normally distributed continuous variables) or Kruskal-Wallis (not normally distributed continuous variables)

[†] Prior cardiac surgery \neq coronary bypass + valve surgery + other cardiac surgery, one patient had multiple surgeries.

		All Patients	Non-POAF	POAF	P* univariate	OR (95% CI) multivariable	P†
Fibrosis (%)		8.21 [5.02–12.65]	8.34 [5.28–12.65]	7.55 [4.51–12.50]	0.356	0.995 (0.959-1.032)	0.793
Myocytolysis (%)		3.9 [1.7-8.4]	4.0 [1.8-8.8]	3.7 [1.3–6.9]	0.277	0.974 (0.914-1.039)	0.427
Cardiomyocytes Diameter (µm)		14.8 ± 2.6	14.9 ± 2.6	14.7 ± 2.6	0.559	0.915 (0.810-1.034	0.155
	None	58 (26.1%)	35 (23.3%)	23 (31.9%)		Reference	
Glycogen accumulation	Mild-Moderate	147 (66.2%)	104 (69.3%)	43 (59.7%)	0.346	0.628 (0.312-1.263)	0.192
	Severe	17 (7.7%)	11 (7.3%)	6 (8.3%)		0.865 (0.256-2.921)	0.815
Cx43 Latarization	Little/Absent	30 (12.9%)	17 (11.3%)	13 (16.1%)	0.300	Reference	e
	Present	202 (87.1%)	134 (88.7%)	68 (83.9%)	0.500	0.460 (0.191-1.104)	0.082
	Control-like	139 (59.9%)	97 (64.2%)	42 (51.8%)		Reference	e
Cx43 Junctional	Depressed	77 (33.2%)	46 (30.5%)	31 (38.3%)	0.141	1.503 (0.785-2.877)	0.219
	Heterogeneous	16 (6.9%)	8 (5.3%)	8 (9.9%)	0.	2.174 (0.682-6.932)	0.189
	Normal + Normal	11 (4.8%)	9 (6.0%)	2 (2.5%)		Reference	e
Connexin 43	Normal + Abnormal	127 (64.8%)	87 (64.9%)	40 (64.5%)	0.566	1.139 (0.222-5.840)	0.876
	Abnormal + Abnormal	58 (29.6%)	38 (28.4%)	20 (32.3%)		1.264 (2.33-6.865)	0.786

Table 2. Histological characteristics at the time of cardiac surgery in relation to subsequent risk of Post-Operative Atrial Fibrillation.

* P-value for Chi² (discrete variables), ANOVA (normally distributed continuous variables) or Kruskal-Wallis (not normally distributed continuous variables). † P-value for multivariable Logistic regression adjusted for age, COPD, atrial fibrillation, congestive HF, valvular surgery and log EuroSCORE. Three classes of Connexin 43: normal pattern (patients with control Cx43 junctional and little/absent Cx43 lateralization); normal + abnormal pattern (patients

with control Cx43 junctional and present Cx43 lateralization or patients with depressed Cx43 junctional and little/absent Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and

Fibrosis (%)		Tertile 1 N=79 (2.00-5.99) Median [IQR]	Tertile 2 N=80 (6.00-10.59) Median [IQR]	Tertile 3 N=80 (10.59-65.83) Median [IQR]	P *	Adjusted unit difference Beta (95%CI)**	P†
plasma	NT-proBNP	271 [120-853]	255 [95 - 615]	463 [188 - 888]	0.152	1.50 (-0.74-3.74)	0.187
	Hs cTnT	10.7 [5.9-18.5]	10.8 [6.3 – 21.1]	14.3 [9.0 - 23.8]	0.109	1.71 (-0.87–4.29)	0.193
	Hs-CRP	2.7 [1.0-5.2]	2.1 [1.0 - 5.9]	2.8 [1.1 - 6.9]	0.426	1.00 (-1.16–3.17)	0.363
	F2-isoprostanes	41 [31-52]	40 [31 - 56]	41 [31 - 51]	0.962	-7.71 (-12.84-2.59)	0.003
	F3-isoprostanes	1.6 [0.5-3.3]	1.0 [0.5 – 3.0]	1.5 [0.5 – 4.0]	0.852	1.94 (-0.73–4.60)	0.154
	Isofurans	37 [28-51]	35 [25 - 50]	36 [20 - 53]	0.578	-0.99 (-4.49–2.52)	0.580
e	F2-isoprostanes	1.16 [0.77-2.03]	1.19 [0.95 - 1.96]	1.15 [0.81 - 1.47]	0.297	-1.81 (-6.71–3.10)	0.468
irin	F3-isoprostanes	0.11 [0.05-0.19]	0.12 [0.07 - 0.23]	0.13 [0.07 - 0.19]	0.748	1.69 (-0.82-4.19)	0.185
C	Isofurans	1.79 [1.19-2.60]	1.74 [1.28 - 2.43]	1.80 [1.08 - 2.29]	0.774	-1.11 (-5.65–3.43) rican	0.629
Myocytolysis (%)		Tertile 1 N=74 (0.00 – 2.22) Median [IQR]	Tertile 2 N=74 (2.23 – 6.05) Median [IQR]	Tertile 3 N=74 (6.05 – 26.85) Median [IQR]	P *	Adjusted unit difference Beta (95%CI)**	P†
	NT-proBNP	403 [175-1036]	292 [140 - 951]	203 [69 - 542]	0.012	-2.24 (-3.59—0.90)	0.001
	Hs cTnT	11.3 [6.9-16.6]	10.0 [5.9 - 20.9]	12.5 [7.7 - 21.6]	0.529	-0.08 (-1.67 - 1.51)	0.919
ŝmŝ	Hs-CRP	2.1 [1.0-5.7]	2.9 [0.9 - 7.0]	2.1 [1.0- 5.7]	0.868	0.07 (-1.27 – 1.41)	0.922
olas	F2-isoprostanes	44 [33-55]	39 [30 - 50]	38 [29 - 52]	0.181	-0.68 (-4.36 - 2.99)	0.714
	F3-isoprostanes	1.0 [0.5–4.0]	2.0 [0.5 - 4.0]	1.0 [0.5 - 3.2]	0.144	-0.45 (-2.16 - 1.25)	0.602
	Isofurans	35 [25-54]	38 [23 - 53]	35 [27 - 49]	0.783	0.09 (-2.09 – 2.28)	0.934
e	F2-isoprostanes	1.16 [0.83-1.57]	1.16 [0.80 - 1.79]	1.18 [0.82 - 1.88]	0.859	1.58 (-1.31 – 4.46)	0.281
irin	F3-isoprostanes	0.12 [0.07-0.22]	0.13 [0.06 - 0.21]	0.10 [0.05 - 0.19]	0.404	-1.25 (-2.68 - 0.19)	0.089
C	Isofurans	1.74 [1.19-2.45]	1.88 [1.45 - 2.60]	1.74 [1.11 - 2.29]	0.672	-0.10 (-2.78 – 2.58)	0.942
Cardiomyocyte diameter, µm		Tertile 1 N=74 (8.44 – 13.75) Median [IQR]	Tertile 2 N=74 (13.76 – 15.67) Median [IQR]	Tertile 3 N=74 (15.72 – 23.42) Median [IQR]	P*	Adjusted unit difference Beta (95%CI)**	P†
	NT-proBNP	246 [124 - 861]	271 [120 - 888]	336 [158 - 713]	0.860	-0.36 (-1.08 - 0.36)	0.325
-	Hs cTnT	10.1 [6.2 - 16.1]	9.3 [5.7 - 19.9]	15.6 [8.3 - 22.4]	0.016	0.78 (-0.04 - 1.60)	0.063
sme	Hs-CRP	1.3 [0.8 - 6.6]	2.1 [1.1 - 5.2]	3.1 [1.7 - 9.2]	0.030	0.68 (-0.02 – 1.37)	0.056
pla	F2-isoprostanes	42 [32 - 63]	40 [31 - 49]	40 [31 - 51]	0.337	-0.35 (-2.27 – 1.56)	0.715
_	F3-isoprostanes	2.0 [0.5 - 4.0]	1.0 [0.5 – 3.0]	1.0 [0.5 - 3.3]	0.214	-0.26 (-1.14 – 0.63)	0.568
	Isofurans	36 [28 - 54]	33 [21 - 50]	37 [26 - 50]	0.428	-0.22 (-1.36 - 0.91)	0.697
e	F2-isoprostanes	1.18 [0.84 - 1.59]	1.14 [0.75 - 1.75]	1.29 [0.94 - 1.92]	0.261	0.94 (-0.60 - 2.48)	0.230
urin∈	F3-isoprostanes	0.13 [0.06 - 0.24]	0.09 [0.05 - 0.15]	0.14 [0.09 - 0.22]	0.056	0.58 (-0.22 – 1.37)	0.153
	Isofurans	1.81 [1.20 - 2.56]	1.56 [1.16 - 2.17]	2.03 [1.46 - 2.61]	0.136	0.55 (-0.88 – 1.99)	0.446

