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1 Molecular mechanisms of diaphragm myopathy in humans with severe heart failure

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25
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Abstract:

2

Rationale: Diaphragm weakness impairs quality-of-life, exercise capacity, and survival in patients with chronic heart failure (CHF) and reduced left ventricular ejection fraction. However, the underlying cellular mechanisms responsible in humans remain poorly resolved.

Objectives: We prospectively evaluated clinical, functional (*in vivo/in vitro*), histological/ultrastructural and molecular alterations of the diaphragm from CHF patients receiving a left ventricular assist device compared to patients without CHF undergoing elective coronary bypass grafting (control) in the observational LIPsia DiaPhrAgm and MUScle Heart Failure Trial (LIPAMUS-HF).

Methods and Results: Participants (Controls=21, CHF=18) underwent cardiopulmonary exercise and spirometry/respiratory muscle testing alongside diaphragm and cardiac imaging. Diaphragm biopsies were phenotyped for mitochondrial respiration, muscle fiber function, histology/ultrastructure, and protein expression.

In vivo respiratory muscle function and diaphragm thickness were reduced in CHF by 38% and 23%. Diaphragm biopsies revealed a fiber-type shift and severe fiber atrophy in CHF alongside elevated proteasome-dependent proteolysis (i.e., MuRF1 expression, ubiquitination, ubiquitin proteasome activity) and myofibrillar protein oxidation, which corresponded to upregulated NADPH oxidase (Nox2/Nox4) signaling. Mitochondria demonstrated severe intrinsic functional and ultrastructural abnormalities in CHF characterized by accumulation of small mitochondria and inhibited autophagy/mitophagy. Single muscle fiber contractile function revealed reduced Ca²⁺ sensitivity in CHF and there was evidence of ryanodine receptor 1 (RyR1) dysfunction indicating Ca²⁺ leak from the sarcoplasmic reticulum. Mitochondrial and Ca²⁺ measures corresponded to upregulated Nox4 isoform NADPH oxidase expression. Molecular markers correlated to whole-body exercise intolerance and diaphragm dysfunction/wasting.

Conclusions: CHF patients demonstrate an obvious diaphragm myopathy independent of disuse or other confounding factors such as ageing, obesity, or hypertension. Diaphragm weakness in CHF was associated with intracellular abnormalities characterized by fiber atrophy, oxidative stress, mitochondrial dysfunction, impaired Ca²⁺ homeostasis, elevated proteasome dependent proteolysis, but inhibited autophagy/mitophagy, which we speculate offers a novel therapeutic molecular target regulated by a Nox-MuRF1/ubiquitin proteasome-mitochondria-RyR1/Ca²⁺ signaling axis.

32

Clinical Trial Registration: URL: <https://clinicaltrials.gov> Unique Identifier: NCT02663115

34

Keywords: diaphragm; humans; heart failure; MuRF1, NADPH

36

1 Non-standard Abbreviations and Acronyms

2

3 ATP – adenosine triphosphate

4 CABG – coronary artery bypass grafting

5 CHF – chronic heart failure

6 ETC – electron transfer chain

7 FEV1 – forced expiratory volume in 1 second

8 ROS – reactive oxygen species

9 SR – sarcoplasmic reticulum

10 UPS – ubiquitin-proteasome system

11 VC – vital capacity

12 VE/VCO₂ – ratio ventilation to carbon dioxide output13 VO_{2peak} – maximal oxygen uptake

14

1 Introduction

2

3 The diaphragm is the main respiratory muscle responsible for normal ventilatory behaviors.¹ Diaphragm
4 weakness is common in patients with chronic heart failure (CHF)² and is closely associated with
5 symptoms and mortality.^{3,4} Yet, the underlying mechanisms in humans remain only partially resolved
6⁵⁻⁸ and most of our knowledge is derived from animal models.⁹⁻¹¹ Of the few human studies performed,
7 most were underpowered, or lacked modern molecular biology approaches⁵⁻⁸, thus limiting our
8 mechanistic understanding and potential treatment options for the current patient.

9 Both acute⁹ and chronic heart failure¹¹ cause diaphragm weakness in CHF mouse models, which is
10 characterized by both fiber contractile dysfunction and fiber atrophy.¹¹⁻¹⁴ Reactive oxygen species
11 (ROS) play a causal role to induce diaphragm weakness^{15,16} since antioxidant interventions such as
12 exercise training¹¹, pharmacological aids¹⁷, or genetic deletion¹⁸ can prevent diaphragm weakness by
13 reducing posttranslational oxidative modifications of contractile proteins and suppressing protein
14 degradation via the ubiquitin-proteasome system¹⁹ in a MuRF1 dependent manner.²⁰ In end-stage CHF
15 patients, the diaphragm shows ultra-structural and myofibrillar alterations that implicate metabolic and
16 contractile abnormalities^{6,8}, which are related to NADPH signaling.⁵ Animal models have identified
17 NADPH oxidase²¹ and mitochondria^{16,22} as major sources of ROS, but their role in the human
18 diaphragm remains poorly defined. In addition, the level of structural and molecular alterations in the
19 diaphragm has never been extensively linked to clinical presentation.

20 We hypothesized that CHF patients would have a distinctive diaphragm myopathy, which would closely
21 correlate to clinical observations. The present study, therefore, evaluated clinical, histological and
22 molecular evidence from diaphragm biopsies in CHF patients scheduled for left ventricular assist device
23 implantation and patients without CHF requiring coronary artery bypass grafting.

1 Methods

2

3 The data that support the findings of this study are available from the corresponding author upon
4 reasonable request. Patients with severe CHF and reduced left ventricular ejection fraction (LV-EF)
5 scheduled for left ventricular assist device implantation and patients with coronary artery disease
6 (control) undergoing elective coronary artery bypass grafting without CHF were eligible for this study.
7 All patients gave written informed consent. The trial was approved by the local ethics committee,
8 complied with the Declaration of Helsinki, and is registered at www.clinicaltrials.gov (Unique
9 Identifier: NCT02663115).

10 Categorical variables are given as numbers and percentage compared between groups by the chi-squared
11 test or Fisher exact test as appropriate. Continuous variables are shown as mean and 95%-confidence
12 interval irrespective of its distribution assessed by the Shapiro-Wilk-Test. Groups were compared using
13 the Students t-test or Mann-Whitney-U-Test according to normal vs non-normal distribution. Due to
14 limited biopsy size, not all parameters were determined in all biopsies, exact numbers and all values
15 depicted in the *Figures* are given in *Online Table I*. For all the representative images in the manuscript,
16 demonstrative examples reflecting the group mean trend were selected. Correlation analyses were
17 performed using Pearson correlation to evaluate the association of both potential biological pathways
18 predominantly known from animal studies and molecular measures with three defined clinical
19 parameters (diaphragm thickness, VO_{2peak} , VE/VCO_2). A two-sided p-value <0.05 was considered
20 significant.

21 For detailed methodological information, please refer to the *Online Supplement* and the *Major*
22 *Resources Table*.

23

1 Results

2

3 Baseline characteristics

4 Between January 2016 and December 2017, 18 CHF patients and 21 controls were included. Baseline
 5 characteristics are shown in *Table 1*. Compared to controls, CHF were slightly younger, had worse renal
 6 function, a lower body mass index, and lower rates of coronary artery disease, but had a higher incidence
 7 of atrial fibrillation and previous cardiac surgery (all $p < 0.05$). One third of CHF had ischemic and two
 8 thirds had dilated cardiomyopathy. With regard to whole body exercise capacity, CHF exhibited a 35%
 9 lower VO_{2peak} and a significantly higher VE/VCO_2 slope than controls, whereas the respiratory exchange
 10 ratio was comparable indicating physical exhaustion in both groups (*Online Table II*). Catheterization
 11 data of CHF are provided in *Online Table III*.

