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# Sodium bicarbonate ingestion prior to training improves mitochondrial adaptations in rats

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We tested the hypothesis that reducing hydrogen ion accumulation during training would result in greater improvements in muscle oxidative capacity and time to exhaustion (TTE). Male Wistar rats were randomly assigned to one of three groups (CON, PLA, and BIC). CON served as a sedentary control, whereas PLA ingested water and BIC ingested sodium bicarbonate 30 min prior to every training session. Training consisted of seven to twelve 2-min intervals performed five times/wk for 5 wk. Following training, TTE was significantly greater in BIC ( $81.2 \pm 24.7$  min) compared with PLA ( $53.5 \pm 30.4$  min), and TTE for both groups was greater than CON ( $6.5 \pm 2.5$  min). Fiber respiration was determined in the soleus (SOL) and extensor digitorum longus (EDL), with either pyruvate (Pyr) or palmitoyl carnitine (PC) as substrates. Compared with CON ( $14.3 \pm 2.6$  nmol O<sub>2</sub> · min<sup>-1</sup> · mg dry wt<sup>-1</sup>), there was a significant greater SOL-Pyr state 3 respiration in both PLA ( $19.6 \pm 3.0$  nmol O<sub>2</sub> · min<sup>-1</sup> · mg dry wt<sup>-1</sup>) and BIC ( $24.4 \pm 2.8$  nmol O<sub>2</sub> · min<sup>-1</sup> · mg dry wt<sup>-1</sup>), with a significantly greater value in BIC. However, state 3 respiration was significantly lower in the EDL from both trained groups compared with CON. These differences remained significant in the SOL, but not the EDL, when respiration was corrected for citrate synthase activity (an indicator of mitochondrial mass). These novel findings suggest that reducing muscle hydrogen ion accumulation during running training is associated with greater improvements in both mitochondrial mass and mitochondrial respiration in the soleus.

mitochondrial respiration; state 3 respiration; muscle pH; citrate synthase; muscle buffer capacity

MITOCHONDRIA ARE CENTRAL TO THE CONVERSION of energy by oxidizing substrates and generating the cell fuel ATP. During steady-state exercise, the rate of mitochondrial ATP production is closely matched to the rate of ATP hydrolysis, and this demonstrates the existence of efficient cellular mechanisms to control mitochondrial ATP synthesis. However, there is evidence that the maximal rate of mitochondrial ATP production ( $V_{\max}$ ) is in excess of that which is required during exercise with large muscle groups (e.g., 2-legged exercise) (52). Nonetheless, despite this apparent overcapacity, there are further

increases in  $V_{\max}$  with endurance training (14, 53, 59). Although this adaptation will have little influence on whole body maximal oxygen utilization, it will play a major role in reducing metabolic perturbations and increasing time to exhaustion (TTE) during submaximal exercise (18). In contrast, reduced mitochondrial respiration appears to provide an important mechanism that links a low aerobic capacity to the pathogenesis of cardiovascular disease (63) and insulin resistance (33). Regularly performed exercise can result in a rapid increase in the activities of oxidative enzymes (26), mitochondrial density (56), and mitochondrial respiration (14, 49, 53, 54, 60). However, given the importance of mitochondrial respiration for both performance and health, further research is required to determine factors that regulate training-induced changes in mitochondrial respiration. One potential factor, unexplored to date, is the degree of acidosis experienced during training. We have reported recently that reducing H<sup>+</sup> accumulation during training (via pretraining ingestion of sodium bicarbonate, NaHCO<sub>3</sub>) resulted in greater improvements in both short-term endurance and the lactate threshold in humans (20). Because the lactate threshold has previously been correlated with mitochondrial respiration (28), we hypothesized that this finding may have been due to the positive effects of reducing H<sup>+</sup> accumulation during training on training-induced changes in mitochondrial respiration.

When a molecular view on training adaptation is taken, it is apparent that adaptations to training are the consequence of changes in gene expression that lead to the accumulation of specific proteins. It has been shown that the muscle environment (e.g., low glycogen) is a determining factor for the transcription of some genes in response to training (40). Recent research suggests that cellular pH may also affect the training-induced expression of some genes, in particular, nuclear-encoded genes that have been proposed to regulate mitochondrial biogenesis [e.g., peroxisome proliferator-activated receptor- $\gamma$  coactivator-1 $\alpha$  (PGC-1 $\alpha$ )] (5, 38). Such findings further suggest that reducing H<sup>+</sup> accumulation during training may promote mitochondrial adaptations.

