	Subchronic nandrolone administration reduces cardiac oxidative markers during restraint
	stress by modulating protein expression patterns
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1 Abstract

2 Nandrolone decanoate (ND), an anabolic-androgenic steroid prohibited in collegiate and

- 3 professional sports, is associated with detrimental cardiovascular effects through redox-dependent
- 4 mechanisms. We previously observed that high-dose short-term ND administration (15 mg/kg for 2
- 5 weeks) did not induce left heart ventricular hypertrophy and, paradoxically, improved postischemic
- 6 response, whereas chronic ND treatment (5 mg/kg twice a week for 10 weeks) significantly reduced
- 7 the cardioprotective effect of postconditioning, with an increase in infarct size and a decrease in
- 8 cardiac performance.
- 9 We wanted to determine whether short-term ND administration could affect the oxidative redox
- status in animals exposed to acute restraint stress. Our hypothesis was that, depending on treatment
- schedule, ND may have a double-edged sword effect.

12 Measurement of malondialdehyde and 4-hydroxynonenal, two oxidative stress markers, in rat

13 plasma and left heart ventricular tissue, revealed that the levels of both markers were increased in

- 14 animals exposed to restraint stress, whereas no increase in marker levels was noted in animals
- 15 pretreated with ND, indicating a possible protective action of ND against stress-induced oxidative
- 16 damage. Furthermore, isolation and identification of proteins extracted from the left heart
- ventricular tissue samples of rats pretreated or not with ND and exposed to acute stress showed a
- 18 prevalent expression of enzymes involved in amino acid synthesis and energy metabolism. Among
- 19 other proteins, peroxiredoxin 6 and alpha B-crystallin, both involved in the oxidative stress
- 20 response, were predominantly expressed in the left heart ventricular tissues of the ND-pretreated
- 21 rats.
- 22 In conclusion, ND seems to reduce oxidative stress by inducing the expression of antioxidant
- 23 proteins in the hearts of restraint-stressed animals, thus contributing to amelioration of postischemic
- 24 heart performance.
- 25

26 Keywords: anabolic-androgenic steroids; heart; proteomics; oxidative stress; restraint stress

27

28 Introduction

Anabolic-androgenic steroids (AASs), synthetic derivatives of testosterone, exert both androgenic and anabolic actions [1] through their binding to the androgen receptor, acting at both the genomic and non genomic levels. AAS abuse, because of their anabolic effects, is detrimental to health [2-5]; nonetheless, AASs are widely used by athletes to increase muscle mass and power or enhance physical endurance [6]. Human and animal studies have reported discordant results regarding the cardiac and metabolic complications of AAS abuse [5,7-11], with some suggesting that testosterone improves myocardial lipid metabolism [12] and may act as a cardioprotectant and coronary
 vasodilator comparable to conventional anti-ischemic drugs [13,14].

In cardiovascular diseases, both protective and damaging mechanisms derive from redox conditions. For example, exercise protects the heart by increasing myocardial resistance to ischemia-reperfusion (I/R) injury [15,16]. Yet, as a consequence of I/R injury, myocardial redox status is altered because of increased reactive oxygen species (ROS) production. This effect can be counteracted by exercise, which improves antioxidant defenses and reduces infarct size [15-17].

Nandrolone decanoate (ND), one of the most extensively studied AASs, may interfere with redox
conditions [18] and its abuse can limit the cardioprotective effects of exercise. Indeed, chronic ND
administration for 6 to 10 weeks at supraphysiological doses causes heart hypertrophy [19-21],
increases cardiac susceptibility to I/R injury [17,22,23], activates the local renin-angiotensin system
[24], and induces ventricular repolarization disturbances [25].