Table 3. Pre-operative levels of biomarkers of cardiac strain, inflammation, and oxidative stress in relation to tertiles of continuous histopathological features of the right atrium at the time of cardiac surgery.

* P for Kruskall Wallis test. † All biomarkers were log-transformed and included in linear regression analysis with histopathological feature as dependent adjusted for age, COPD, history of atrial fibrillation, heart failure, valve surgery and EuroSCORE log. Beta represents the % unit difference in histopathologic features for continuous outcomes for a doubling in levels of each biomarker.

Table 4. Pre-operative levels of biomarkers of cardiac strain, inflammation, and oxidative stress in relation to categories of histopathological features of the right atrium at the time of cardiac surgery.

						Adjusted OR (95%	
Glyc	ogen accumulation	None N=58	Mild-Moderate N=147	Severe N=17	P*	CI) Mild- Moderate+Severe vs	P†
	NT-proBNP	308 [132 - 674]	266 [123 - 853]	625 [77 – 1132]	0.966	0.85 (0.47-1.54)	0.600
	HscTnT	9.3 [5.7 - 14.5]	14.0 [7.4 - 23.2]	9.0 [5.2 - 45.9]	0.023	2.46 (1.02-5.92)	0.045
na	Hs-CRP	1.4 [0.9 - 5.4]	2.7 [1.1 - 6.7]	2.6 [1.6 - 6.6]	0.118	1.76 (0.96-3.22)	0.068
plasr	F2-isoprostanes	44 [32 - 55]	39 [30 - 51]	42 [34 - 52]	0.565	0.36 (0.07-1.80)	0.214
	F3-isoprostanes	1.0 [0.5 – 4.0]	1.6 [0.5 - 3.6]	2.0 [0.5 - 4.0]	0.865	1.22 (0.59-2.55)	0.591
	Isofurans	34 [20 - 59]	36 [27 - 50]	45 [32 - 53]	0.371	1.57 (0.59-4.22)	0.371
	F2-isoprostanes	1.18 [0.88 - 1.59]	1.18 [0.81 - 1.66]	1.12 [0.88 - 2.02]	0.960	0.78 (0.22-2.76)	0.695
ine	F3-isoprostanes	0.13 [0.08 - 0.24]	0.11 [0.05 - 0.19]	0.07 [0.01 - 0.12]	0.054	0.29 (0.13-0.67)	0.004
m	Isofurans	1.77 [1.36 - 2.25]	1.82 [1.17 - 2.53]	1.75 [1.48 - 2.47]	0.908	1.05 (0.32-3.44)	0.940
C-41	T - 4 4 ¹	Little / absent	Pres	ent	D¥	Adjusted OR (95%	D4
CX43	Laterization	N=30	N=2	202	P*	CI)	Pf
	NT-proBNP	459 [237 - 760]	267 [125	5 - 847]	0.368	0.78 (0.21 -1.11)	0.087
ਰ	Hs cTnT	12.7 [5.9 - 15.7]	11.3 [6.9	- 21.8]	0.269	1.47 (0.51-4.21)	0.471
sma	Hs-CRP	3.1 [1.0 - 5.7]	2.3 [1.0	9 - 6.3]	0.977	1.05 (0.47-2.35)	0.904
pla	F2-isoprostanes	39 [33 - 53]	41 [31	- 52]	0.683	0.27 (0.04-1.78)	0.174
	F3-isoprostanes	3.0 [0.5 – 4.0]	1.2 [0.5	5 - 3.2]	0.191	0.67 (0.26-1.70)	0.400
	Isofurans	39 [28 - 59]	36 [25	- 50]	0.233	0.42 (0.13-1.36)	0.145
е	F2-isoprostanes	1.21 [0.8 - 2.02]	1.18 [0.83 - 1.64]		0.751	1.13 (0.20-6.44)	0.888
urin	F3-isoprostanes	0.19 [0.1 - 0.34]	0.12 [0.06 - 0.19]		0.039	0.31 (0.08-1.16)	0.081
Isofurans		1.61 [1.01 - 2.60]	1.78 [1.23 - 2.30]		0.330	2.68 (0.59-12.27)	0.205
		Control like	Depressed	Heterogeneous		Adjusted OR (95%	
Cx43	Junctional	Control like N=139	Depressed N=77	Heterogeneous N=16	P*	Adjusted OR (95% CI) Depressed vs Control-like	₽†
Cx43	Junctional NT-proBNP	Control like N=139 273 [108 - 759]	Depressed N=77 323 [158 - 875]	Heterogeneous N=16 304 [180 – 1024]	P *	Adjusted OR (95% CI) Depressed vs Control-like 1.41 (0.77-2.56)	P †
Cx43	B Junctional NT-proBNP Hs cTnT	Control like N=139 273 [108 - 759] 12.7 [7.0 - 22.4]	Depressed N=77 323 [158 - 875] 10.9 [7.1 - 20.3]	Heterogeneous N=16 304 [180 – 1024] 5.8 [5.0 - 16.5]	P * 0.406 0.351	Adjusted OR (95% CI) Depressed vs Control-like 1.41 (0.77-2.56) 0.90 (0.46-1.78)	P † 0.262 0.767
Cx43	B Junctional NT-proBNP Hs cTnT Hs-CRP	Control like N=139 273 [108 - 759] 12.7 [7.0 - 22.4] 2.6 [1.1 - 7.0]	Depressed N=77 323 [158 - 875] 10.9 [7.1 - 20.3] 2.6 [1.0 - 6.5]	Heterogeneous N=16 304 [180 - 1024] 5.8 [5.0 - 16.5] 1.4 [0.7 - 2.4]	P * 0.406 0.351 0.142	Adjusted OR (95% CI) Depressed vs Control-like 1.41 (0.77-2.56) 0.90 (0.46-1.78) 0.85 (0.48-1.48)	P † 0.262 0.767 0.558
Cx43 emseld	B Junctional NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes	Control like N=139 273 [108 - 759] 12.7 [7.0 - 22.4] 2.6 [1.1 - 7.0] 40 [31 - 51]	Depressed N=77 323 [158 - 875] 10.9 [7.1 - 20.3] 2.6 [1.0 - 6.5] 41 [31 - 57]	Heterogeneous N=16 304 [180 - 1024] 5.8 [5.0 - 16.5] 1.4 [0.7 - 2.4] 41 [30 - 44]	P* 0.406 0.351 0.142 0.479	Adjusted OR (95% CI) Depressed vs Control-like 1.41 (0.77-2.56) 0.90 (0.46-1.78) 0.85 (0.48-1.48) 2.45 (0.56-10.80)	P † 0.262 0.767 0.558 0.235
Cx43 emseld	B Junctional NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes F3-isoprostanes	Control like N=139 273 [108 - 759] 12.7 [7.0 - 22.4] 2.6 [1.1 - 7.0] 40 [31 - 51] 1.0 [0.5 - 4.0]	Depressed N=77 323 [158 - 875] 10.9 [7.1 - 20.3] 2.6 [1.0 - 6.5] 41 [31 - 57] 2.0 [0.5 - 4.0]	Heterogeneous N=16 304 [180 - 1024] 5.8 [5.0 - 16.5] 1.4 [0.7 - 2.4] 41 [30 - 44] 0.5 [0.5 - 3.0]	P * 0.406 0.351 0.142 0.479 0.125	Adjusted OR (95% CI) Depressed vs Control-like 1.41 (0.77-2.56) 0.90 (0.46-1.78) 0.85 (0.48-1.48) 2.45 (0.56-10.80) 1.50 (0.75-3.02)	P† 0.262 0.767 0.558 0.235 0.251