Therefore, the present study investigated for the first time the effects of altering muscle pH during training (via NaHCO<sub>3</sub> ingestion) on citrate synthase (CS) activity (a marker of mitochondrial mass) and mitochondrial respiration in rat skeletal muscle. Although many previous studies have investigated the effects of training on the function of isolated mitochondria (7, 30, 43), the structure of the mitochondrial membrane and the

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Table 1. Summary of the training performed by both training groups during the experimental period

Training Weeks	1	2	3	4	5
No. of intervals/training session	7	7	9	10	12
Mean speed/wk, m/min	33	35	39	44	48
Minimal speed at the beginning of the training session, m/min	28	28	31	31	37
Maximal speed reached during the training session, m/min	37	40	43	49	52
Mean running distance/training session, m	435	489	731	962	1,202

function of the isolated mitochondria may be altered by the isolation procedure (45). Therefore, we chose to investigate the effects of training on mitochondrial respiration in saponin-permeabilized fibers, since it is possible to study mitochondria in situ, keeping their natural environment (46). We hypothesized that rats ingesting  $\text{NaHCO}_3$  prior to every training session (BIC) would have a significantly greater maximal, ADP-stimulated (state 3) respiration than either sedentary rats (CON) or rats ingesting water prior to training (PLA). In addition, we hypothesized that, despite performing identical training, BIC would have a greater submaximal TTE than PLA.

## METHODS

### Ethical Approval

The experimental protocol was approved by the Research Ethics Committee of the University of Montpellier and conducted according to the guidelines of the National Research Council for the Care and Use of Laboratory Animals and French law on animal handling and protection.

### Animals

Twenty-one male Wistar rats (Charles River Laboratories, St. Germain sur l'Arbresle, France) were housed individually in a temperature-controlled room and maintained with food and drink ad libitum in a 12:12-h light-dark cycle (lights on at 7 PM), thus allowing exercise during their active phase. All exercise tests were conducted in the daytime, but a dark cover was placed over the treadmill to ensure that the rats exercised in darkness. The body mass of the rats was monitored throughout the experimental period.

### Experimental Design

All animals were familiarized with a motor-driven treadmill for 3–4 days, 5 min/day, on a 10% grade. The running speed was initially 15 m/min and was gradually increased to 25 m/min within 2 days. Following the familiarization, all animals performed an incremental test to exhaustion on a treadmill inclined to  $15^\circ$  [starting at 10 m/min with increments of 3 m/min every 2 min (24)]. At least 72 h later, the rats performed a TTE test at the individually determined maximal velocity reached during the incremental test. Exhaustion in both tests was defined as the inability of the rats to run on the treadmill despite small electric shocks and an inability to upright themselves when placed on their back.

Rats were then randomly assigned to one of the following three groups: a control group that performed no training (CON;  $n = 7$ ), a high-intensity training group supplemented with a placebo (PLA;  $n = 7$ ), or a high-intensity training group supplemented with sodium bicarbonate (BIC;  $n = 7$ ). At least 72 h after the last training session, rats performed a TTE test at the pretraining maximal velocity reached during the incremental test (without supplementation). Seventy-two hours after the last exercise, all rats were euthanized by cervical dislocation. One of the soleus (SOL) and extensor digitorum longus (EDL) muscles was quickly removed, frozen in liquid nitrogen, and stored at  $-80^\circ\text{C}$  until biochemical analysis. Before freezing, a section of the sample was mounted in embedding medium and frozen in

isopentane cooled to its freezing point in liquid nitrogen. This sample was stored at  $-80^\circ\text{C}$  until later histochemical analysis. The same muscles were removed from the contralateral hindlimb, placed in an ice-cold preservation buffer on ice, and immediately prepared for fiber respiration measurements.

### Training Intervention

As shown in Table 1, training consisted of seven (1st wk) to 12 (5th wk) 2-min intervals (interspersed with 1 min of rest) performed five times/wk for 5 wk. The intensity of the intervals was initially set at 80% of the peak speed reached for each rat during the incremental test and was increased by 10% each week. Rats were matched, on the basis of the peak speed reached during the incremental test, with a rat from the alternate training group, and both rats trained concurrently on the treadmill for each training session to ensure that all pairs performed identical amounts of training.

### Supplementation

Thirty minutes prior to each training session, PLA received water, whereas BIC received  $\text{NaHCO}_3$  (0.05 g/kg body mass) administered by esophageal catheter. The timing and dose were based on our pilot work, which showed that there was a significant increase in blood pH 15–30 min after the administration of  $\text{NaHCO}_3$ , which was maintained to a steady state for  $>60$  min (Fig. 1) (51). In addition, our further pilot work demonstrated that the decrease in muscle pH immediately following a typical training session (6 X 2-min intervals interspersed with 1 min of rest) was significantly less following the administration of  $\text{NaHCO}_3$  than without in the SOL [ $6.83 \pm 0.04$  (range: 6.76–6.87) vs.  $6.94 \pm 0.03$  (range: 6.90–6.97) pH units,  $P < 0.05$ ] and tended to be less in the EDL [ $6.60 \pm 0.04$  (range: 6.54–6.66) vs.  $6.65 \pm 0.04$  (range: 6.58–6.70) pH units in the EDL,  $P > 0.05$ ]. Although the range of pH values is greater than that which we have observed in our previously published human studies, we believe that this can be explained by the time required to remove the rat from the treadmill, perform the cervical dislocation, remove the fur, extract the muscles, and place them in liquid nitrogen. Because muscle pH can increase by 0.1 of a pH unit within 30 s postexercise (6), small differences in any of these postexercise procedures will obviously contribute to the variability of the reported measurements.