12

13 **Table 1: Baseline characteristics**

14

	Control n=21	CHF n=18	P-value
Age [years]	62 (60; 65)	57 (53; 61)	0.014*
Male Gender, n (%)	17 (81.0)	17 (94.4)	0.349 [†]
Caucasian ethnicity, n (%)	21 (100)	18 (100)	n.a.
Body Mass Index [kg/m ²]	32.0 (29.5; 34.6)	28.6 (25.9; 31.3)	0.037*
Systolic blood pressure [mmHg]	133 (124; 142)	109 (103; 115)	8.3E-5 [‡]
Heart rate [min ⁻¹]	76 (69; 82)	82 (75; 89)	0.180 [‡]
New York Heart Association class			3.4E-4 [§]
I	2 (9.5)	0 (0)	
II	14 (66.7)	2 (11.1)	
III	5 (23.8)	10 (55.6)	
IV	0 (0)	6 (33.3)	
STS-Score (for CABG) [%]	0.42 (0.36; 0.49)	1.58 (1.02; 2.13)	2.3E-7*
Medical history			
Coronary artery disease, n (%)	21 (100)	10 (55.6)	7.1E-4 [†]
Previous Myocardial Infarction, n (%)	1 (4.8)	5 (27.8)	0.077 [†]
Previous percutaneous coronary intervention, n (%)	5 (23.8)	8 (44.4)	0.173 [§]
Previous cardiac surgery, n (%)	0 (0)	4 (22.2)	0.037 [†]
Atrial fibrillation/flutter, n (%)	2 (9.5)	11 (61.1)	6.6E-4 [§]
Arterial hypertension, n (%)	20 (95.2)	17 (94.4)	1.000 [†]
Diabetes mellitus, n (%)	10 (47.6)	10 (55.6)	0.521 [§]
Previous stroke, n (%)	0 (0)	1 (5.6)	0.462 [†]
Peripheral artery disease, n (%)	1 (4.8)	2 (11.1)	0.586 [†]
Chronic kidney disease, stage $\geq 3b$, n (%)	0 (0)	9 (50)	2.3E-4 [†]
Serum creatinine [mmol/l]	78 (71; 84)	122 (100; 144)	1.3E-4 [‡]
Heart failure features			
Ischemic cardiomyopathy, n (%)	n.a.	6 (33.3)	n.a.
Dilated cardiomyopathy, n (%)	n.a.	12 (66.7)	
Number of cardiac decompensation	0 (0; 0)	1.39 (0.93; 1.84)	1.1E-7*
NT-proBNP [pg/ml]	244 (84; 405)	3449 (1791; 5108)	1.6E-8*
Echocardiography			
Left ventricular-ejection fraction [%]	61 (58; 64)	19 (17; 22)	8.6E-23 [‡]
Left ventricular enddiastolic diameter [mm]	51 (49; 53)	73 (69; 77)	5.2E-11 [‡]

Left atrial volume index [ml/m ²]	34 (28; 41)	63 (54; 72)	6.2E-6*
Right ventricular enddiastolic parameter [mm]	29 (27; 32)	38 (34; 41)	1.4E-4‡
Right atrial area [mm ²]	13 (12; 15)	23 (19; 28)	2.8E-4‡
Tricuspid annular plane excursion [mm]	23 (20; 25)	14 (12; 15)	2.6E-7‡
Diameter inferior vena cava [mm]	14 (13; 16)	19 (16; 22)	1.1E-3‡
Lung function			
VC [l]	3.9 (3.4; 4.3)	3.2 (2.9; 3.4)	8.0E-3‡
FEV1 [l]	3.1 (2.8; 3.5)	2.6 (2.3; 2.9)	0.022‡
FEV1/VC [%]	81 (78; 85)	79 (75; 83)	0.250‡
Residual volume [l]	2.7 (2.2; 3.2)	2.5 (2.1; 2.8)	0.466‡
Resistance [kPa/l/s]	0.24 (0.19; 0.29)	0.26 (0.21; 0.32)	0.415*
Transfer coefficient [%]	99 (92; 106)	80 (73; 87)	3.8E-4‡
Treatments at inclusion			
Beta-blocker, n (%)	17 (81.0)	17 (94.4)	0.349†
ACE-inhibitor, n (%)	8 (38.1)	7 (38.9)	0.959§
Angiotensin receptor blocker, n (%)	8 (38.1)	0 (0)	4.0E-3†
Neprylisin inhibitor/angiotensin receptor blocker, n (%)	0 (0)	6 (33.3)	5.7E-3†
Mineral receptor antagonist, n (%)	1 (4.8)	16 (88.9)	1.3E-7§
Digitoxin, n (%)	0 (0)	4 (22.2)	0.037†
Diuretics, n (%)	10 (47.6)	16 (88.9)	6.4E-3§
Intravenous Phosphodiesterase inhibitor, n (%)	0 (0)	15 (83.3)	9.7E-8§
Implantable cardioverter defibrillator, n (%)	0 (0)	16 (88.9)	1.8E-8§
Cardiac resynchronization therapy, n (%)	0 (0)	4 (22.2)	0.037†

1
2 Values are expressed as mean with 95%-confidence interval or numbers and percentages. STS indicates
3 Society of Thoracic Surgeons score; CABG, coronary artery bypass grafting; VC, vital capacity; FEV₁,
4 forced expiratory volume in 1 second; ACE, angiotensin converting enzyme.

5 All p-values are two-sided and were not corrected for multiple testing. Continuous variables were tested
6 for normal distribution applying the Shapiro-Wilk-test.

7 * Mann-Whitney-U-Test

8 † Fisher exact test

9 ‡ unpaired Students t-test

10 § chi-squared test

11

12 *In vivo respiratory function*

13 P_{i,max} and P_{e,max} were significantly reduced by 38% and 25% in CHF compared to controls (*Figure 1 A*).
14 P_{0.1} was not significantly different between groups; however, P_{0.1}/P_{i,max} ratio was higher in CHF
15 indicating increased ventilatory drive (*Figure 1 B, Online Table II*). Diaphragm wasting was obvious in
16 CHF, with mean diaphragm thickness 23% lower compared to control (*Figure 1 C, Online Table II*).
17 Diaphragm thickness was moderately correlated to P_{i,max} (R=0.44, p=5.0E-3), the latter also correlated
18 to vital capacity (r=0.47; p=3.0E-3) (*Figure 1 D-E*). Spirometry showed vital capacity, absolute FEV₁,
19 and transfer coefficient were lower in CHF, but FEV₁/VC, residual volume und resistance were
20 comparable between groups (*Table 1*). No patient had a FEV₁/VC <70% excluding obstructive lung
21 disease.

22

23 *Diaphragm sampling*

24 No serious adverse events occurred during baseline or diaphragm sampling in either group. The time
25 from intubation to sampling of the diaphragm probes was comparable between both groups (CHF 69
26 min (95%-CI 53; 85) vs control 64 min (95%-CI 55; 74), p=0.576), which suggests any differences in
27 diaphragm measures were unrelated to mechanical ventilation.

28

29

1 *Fiber remodeling*

2 Fiber remodeling involving isoform shift and atrophy are suggested to underpin diaphragm weakness in
 3 CHF. We confirmed a fiber-type shift and atrophy in CHF compared to controls: a fast type II to slow
 4 type I was observed ($p=0.047$) alongside atrophy in both type I and type II fibers with lower cross
 5 sectional areas of 35% and 51% respectively (*Figure 2 A-D*). The ubiquitin-proteasome system (UPS)
 6 is the major proteolytic pathway responsible for protein degradation, which is rate-limited in wasting
 7 conditions by upregulation of muscle-specific E3 ligases that include MuRF1 and MAFbx. MuRF1
 8 protein expression was higher by 46% ($p=3.9E-4$) (*Figure 2 E*), whereas MAFbx protein expression was
 9 lower by 35% in CHF vs control ($p=2.0E-3$) (*Figure 2 F*). However, we assume a higher overall E3
 10 ligase activity in CHF that is supported by a 54% higher protein ubiquitination at lysine 48 residues,
 11 with proteasome activity elevated by 54% ($p=0.026$) (*Figure 2 G-H*). MuRF1 expression was correlated
 12 with protein ubiquitination at lysine residues (Ubi-K48) ($R=0.46$, $p=0.017$) and Ubi-K48 correlated with
 13 proteasome activity ($R=0.58$, $p=2.9E-3$) supporting the biological pathway of E3 Ligase dependent
 14 proteasome activation. Moreover, MuRF1 protein expression was correlated with clinical parameters
 15 (diaphragm thickness ($R=-0.46$, $p=0.015$), VO_{2peak} ($R=-0.43$, $p=0.024$), VE/VCO_2 ($R=0.40$, $p=0.045$)).
 16 These data implicate diaphragm weakness and atrophy in CHF is underpinned by MuRF1-dependent
 17 ubiquitin-proteasome degradation.

18

19 *Oxidative stress*

20 Increased oxidative stress leads to 1) activation of UPS components to elevate protein degradation and
 21 2) contractile dysfunction by damaging sarcomeric proteins.²¹ As NADPH oxidases are a major source
 22 of muscle-derived ROS, we confirmed that protein expression of Nox2 and Nox4 isoforms were 2.8-
 23 ($p=4.2E-3$) and 10.2-fold ($p=1.4E-3$) higher in CHF vs controls (*Figure 2 I-J*). ROS-induced protein
 24 damage was evident in CHF as indicated by a 33% higher carbonylation of myosin heavy chain in CHF
 25 vs controls ($p=0.015$) (*Figure 2 K*), supporting higher oxidation of myofibrillar proteins. Nox2 was
 26 correlated to MuRF1 expression ($R=0.40$, $p=0.049$) and Ubi-K48 ($R=0.67$, $p<1.0E-4$) (*Figure 2 L*).
 27 Nox2 also correlated with *in vivo* measures as diaphragm thickness ($R=-0.52$, $p=3.3E-3$), VO_{2peak} ($R=-$
 28 0.72 , $p=1.1E-5$) and VE/VCO_2 ($R=0.56$, $p=1.5E-3$) (*Online Table IV*). Overall, these data suggest that
 29 NADPH oxidases are likely a major upstream mechanism inducing oxidative stress causing diaphragm
 30 weakness in human CHF⁵, at least partially via 1) activating components of the UPS to elevate protein
 31 degradation, leading to muscle atrophy, and 2) inducing contractile dysfunction by damaging sarcomeric
 32 proteins. Those molecular mechanisms are linked to whole-body exercise intolerance.