13 In two previous studies we reported that subchronic treatment with high-dose ND (15 mg/kg for 14 days) does not induce evident left ventricular hypertrophy but, paradoxically, improves cardiac 14 postischemic response [26], whereas chronic ND treatment (10 weeks) significantly reduces the 15 cardioprotective effect of postconditioning, with an increase in infarct size and decrease in cardiac 16 performance [27]. We hypothesized that the double-edged sword effect of ND may be related to 17 treatment schedule and redox conditions. To test this hypothesis, we investigated the effects of 18 subchronic ND treatment on left heart ventricular tissue and on systemic and cardiac redox status in 19 rats exposed or not to restraint stress. Experimental evidence supports the notion that stressful 20 conditions produce oxidative damage and the rationale for using restraint to investigate alterations 21 in redox activities in different tissues. Restraint conditions have been associated with impairment of 22 antioxidant enzyme activity (e.g., superoxide dismutase) and depletion of reduced glutathione 23 [28,29]. Moreover, we previously reported that restraint conditions induce downregulation of 24 25 myocardial contractility response to isoproterenol stimulation which is reversed by ND pretreatment [30]. 26

In this experimental model of subchronic ND treatment in modulating redox status, we measured two oxidative stress markers in plasma and left heart ventricular tissue samples: malondialdehyde (MDA) and 4-hydroxynonenal (4-HNE), the aldehydes resulting from polyunsaturated fatty acid lipid peroxidation. Using a proteomic approach, we identified the protein profile signatures in the left heart ventricular tissues of rats pretreated or not with ND and exposed to acute restraint stress .

32

33 Methods

34 Animals and treatments

1 Treatment schedule: 70 two-month-old male Sprague–Dawley rats weighing 300 g (Harlan

- 2 Laboratories, San Pietro al Natisone, Udine, Italy) were housed in groups of two in polycarbonate
- 3 cages with ad libitum access to food and tap water and maintained in a controlled environment

4 (12:12 light–dark cycle, room temperature 20-24° C, and humidity 50-60%). They were allowed 1

5 week of acclimatization before the experiment began and were handled daily during this period. To

6 avoid circadian variability, all tests were performed between 10:00 and 12:00, when plasma

7 hormones levels are relatively low. Institutional ethical committee approval was obtained for this

8 study; all experimental procedures were performed in accordance with EC Directive 86/609/EEC

9 and Italian law regulating experiments on animals.

10 One group of rats (n=32) received once a day for 14 days an intramuscular (i.m.) injection of 0.5

11 mL of peanut oil (vehicle) containing 15 mg/kg of ND (DECA-Durabolin®, Organon), which is

equivalent to the abuse dose of AAS in humans [31]. Sixteen of these rats (ND group) remained in

their cages until sacrifice, and the 16 other animals were exposed to 1 h of restraint stress 24 h after

the last injection and immediately before sacrifice (ND+S group) [32]. A second group (n=32)

received 0.5 mL peanut oil i.m. injection once a day for 14 days. This group was divided into two

subgroups: 16 were treated with the vehicle only (V group) and were sacrificed, and the 16 other

animals were exposed to the same acute restraint stress protocol described above (V+S group). The

control group (n=6) received i.m. injections of normal saline solution daily for 14 days to check for
possible peanut oil effects. The experimental design is presented in Table 1. At the end of 14 days

20 of treatment, all animals were sacrificed by decapitation and trunk blood was collected. The hearts

21 were rapidly removed, rinsed with saline buffer solution, and dissected to collect the left heart

ventricle. All samples were snap frozen and stored at -80° C until use.

23

24 Oxidative stress marker measurement in plasma and left heart ventricles

Plasma oxidative stress damage was determined by measuring the fluorescent adducts formed 25 between lipid peroxidation-derived aldehydes and plasma proteins. Plasma samples were extracted 26 in ethanol/ether (3:1 v/v) and then centrifuged at 1000 g for 10 min. Pellets were resuspended in 27 ethanol/ether (3:1 v/v); this procedure was repeated three times. Protein adducts were dried under 28 nitrogen and resuspended in a 5% SDS solution. Hydroxynoneal (4-HNE) protein adducts (355 nm 29 excitation/460 nm emission) and malondialdehyde (MDA) protein adducts (390 nm excitation/460 30 nm emission) were measured using a spectrofluorometer (SFM 25 Kontron Instruments AG, 31 Zurich, Switzerland). Data are expressed as arbitrary unit fluorescence (AUF)/mg plasma proteins 32

33 [33].