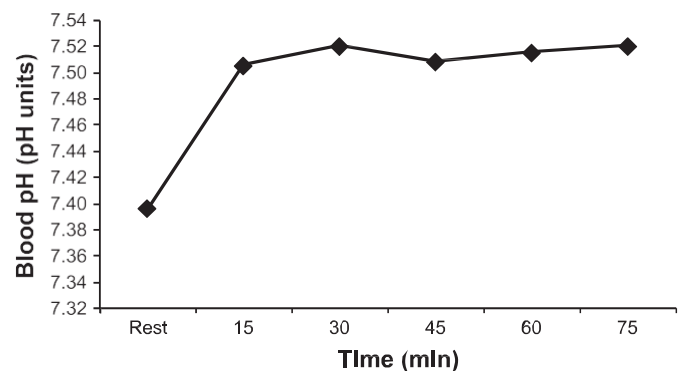


Fig. 1. Evolution of blood pH after administration of sodium bicarbonate ( $\text{NaHCO}_3$ ) by esophageal catheter.

**Histochemical determination of fiber type.** Serial transverse sections (10- $\mu$ m-thick cross-sections) were cut in a cryostat at  $-20^{\circ}\text{C}$  and mounted on glass coverslips. The samples were stained for myofibrillar ATPase according to the procedure of Mabuchi and Sreter (34). The stained slides were photographed with a CMOS Pro camera (Sound Vision) connected to Nikon eclipse 400 microscope (Nikon, Tokyo, Japan). Computer image analysis was performed using the National Institutes of Health Image Analysis software (National Institutes of Health, Bethesda, MD), and muscle samples were classified as types I, Ila, and I Ib.

**Muscle buffer capacity.** Resting muscle samples (45–50 mg) were homogenized on ice for 2 min in a solution containing sodium fl (10 mM) at a dilution of 25  $\mu$ l homogenizing solution/mg wet wt muscle (based on Ref. 35). The muscle homogenate was then placed in a circulating water bath at  $37^{\circ}\text{C}$  for 5 min prior to and during the measurement of pH. The pH measurements were made with a microelectrode (MI-415; Microelectrodes, Bedford, NH) connected to a pH meter (SA 520; Orion Research, Cambridge, MA). After initial pH measurement, muscle homogenates were adjusted to a pH of 7.2 with a sodium hydroxide (0.1 M) solution and then titrated to a pH of 6.2 by the serial addition of 10  $\mu$ l of hydrochloric acid (0.01 M). From the titration trend line, the number of moles of  $\text{H}^{+}$  (per gram of wet muscle) required to change the pH from 7.1 to 6.5 was interpolated. This value was then normalized to the whole pH unit for fi display as micromoles  $\text{H}^{+}$  per gram wet muscle per unit pH ( $\mu\text{mol H}^{+} \cdot \text{g muscle wet wt}^{-1} \cdot \text{pH}^{-1}$ ) and determined as the rat's muscle buffer capacity ( $r3m_{\text{in vitro}}$ ).

**CS and 6-phosphofructokinase activities.** Homogenates for CS activity were prepared in buffer (in mM: 210 sucrose, 2 EGTA, 40 NaCl, 30 HEPES, 5 EDTA, and 2 phenylmethylsulfonyl fl pH 7.4) and stored at  $-80^{\circ}\text{C}$ . CS activity was assayed by a spectrophotometric method according to Sreter (48). Changes in absorbance were recorded over 3 min at 412 nm and at  $25^{\circ}\text{C}$ . Phosphofructokinase (PFK) activity was determined on fresh SOL and EDL homogenates according to Opie and Newsholme's (37) method and followed the rate of decrease in NADH at 340 nm and at  $25^{\circ}\text{C}$ .

#### Fiber Respiration

**Preparation of permeabilized fibers.** Fibers were isolated from the SOL and the EDL muscles and each placed into 5 ml of buffer A [10 mM EGTA, 5 mM  $\text{MgCl}_2$ , 20 mM taurine, 0.5 mM DTT, 5 mM ATP, 20 mM imidazole, 0.1 M MES potassium salt (KMES), 15 mM phosphocreatine, pH 7.0] with 50  $\mu\text{g/ml}$  saponin for 20 min at  $4^{\circ}\text{C}$ . The saponin was then removed by rinsing the fi at two 10-min intervals at  $4^{\circ}\text{C}$  using 2 X 5 ml of ice-cold buffer B (10 mM EGTA, 5 mM  $\text{MgCl}_2$ , 20 mM taurine, 0.5 mM DTT, 20 mM imidazole, 0.1 M KMES, 3 mM phosphate, 5 mg/ml bovine serum albumin, pH 7.0).