33

34 *Mitochondrial function and morphology*

35 Since mitochondrial dysfunction is known to play a key role in fiber atrophy, and Nox4 interacts with
 36 mitochondria, we next examined the mitochondria. Mitochondria of CHF exhibited a lower citrate
 37 synthase activity per mg mitochondrial protein (*Figure 3 A*) and were smaller according to a FACS
 38 analysis of all undamaged mitochondria (*Figure 3 B*). In addition, state 3 and state 4 respiration were
 39 significantly lower across different substrates for complex I, II, III, and IV in CHF patients compared to
 40 controls (*Online Table V*). Respiratory control index and ADP/O ratio were not different between groups
 41 irrespective of the substrate indicating comparable phosphorylation efficiency. Representative graphs
 42 are shown for palmitoylcarnitine/malate as complex I substrate in *Figure 3 C-F*. State 3 respiration was
 43 correlated to *in vivo* measures of ventilatory efficiency ($R=-0.58$, $p=0.012$) and diaphragm thickness
 44 ($r=0.60$, $p=6.4E-3$) (*Figure 3 G-H*) and VO_{2peak} ($R=0.57$, $p=0.014$). We used electron microscopy to
 45 assess diaphragm mitochondrial morphology in CHF and control samples, which revealed accumulation
 46 of intermyofibrillar and subsarcolemmal mitochondria characterized by variable size, but predominately
 47 small-sized mitochondria in CHF (*Figure 3 I-L*).

48 Given that mitochondrial function depends on both intrinsic function and amount, we assessed
 49 mitochondrial content in patients by examining different mitochondrial markers and enzyme activities
 50 of proteins related to the electron transfer chain (ETC). Mitochondrial quantity was higher in CHF vs
 51 controls indicated by a 2.2-fold elevated porin protein expression (*Figure 4 A*) and 1.3-fold higher citrate
 52 synthase activity in whole muscle homogenates (*Figure 4 B*). Succinate dehydrogenase activity
 53 remained unchanged (*Figure 4 C*). However, after normalizing these values for mitochondrial amount
 54 to provide an index of intrinsic mitochondrial function, both ratios for citrate synthase and succinate
 55 dehydrogenase activity were significantly lower in CHF (*Figure 4 B-C*). In addition, enzymatic

1 complex-I activity was reduced by 41% in CHF, which is in line with the reduced complex-I respiration
2 found in isolated mitochondria (*Figure 4 D*). The protein expression of uncoupling protein 3 (UCP3)
3 was higher in CHF compared to controls ($p=0.052$) (*Figure 4 E*), whereas protein expression of
4 uncoupling protein 2 was not detectable in both CHF and controls (data not shown).
5 To provide further insights into mitochondrial pathology, we examined parameters of fusion and fission.
6 The protein expression of the fusion parameter Mitofusin-2 (Mfn2) and the fission parameter cytosolic
7 GTPase dynamin-related protein 1 (Drp1) was 2.3-fold ($p=1.4E-3$) and 1.6-fold ($p=0.041$) higher in
8 CHF than in controls (*Figure 4 F-G*) suggesting both increased fusion and fission. To assess
9 mitochondria biogenesis, we investigated the protein expression of PGC1alpha found to be 1.5-fold
10 higher in CHF ($p=0.032$). Autophagy/mitophagy was assessed by evaluating the protein expression of
11 LC3-I/LC3-II and p62. The ratio LC3-I/LC3-II was 1.4-fold higher in CHF ($p=9.0E-3$) and p62 protein
12 expression was 2-fold higher in CHF ($p=1.4E-3$). The combination of a reduced lipidation of LC3 and
13 accumulation of p62 indicates inhibited autophagy/mitophagy in CHF.²³
14 Overall, these data reveal that despite increased measures of overall number and content, intrinsic
15 dysfunction of mitochondria is present in the CHF diaphragm, potentially caused by impaired
16 mitophagy.

17 18 *Calcium homeostasis*

19 The observed mitochondrial alterations closely mirror those reported to occur when Ca^{2+} overload is
20 present consequent to “leaky” ryanodine receptor (RyR1).²⁴ We investigated the binding status of
21 FKBP12 to RyR1, which when reduced is a proxy for “leaky” RyR1 to indicate altered Ca^{2+} homeostasis.
22 Binding of FKBP12 to RyR1 was lower in CHF compared to controls ($p=0.050$), which indicated Ca^{2+}
23 efflux from the sarcoplasmic reticulum (SR) into the cytosol (*Figure 5 A*). There was a parallel
24 significant increase in protein expression of Ca^{2+} re-uptake proteins in CHF including SERCA-1 and
25 SERCA-2A (*Figure 5 B-C*). Elevated Ca^{2+} levels can also activate Ca^{2+} -dependent proteases such as
26 calpains to promote fiber atrophy.²⁵ We therefore measured calpain activity but did not find a significant
27 difference between groups (*Figure 5 D*), which suggests that catabolic activity in the diaphragm of
28 humans with CHF is potentially driven by the UPS. Impaired Ca^{2+} homeostasis can also limit excitation-
29 contraction coupling to impair diaphragm contractile function and induce weakness. As such, we next
30 assessed Ca^{2+} -activated single-fiber contractile function in isolated diaphragm type II fibers, which
31 revealed significantly reduced myofilament Ca^{2+} sensitivity in CHF vs controls (*Figure 5 E-F*).
32

1 Discussion

2
3 Our data show that CHF patients suffer from a diaphragm myopathy that occurs independent of disuse
4 or other confounding factors such as ageing, obesity, or hypertension. Specifically, we confirm using *in*
5 *vivo* and *in vitro* approaches that diaphragm weakness and wasting is present in CHF, which is
6 underpinned by severe cellular remodeling and molecular alterations that include fiber atrophy, NOX-
7 related oxidative stress, proteasome-linked catabolic activation, mitochondrial dysfunction, and
8 impaired Ca²⁺ homeostasis. Clinically, the molecular alterations closely correlate to both whole-body
9 exercise capacity and respiratory function. Thus our data reveal, for the first time in humans with heart
10 failure, a novel molecular mechanism and potential therapeutic target for diaphragm weakness that is
11 centered around a Nox-MuRF1/ubiquitin proteasome-mitochondria-RyR1/Ca²⁺ dependent signaling
12 axis.

13 14 *Patient population and clinical data*

15 The CHF group comprised patients with exercise capacity reduced to 57% of the age- and sex-specific
16 predicted norm. CPX was performed in part under inotropic support, which might explain the slightly
17 higher VO_{2peak} compared to other left ventricular assist device studies.^{26, 27} In the control group, we
18 included non-CHF patients undergoing elective coronary artery bypass grafting. Clinically, control
19 patients did not have CHF and presented with normal measures of LV-EF, nt-proBNP and VO_{2peak}.
20 Moreover, we excluded patients in both groups with comorbidities known to influence the morphology
21 and function of the diaphragm, in particular recent mechanical ventilation, COPD, severe chronic and/or
22 acute renal failure.

23 The magnitude of diaphragm dysfunction in our study was comparable to other studies including those
24 examining Pi_{max} as a prognostic marker^{3, 4} and including NYHA III/IV patients who have pronounced
25 diaphragm weakness.²⁸ The ratio P_{0.1}/Pi_{max}, commonly interpreted as an index for ventilatory drive, was
26 higher in CHF compared to controls and comparable to published data.²⁸ This observation is consistent
27 with rapid shallow breathing commonly found in advanced CHF. Despite this increased workload, we
28 found reduced diaphragm thickness in CHF, which is an index of atrophy closely associated with
29 exercise intolerance in CHF.²⁹ Structural changes in CHF were associated with *in vivo* respiratory
30 dysfunction and cellular markers of proteolysis (i.e. MuRF1) and mitochondrial derangements.

31 32 *NADPH oxidase as a major mechanism triggering diaphragm weakness*

33 NADPH oxidases have emerged as one major source of ROS in skeletal muscle cells.²¹ An increased
34 NADPH expression has been observed in both acute⁹ and chronic animal models of heart failure.¹⁸
35 Increased Nox2 protein expression in the diaphragm of CHF patients is in line with a former study
36 examining Nox2 and its downstream mediators.⁵ An ~40% increase of ROS induced protein damage
37 was also evident⁵, which is similar to the increased carbonylation of MHC in our study.