1 In the left heart ventricle lysates, free MDA levels were determined by measuring thiobarbituric

- 2 acid-reactive substances [34]. Briefly, left heart ventricle tissues were disintegrated with a Mikro-
- 3 Dismembrator II (B. Braun, Melsungen, Germany) and dissolved in Tris-HCl buffer (0.1 M, pH
- 4 7.4) using a Potter-Elvehjem homogenizer (Sigma-Aldrich, St. Louis, MO, USA) The MDA
- 5 concentrations determined by the thiobarbituric acid reaction are expressed as nmol/g tissue.
- 6

7 Tissue damage evaluation

- 8 Tissue damage was evaluated by measuring two cytotoxic markers, lactate dehydrogenase (LDH)
- 9 (Abbott Laboratories, Abbott Park, IL, USA, 2P56-21) and creatine kinase MB isozyme (CK-MB)
- 10 (Abbott, 6K25-30), in blood samples according to the manufacturer's instructions with the aid of a
- 11 clinical chemistry analyzer (Abbott, Architect C8000 Plus).
- 12

13 Protein sample preparation

Frozen left heart ventricular tissue samples were disintegrated with a Mikro-Dismembrator II and
lysed in RIPA buffer (20 mM HEPES, 150 mM NaCl, 5m M EDTA, 1 mM DTT, 1% Triton X-100,
10% glycerol, 1 µg/mL leupeptin, 1 µg/mL aprotinin, 1 µg/mL phenylmethyl–sulfonyl fluoride, 1
mM sodium orthovanadate) at 4° C. The lysates were then spun-down to remove insoluble cellular
debris. Protein concentration was measured by the Lowry method [35].

19

20 Two-dimensional gel electrophoresis (2-DE)

- 21 Two-dimensional gel electrophoresis (2-DE) was performed according to the manufacturer's
- instructions (GE Healthcare, Milan, Italy) with slight modifications. Samples containing 0.125 mg
- proteins were in-gel rehydrated for 12 h onto 7-cm immobilized non-linear pH gradient strips (pH
- 24 3-10), electrofocused at 200 V, 500 V, and 1000 V each for 1 h (step-N-hold), 1000-5000 V
- 25 (gradient) for 30 min, and 5000 V (step-N-hold) for 3 h using an IPGphor system (GE Healthcare).
- 26 The strips were then reduced for 30 min in SDS equilibration buffer (50 mM Tris-HCl pH 8.8, 6 M
- urea, 30% glycerol, 2% SDS) containing 1% (w/v) DTT, followed by 30 min alkylation in the same
- equilibration buffer with 2.5% (w/v) iodoacetamide instead of DTT. Electrophoresis was run on an
- 29 Ettan DALT II system (GE Healthcare). The gels were stained with colloidal Coomassie Blue G250
- 30 (Bio-Rad, Hercules, CA, USA), scanned using image scanner and the gel images were analyzed
- 31 with ImageMaster 2D Platinum 6.0 software (GE Healthcare).
- 32

33 Mass spectrometry and protein identification

Protein spots were excised and in-gel trypsin (Sigma-Aldrich) digested according to the 1 manufacturer's instructions. Aliquots of 1 µL of each sample were mixed with an equal volume of 2 matrix (α -cyano-4-hydroxycinnamic acid), saturated in 50% acetonitrile (Sigma Aldrich), and the 3 mixture was dropped onto the metal target plate of a Microflex LRF MALDI-TOF mass 4 spectrometer (Bruker Daltonics, Germany). Mass spectrometry analysis was carried out in reflector 5 mode. The adrenocorticotropic hormone (2464.1989 m/z) and the angiotensin II fragments 6 (1045.5423 m/z) were used for internal mass calibration to maximize mass accuracy. The peptide 7 mass fingerprint spectra were searched against the MASCOT (Matrix Science) and National Center 8 9 for Biotechnology Information (NCBI, www.ncbi.nlm.gov) protein databases.