Cx43 emseld	B Junctional NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes F3-isoprostanes Isofurans	Control like N=139 273 [108 - 759] 12.7 [7.0 - 22.4] 2.6 [1.1 - 7.0] 40 [31 - 51] 1.0 [0.5 - 4.0] 36 [23 - 50]	Depressed N=77 323 [158 - 875] 10.9 [7.1 - 20.3] 2.6 [1.0 - 6.5] 41 [31 - 57] 2.0 [0.5 - 4.0] 36 [29 - 56]	Heterogeneous N=16 304 [180 - 1024] 5.8 [5.0 - 16.5] 1.4 [0.7 - 2.4] 41 [30 - 44] 0.5 [0.5 - 3.0] 28 [22 - 43]	P* 0.406 0.351 0.142 0.479 0.125 0.371	Adjusted OR (95% CI) Depressed vs Control-like 1.41 (0.77-2.56) 0.90 (0.46-1.78) 0.85 (0.48-1.48) 2.45 (0.56-10.80) 1.50 (0.75-3.02) 1.30 (0.49-3.42)	P † 0.262 0.767 0.558 0.235 0.251 0.598
Cx43 plasma	B Junctional NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes F3-isoprostanes Isofurans F2-isoprostanes	Control like N=139 273 [108 - 759] 12.7 [7.0 - 22.4] 2.6 [1.1 - 7.0] 40 [31 - 51] 1.0 [0.5 - 4.0] 36 [23 - 50] 1.23 [0.94 - 1.72]	Depressed N=77 323 [158 - 875] 10.9 [7.1 - 20.3] 2.6 [1.0 - 6.5] 41 [31 - 57] 2.0 [0.5 - 4.0] 36 [29 - 56] 1.08 [0.64 - 1.94]	Heterogeneous N=16 304 [180 - 1024] 5.8 [5.0 - 16.5] 1.4 [0.7 - 2.4] 41 [30 - 44] 0.5 [0.5 - 3.0] 28 [22 - 43] 0.99 [0.55 - 1.52]	P* 0.406 0.351 0.142 0.479 0.125 0.371 0.133	Adjusted OR (95% CI) Depressed vs Control-like 1.41 (0.77-2.56) 0.90 (0.46-1.78) 0.85 (0.48-1.48) 2.45 (0.56-10.80) 1.50 (0.75-3.02) 1.30 (0.49-3.42) 0.32 (0.09-1.17)	P† 0.262 0.767 0.558 0.235 0.251 0.598 0.084
Cx43 plasma	B Junctional NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes F3-isoprostanes Isofurans F2-isoprostanes F3-isoprostanes	Control like N=139 273 [108 - 759] 12.7 [7.0 - 22.4] 2.6 [1.1 - 7.0] 40 [31 - 51] 1.0 [0.5 - 4.0] 36 [23 - 50] 1.23 [0.94 - 1.72] 0.12 [0.07 - 0.20]	Depressed N=77 323 [158 - 875] 10.9 [7.1 - 20.3] 2.6 [1.0 - 6.5] 41 [31 - 57] 2.0 [0.5 - 4.0] 36 [29 - 56] 1.08 [0.64 - 1.94] 0.12 [0.06 - 0.24]	Heterogeneous N=16 304 [180 - 1024] 5.8 [5.0 - 16.5] 1.4 [0.7 - 2.4] 41 [30 - 44] 0.5 [0.5 - 3.0] 28 [22 - 43] 0.99 [0.55 - 1.52] 0.12 [0.07 - 0.16]	P* 0.406 0.351 0.142 0.479 0.125 0.371 0.133 0.943	Adjusted OR (95% CI) Depressed vs Control-like 1.41 (0.77-2.56) 0.90 (0.46-1.78) 0.85 (0.48-1.48) 2.45 (0.56-10.80) 1.50 (0.75-3.02) 1.30 (0.49-3.42) 0.32 (0.09-1.17) 0.95 (0.49-1.86)	P† 0.262 0.767 0.558 0.235 0.251 0.598 0.084 0.889
Cx43	Junctional NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes F3-isoprostanes Isofurans F2-isoprostanes F3-isoprostanes F3-isoprostanes Isofurans	Control like N=139 273 [108 - 759] 12.7 [7.0 - 22.4] 2.6 [1.1 - 7.0] 40 [31 - 51] 1.0 [0.5 - 4.0] 36 [23 - 50] 1.23 [0.94 - 1.72] 0.12 [0.07 - 0.20] 1.86 [1.31 - 2.31]	Depressed N=77 323 [158 - 875] 10.9 [7.1 - 20.3] 2.6 [1.0 - 6.5] 41 [31 - 57] 2.0 [0.5 - 4.0] 36 [29 - 56] 1.08 [0.64 - 1.94] 0.12 [0.06 - 0.24] 1.73 [1.00 - 2.74]	Heterogeneous N=16 304 [180 - 1024] 5.8 [5.0 - 16.5] 1.4 [0.7 - 2.4] 41 [30 - 44] 0.5 [0.5 - 3.0] 28 [22 - 43] 0.99 [0.55 - 1.52] 0.12 [0.07 - 0.16] 1.46 [1.05 - 1.77]	P* 0.406 0.351 0.142 0.479 0.125 0.371 0.133 0.943 0.211	Adjusted OR (95% CI) Depressed vs Control-like 1.41 (0.77-2.56) 0.90 (0.46-1.78) 0.85 (0.48-1.48) 2.45 (0.56-10.80) 1.50 (0.75-3.02) 1.30 (0.49-3.42) 0.32 (0.09-1.17) 0.95 (0.49-1.86) 0.81 (0.25-2.58)	P† 0.262 0.767 0.558 0.235 0.251 0.598 0.084 0.889 0.717
Cx43	B Junctional NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes F3-isoprostanes Isofurans F2-isoprostanes F3-isoprostanes F3-isoprostanes Isofurans	Control like N=139 273 [108 - 759] 12.7 [7.0 - 22.4] 2.6 [1.1 - 7.0] 40 [31 - 51] 1.0 [0.5 - 4.0] 36 [23 - 50] 1.23 [0.94 - 1.72] 0.12 [0.07 - 0.20] 1.86 [1.31 - 2.31]	Depressed N=77 323 [158 - 875] 10.9 [7.1 - 20.3] 2.6 [1.0 - 6.5] 41 [31 - 57] 2.0 [0.5 - 4.0] 36 [29 - 56] 1.08 [0.64 - 1.94] 0.12 [0.06 - 0.24] 1.73 [1.00 - 2.74]	Heterogeneous N=16 304 [180 - 1024] 5.8 [5.0 - 16.5] 1.4 [0.7 - 2.4] 41 [30 - 44] 0.5 [0.5 - 3.0] 28 [22 - 43] 0.99 [0.55 - 1.52] 0.12 [0.07 - 0.16] 1.46 [1.05 - 1.77]	P* 0.406 0.351 0.142 0.479 0.125 0.371 0.133 0.943 0.211	Adjusted OR (95% CI) Depressed vs Control-like 1.41 (0.77-2.56) 0.90 (0.46-1.78) 0.85 (0.48-1.48) 2.45 (0.56-10.80) 1.50 (0.75-3.02) 1.30 (0.49-3.42) 0.32 (0.09-1.17) 0.95 (0.49-1.86) 0.81 (0.25-2.58) Adjusted OR (95%	P† 0.262 0.767 0.558 0.235 0.251 0.598 0.084 0.889 0.717