38 ROS, but also cytokines, angiotensin II and sphingomyelinase, are able to activate proteolytic pathways
39 in skeletal muscle including the UPS.³⁰ In the diaphragm of CHF, the UPS appeared to be the dominant
40 proteolytic system since calpain activity was not significantly different between groups. Moreover,
41 Nox2 and Nox 4 protein expression was correlated to activation of the key atrogene, muscle-specific E3
42 ligase MuRF1, and protein ubiquitination, thus supporting the biological pathway of MuRF1-dependent
43 proteasome degradation of proteins. Both Nox2 and the UPS are linked to diaphragm weakness in animal
44 models^{18, 31} and the genetic knock out of the p47(phox)-Nox2 subunit is associated with reduced ROS
45 production and preservation of diaphragm function.¹⁸ In humans, the correlation of Nox2 and MuRF1
46 to *in vivo* measures of diaphragm morphology and function supports the association between increased
47 ROS/catabolic activity and whole-body symptoms related to exercise intolerance and respiratory
48 dysfunction. While we saw an increase in MuRF1 expression, MAFbx was lower in CHF than in
49 controls. This phenomenon has been observed before with unchanged or reduced expression levels in
50 humans and animals with heart failure and/or after exercise training.^{10, 32, 33} However, we assume a
51 higher overall E3 ligase activity in CHF supported by increased polyubiquitination at lysine 48 residues
52 and proteasome activity in CHF, with the former one being the typical polyubiquitination signal for
53 proteasome dependent degradation³⁴. Reduction of MAFbx, which predominantly leads to degradation
54 of initiation factor 3 subunit 5 (eIF3-f) and MyoD³⁵, might be a compensation to the increased
55 proteolysis of structural proteins by MuRF1 since an increase in eIF3-f leads to increased expression of

1 structural proteins and causes muscle hypertrophy.³⁶ Further research is necessary to investigate the
2 interaction between these two E3 ligases.

3 Many studies have focused on Nox2, however, the role of Nox4 in skeletal muscle and the diaphragm
4 of CHF patients is not well defined. While Nox2 is regulated by phosphorylation of p47(phox) or
5 p67(phox), and activation of Rac1, Nox4 is constitutively active and, therefore, transcriptionally
6 regulated.²¹ We found an astonishing 10.2-fold higher Nox4 protein expression in the diaphragm of
7 CHF indicating increased Nox4 activity. While Nox2 is a sarcolemmal protein, Nox4 contains a
8 mitochondria localization sequence in its N-terminal region and is found in muscle mitochondria and
9 sarcoplasmic reticulum (SR).²¹ Therefore, we assessed mitochondrial function and parameters of
10 calcium handling to provide further mechanistic insight.

11 *Mitochondrial function and calcium handling*

12 In isolated mitochondria, we found reduced state 3 and state 4 respiration across different substrates for
13 complex I-IV in CHF. Electron microscopy revealed accumulation of intermyofibrillar and
14 subsarcolemmal mitochondria in CHF characterized by variable size, but predominately small-sized
15 mitochondria, which is in line with the results of the FACS analysis of isolated mitochondria. This might
16 be the consequence of inhibited autophagy/mitophagy with an imbalance between fusion and fission,
17 which is discussed below. Increased markers of mitochondria quantity (citrate synthase, porin)
18 supported a higher amount of mitochondria in CHF. The reduced respiratory capacity across all four
19 electron transport chain (ETC) complexes suggests a defect in complex IV and/or assembly of
20 supercomplexes.³⁷ However, this does not exclude impairments in other complex activities. Indeed, we
21 proved a reduction in enzymatic complex-I- and succinate dehydrogenase activity suggesting that
22 reduced respiration of complex I and II is not only due to a defect in complex IV but also caused by
23 complex specific alterations. This may finally lead to a reduced proton gradient across the inner
24 membrane with reduced capacity for ATP production contributing to impaired muscle function and
25 exercise capacity.³⁸ From a clinical point of view, this hypothesis is supported by our correlative
26 analysis between a reduced state 3 respiration vs ventilatory efficiency and $\text{VO}_{2\text{peak}}$ derived from whole
27 body exercise testing.

28 To elucidate potential reasons for the impaired mitochondrial function in CHF, we evaluated markers
29 controlling the quality of the mitochondrial network. A higher protein expression of Mfn2 and Drp1 in
30 CHF indicates increased fusion and fission. Both parameters are sensitive to exercise³⁹ and the observed
31 higher expression might be the consequence of a higher ventilatory workload in CHF. The same could
32 be the case for PGC1alpha known to be very sensitive to exercise.³⁹ In contrast, the combination of a
33 reduced lipidation of LC3 and accumulation of p62 indicates inhibition of autophagy/mitophagy in CHF.
34²³ Blocked mitophagy is associated with a diminished mitochondrial function⁴⁰ and protein expression
35 of ETC enzymes.³⁹ Therefore, this finding might serve as an explanation for the reduced mitochondrial
36 function observed in the diaphragm of CHF patients. Further research is necessary to elucidate the
37 reasons for autophagy inhibition.

38 Moreover, we found an increased UCP3 expression, which can be the consequence of elevated ROS.
39 Increased UCP3 is found after acute exercise and is thought to alleviate the proton gradient across the
40 inner membrane, thereby reducing excessive ROS production but also ATP production by the ETC.⁴¹
41 In CHF, a similar scenario may occur in response to the increased work of breathing and acute episodes
42 of dyspnea⁷, potentially aggravated by hypoxia.⁴² Another hypothesis is Nox4 induced mitochondrial
43 stress, which is supported by the correlation between Nox4 and UCP3 in our patient cohort. Cardiac-
44 specific Nox4-deletion attenuated mitochondrial dysfunction in response to pressure overload in mice
45 being associated with reduced ROS production and oxidative damage to mitochondrial proteins.⁴³

46 Several mechanisms can account for the cross talk between Nox4, mitochondria and the SR including
47 oxidation of mitochondrial ETC proteins or oxidation of the RyR within the SR.²¹ The latter one causes
48 a Ca^{2+} leak from the SR.²⁴ Mitochondria act as a buffer with the result of Ca^{2+} overload, which in return
49 induces mitochondrial dysfunction.²⁴ Additionally, impaired mitophagy is associated with increased
50 mitochondrial Ca^{2+} uptake and a reduced Ca^{2+} availability for contraction.⁴⁰ Such a feedback-loop
51 between the SR and mitochondria in which SR Ca^{2+} leak triggers mitochondrial dysfunction has been
52 described in failing myocytes²⁴ and in the diaphragm during mechanical ventilation.⁴⁴ Against this
53 background, we examined the binding status of the subunit FKBP12 to RyR1, which if reduced is a
54 molecular marker of a “leaky” receptor state. Binding of FKBP12 to RyR1 was lower in CHF compared
55

1 to controls suggesting increased Ca^{2+} efflux from the SR. We also found an increased protein expression
2 of calcium re-uptake handling proteins in CHF, which likely indicates a compensatory response to
3 elevated Ca^{2+} levels by increasing reuptake pumping. Moreover, single-fiber contractile function from
4 CHF patients was impaired, showing reduced Ca^{2+} sensitivity. The rightward shift of the pCa^{2+} -force
5 curve indicates that diaphragm fibers of CHF patients generate a lower force of their maximal for a
6 given Ca^{2+} concentration.¹⁴ This suggests that contractile function in the CHF diaphragm is even more
7 affected at submaximal than at maximal activation, which is clinically relevant as the diaphragm most
8 often works at submaximal workloads. Overall, our data confirm impaired Ca^{2+} homeostasis in the
9 diaphragm of CHF patients, which is caused by leaky RyR1 coupled to reduced myofibrillar sensitivity
10 and mitochondrial dysfunction.