10

11 Western blot analysis (WB)

12 Total cellular proteins (25 μg) were resolved by 10% SDS-PAGE and transferred onto

13 nitrocellulose membrane (HybondTM ECLTM, GE Healthcare) for immunoblotting. The membranes

14 were probed with the following antibodies: rabbit polyclonal anti-glutathione-S-transferase (GST)

15 (1:2000, Alpha Diagnostic International, San Antonio, TX, USA, GSTM11-S), rabbit monoclonal

16 anti-CRYAB (1:5000, GeneTex, Irvine CA, USA, GTX62094), rabbit monoclonal anti-PRDX6

17 (1:1000, GeneTex GTX62281), mouse monoclonal anti-alpha tubulin (TUB) (1:10000, Santa Cruz

18 Biotechnology, Dallas, TX, USA, SC-23948) and goat polyclonal anti-vinculin (VCL) (1:1000,

19 Santa Cruz Biotechnology sc-7649). The blots were then incubated with their appropriate secondary

20 antibodies. Protein signals were detected using enhanced chemiluminescence ECLTM WB detection

21 reagents (GE Healthcare). WB images were analyzed using ImageJ software

22 (http://imagej.nih.gov/ij/). CRYAB and PRDX6 were quantified by normalizing them against TUB

selected as the reference housekeeping protein; GST was normalized using VCL.

24

25 Statistical analysis

All data are presented as the mean and the median as a central measure and interquartile range

27 (IQR) as a measure of spread. The Shapiro-Wilk test was applied to check the normality

assumption. Since the MDA concentrations in the left heart ventricular tissue samples had skewed

29 distribution, natural log transformation was applied to achieve normality assumption for standard

- 30 parametric analysis. Two-way analysis of variance (ANOVA) and Tukey's post hoc test were
- carried out in the event of a significant F ratio. Statistical significance was set at p < 0.05. Analyses
- were performed using R version 3.02 (www.r-project.org).
- 33
- 34 **Results**

1 Nandrolone prevents HNE and MDA accumulation

- 2 We measured HNE and MDA adducts in plasma and MDA in its free form in left heart ventricular
- 3 specimens. Since HNE is quickly metabolized to less cytotoxic compounds in many tissues, and
- 4 thus scantly detectable by photometric methods, we measured in left heart ventricular tissue
- 5 samples the protein expression levels of GST, an enzyme involved in its detoxification. Indeed,
- 6 through GST, HNE is rapidly and covalently conjugated to GSH, giving rise to GSH-HNE, which is
- 7 more water soluble and thus more easily excreted.
- 8 A significant increase in the plasma levels of both HNE and MDA adducts was observed in the rats
- 9 exposed to restraint stress (Fig. 1 and Table 2; p=0.04 and p=0.02, respectively). The aldehyde-
- 10 protein adduct levels were lower in the ND+S group than in the V+S group, suggesting that ND
- 11 may prevent against restraint-induced stress effects. However, two-way ANOVA found no
- 12 significant interaction (Table 2). MDA in its free form in the left heart ventricular tissues was
- increased in the vehicle-treated group exposed to restraint stress (V vs V+S, p=0.002), whereas the
- stress-induced MDA increase was significantly lower in the ND+S group (Fig. 2).
- 15 Western blotting indicated a significant increase in GST protein levels after ND treatment in the left
- 16 heart ventricular tissue samples from animals exposed or not to restraint stress, as compared to the
- 17 tissues from animals that had not received ND (V and V+S groups) (Fig. 3).
- 18 The MDA levels in the control group and the V group were similar (data not shown).
- 19

