

Easily performed interval exercise induces to increase in skeletal muscle PGC-1 α gene expression

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Abstract

Introduction: Understanding the relationship between PGC-1 α expression and exercise is important for developing therapeutic exercise programs focusing on the prevention of lifestyle diseases. The current study examined whether easily performed moderate intensity interval exercise can induce PGC-1 α gene expression. **Methods:** Nine subjects performed cycling in one of three protocols: maximal intensity [MIE: 20 \times 1 min with 4 min recovery], high intensity [HIE] and a moderate intensity at anaerobic threshold interval exercise [AIE]. Both HIE and AIE were adjusted to obtain the same exercise volume as MIT performed for 5min including recovery time and repeated 20 times. **Results:** Increase in PGC-1 α mRNA expression was observed in all conditions, with a significant increase in plasma epinephrine. ACC phosphorylation also increased in all condition. **Conclusion:** These findings suggest that easily performed interval exercise at anaerobic threshold induces PGC-1 α expression which induce aerobic training adaptation.

Introduction

Physical inactivity has resulted in a drastic increase in the prevalence of a number of chronic diseases [1]. Given its health consequences, understanding the relationship between PGC-1 α expression and exercise is important for developing therapeutic exercise programs focusing on the prevention of lifestyle diseases [2]. Numerous studies have shown that prolonged endurance exercise results in PGC-1 α expression and improves aerobic capacity after several weeks of training [3]. Moreover, there is evidence that low volume sprint interval exercise has similar effects to traditional endurance training, eliciting similar levels of PGC-1 α and aerobic adaptation as typical endurance exercise training [4] [5]. Moreover, it has been shown that practical aerobic interval exercise (10 \times 1 min at 85-95% heart rate reserve interspersed with 1 min recovery) increases markers of skeletal muscle mitochondrial content and glucose transport capacity in sedentary middle-aged adults [6].

Accessibility and ease of completion are major considerations in creating therapeutic exercise programs for the general public. As such, short periods of higher-intensity interval exercise that have the same effect on PGC-1 α expression as long durations of continuous exercise presents an attractive prospect. However, for participants in exercise therapy programs, who often suffer from illnesses or obesity and have low levels of fitness [e.g., sedentary middle-aged or elderly people], even the 85-95%HR exercise is hard to practice in the longer term. In addition, short bursts of exercise are ineffective in weight loss because of the relatively small energy expenditure involved.

We already showed that even the moderate intensity exercise for 1hour duration could induce PGC-1 α expression [7]. We hypothesis that the easily performed interval exercise with short duration for several min which might easily performed even in elderly patients could induce PGC-1 α expression.

The current study sought was to verify whether moderate intensity exercise at anaerobic threshold performed in intervals (AIE) leads to a

PGC-1 α mRNA expression increase similar to that observed in typical endurance exercise and high intensity interval exercise (HIE).

Methods

Subjects

Subjects were nine active males (mean age: 27 \pm 12 years, mean height: 169.6 \pm 4.4 cm, mean weight: 71.9 \pm 16.6 kg, Vo₂max: 47.2 \pm 10.5 ml/kg.min) including 65 years old. All subjects were informed of the procedure and any risks, and gave written informed consent before participation. The study was approved by the Ethics committee of Fukuoka University.

Exercise load settings

We used cycle ergometers (Rehcor, Lode, Groningen, Netherlands) while subjects performed a graded ramp exercise test. After 15 min rest in a sitting position, the subjects warmed up for 4 min at 10 watts. During the test, 1 watt of load was added every 4 seconds until volitional fatigue. Gas analysis and electrocardiogram were measured during the test. Throughout the procedure, gas exchange was monitored continuously using the metabolic cart with mass spectrometry (ARCO-2000, Arco System, Chiba, Japan). The measures of gas concentration and amount of ventilation were converted to values per minute and the oxygen uptake was calculated. At 3 and 5 min after the exercise, blood was collected from the earlobes and the concentration of lactate was measured.

From the changes in oxygen inhalation and carbon dioxide exhalation during exercise, anaerobic threshold (an indirect measure of the lactate threshold (LT), moderate intensity) was determined

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using the V slope method. VO₂ max intensity (maximal intensity) was determined from the regression equation of exercise load and oxygen uptake, and high intensity was calculated (between maximal and moderate intensities) as follows. high intensity = (maximal Intensity + moderate Intensity) \div 2.