11 *Limitations*

12 Our results should be interpreted against the background of the following limitations. First, the control
13 group was not a healthy comparator since patients had coronary artery disease requiring coronary artery
14 bypass grafting; however, for reasons of gathering both clinical data and diaphragm biopsy, this was the
15 most appropriate group in our setting. Second, CHF were slightly younger than controls. We believe
16 that the difference of 7 years was not clinically meaningful. It even underlines the robustness of our
17 findings since respiratory function decreases with age. Third, CHF comprised both ischemic and dilated
18 cardiomyopathy, but number of patients was not sufficient to make meaningful comparisons between
19 both entities. Fourth, Nox2 and Nox4 subunits are expressed in several other cell types (e.g.,
20 macrophages, endothelium, and smooth muscle cells), which may have made minor contribution to our
21 muscle homogenate data. Fifth, single fiber measurement could only be done in type II fibers due to
22 contracted type I fibers. Preserving techniques are well established for those contractile measurements
23^{45, 46}; therefore, this phenomenon remains unclear. However, we do not believe that it was a disease
24 specific problem because it was found in both CHF and control probes. Moreover, we cannot rule out
25 that a fiber selection bias affected the outcomes of our contractility assays. To minimize this effect, we
26 randomly selected fibers for the contractile assays and excluded only those exhibiting injury patterns
27 that are associated with impossible or unreliable single fiber measurement. Sixth, our data are descriptive
28 and despite correlative evidence, we cannot provide a cause-relationship despite our assumptions being
29 supported in some instances by genetic knock out models. We were also not able to examine all the
30 downstream mediators known to be responsible for ROS production by NADPH oxidases due to tissue
31 limitation.²¹ Moreover, we are not able to distinguish between causative and compensatory molecular
32 events since it is a single time point analysis in humans and correlation analysis does not mean causality.
33 To elucidate cause and compensation, longitudinal experiments with specific therapies aimed at specific
34 molecular components would be necessary. However, it is important to recognize that the
35 pathophysiology of diaphragm dysfunction seen in different animal models (e.g. catabolic activation,
36 mitochondrial dysfunction, Nox activation) is also present in human specimens. Those results form a
37 solid base for development and testing of different treatment strategies in CHF to address respiratory
38 and skeletal muscle dysfunction including MuRF1 inhibition^{47, 48}, increasing Ca^{2+} sensitivity⁴⁹ or Nox
39 inhibition.⁵⁰

41 *Conclusions*

42 CHF patients demonstrate a diaphragm myopathy independent of disuse or other confounding factors
43 such as ageing, obesity, or hypertension. *In vivo* diaphragm weakness and wasting induced by CHF were
44 associated with intracellular abnormalities characterized by fiber atrophy, oxidative stress,
45 mitochondrial dysfunction, impaired Ca^{2+} homeostasis, elevated proteasome dependent proteolysis, but
46 inhibited autophagy/mitophagy, which we speculate is regulated via a Nox-MuRF1/ubiquitin
47 proteasome-mitochondria-RyR1/ Ca^{2+} signaling axis to offer a novel therapeutic molecular target
48 (*Figure 6*).

49
50

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5

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4

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- 2
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- 23 Volker Adams has nothing to disclose.
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- 1 **Supplemental Materials**
- 2 Expanded Materials & Methods
- 3 Online Tables I - V
- 4 References only in the Supplement ⁵¹⁻⁶¹
- 5
- 6

1 **Figure Legend**

2
3 **Figure 1 – Clinical assessment of respiratory function.** Reduced inspiratory ($P_{i_{max}}$) and expiratory
4 function ($P_{e_{max}}$) in patients with heart failure (CHF) compared to control patients (A). $P_{0.1}$ was equal
5 between groups, but the ratio $P_{0.1}/P_{i_{max}}$ was higher in CHF indicating increased respiratory drive (B).
6 Diaphragm thickness was lower in CHF for both right and left hemi-diaphragm (C). Reduced diaphragm
7 thickness was associated with reduced $P_{i_{max}}$ (D). $P_{i_{max}}$ was associated with reduced vital capacity (E).
8 Number of patients: A-D) control: 21, CHF: 18; E) control: 20, CHF: 17.

9 All p-values are two-sided and were not corrected for multiple testing.

10 Continuous variables were tested for normal distribution applying the Shapiro-Wilk-test.

11 * unpaired Students t-test.

12 † Mann-Whitney-U-Test.

13
14 **Figure 2 – Diaphragm fiber phenotype, ubiquitin-proteasome system, and NADPH oxidase.**

15 Immunostaining of type-1 fibers (brown) in the diaphragm of control (A) and CHF (B). In seven control
16 and six CHF patients, an average of 288 type-1 and 164 type-2 fibers per patient were analyzed;
17 statistical comparisons were made between control and CHF for each fiber type regarding distribution
18 and cross-sectional area (CSA). CHF had a shift towards type-1 fibers (C) with a reduction of CSA in
19 both type-1 (slow) and type-2 (fast) fibers (D). Protein expression of the muscle-specific E3-ligase
20 MuRF1 (E) was higher, whereas the protein expression of another E3-ligase MafBx (F) was lower in
21 CHF vs controls. Protein ubiquitination (G) and proteasome activity (H) were higher in CHF vs controls.
22 Protein expression of the NADPH oxidase isoforms Nox2 (I) and Nox4 (J) were higher in CHF vs
23 controls. The amount of carbonylated myosin heavy chain (MHC) was higher in CHF vs controls (J).
24 Nox2 protein expression was correlated to protein ubiquitination (K).

25 Number of patients: E) control: 16, CHF: 11; F) control: 19, CHF: 12; G) control: 18, CHF: 12; H)
26 control: 16, CHF: 9; I) control: 18, CHF: 12; J) control: 15, CHF: 12; K) control: 6, CHF: 6; L) control:
27 16, CHF: 11.

28 All p-values are two-sided and were not corrected for multiple testing.

29 Continuous variables were tested for normal distribution applying the Shapiro-Wilk-test.

30 * unpaired Students t-test.

31 † Mann-Whitney-U-Test.

32
33 **Figure 3 – Isolated mitochondrial measures.** The activity of citrate synthase (relative to mitochondrial
34 protein content) (A) and mitochondrial size (B) was lower in CHF vs controls. State 3 (C) and state 4
35 respiration (D) of complex I using palmitoylcarnitine/malate as substrate with significantly lower values
36 in CHF vs controls. The respiratory control index (RCI, E) and ADP/O ratio (F) were comparable
37 between both groups. State 3 respiration (glutamate/malate) was associated with exercise capacity (G)
38 and diaphragm thickness (H). Electron micrograph showing normal mitochondria localized in
39 correspondence of sarcomeric Z-disk in a control patient (I, J). In CHF, we found intermyofibrillar
40 mitochondria with small size (K) and prominent accumulation of mitochondria with variable size in
41 subsarcolemmal areas (L).

42 Number of patients: A) control: 12, CHF: 9; B) control: 10, CHF: 5; C) control: 11, CHF: 6; D) control:
43 11, CHF: 6; E) control: 11, CHF: 6; F) control: 11, CHF: 6; G) control: 12, CHF: 6; H) control: 12,
44 CHF: 6; I-L) control: 5, CHF: 5.

45 All p-values are two-sided and were not corrected for multiple testing.

46 Continuous variables were tested for normal distribution applying the Shapiro-Wilk-test.

47 * unpaired Students t-test.

48 † Mann-Whitney-U-Test.

49
50 **Figure 4 – Mitochondrial protein, enzyme activities and morphology.** Protein expression of porin
51 (A) and citrate synthase activity (B, left side) was higher in CHF vs controls indicating increased
52 mitochondrial content. Citrate synthase activity relative to porin expression (B, right side) and succinate
53 dehydrogenase activity relative to porin expression (C, right side) were lower in CHF vs controls,
54 whereas absolute succinate dehydrogenase activity was not different (C, left side). Enzymatic complex
55 I activity was significantly lower in CHF vs controls (D). Protein expression of uncoupling protein 3

1 (UCP3) was higher in CHF (E). Higher protein expression of Mfn2 (F) and Drp1 (G) in CHF vs controls.
2 Protein expression of PGC1alpha was higher in CHF vs controls (H). Both LC3-I/LC3-II ratio (I) and
3 p62 protein expression (J) were higher in CHF vs controls.

4 Number of patients: A) control: 16, CHF: 12; B) control: 17/15, CHF: 9; C) control: 16, CHF: 12; D)
5 control: 14, CHF: 9; E) control: 17, CHF: 12; F) control: 17, CHF: 11; G) control: 15, CHF: 12; H)
6 control: 19, CHF: 12; I) control: 16, CHF: 12; J) control: 17, CHF: 11.

7 All p-values are two-sided and were not corrected for multiple testing.

8 Continuous variables were tested for normal distribution applying the Shapiro-Wilk-test.

9 * unpaired Students t-test.

10 † Mann-Whitney-U-Test.

11

12 **Figure 5 – Markers of calcium homeostasis in the diaphragm.** Immunoprecipitation for RyR1
13 showing reduced binding of FKBP12 in CHF vs controls (A). Protein expression of SERCA-1 (B) and
14 SERCA-2A (C) were higher in CHF vs controls. Calpain activity was not different between CHF and
15 controls (D). Calcium sensitivity was lower in CHF when measured in isolated single type II fibers (E-
16 F).

17 Number of patients: A) control: 4, CHF: 4; B) control: 17, CHF: 13; C) control: 18, CHF: 11; D) control:
18 14, CHF: 9; E-F) control: 12, CHF: 9 (an average of 12 fibers per subject were evaluated).

19 All p-values are two-sided and were not corrected for multiple testing.

20 Continuous variables were tested for normal distribution applying the Shapiro-Wilk-test.

21 * unpaired Students t-test.