20 Nandrolone treatment impairs blood LDH and CK-MB activity levels

- 21 HNE and MDA are highly toxic side products of lipid peroxidation. To determine whether these
- 22 catabolites might have damaged or altered heart muscle viability, we evaluated the activity of
- 23 lactate dehydrogenase (LDH), a classic indicator of tissue cytotoxicity, and creatine kinase MB
- 24 isozyme (CK-MB), a cardiac-specific cytotoxicity marker. LDH and CK-MB levels were increased
- in the control group exposed to restraint stress as compared to the unstressed group (V vs. V+S
- 26 groups), though the difference between-groups was not statistically significant. Interestingly, LDH
- and CK-MB levels were significantly lower in both ND-treated groups than in the control group
- exposed to restraint stress (V+S) (Fig. 4). Overall, these data strengthen the hypothesis that the
- 29 changes in oxidative stress after exposure to restraint stress did not damage heart muscle integrity.
- 30

31 Left heart ventricular tissue from V+S and ND+S animals displays different proteomic

- 32 signatures
- 33 To gain further insights into the molecular components involved in triggering differential responses
- to ND treatment in the animals exposed to restraint stress, 2-DE was performed on lysates of left

heart ventricular tissue samples from both groups (V+S and ND+S). Representative images are
shown in Fig.5. Comparison between the two groups showed a higher number of protein spots in

the ND+S than in the V+S samples (282 spots in ND+S vs 201 spots in V+S), especially within the

4 range of 20-45 kDa. Eighty-five visible spots were manually excised from the gels and subjected to

5 MALDI-TOF-MS analysis to identify the corresponding proteins. The spots unambiguously

6 identified via MALDI-TOF-MS are numbered in Fig. 5 and the proteins identified (together with

7 their relative scores, sequence coverage, and functional categories) are reported in Table 3. Several

8 spots were identified by MALDI-TOF-MS analysis as being either different isoforms of the same

9 protein or the same protein that migrated differently in other cases, suggesting post-translational

- 10 modifications. Functional category analysis highlighted that the proteins involved in energy
- 11 metabolism were the major components detected.

12 We identified three non-overlapping spots detectable only in the protein profile of the left heart

13 ventricular tissue samples from the ND+S group. Mass spectrometry analysis identified these spots

14 as succinyl–CoA transferase, alpha B-crystallin (CRYAB), and peroxiredoxin 6 (PRDX6), the last

two of which are involved in cellular response to oxidative stress [36,37]. Indeed, CRYAB is a

16 chaperone protein that prevents apoptosis induced by OS and PRDX6 protects against oxidative

17 injury through its peroxidase activity.

These data were corroborated by WB analysis carried-out onof left heart ventricular tissue samples from the four groups (V, V+S, ND, ND+S). The comparison showed that PRDX6 expression levels were higher in both ND-treated groups, whether exposed to restraint stress or not, whereas CRYAB expression was increased in the two groups exposed to restraint stress, with a greater increase in the ND-treated group (Fig. 6).

23

24 Discussion

Our findings indicate that restraint stress impacts on oxidative stress and that ND administration can alter the oxidative status of cardiac tissue in rats exposed to restraint stress. Previous experimental evidence has shown that short-term ND pretreatment can protect against pro-oxidant conditions, such as I/R injury [26]; therefore, we hypothesized that it may exert a protective action against oxidative stress. To settle this issue, and because stressful conditions have been found to cause

30 oxidative damage in diverse tissues, we employed an *in vivo* model in which rats were exposed to

31 restraint stress [28,29,38,39].

32 Our assumption seems to contrast with that of authors reporting a ND-mediated impairment of

exercise-induced cardioprotection [17,24,22,40]. A possible explanation for the protective effect of

1 ND we observed in left heart ventricular tissue is that ND exerts a differential effect in relation to

2 duration of treatment (14 days in our study vs 6 to 8 weeks in those mentioned above).