Cx43 nuine Conr	Junctional NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes F3-isoprostanes Isofurans F2-isoprostanes F3-isoprostanes F3-isoprostanes Isofurans	Control like N=139 273 [108 - 759] 12.7 [7.0 - 22.4] 2.6 [1.1 - 7.0] 40 [31 - 51] 1.0 [0.5 - 4.0] 36 [23 - 50] 1.23 [0.94 - 1.72] 0.12 [0.07 - 0.20] 1.86 [1.31 - 2.31] Normal+normal	Depressed N=77 323 [158 - 875] 10.9 [7.1 - 20.3] 2.6 [1.0 - 6.5] 41 [31 - 57] 2.0 [0.5 - 4.0] 36 [29 - 56] 1.08 [0.64 - 1.94] 0.12 [0.06 - 0.24] 1.73 [1.00 - 2.74] Normal + Abnor	Heterogeneous N=16 304 [180 - 1024] 5.8 [5.0 - 16.5] 1.4 [0.7 - 2.4] 41 [30 - 44] 0.5 [0.5 - 3.0] 28 [22 - 43] 0.99 [0.55 - 1.52] 0.12 [0.07 - 0.16] 1.46 [1.05 - 1.77]	P* 0.406 0.351 0.142 0.479 0.125 0.371 0.133 0.943 0.211 P*	Adjusted OR (95% CI) Depressed vs Control-like 1.41 (0.77-2.56) 0.90 (0.46-1.78) 0.85 (0.48-1.48) 2.45 (0.56-10.80) 1.50 (0.75-3.02) 1.30 (0.49-3.42) 0.32 (0.09-1.17) 0.95 (0.49-1.86) 0.81 (0.25-2.58) Adjusted OR (95% CI)	P† 0.262 0.767 0.558 0.235 0.251 0.598 0.084 0.889 0.717 P†
Cx43 nuine Conr	Junctional NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes F3-isoprostanes Isofurans F2-isoprostanes F3-isoprostanes F3-isoprostanes Isofurans	Control like N=139 273 [108 - 759] 12.7 [7.0 - 22.4] 2.6 [1.1 - 7.0] 40 [31 - 51] 1.0 [0.5 - 4.0] 36 [23 - 50] 1.23 [0.94 - 1.72] 0.12 [0.07 - 0.20] 1.86 [1.31 - 2.31] Normal+normal N=11	Depressed N=77 323 [158 - 875] 10.9 [7.1 - 20.3] 2.6 [1.0 - 6.5] 41 [31 - 57] 2.0 [0.5 - 4.0] 36 [29 - 56] 1.08 [0.64 - 1.94] 0.12 [0.06 - 0.24] 1.73 [1.00 - 2.74] Normal + Abnor N=146	Heterogeneous N=16 304 [180 - 1024] 5.8 [5.0 - 16.5] 1.4 [0.7 - 2.4] 41 [30 - 44] 0.5 [0.5 - 3.0] 28 [22 - 43] 0.99 [0.55 - 1.52] 0.12 [0.07 - 0.16] 1.46 [1.05 - 1.77] Ab + Abnormal N=74	P* 0.406 0.351 0.142 0.479 0.125 0.371 0.133 0.943 0.211 P*	Adjusted OR (95% CI) Depressed vs Control-like 1.41 (0.77-2.56) 0.90 (0.46-1.78) 0.85 (0.48-1.48) 2.45 (0.56-10.80) 1.50 (0.75-3.02) 1.30 (0.49-3.42) 0.32 (0.09-1.17) 0.95 (0.49-1.86) 0.81 (0.25-2.58) Adjusted OR (95% CI) Normal+Abnor vs	P† 0.262 0.767 0.558 0.235 0.251 0.598 0.084 0.889 0.717
Cx43 enrine Cour	3 Junctional NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes Isofurans hexin 43	Control like N=139 273 [108 - 759] 12.7 [7.0 - 22.4] 2.6 [1.1 - 7.0] 40 [31 - 51] 1.0 [0.5 - 4.0] 36 [23 - 50] 1.23 [0.94 - 1.72] 0.12 [0.07 - 0.20] 1.86 [1.31 - 2.31] Normal+normal N=11	Depressed N=77 323 [158 - 875] 10.9 [7.1 - 20.3] 2.6 [1.0 - 6.5] 41 [31 - 57] 2.0 [0.5 - 4.0] 36 [29 - 56] 1.08 [0.64 - 1.94] 0.12 [0.06 - 0.24] 1.73 [1.00 - 2.74] Normal + Abnor N=146 255 [115 - 791]	Heterogeneous N=16 304 [180 - 1024] 5.8 [5.0 - 16.5] 1.4 [0.7 - 2.4] 41 [30 - 44] 0.5 [0.5 - 3.0] 28 [22 - 43] 0.99 [0.55 - 1.52] 0.12 [0.07 - 0.16] 1.46 [1.05 - 1.77] Ab + Abnormal N=74 308 [174 - 1012]	P* 0.406 0.351 0.142 0.479 0.125 0.371 0.133 0.943 0.211 P* 0.408	Adjusted OR (95% CI) Depressed vs Control-like 1.41 (0.77-2.56) 0.90 (0.46-1.78) 0.85 (0.48-1.48) 2.45 (0.56-10.80) 1.50 (0.75-3.02) 1.30 (0.49-3.42) 0.32 (0.09-1.17) 0.95 (0.49-1.86) 0.81 (0.25-2.58) Adjusted OR (95% CI) Normal+Abnor vs Ab+Abnormal 1.41 (0.72-2.75)	P† 0.262 0.767 0.558 0.235 0.251 0.598 0.084 0.889 0.717 P† 0.310
Cx43 enrine Cour	Junctional NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes Isofurans rexin 43 NT-proBNP Hs cTnT	Control like N=139 273 [108 - 759] 12.7 [7.0 - 22.4] 2.6 [1.1 - 7.0] 40 [31 - 51] 1.0 [0.5 - 4.0] 36 [23 - 50] 1.23 [0.94 - 1.72] 0.12 [0.07 - 0.20] 1.86 [1.31 - 2.31] Normal+normal N=11 459 [273 - 674] 7 6 [4 9 - 16 0]	Depressed N=77 323 [158 - 875] 10.9 [7.1 - 20.3] 2.6 [1.0 - 6.5] 41 [31 - 57] 2.0 [0.5 - 4.0] 36 [29 - 56] 1.08 [0.64 - 1.94] 0.12 [0.06 - 0.24] 1.73 [1.00 - 2.74] Normal + Abnor N=146 255 [115 - 791] 12 7 [7 0 - 21 6]	Heterogeneous N=16 304 [180 - 1024] 5.8 [5.0 - 16.5] 1.4 [0.7 - 2.4] 41 [30 - 44] 0.5 [0.5 - 3.0] 28 [22 - 43] 0.99 [0.55 - 1.52] 0.12 [0.07 - 0.16] 1.46 [1.05 - 1.77] Ab + Abnormal N=74 308 [174 - 1012] 10.1 [6.4 - 20.7]	P* 0.406 0.351 0.142 0.479 0.125 0.371 0.133 0.943 0.211 P* 0.408 0.564	Adjusted OR (95% CI) Depressed vs Control-like 1.41 (0.77-2.56) 0.90 (0.46-1.78) 0.85 (0.48-1.48) 2.45 (0.56-10.80) 1.50 (0.75-3.02) 1.30 (0.49-3.42) 0.32 (0.09-1.17) 0.95 (0.49-1.86) 0.81 (0.25-2.58) Adjusted OR (95% CI) Normal+Abnor vs Ab+Abnormal 1.41 (0.72-2.75) 1.11 (0.51-2.45)	P† 0.262 0.767 0.558 0.235 0.251 0.598 0.084 0.889 0.717 P† 0.310 0.788