22 † Mann-Whitney-U-Test.

23

24

25 **Figure 6 – Proposed mechanisms in Heart Failure associated Diaphragm Dysfunction (HFaDD)**
26 **based on the presented data.** Sarcolemmal-bound Nox2 and sarcoplasmic reticulum (SR) and
27 mitochondrial associated Nox4 are two major sources of reactive oxygen species (ROS) that drive
28 diaphragm weakness via promoting fiber atrophy and contractile dysfunction. ROS-mediated activation
29 of the ubiquitin-proteasome system (UPS) to induce atrophy in a MuRF1-dependent manner, while also
30 inducing protein oxidation to impair key contractile proteins such as myosin heavy chain. Inhibited
31 mitophagy, ROS-mediated mitochondrial dysfunction and damage of the SR to induce calcium efflux
32 via “leaky” ryanodine receptors, which proceeds as a vicious circle to exacerbate mitochondrial and
33 contractile dysfunction.

34

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1 **Novelty and Significance**

3 **What Is Known?**

- 4 • The diaphragm is the main inspiratory muscle responsible for normal ventilation.
- 5 • In chronic heart failure (CHF), diaphragm weakness is associated with a worse prognosis;
- 6 however, we have limited understanding of the basic molecular mechanisms with most of the
- 7 data derived from animal studies.

9 **What New Information Does This Article Contribute?**

- 10 • Clinical, functional (*in vivo/in vitro*), histological/ultrastructural and molecular alterations of
- 11 diaphragm biopsies from CHF patients receiving a left ventricular assist device were compared
- 12 to biopsies of patients without CHF undergoing elective coronary bypass grafting (control).
- 13 • NADPH oxidase isoform Nox2 and Nox4 expression was increased in CHF accompanied by
- 14 catabolic activation and reduced cross-sectional area of type-I and type-II fibers, whereas
- 15 mitochondria exhibited functional and ultrastructural abnormalities characterized by inhibited
- 16 autophagy/mitophagy.
- 17 • Molecular measures were associated with a clinically evident respiratory muscle dysfunction
- 18 and reduced whole body exercise capacity in CHF.

19
20 Diaphragm weakness in CHF is both responsible for aggravated symptoms and is associated with a
21 worse prognosis. The molecular mechanisms in humans with CHF are poorly understood and, therefore,
22 specific treatment options are lacking. Our data derived from *in vivo* measures and molecular analyses
23 of diaphragm biopsies revealed that CHF patients suffers from a diaphragm myopathy that occurs
24 independent of disuse or other confounding factors such as ageing, obesity, or hypertension.
25 Specifically, we confirm that severe cellular remodeling and molecular alterations including fiber
26 atrophy, oxidative stress, proteasome-linked catabolic activation, mitochondrial dysfunction, and
27 impaired Ca²⁺ homeostasis underpin diaphragm weakness and wasting in CHF. Clinically, the molecular
28 alterations closely correlate to both whole-body exercise capacity and respiratory function. Thus, our
29 data reveal, for the first time in humans with heart failure, a molecular mechanism and potential
30 therapeutic target for diaphragm weakness that is centered around a cross-talk of NADPH oxidase
31 derived oxidative stress, catabolic activation via the ubiquitin-proteasome system, quantitative and
32 qualitative mitochondrial disturbances and altered Ca²⁺ signaling.