3 With the present study, we report that oxidative stress marker levels in plasma and left heart

4 ventricular tissues are increased in rats exposed to restraint stress. Although ND per se does not

5 directly modify oxidative stress markers, it appears to prevent an increase in free MDA in the left

6 heart ventricular tissue of restrained animals and that it enables HNE detoxification by means of

7 increased levels of GST. This observation is shared by previous results reporting that ND enhances

8 GST activity [41]. A similar trend was observed for the accumulation of HNE and MDA adducts in

9 the plasma samples. The lack of significant differences between the oxidative stress marker levels

10 in the plasma samples may be due to several reasons; indeed, it cannot be ruled out that oxidative

stress varies in different tissues of the body. Two data sets indicate restraint stress-induced lipid

12 peroxidation: the increased HNE and MDA levels and the appreciable, though statistically not

13 significant, blood accumulation of LDH and CK-MB detected in the V+S group.

14 Overall, the data indicate that, albeit restraint stress led to oxidative damage, it apparently does not

15 affect tissue viability. Interestingly, the opposite behavior of the oxidative stress and cytotoxic

16 markers detected in the ND+S group suggests that ND plays a protective role against the negative

17 effects of oxidative stress.

18 The effects of ND on oxidative stress in the animals exposed to restraint stress may be related to its

19 effect on the corticosterone (CORT) response to stressful conditions. In a previous study, we

20 reported that in rats that received ND treatment identical to the schedule used in the present study

21 CORT levels were reduced in both basal condition and after stress exposure [42]. Indeed,

22 glucocorticoids may affect the redox status of different tissues (e.g., endothelial cells and brain) via

23 different mechanisms, including an increase in superoxide cell production and impairment of tissue

antioxidant capacity [43,44]. Therefore, one can assume that the increase in the two oxidative stress

25 markers in the V+S group is a consequence of the high CORT levels induced by stress and because

ND decreases the CORT levels in plasma after stress, it could conceivably exert a protective action

27 against oxidative stress.

28 We identified two additional novel players implicated in oxidative stress response to subchronic ND

29 pretreatment: CRYAB and PRDX6. These two proteins, which were expressed at higher levels in

- the ND+S as compared to the V+S samples, are both involved in oxidative response. PRDX6, a
- class of thiol-specific antioxidant enzymes, exerts peroxidase activity against a broad range of
- peroxides [36,45-47]. CRYAB is activated by phosphorylation during oxidative stress, thus
- protecting cells from apoptosis induced by oxidative damage [48-50]. Previous studies have shown
- that transgenic mice that ubiquitously overexpress CRYAB have increased tolerance to I/R injury

1 [51], whereas CRYAB/HSPB2-null mouse hearts display poorer functional recovery, a higher cell

- 2 death rate [52], increased stiffness, and, hence, poor myocardial relaxation following I/R injury [53]
- 3 as compared with wild-type controls. Of note, we previously observed an improvement in

4 postischemic diastolic and systolic cardiac functions after subchronic ND treatment [26].

5 To elucidate to what extent the effect on PRDX6 and CRYAB expression was due to ND

6 administration by itself or ND coupled to stress, we compared their expression in samples from the

7 four groups (V, V+S, ND, ND+S) by WB analysis. ND treatment alone was enough to induce up-

- 8 modulation of both PRDX6 and CRYAB.
- 9 We previously reported that subchronic ND treatment improves systolic function induced by
- 10 postconditioning cardioprotection via β 2-adrenoreceptor activation [26] and probably, via the same
- 11 pathway, prevents the decrease induced by restraint stress in heart contractile response to
- isoproterenol [30]. These findings appear to be in line with results from studies on other antioxidant
- 13 substances [54,55]. Vitamin C is known to enhance the positive inotropic response to the β 1-
- adrenoreceptor agonist dobutamine in the presence of high levels of reactive oxygen species [54]
- and, in the same condition, the free radical scavenger N-acetylcysteine was found to strengthen the
- 16 inotropic effect of dobutamine in isolated heart tissue [55]. In our experimental conditions, it is
- 17 unlikely that ND had a direct antioxidant action. ND-pretreatment affects the protein expression
- 18 patterns, as revealed by 2-DE mapping.
- 19 In conclusion, our study provides further evidence for an enhanced oxidative response in animals
- 20 exposed to restraint stress and reports the novel observation that subchronic ND-pretreatment limits
- 21 myocardial redox response induced by restraint *via* the modification of anti-oxidant protein