cx43 crine plasma	Junctional NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes F3-isoprostanes Isofurans F3-isoprostanes Isofurans rexin 43 NT-proBNP Hs cTnT Hs-CRP	Control like N=139 273 [108 - 759] 12.7 [7.0 - 22.4] 2.6 [1.1 - 7.0] 40 [31 - 51] 1.0 [0.5 - 4.0] 36 [23 - 50] 1.23 [0.94 - 1.72] 0.12 [0.07 - 0.20] 1.86 [1.31 - 2.31] Normal+normal N=11 459 [273 - 674] 7.6 [4.9 - 16.0] 5.2 [0.7 - 5.7]	Depressed N=77 323 [158 - 875] 10.9 [7.1 - 20.3] 2.6 [1.0 - 6.5] 41 [31 - 57] 2.0 [0.5 - 4.0] 36 [29 - 56] 1.08 [0.64 - 1.94] 0.12 [0.06 - 0.24] 1.73 [1.00 - 2.74] Normal + Abnor N=146 255 [115 - 791] 12.7 [7.0 - 21.6] 2.3 [1.1 - 7 0]	Heterogeneous N=16 304 [180 - 1024] 5.8 [5.0 - 16.5] 1.4 [0.7 - 2.4] 41 [30 - 44] 0.5 [0.5 - 3.0] 28 [22 - 43] 0.99 [0.55 - 1.52] 0.12 [0.07 - 0.16] 1.46 [1.05 - 1.77] Ab + Abnormal N=74 308 [174 - 1012] 10.1 [6.4 - 20.7] 2.3 [1.0 - 5 2]	P* 0.406 0.351 0.142 0.479 0.125 0.371 0.133 0.943 0.211 P* 0.408 0.564 0.720	Adjusted OR (95% CI) Depressed vs Control-like 1.41 (0.77-2.56) 0.90 (0.46-1.78) 0.85 (0.48-1.48) 2.45 (0.56-10.80) 1.50 (0.75-3.02) 1.30 (0.49-3.42) 0.32 (0.09-1.17) 0.95 (0.49-1.86) 0.81 (0.25-2.58) Adjusted OR (95% CI) Normal+Abnor vs Ab+Abnormal 1.41 (0.72-2.75) 1.11 (0.51-2.45) 0.88 (048-1.64)	P† 0.262 0.767 0.558 0.235 0.251 0.598 0.084 0.889 0.717 P† 0.310 0.788 0.696
Cx43 email Court court court	B Junctional NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes F3-isoprostanes F3-isoprostanes Isofurans F2-isoprostanes Isofurans Rexin 43 NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes	Control like N=139 273 [108 - 759] 12.7 [7.0 - 22.4] 2.6 [1.1 - 7.0] 40 [31 - 51] 1.0 [0.5 - 4.0] 36 [23 - 50] 1.23 [0.94 - 1.72] 0.12 [0.07 - 0.20] 1.86 [1.31 - 2.31] Normal+normal N=11 459 [273 - 674] 7.6 [4.9 - 16.0] 5.2 [0.7 - 5.7] 39 [35 - 51]	Depressed N=77 323 [158 - 875] 10.9 [7.1 - 20.3] 2.6 [1.0 - 6.5] 41 [31 - 57] 2.0 [0.5 - 4.0] 36 [29 - 56] 1.08 [0.64 - 1.94] 0.12 [0.06 - 0.24] 1.73 [1.00 - 2.74] Normal + Abnor N=146 255 [115 - 791] 12.7 [7.0 - 21.6] 2.3 [1.1 - 7.0] 41 [31 - 54]	Heterogeneous N=16 304 [180 - 1024] 5.8 [5.0 - 16.5] 1.4 [0.7 - 2.4] 41 [30 - 44] 0.5 [0.5 - 3.0] 28 [22 - 43] 0.99 [0.55 - 1.52] 0.12 [0.07 - 0.16] 1.46 [1.05 - 1.77] Ab + Abnormal N=74 308 [174 - 1012] 10.1 [6.4 - 20.7] 2.3 [1.0 - 5.2] 42 [31 - 52]	P* 0.406 0.351 0.142 0.479 0.125 0.371 0.133 0.943 0.211 P* 0.408 0.564 0.720 0.928	Adjusted OR (95% CI) Depressed vs Control-like 1.41 (0.77-2.56) 0.90 (0.46-1.78) 0.85 (0.48-1.48) 2.45 (0.56-10.80) 1.50 (0.75-3.02) 1.30 (0.49-3.42) 0.32 (0.09-1.17) 0.95 (0.49-1.86) 0.81 (0.25-2.58) Adjusted OR (95% CI) Normal+Abnor vs Ab+Abnormal 1.41 (0.72-2.75) 1.11 (0.51-2.45) 0.88 (.048-1.64) 2.34 (0.40-13.86)	P† 0.262 0.767 0.558 0.235 0.251 0.598 0.084 0.889 0.717 P† 0.310 0.788 0.696 0.349
Cx43 enrine Court Court	B Junctional NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes Isofurans F2-isoprostanes Isofurans Rexin 43 NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes	Control like N=139 273 [108 - 759] 12.7 [7.0 - 22.4] 2.6 [1.1 - 7.0] 40 [31 - 51] 1.0 [0.5 - 4.0] 36 [23 - 50] 1.23 [0.94 - 1.72] 0.12 [0.07 - 0.20] 1.86 [1.31 - 2.31] Normal+normal N=11 459 [273 - 674] 7.6 [4.9 - 16.0] 5.2 [0.7 - 5.7] 39 [35 - 51] 3.0 [0.5 - 4.0]	Depressed N=77 323 [158 - 875] 10.9 [7.1 - 20.3] 2.6 [1.0 - 6.5] 41 [31 - 57] 2.0 [0.5 - 4.0] 36 [29 - 56] 1.08 [0.64 - 1.94] 0.12 [0.06 - 0.24] 1.73 [1.00 - 2.74] Normal + Abnor N=146 255 [115 - 791] 12.7 [7.0 - 21.6] 2.3 [1.1 - 7.0] 41 [31 - 54] 1.0 [0.5 - 4.0]	Heterogeneous N=16 304 [180 - 1024] 5.8 [5.0 - 16.5] 1.4 [0.7 - 2.4] 41 [30 - 44] 0.5 [0.5 - 3.0] 28 [22 - 43] 0.99 [0.55 - 1.52] 0.12 [0.07 - 0.16] 1.46 [1.05 - 1.77] Ab + Abnormal N=74 308 [174 - 1012] 10.1 [6.4 - 20.7] 2.3 [1.0 - 5.2] 42 [31 - 52] 2.0 [0.5 - 3 3]	P* 0.406 0.351 0.142 0.479 0.125 0.371 0.133 0.943 0.211 P* 0.408 0.564 0.720 0.928 0.764	Adjusted OR (95% CI) Depressed vs Control-like 1.41 (0.77-2.56) 0.90 (0.46-1.78) 0.85 (0.48-1.48) 2.45 (0.56-10.80) 1.50 (0.75-3.02) 1.30 (0.49-3.42) 0.32 (0.09-1.17) 0.95 (0.49-1.86) 0.81 (0.25-2.58) Adjusted OR (95% CI) Normal+Abnor vs Ab+Abnormal 1.41 (0.72-2.75) 1.11 (0.51-2.45) 0.88 (.048-1.64) 2.34 (0.40-13.86) 1.61 (0.74-3.48)	P† 0.262 0.767 0.558 0.235 0.251 0.598 0.084 0.889 0.717 P† 0.310 0.788 0.696 0.349 0.231
Cx43 anine plasma	B Junctional NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes F3-isoprostanes Isofurans F2-isoprostanes Isofurans NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes Isofurans Isofurans	Control like N=139 273 [108 - 759] 12.7 [7.0 - 22.4] 2.6 [1.1 - 7.0] 40 [31 - 51] 1.0 [0.5 - 4.0] 36 [23 - 50] 1.23 [0.94 - 1.72] 0.12 [0.07 - 0.20] 1.86 [1.31 - 2.31] Normal+normal N=11 459 [273 - 674] 7.6 [4.9 - 16.0] 5.2 [0.7 - 5.7] 39 [35 - 51] 3.0 [0.5 - 4.0] 35 [20 - 55]	Depressed N=77 323 [158 - 875] 10.9 [7.1 - 20.3] 2.6 [1.0 - 6.5] 41 [31 - 57] 2.0 [0.5 - 4.0] 36 [29 - 56] 1.08 [0.64 - 1.94] 0.12 [0.06 - 0.24] 1.73 [1.00 - 2.74] Normal + Abnor N=146 255 [115 - 791] 12.7 [7.0 - 21.6] 2.3 [1.1 - 7.0] 41 [31 - 54] 1.0 [0.5 - 4.0] 37 [25 - 53]	Heterogeneous N=16 304 [180 - 1024] 5.8 [5.0 - 16.5] 1.4 [0.7 - 2.4] 41 [30 - 44] 0.5 [0.5 - 