**Assessing permeabilized fiber oxygen consumption.** For all animals, the respiratory parameters of the total mitochondrial population of SOL and EDL muscle fibers were studied in situ using a Clark electrode (Strathkelvin Instruments, Glasgow, UK). Measurements were carried out at  $30^{\circ}\text{C}$  with continuous stirring in 2 ml of the oxygraph solution with different respiratory substrates. The oxygen concentration in the medium used was 150 nmol/ml. It consisted of buffer B supplemented with 20  $\mu\text{M}$  EDTA, 50  $\mu\text{M}$  AP<sub>5</sub>A (an inhibitor of adenylate kinase), and 1 mM iodoacetate and is referred to here as buffer B=. Eight to 10 mg wet weight of permeabilized SOL and EDL fibers was placed into their respective oxygen chambers along with 2 ml of buffer B= and respiratory substrates. Two metabolic pathways were investigated by using two pairs of respiratory substrates. One set of measurements was made using 10 mM of pyruvate plus 10 mM of malate (Pyr); the procedure was then repeated using 10 mM of

palmitoyl carnitine plus 10 mM malate (PC), producing a second set of measurements for each muscle. A low oxygen consumption rate (state 2) was left to establish over a 5-min period, and then 2 mM of ADP was injected into the chambers, resulting in an increase in respiration (state 3). After 5 min of state 3 respiration, respiration was inhibited using 70  $\mu\text{M}$  atractyloside (inhibitor of mitochondrial inner membrane ATP/ADP exchange) and 1  $\mu\text{M}$  oligomycin (ATP synthase inhibitor). The respiratory control ratio (RCR) was also calculated and is the quotient of state 3 respiration (with ADP in excess) over that of stage 4 (respiration inhibited).

#### Statistical Analysis

Descriptive statistics are expressed as means  $\pm$  SD. One-way ANOVA was performed to compare the effects of high-intensity training (comparison with CON) and the effects of  $\text{NaHCO}_3$  administration (PLA vs. BIC). Individual relationships between variables were studied by means of linear regressions. SigmaStat software was used for all tests, and the level of significance was set at  $P < 0.05$ .

## RESULTS

### Body Mass

The mean body mass of the three groups of rats before training was  $235 \pm 17$  g. After 5 wk of treadmill training, the body masses of the PLA and BIC rats were  $376 \pm 12$  and  $375 \pm 18$  g, respectively, with no significant difference between groups. After 5 wk of caged living, the control rats ( $426 \pm 24$  g) weighed significantly more than both training groups.

### CS and PFK Enzyme Activities

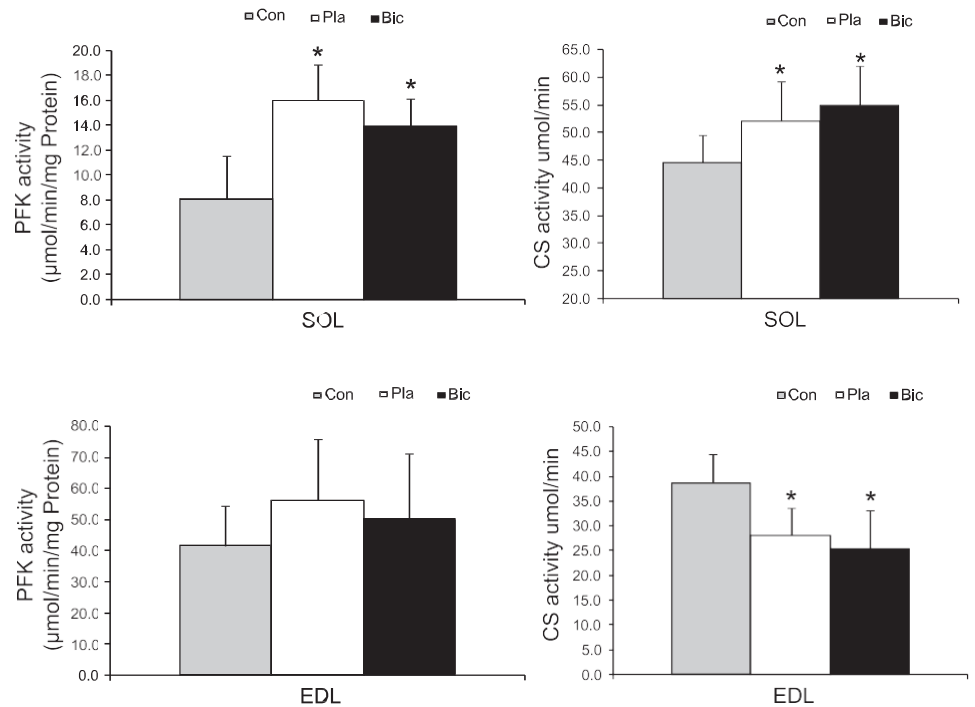
As shown in Fig. 2, compared with CON, the activity of CS in the trained groups was higher for both PLA and BIC in the SOL ( $P < 0.05$ ) and significantly lower for both training groups in the EDL. There was also significantly greater PFK activity ( $P < 0.05$ ) in the SOL only for both trained groups, without a significant main effect for groups in the EDL ( $P = 0.35$ ).

### Fiber Respiration

**SOL.** Compared with CON, both training groups had a significantly greater SOL-Pyr state 3 respiration ( $+39 \pm 26$  and  $+74 \pm 29\%$  for PLA and BIC, respectively,  $P < 0.05$ ), with a significantly greater value in BIC compared with PLA ( $P = 0.009$ ) (Fig. 3). When the respiration was related to CS activity, a marker of mitochondrial mass, respiration remained significantly higher for BIC compared with CON ( $P = 0.006$ ). Compared with CON, the RCR in SOL-Pyr was significantly greater in both the PLA ( $+43 \pm 25\%$ ,  $P < 0.001$ ) and the BIC ( $+77 \pm 22\%$ ,  $P < 0.001$ ), with a significant difference between the two training groups ( $P = 0.016$ ). There were no significant differences between the groups for any of the respiration measurements for SOL-PC.