22 expression. These effects may explain the amelioration of postischemic heart performance observed

- after ND administration [26]. It can be argued that ND acts as a double-edged sword, with positive
- 24 or negative effects on the heart depending on the treatment schedule.
- 25

26 **Conflict of interest**

- 27 The Authors declare that they have no conflict of interest.
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- Fig. 1 Restraint stress increases plasma HNE and MDA adduct levels. The boxes indicate the lower
 and upper quartiles, and the central line is the median. The dashed line represents the mean. The
- 3 whiskers above and below the boxes indicate a distance of 1.5 IQR. *p<0.05.
- 4 HNE denotes 4-hydroxynoneal; MDA malondialdehyde; IQR interquartile range.
- 5
- Fig. 2 Free MDA levels in left heart ventricle are lowered by ND treatment upon restraint stress. 6 Box plot of MDA levels in left heart ventricular tissue (log(nmol/g)) in ND-treated and control 7 animals. The boxes indicate the lower and upper quartiles, and the central line is the median. The 8 9 dashed line represents the mean. The whiskers above and below the boxes indicate a distance of 1.5 IQR. Two-way ANOVA and post hoc Tukey test showed significant differences between ND and 10 ND+S (p=0.027), V and V+S (p=0.002), ND+S and V+S (p=0.001). 11 MDA denotes malondialdehyde; ND nandrolone decanoate; V vehicle-treated group; V+S vehicle-12 13 treated group exposed to restraint stress; ND nandrolone-treated group; ND+S nandrolone-treated group exposed to restraint stress; IQR interquartile range. 14 15 Fig. 3 Increased GST protein levels in left heart ventricular tissue samples from ND-treated animals. 16 Representative Western blot image showing quantification of GST protein levels. 17 Fig. 4 Measurement of blood LDH and CK-MB activity. Box plot of LDH (panel A) and CK-MB 18 isozyme (panel B) activity levels (U/L). The boxes indicate the lower and upper quartiles. The 19 central and the dashed lines represent the median and the mean values, respectively. The whiskers 20 above and below the boxes indicate a distance of 1.5 IQR. Two-way ANOVA and post hoc Tukey 21 test showed significant differences between the V+S vs ND groups and the V+S vs ND+S 22 groups.Fig. 5 Two-dimensional gel electrophoresis (2-DE) displaying different patterns in the 23 lysates obtained from left heart ventricular tissue samples from the two groups exposed to restraint 24 stress (V+S and ND+S). Proteins from V+S (panel A) and ND+S (panel B) left heart ventricular 25 tissues were separated by isoelectric focusing on 7-cm strips with a non linear 3 to 10 pH gradient 26 and subjected to SDS-polyacrylamide gel electrophoresis. Gels were stained with Coomassie Blue 27 28 (G250). The numbers indicate the spots analyzed by mass spectrometry. Molecular size markers are
- indicated on the left.
- 30

Fig. 6 Immunoblots of PRDX6 and CRYAB proteins in left heart ventricular tissue. Histograms
show the relative concentrations of the proteins normalized to the house-keeping protein tubulin
(TUB).

- 1 PRDX6 denotes peroxiredoxin 6; CRYAB alpha B-crystallin; V vehicle-treated group; V+S
- 2 vehicle-treated group exposed to restraint stress; ND nandrolone-treated group; ND+S nandrolone-
- 3 treated group exposed to restraint stress. Two-way ANOVA and post hoc Tukey test showed
- 4 significant differences between the groups. PRDX6 *p<0.05 vs V and V+S, n=3; CRYAB *p<0.05
- 5 *vs* V; #p<0.05 *vs* V+S, n=3.
- 6 The immunoblots are from one representative experiment.
- 7
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