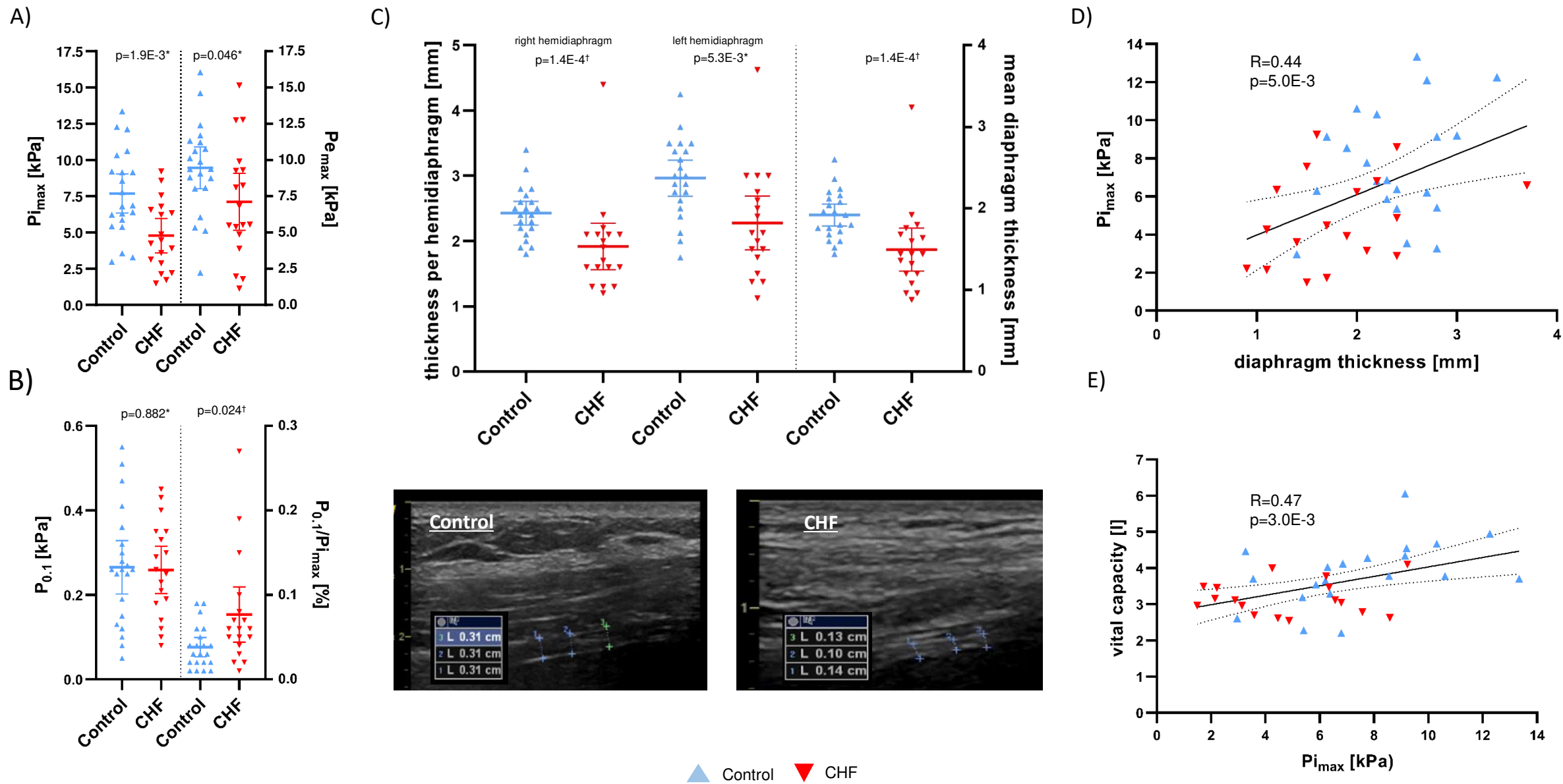


Figure 1

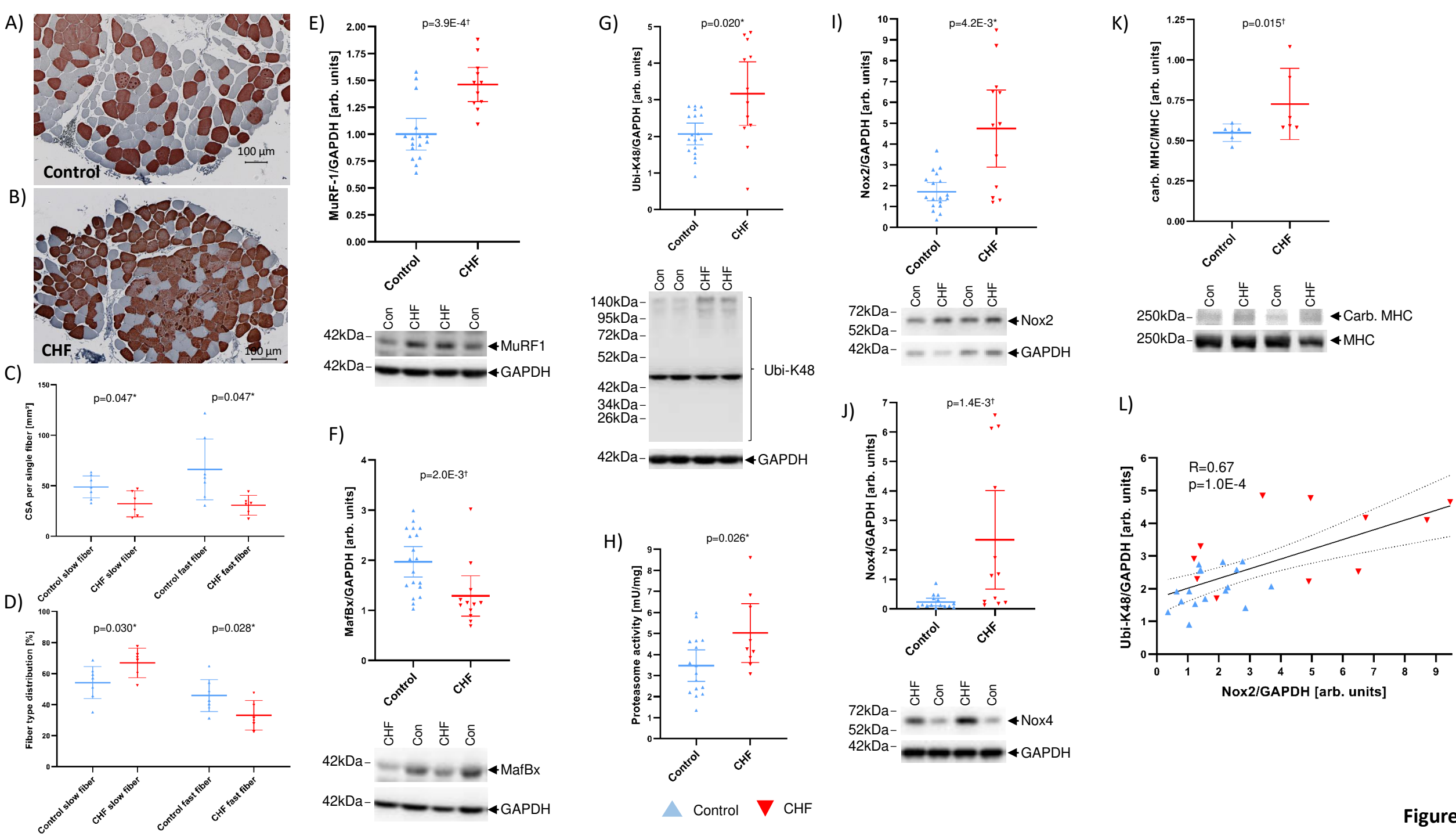


Figure 2

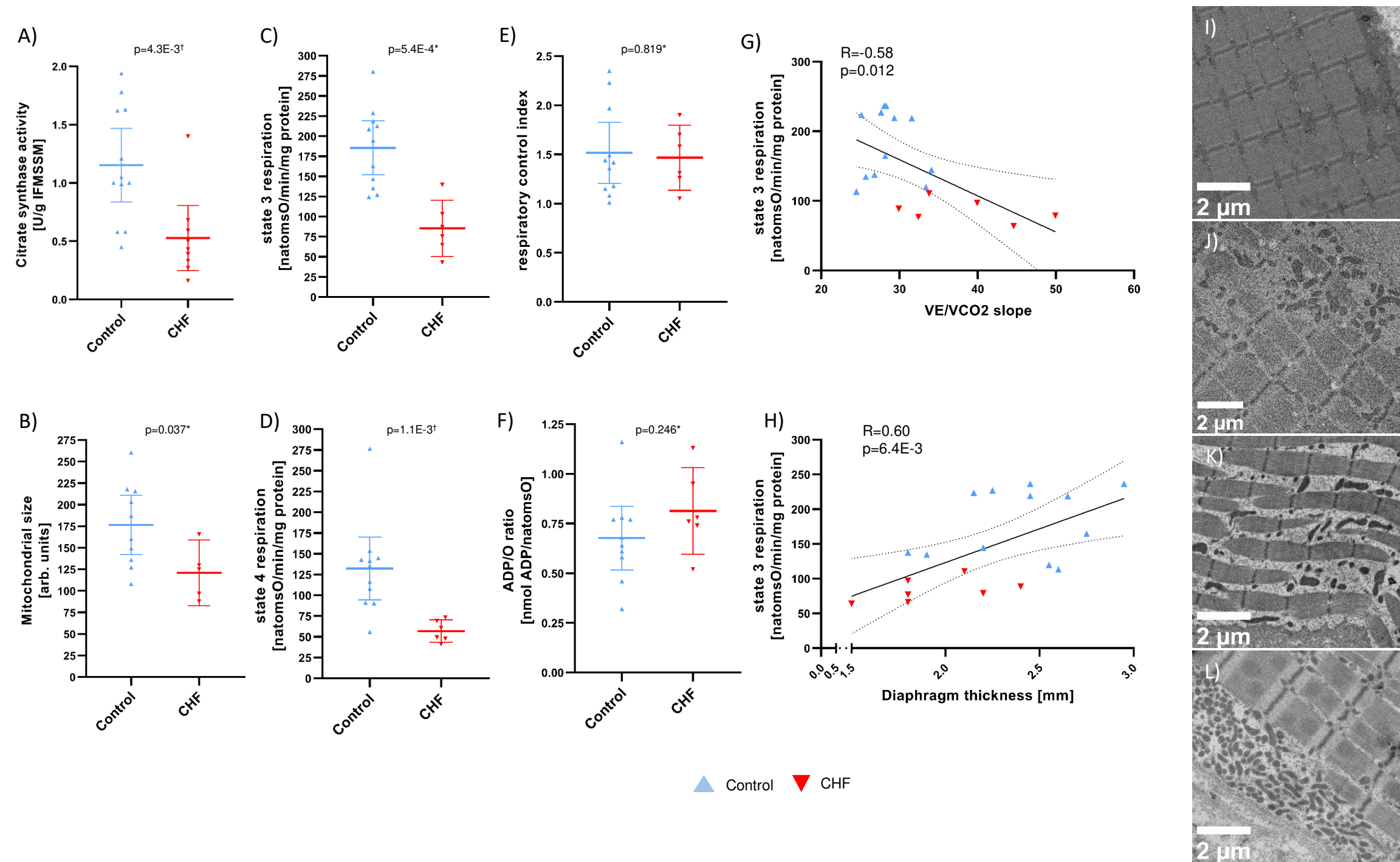
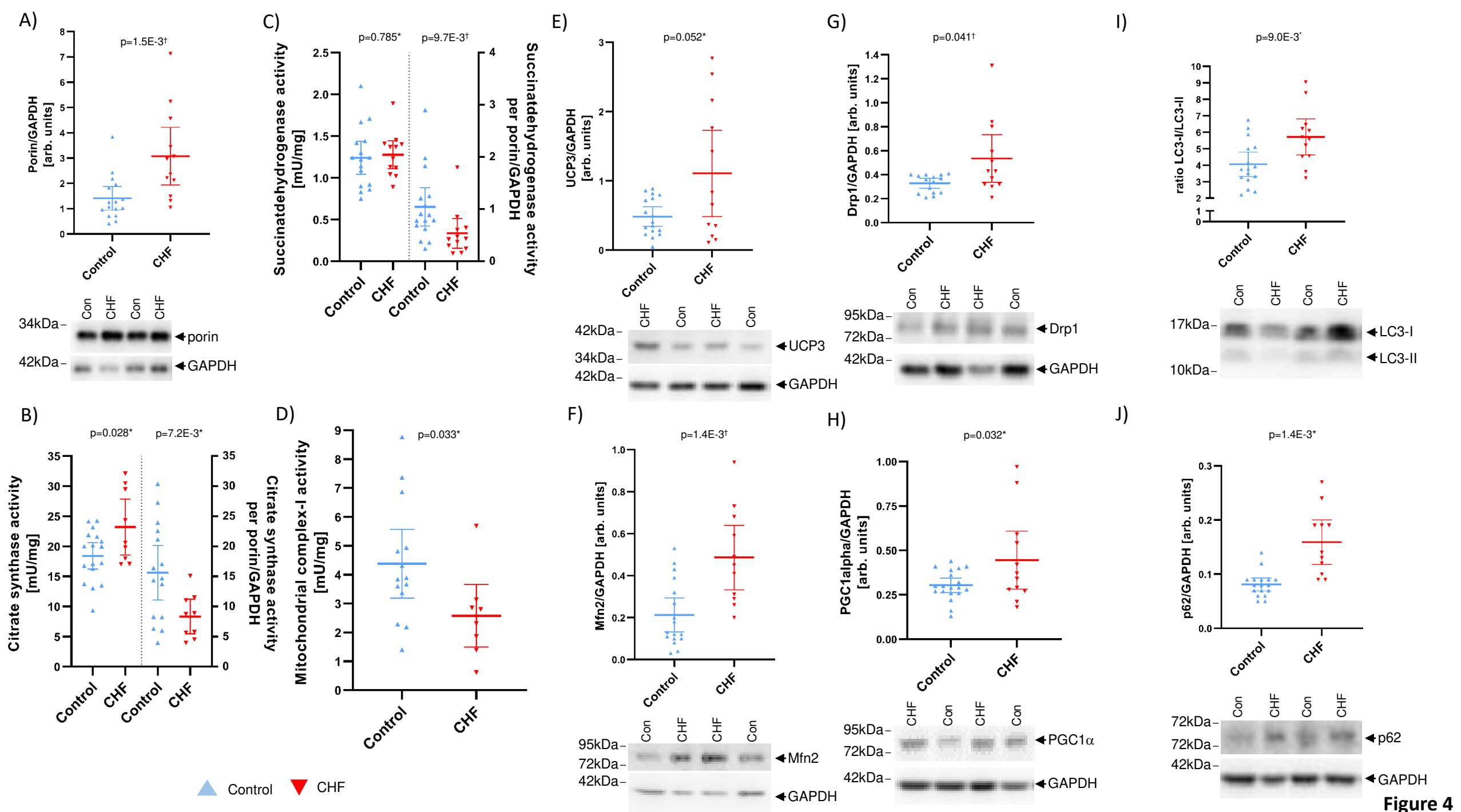