**EDL.** Compared with CON, there was a significantly lower state 3 respiration in the two training groups when either Pyr ( $-32 \pm 11$  and  $-33 \pm 11\%$  for PLA and BIC, respectively,  $P = 0.002$ ) or PC ( $-23 \pm 23$  and  $-32 \pm 12\%$  for PLA and BIC, respectively,  $P = 0.009$ ) was used as substrate; there were no significant differences between the two training groups for either substrate. A similar pattern was observed for state 4 respiration. However, there were no

Fig. 2. Citrate synthase (CS) and phosphofructokinase (PFK) enzyme activities in the 3 groups [a control group that performed no training (CON), a high-intensity training group supplemented with a placebo (PLA), and a high-intensity training group supplemented with sodium bicarbonate (BIC)] for both the soleus (SOL) and the extensor digitorum longus (EDL) muscles. Data are means  $\pm$  SD;  $n = 7$  for each group. \*Signifi different from CON.



signifi differences between the three groups for state 3 respiration expressed relative to CS activity, or for the RCR, using either substrate.

13 $m_{in vitro}$ . Compared win CON, r3 $m_{in vitro}$  was significantly greater in both training groups (PLA and BIC) in both the soleus ( $+40 \pm 29$  and  $+34 \pm 19\%$  for PLA and BIC, respectively,  $P = 0.001$ ) and in the EDL ( $+22 \pm 15$  and  $+22 \pm 29\%$  for PLA and BIC, respectively,  $P = 0.02$ ) (Fig. 4).

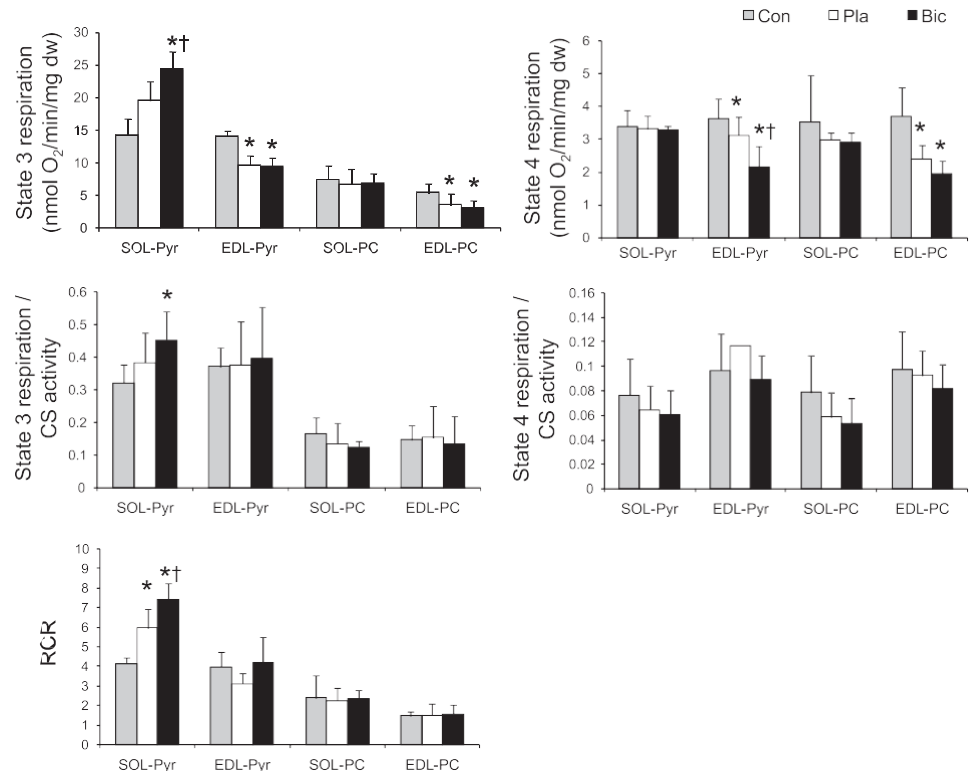
#### Fiber Composition

There were no significant differences in fiber composition between the three groups (Fig. 5).

#### TTE

Following training, TTE was significantly greater in BIC ( $81.2 \pm 24.7$  min) compared with PLA ( $53.5 \pm 30.4$  min,  $P =$

Fig. 3. State 3 respiration, state 3 respiration expressed relative to CS activity, state 4 respiration, state 4 respiration expressed relative to CS activity, and the respiratory control ratio (RCR) in the 3 groups in both the SOL and the EDL when either pyruvate + malate (Pyr) or palmitoyl carnitine + malate (PC) is used as substrates. \*Significantly different from CON; †significantly different from PLA. Data are means  $\pm$  SD;  $n = 7$  for each group.