3.0] 28 [22 - 43] 0.99 [0.55 - 1.52] 0.12 [0.07 - 0.16] 1.46 [1.05 - 1.77] Ab + Abnormal N=74 308 [174 - 1012] 10.1 [6.4 - 20.7] 2.3 [1.0 - 5.2] 42 [31 - 52] 2.0 [0.5 - 3.3] 35 [25 - 51]	P* 0.406 0.351 0.142 0.479 0.125 0.371 0.133 0.943 0.211 P* 0.408 0.564 0.720 0.928 0.764 0.872	Adjusted OR (95% CI) Depressed vs Control-like 1.41 (0.77-2.56) 0.90 (0.46-1.78) 0.85 (0.48-1.48) 2.45 (0.56-10.80) 1.50 (0.75-3.02) 1.30 (0.49-3.42) 0.32 (0.09-1.17) 0.95 (0.49-1.86) 0.81 (0.25-2.58) Adjusted OR (95% CI) Normal+Abnor vs Ab+Abnormal 1.41 (0.72-2.75) 1.11 (0.51-2.45) 0.88 (.048-1.64) 2.34 (0.40-13.86) 1.61 (0.74-3.48) 1.11 (0.36-3.40)	P† 0.262 0.767 0.558 0.235 0.251 0.598 0.084 0.889 0.717 P† 0.310 0.788 0.696 0.349 0.231 0.862
Cx43 blasma Dasma	B Junctional NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes F3-isoprostanes Isofurans F2-isoprostanes Isofurans Rexin 43 NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes Isofurans F3-isoprostanes F3-isoprosta	$\begin{tabular}{ c c c c c c } \hline Control like \\ N=139 \\ \hline $273 [108 - 759] \\ 12.7 [7.0 - 22.4] \\ 2.6 [1.1 - 7.0] \\ 40 [31 - 51] \\ 1.0 [0.5 - 4.0] \\ 36 [23 - 50] \\ 1.23 [0.94 - 1.72] \\ 0.12 [0.07 - 0.20] \\ 1.86 [1.31 - 2.31] \\ \hline $1.23 [0.94 - 1.72] \\ 0.12 [0.07 - 0.20] \\ 1.86 [1.31 - 2.31] \\ \hline $1.23 [0.94 - 1.72] \\ 0.12 [0.07 - 0.20] \\ 1.86 [1.31 - 2.31] \\ \hline $1.23 [0.74 - 1.72] \\ 0.12 [0.07 - 0.20] \\ 1.86 [1.31 - 2.31] \\ \hline $1.23 [0.94 - 1.72] \\ 0.12 [0.07 - 0.20] \\ 1.86 [1.31 - 2.31] \\ \hline $1.23 [0.75 - 0.20] \\ 1.85 [1.21 - 2.02] \\ 1.55 [1.21 - 2.02] \\ \hline \end{tabular}$	Depressed N=77 323 [158 - 875] 10.9 [7.1 - 20.3] 2.6 [1.0 - 6.5] 41 [31 - 57] 2.0 [0.5 - 4.0] 36 [29 - 56] 1.08 [0.64 - 1.94] 0.12 [0.06 - 0.24] 1.73 [1.00 - 2.74] Normal + Abnor N=146 255 [115 - 791] 12.7 [7.0 - 21.6] 2.3 [1.1 - 7.0] 41 [31 - 54] 1.0 [0.5 - 4.0] 37 [25 - 53] 1.18 [0.85 - 1 58]	Heterogeneous N=16 304 [180 - 1024] 5.8 [5.0 - 16.5] 1.4 [0.7 - 2.4] 41 [30 - 44] 0.5 [0.5 - 3.0] 28 [22 - 43] 0.99 [0.55 - 1.52] 0.12 [0.07 - 0.16] 1.46 [1.05 - 1.77] Ab + Abnormal N=74 308 [174 - 1012] 10.1 [6.4 - 20.7] 2.3 [1.0 - 5.2] 42 [31 - 52] 2.0 [0.5 - 3.3] 35 [25 - 51] 1.11 [0.69 - 1 75]	P* 0.406 0.351 0.142 0.479 0.125 0.371 0.133 0.943 0.211 P* 0.408 0.564 0.720 0.928 0.764 0.872 0.131	Adjusted OR (95% CI) Depressed vs Control-like 1.41 (0.77-2.56) 0.90 (0.46-1.78) 0.85 (0.48-1.48) 2.45 (0.56-10.80) 1.50 (0.75-3.02) 1.30 (0.49-3.42) 0.32 (0.09-1.17) 0.95 (0.49-1.86) 0.81 (0.25-2.58) Adjusted OR (95% CI) Normal+Abnor vs Ab+Abnormal 1.41 (0.72-2.75) 1.11 (0.51-2.45) 0.88 (.048-1.64) 2.34 (0.40-13.86) 1.61 (0.74-3.48) 1.11 (0.36-3.40) 0.46 (0.11-1 97)	P† 0.262 0.767 0.558 0.235 0.251 0.598 0.084 0.889 0.717 P† 0.310 0.788 0.696 0.349 0.231 0.862 0.296
cx43 plasma plasma	B Junctional NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes F3-isoprostanes Isofurans F2-isoprostanes Isofurans Rexin 43 NT-proBNP Hs cTnT Hs-CRP F2-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes F3-isoprostanes	Control like N=139 $273 [108 - 759]$ $12.7 [7.0 - 22.4]$ $2.6 [1.1 - 7.0]$ $40 [31 - 51]$ $1.0 [0.5 - 4.0]$ $36 [23 - 50]$ $1.23 [0.94 - 1.72]$ $0.12 [0.07 - 0.20]$ $1.86 [1.31 - 2.31]$ Normal+normal N=11 $459 [273 - 674]$ $7.6 [4.9 - 16.0]$ $5.2 [0.7 - 5.7]$ $39 [35 - 51]$ $3.0 [0.5 - 4.0]$ $35 [20 - 55]$ $1.55 [1.21 - 2.02]$ $0.32 [0.16 - 0.341]$	Depressed N=77 323 [158 - 875] 10.9 [7.1 - 20.3] 2.6 [1.0 - 6.5] 41 [31 - 57] 2.0 [0.5 - 4.0] 36 [29 - 56] 1.08 [0.64 - 1.94] 0.12 [0.06 - 0.24] 1.73 [1.00 - 2.74] Normal + Abnor N=146 255 [115 - 791] 12.7 [7.0 - 21.6] 2.3 [1.1 - 7.0] 41 [31 - 54] 1.0 [0.5 - 4.0] 37 [25 - 53] 1.18 [0.85 - 1.58] 0.11 [0.07 - 0.19]	Heterogeneous N=16 $304 [180 - 1024]$ $5.8 [5.0 - 16.5]$ $1.4 [0.7 - 2.4]$ $41 [30 - 44]$ $0.5 [0.5 - 3.0]$ $28 [22 - 43]$ $0.99 [0.55 - 1.52]$ $0.12 [0.07 - 0.16]$ $1.46 [1.05 - 1.77]$ Ab + Abnormal N=74 $308 [174 - 1012]$ $10.1 [6.4 - 20.7]$ $2.3 [1.0 - 5.2]$ $42 [31 - 52]$ $2.0 [0.5 - 3.3]$ $35 [25 - 51]$ $1.11 [0.69 - 1.75]$ $0.12 [0.06 - 0.19]$	P* 0.406 0.351 0.142 0.479 0.125 0.371 0.133 0.943 0.211 P* 0.408 0.564 0.720 0.928 0.764 0.872 0.131 0.064	Adjusted OR (95% CI) Depressed vs Control-like 1.41 (0.77-2.56) 0.90 (0.46-1.78) 0.85 (0.48-1.48) 2.45 (0.56-10.80) 1.50 (0.75-3.02) 1.30 (0.49-3.42) 0.32 (0.09-1.17) 0.95 (0.49-1.86) 0.81 (0.25-2.58) Adjusted OR (95% CI) Normal+Abnor vs Ab+Abnormal 1.41 (0.72-2.75) 1.11 (0.51-2.45) 0.88 (.048-1.64) 2.34 (0.40-13.86) 1.61 (0.74-3.48) 1.11 (0.36-3.40) 0.46 (0.11-1.97) 0.97 (0.48-1.98)	P† 0.262 0.767 0.558 0.235 0.251 0.598 0.084 0.889 0.717 P† 0.310 0.788 0.696 0.349 0.231 0.862 0.296