Figure 3



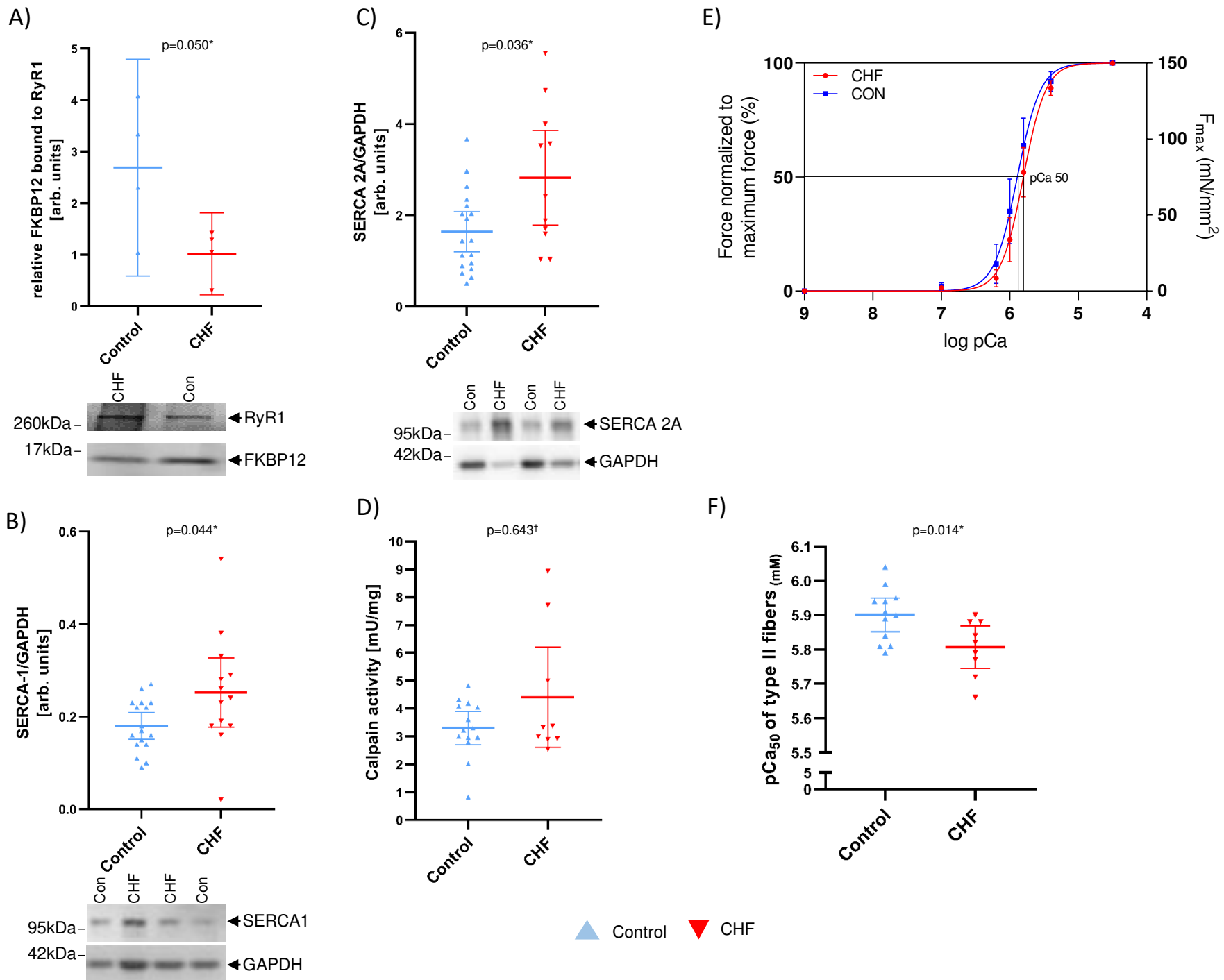


Figure 5

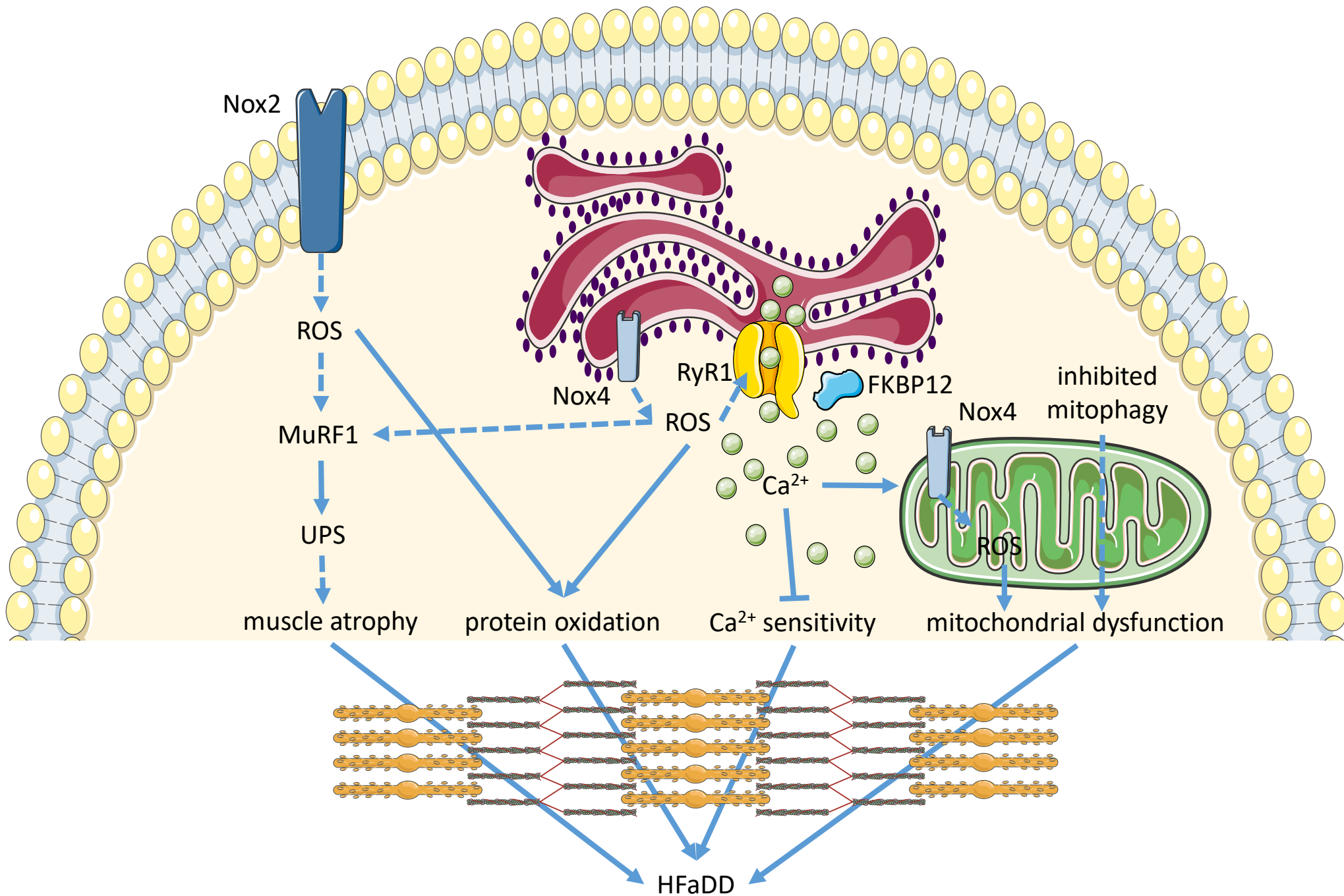


Figure 6