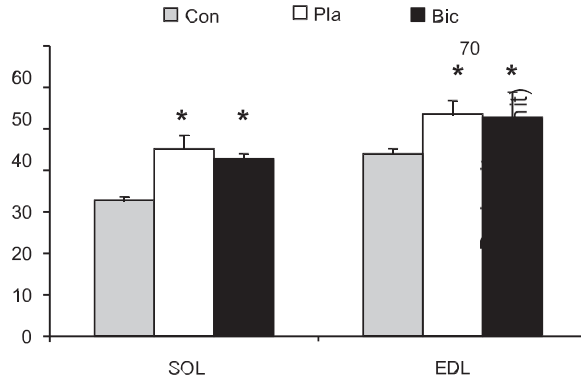


Fig. 4. Muscle buffer capacity ( $r3_{\text{in vitro}}$ ) in the 3 groups for both the SOL and the EDL muscles. \*Significantly different from CON. Data are means  $\pm$  SD;  $n = 7$  for each group.

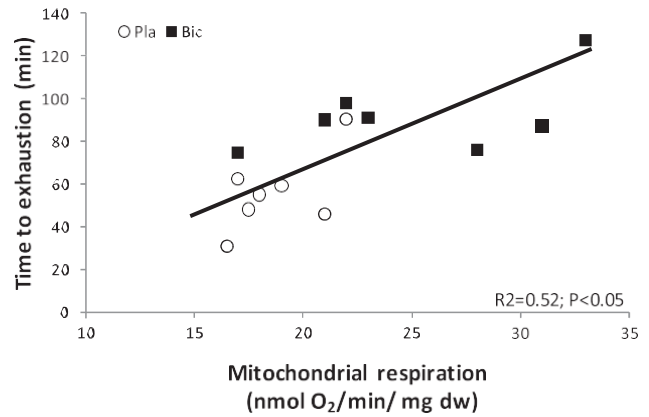


Fig. 6. Scatter plot showing the relationship between state 3 respiration in the SOL (with pyruvate and malate as substrates) and time to exhaustion in trained rats ( $n = 14$ ).

0.038), and TTE for both groups was greater than CON ( $6.5 \pm 2.5$  min,  $P < 0.05$ ). There was a significant correlation between state 3 respiration in the SOL (using PC as the substrate) and TTE in the two training groups (Fig. 6).

## DISCUSSION

In the present study, training clearly increased muscle oxidative capacity in the soleus, as indicated by the greater state 3 fiber respiration (only with pyruvate as a substrate) and by the greater CS activity in trained rats. However, the main finding was that there was a significantly greater state 3 respiration in the soleus muscle of rats provided with  $\text{NaHCO}_3$  before every training session. We also report that there was a significant correlation between the state 3 respiration in the soleus and time to exhaustion.

In this study, we wanted to test specifically the hypothesis that reducing muscle  $\text{H}^+$  accumulation during training would result in greater improvements in mitochondrial respiration. Thus, it was important to employ a training program that would result in large increases in mitochondrial respiration and a supplement protocol that would reduce the  $\text{H}^+$  accumulation during training. Our training was based on previous research that has demonstrated that high-intensity interval training is required to increase the oxidative capacity of both the slow- and fast-twitch fibers (white vastus lateralis) (19). Furthermore, to maximize adaptations in all rats, the imposed training intensity was based on the individually determined maximal running velocities of all rats, as recommended by others (62).

Our supplement protocol was based on previous research in humans demonstrating that  $\text{NaHCO}_3$  ingestion leads to a smaller decrease in both blood (20) and muscle (13) pH following high-intensity interval training. In addition, our pilot work demonstrated that our  $\text{NaHCO}_3$  ingestion protocol significantly increased resting blood pH (Fig. 1) and reduced the decrease in muscle pH following a typical training session.

### Adaptations to the SOL

The 25% greater CS activity in the soleus of trained rats (PLA and BIC; Fig. 2) following 5 wk of high-intensity interval training is consistent with previous reports (16–69%) (3, 32, 41, 55, 64). Because CS is an enzyme located exclusively in the mitochondria, it has been suggested to provide a good indication of directional changes in the mitochondrial mass (52). Thus, our results suggest that there was a greater mitochondrial mass in the soleus of our trained rats (with no significant difference between PLA and BIC).

Because the maximal activity of CS has been shown to have no correlation with the maximal flux through the tricarboxylic acid cycle (8, 12), it is difficult to infer changes in mitochondrial function with training from changes in the activity of this enzyme. Therefore, we measured maximal ADP-stimulated respiration and observed a significantly greater state 3 respiration in the soleus of the two trained groups (with pyruvate as a substrate) compared with the control group (Fig. 3). When considering the PLA and CON groups, these differences