*P for Kruskal Wallis test. † Biomarkers were log-transformed and included in logistic regression analysis with the histopathological feature dependent adjusted for age, COPD, history of Atrial Fibrillation, heart failure, valve surgery and EuroSCORE log. Three classes of Connexin 43: normal pattern (patients with control Cx43 junctional and little/absent Cx43 lateralization); normal + abnormal pattern (patients with control Cx43 junctional and present Cx43 lateralization or patients with depressed Cx43 junctional and little/absent Cx43 lateralization); abnormal pattern (patients with depressed Cx43 junctional and present Cx43 lateralization; or patients with heterogeneous Cx43 junctional and present Cx43 lateralization).

		Methods of histological analysis		
	Total	Automatic	Semi- automatic	No Automatic
Nr of papers, n	19	4	2	13
Nr of patients, n, mean±SD (total)	96±55 (1828)	64±34 (257)	84±72 (168)	108±58 (1403)
Nr of patients with POAF, n (%), mean±SD	530 (29), 28±17	81 (32) 20±8	98 (58) 49±48	351 (25) 27±12
Age overall (years), mean±SD	63±7	62±13	58±12	65±3
Age of POAF (years), mean±SD	68±7	74±0	61±17	69±4
Sex (Male), n (%)	1200 (66)	189 (74)	80 (48)	931 (66)
Type of surgery, Coronary bypass, n/n Valve surgery, n/n	17/19 10/19	3/4 3/4	1/2 2/2	13/13 5/13
Multivariate analysis, n/n	13/19	2/4	2/2	9/13 American Association
Biomarkers, n	12/19	1/12	2/2	9/13
Histological variables analyzed				
Fibrosis, n/n	15/19	3/4	2/2	10/13
Fibrosis association with POAF, n/n	8/15 Yes 6/15 No 1/15 not clear	2/3 Yes 1/3 No	2/2 Yes	4/10 Yes 5/10 No 1/10 not clear
Myocytolysis, n/n	8/19	1/4	1/2	6/13
Myocytolysis association with POAF, n/n	7/8 Yes 1/8 No	1/1 Yes 0/1 No	1/1 Yes 0/1 No	5/6 Yes 1/6 No
Hypertrophy of cardiomyocytes, n/n	6/19	1/4	0/2	5/13
Hypertrophy of cardiomyocytes association with POAF, n/n	1/6 Yes 5/6 No	1/1 No	-	4/5 Yes 1/5 No
Cx43, n/n	3/19	2/4	0/2	1/13
Cx43 association with POAF, n/n	1/3 Yes 2/3 No	1/2 Yes 1/2 No	-	0/1 Yes 1/1 No