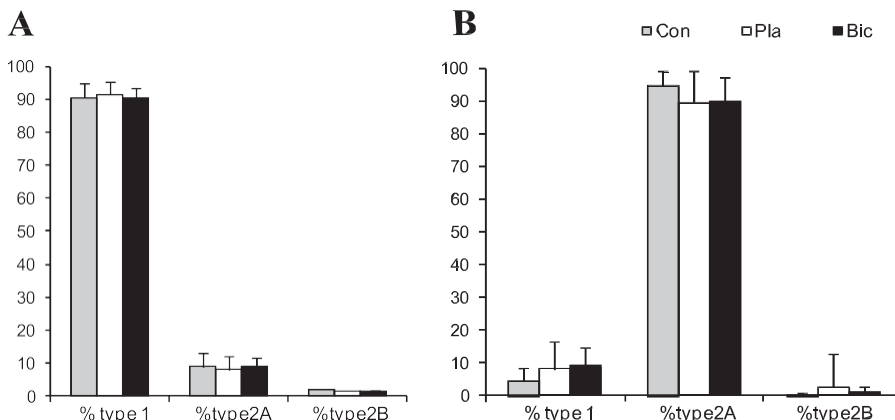


Fig. 5. % Muscle fiber composition in the 3 groups for both the SOL (A) and the EDL (B) muscles. Data are means  $\pm$  SD;  $n = 7$  for each group.

(89%) were similar to those reported in previous training studies using permeabilized fibers from both rats (20–36%) (7, 9) and humans (36–38%) (15, 59).

However, the most important finding of this study is that, compared with CON, state 3 respiration was significantly greater in rats that were provided with NaHCO<sub>3</sub> prior to every training session (+74%; BIC) than in rats that were supplied with the same volume of water prior to every training session (+39%; PLA) (Fig. 3). This increase in BIC is nearly double that reported in previous studies that have reported changes in mitochondrial respiration in saponin-permeabilized fibers following training (20–38%) (9, 15, 59). Furthermore, this difference remained significant even when corrected for CS activity. This suggests that both mitochondrial mass and mitochondrial function were significantly greater in BIC. Possible mechanisms are discussed below, but these novel findings indicate that reducing muscle H<sup>+</sup> accumulation during training is associated with greater improvements in mitochondrial respiration in the rat soleus.

#### Potential Mechanisms

Mitochondrial proteins are continuously degraded and renewed by de novo synthesis. This means that greater training-induced changes in mitochondrial protein (mass) could occur as a result of reduced protein degradation and/or greater protein synthesis. Although we are unaware of any studies investigating the effects of acidosis on mitochondrial protein, skeletal muscle protein synthesis has been reported to be inhibited when the medium pH is lowered in a variety of cell and tissue culture systems (21, 22, 58). In addition, acute metabolic acidosis has been reported to inhibit muscle protein synthesis (11) and to increase protein breakdown (42) in rat skeletal muscle. The protein breakdown pathway differs in mitochondria (ATP-dependent pathway involving mitochondrial proteases) vs. skeletal muscle (ATP-dependent pathway involving ubiquitin and proteasomes) (31). Nonetheless, it is possible that the greater state 3 respiration in BIC vs. PLA may be related to a smaller increase in mitochondrial protein degradation and/or reduced depression of mitochondrial protein synthesis as a result of smaller training-induced decreases in muscle pH.

The degree of acidosis during training may also affect the expression of molecular signaling proteins and the subsequent transcription of nuclear-encoded genes that regulate mitochondrial biogenesis. To date, acidosis has been reported to affect protein kinase B/Akt phosphorylation (2), to increase mRNAs encoding ubiquitin and protease subunits (1, 27), and to activate MAPK/ERK signaling pathways (4). In addition, it has been reported recently that NaHCO<sub>3</sub>-induced chronic alkalosis promotes upregulation of PGC-1 $\alpha$  and some of its downstream targets (COX-II, COX-IV, and cytochrome *c*) in C<sub>2</sub>C<sub>12</sub> myotubes (38). Furthermore, we have demonstrated recently that metabolic acidosis reduced exercise-induced upregulation of PGC-1 $\alpha$  mRNA (5). Thus, these latest findings suggest that our results may be explained, at least in part, by the effects of metabolic acidosis on the upregulation of important genes associated with mitochondrial biogenesis.

Although we have concentrated on the potential effects of acidosis on mitochondrial adaptations, NaHCO<sub>3</sub> ingestion may exert its effects by mechanisms in addition to, or instead of, alterations in exercise-induced muscle pH. For example, it has

been reported that the lactate anion also affects muscle-signaling pathways, leading to an increase in cytochrome *c* oxidase (COX) mRNA and protein, and also PGC-1 $\alpha$  expression (23). An imposed metabolic alkalosis (following NaHCO<sub>3</sub> ingestion) during high-intensity exercise produces an acceleration of glycogenolysis at the level of glycogen phosphorylase relative to maximal PDH activation, resulting in a mismatch between the rates of pyruvate production and oxidation, leading to an increase in lactate production (25). Thus, if lactate is indeed an important signaling molecule, greater increases in muscle lactate production during training, following NaHCO<sub>3</sub> ingestion, may serve to promote even greater trained-induced changes in mitochondrial biogenesis.

#### Adaptations to the EDL

An unexpected finding was that there was a significantly lower (85%) CS activity in the predominantly fast-twitch EDL muscle of our trained rats (Fig. 2). Other researchers have also reported lower (but not always significant) CS activity in the EDL (3–25%) in response to training that was associated with significantly greater CS activity in the soleus (3, 32, 41, 57, 64). It is possible that this can be attributed to a limited recruitment of the EDL during treadmill running in rats. However, if such an explanation is correct, one would expect no difference, rather than a lower CS activity, in the muscle of trained rats. The significantly greater  $r_{3\text{in vitro}}$  in both the soleus and EDL muscle of the trained rats also argues in favor of both muscles being recruited during our training protocol.

An alternative explanation for the reported decrease in CS activity in the EDL is that our chosen training intensity was excessive for the EDL muscle. It has previously been reported that the optimal training intensity varies among different fiber types and that improvements in oxidative capacity (cytochrome *c* activity) can be reduced with increases in training intensity (19). Furthermore, previous studies incorporating more moderate-intensity treadmill training in rats have reported significant increases in the CS of the EDL muscle (47, 50). Regardless of the explanation, our results suggest that high-intensity interval training can lead to a decrease in the mitochondrial mass of the EDL (a predominantly fast-twitch fiber), as indicated by the significantly lower CS activity in the EDL of trained rats.

Consistent with our CS results, we report for the first time that both state 3 and state 4 respiration were significantly lower in the EDL muscles from our trained rats compared with CON (Fig. 2). These differences were no longer apparent when mitochondrial respiration was corrected for CS activity. This suggests that the observed decreases in respiration were attributable to a lower mitochondrial mass rather than lower specific activity of the mitochondria. Furthermore, our observed decrease in CS is unlikely to be an artifact attributable to muscle hypertrophy (and hence, an apparent dilution of an unchanged total mitochondrial mass), since similar training in rats has not been reported to increase the mass of the EDL (168  $\pm$  26 vs. 170  $\pm$  17 g pre- and posttraining, respectively) (64). The absence of significant differences between BIC and PLA may be due to the observation that NaHCO<sub>3</sub> ingestion was less effective in reducing the change in pH in the EDL.



## Mitochondrial Respiration with Pyruvate vs. Palmitoyl Carnitine

Endurance exercise training is typically associated with an increased contribution of lipids to energy metabolism during submaximal work (10). The mechanisms for this are not clear, but this adaptation has been attributed to increased mitochondrial volume (29), higher  $\alpha$ -oxidation enzyme capacity (39), and greater lipid oxidation rates by the mitochondria (61). Therefore, it is surprising that we did not observe an increase in the maximal ADP-stimulated respiration of the soleus when using palmitoyl carnitine as a substrate. However, to our knowledge, all previous studies that have observed training-induced increases in mitochondrial respiration with palmitoyl carnitine as a substrate have used isolated mitochondria. An important difference when working with permeabilized skeletal muscle fibers is that reaching a concentration that is saturating requires a much greater palmitoyl carnitine concentration, and this high concentration may cause uncoupling in some mitochondria. This concern is supported by our respiration values for palmitoyl carnitine, which are less than one-half of the values for pyruvate, whereas these values are very similar when isolated mitochondria are used (36).

### Time to Exhaustion

A final novel finding of the present study is that, in addition to having a greater state 3 respiration, rats in the BIC group also had a significantly greater time to exhaustion than PLA. Furthermore, there was a significant correlation ( $r = 0.72$ ,  $P < 0.05$ ) between state 3 respiration in the soleus (with pyruvate and malate as substrates) and time to exhaustion in trained rats (Fig. 6). This is consistent with previous research, which has suggested that muscle oxidative capacity is a major determinant of endurance performance (16–18). For example, it has been reported previously that there is a temporal association between improvements in muscle oxidative capacity and improvements in endurance time during iron repletion in iron-deficient rats (18). Thus, the results of the present study provide further support for the hypothesis that muscle oxidative capacity is a major determinant of endurance performance. Furthermore, our results suggest that it is the oxidative capacity of the slow-twitch fibers that is a particularly important predictor of submaximal exercise time to fatigue.

### Summary

The main purpose of the present study was to investigate whether reducing muscle  $H^+$  accumulation during training would result in greater improvements in mitochondrial respiration. The success of our endurance training program was confirmed by the significantly greater CS activity in the soleus, but not the EDL, of trained rats. Thus, our results suggest that there was a greater mitochondrial mass in the soleus of our trained rats (with no significant difference between PLA and BIC). In contrast, it appears that high-intensity interval training can also lead to a decrease in the mitochondrial mass of the EDL (a predominantly fast-twitch fiber), as indicated by the significantly lower CS activity in the EDL of trained rats. However, the most important finding of this study is that, compared with CON, state 3 respiration was significantly greater in the rats that took  $NaHCO_3$  prior to every training

session (+74%; BIC) than in the rats that were supplied with water prior to every training session (+39%; PLA). This difference remained significant even when CS activity was used as a reference base, which suggests that both mitochondrial mass and mitochondrial function were significantly greater in BIC. Further research is required to determine whether these findings can be explained by the effects of acidosis on protein degradation, protein synthesis, or the expression of molecular signaling proteins and the subsequent transcription of nuclear-encoded genes that regulate mitochondrial biogenesis.

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### DISCLOSURES

No conflicts of interest, financial or otherwise, are declared by the author(s).

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