Table 5. Literature analysis of papers focused on Post-Operative Atrial Fibrillation, subdivided by methods of histological evaluations.

Automatic: the measurements were performed by using a computer image analyzer; semi-automatic: the measurements were performed manual and with a computer-assisted image analyzer; No Automatic: the measurements were performed manual by the operators.

		Methods of histological analysis			
	Total	Automatic	Semi- automatic	No Automatic	
Nr of papers, n	30	15	6	9	
Nr of patients (without animals), n, mean±SD (total)	72±59 (1877)	69±43 (831)	73±58 (367)	75±82 (679)	
Nr of animals, n, mean±SD	123, 21±13	87 17±12	36	-	
Age(years) of patients, mean±SD	56±9	53±9	57±10	60±7	
Age of AF (years), mean±SD	61±11	57±8	70±18	62±9	
Sex (Male), n (%)	1061 (57)	486 (58)	229 (62)	346 (51)	
Type of surgery Coronary bypass, n/n Valve surgery Others	6/30 19/30 11/30	1/15 7/15 6/15	4/6 2/6	5/9 8/9 3/9	
Multivariate analysis, n/n	6/30	3/15	2/6	2/9	
Biomarkers, n	7/30	4/15	2/6	1/9	
Histological variables analyzed					
Fibrosis, n/n	20/30	11/15	3/6	6/9	
Fibrosis association with POAF, n/n	18/20 Yes 2/20 No	11/11 Yes	3/3 Yes	4/6 Yes 2/6 No	
Myocytolysis, n/n	6/30	-	2/6	4/9	
Myocytolysis association with POAF, n/n	5/6 Yes 1/6 not clear	-	2/2 Yes	3/4 Yes 1/4 not clear	
Hypertrophy of cardiomyocytes, n/n	5/30	1/15	1/6	3/9	
Hypertrophy of cardiomyocytes association with POAF, n/n	5/5 Yes	1/1 Yes	1/1 Yes	3/3 Yes	
Cx43, n/n	1/30	-	1/6	-	
Cx43 association with POAF, n/n	1/1 No	-	1/1 No	-	

Table 6. Literature analysis of papers focused on Other Types of Atrial Fibrillation, subdivided by methods of histological evaluations.

Automatic: the measurements were performed by using a computer image analyzer; semi-automatic: the measurements were performed manual and with a computer-assisted image analyzer; No Automatic: the measurements were performed manual by the operators.

Figure Legends:

Figure 1. A. Low-power histopathological view of a RAA biopsy with mild-to-moderate hypertrophy of the pectinate muscles (arrow) and moderate subepicardial fibro-adipose thickening (arrowhead). B. A detail of the region in A marked by an asterisk, with moderate-to-severe interstitial fibrosis (arrow). C. Intermediate-power view of a RAA biopsy with moderate interstitial fibrosis and cardiomyocytes with transverse diameters within the normal limits. D. C. Intermediate-power view of a RAA biopsy and cardiomyocytes with moderate interstitial fibrosis and cardiomyocytes with enlarged transverse diameters. Stainings: A-B hematoxylin-eosin; C-D Periodic acid–Schiff. Original magnifications: A x2 (scale bar: 2mm), B-D x20 (scale bar: 150µm).

Figure 2. RAA biopsies including cardiomyocytes (arrows) A. without glycogen deposition, B. with mild glycogen deposition, C. with moderate glycogen deposition, and D. with severe glycogen deposition. Staining: A-D Periodic acid–Schiff (PAS). Original magnifications: A-D x40 (scale bar: 100µm).

Figure 3. RAA myocardial fibrosis. A. Interstitial fibrosis within the normal limits and sparse myocytolytic cardiomyocytes (arrow). B. Mildly increased perivascular fibrosis around deeppenetrating myocardial blood vessels (arrow). C. Mild-to-moderate myocardial interstitial fibrosis (arrow). D. Severe myocardial interstitial fibrosis (arrow). Staining: A-D Van Gieson staining method. Original magnifications: A-B x10 (scale bar: 300μm); C-D x20 (scale bar: 150μm).

Figure 4. Connexin 43 immunosignal. A. Control distribution of Connexin 43 at the intercalated disks. B. Depressed signal at the intercalated disks. C. Profound lateralization of Connexin 43. D. Near-absence Connexin 43 immunosignal. A-D confocal microscopy. Original magnifications: A-D x40 (scale bar: 100μm).

American Heart Association.

Circulation: Arrhythmia and Electrophysiology

What is Known:

- Post-operative atrial fibrillation (POAF) occurs in 30-50% of patients undergoing cardiac surgery and is associated with increased morbidity and mortality.
- The lack of understanding of the specific preceding anatomic and molecular changes in the atrial myocardium at the time of cardiac surgery that might influence the risk of POAF is a major limitation in developing preventive strategies.

What the Study Adds:

- In 239 patients undergoing cardiac surgery (35.2% developed POAF), right atrial appendage (RAA) was excised, fixed, paraffin embedded and analyzed.
- On top of well-known risk factors for POAF (i.e. advanced age, heart failure, COPD, higher EuroSCORE, valve surgery), none of the histological features (i.e. fibrosis, myocytolysis, cardiomyocyte diameter, glycogen score, Connexin 43 distribution) was independently associated with risk of POAF.
- RAA structural remodeling does not reflect changes in other parts of the atria, shown to contribute to the pathogenesis of POAF.











Graphic Abstract

ClinicalTrials.gov Identifier: NCT00970489

TO INVESTIGATE THE HISTOPATHOLOGY OF RIGHT ATRIAL APPENDAGE (RAA) THAT MIGHT PREDICT POST-OPERATIVE ATRIAL FIBRILLATION (POAF)



239 RAA BIOPSIES AND BLOOD SAMPLES FROM PATIENTS IN SINUS RHYTHM UNDERGOING CARDIAC SURGERY

HISTOPATHOLOGICAL ANALYSIS

INTERSTITIAL

FIBROSIS

CONNEXIN 43

GENERAL MORPHOLOGY



HYPERTROPHY MYOCYTOLYSIS



CIRCULATING BIOMARKERS



HISTOPATHOLOGICAL CHANGES IN THE RAA AND CIRCULATING BIOMARKERS DO NOT PREDICT POAF OCCURRENCE