Intermittent exercise test at determined intensity

Subjects arrived at 07:00–08:30 and consumed a standardized breakfast [(energy: 476 kcal; carbohydrates: 57.1%; fat: 28.7%; protein: 14.1%). Approximately 1 hour later, they performed 20 sets of intermittent exercise using the power ergometer (Powermax-VIII, Combi Wellness, Tokyo). The experimental protocol was shown in Figure 2. The intensity of exercise was at three levels (Maximal, High and Moderate). Participants performed each protocol for one week, in a random order. One set at Maximal intensity lasted 60 s, and the exercise time for the High and Moderate intensity exercise types was adjusted to obtain the same exercise volume. The resting time for the Maximal condition was 240 s, and the resting time for High and Moderate conditions was calculated as follows: 300 s – [exercise time].

Needle muscle biopsy samples were taken during resting time from the vastus lateralis of one thigh under local anesthesia, as previously described [7], directly after exercise and 3 h after exercise. Muscle samples were collected from the vastus lateralis in the dominant foot. We collected samples of 5–10 mg at a time, and the same sampling method was used before and after exercise. Collected samples of skeletal muscle were promptly separated from blood using a sterilized gauze, frozen with liquid nitrogen and stored at –80 °C until the analysis.

The lactate concentration in the blood was measured before and after exercise. We asked subjects to estimate their rate of perceived exertion [RPE] after 5, 10, 15 and 20 sets of exercise.

Muscle analysis

Gene expression analysis: Total RNA was extracted from muscle sample [5mg] by the acid guanidine phenol chloroform [AGPC] method [ISOGEN, Nippon Gene Co Ltd, Toyama, Japan] with DNase treatment [Deoxyribonuclease I, Invitrogen Corp., Carlsbad, CA, USA]. cDNA was synthesized from 1 μ g of total RNA by reverse transcriptional reagents [Transcriptor First Stand cDNA synthesis kit, Roche Applied Science, Penzberg, Germany]. The PGC-1 α and β -actin gene expressions levels were analyzed by real time RT-PCR [LightCycler and FastStart DNA MasterPLUS SYBR Green I, Roche Applied Science, Penzberg, Germany] and those gene expressions were quantified by the relative standard curve method. A prior analysis was carried out to determine the PCR conditions and the PCR period was stored as a calibrator sample.

The prior PGC product was extracted using a PCR product purification kit [High pure PCR product purification kit, Roche Applied Science, Penzberg, Germany] and the sample was stored at –30 °C. Diluted calibrator samples were included in each analysis to make a standard curve. The PGC- α and PDK gene expression were normalized by β -actin. The intra and interassay coefficient variations were 12% and 15%. The specificity of the PCR product was identified by a melting curve analysis, which determined the temperature of denaturation of the PCR product by heating.

Protein quantification: Skeletal muscles was homogenized in ice-cold homogenization buffer [50 mM Tris [PH 7.5] 137 mM NaCl, 2.7 mM KCl, 1 mM MgCl₂, 1% Triton X-100, 10% [w/v] glycerol, 10mM NaF, 1 mM EDTA, 5 mM sodium pyrophosphate, 0.5 mM Na₃VO₄, 1 μ g ml⁻¹ leupatin, 0.2 mM phenylmethyl sulfonyl fluoride, 1 μ g ml⁻¹ aprotinin, 1 mM dithiothreitol, 1 mM benzamide, 1 μ M microcystin]

using Polytron homogenizer. Homogenates were rotated for 60 min at 4°C and centrifuged [12,000g for 10 min at 4°C], and the protein content of supernatant was determined using commercial kit assay [Bio-Rad, Richmond, CA, USA]. An aliquot of muscle homogenate [50 μ g protein] was mixed with Laemmli buffer.

The sample were separated and identified using SDS-PAGE and transferred to polyvinylidene difluoride membranes. Non-specific binding was blocked in TBST [10mM Tris, 100mM NaCl, 0.02% Tween 20] for 2h at room temperature. Membranes were incubated overnight with primary antibodies directed towards phospho-acetyl-CoA carboxylase [ACC] Ser⁷⁹ [1:500; Cell Signaling no, 9221]. Membranes were washed in TBS-t and incubated with appropriate secondary antibody for 1h at room temperature. Antibody binding was visualized by enhanced chemiluminescence [ECL; GE Healthcare, Arlington Heights, IL, USA] and quantified by densitometry [CS800 Calibrated Imaging Densitometer, Bio-Rad].

Statistical analysis

The data values in the study were expressed as means \pm SD. For all statistical processing IBM SPSS Statistic Ver 18 (SPSS Inc., Chicago, USA) was used. P < 0.05 was used as the threshold for statistical significance.

For the purposes of analysis, the amount of mRNA at rest was taken to have a value of 1, and amplification was estimated for PGC-1 α and PDK. For positively skewed distributed variables, log-transformation was performed. A two-way repeated measures ANOVA method [time and exercise model] was used. When a significant main effect was observed, Bonferroni post hoc analysis was used to examine the difference.

Paired t-tests were used to compare within-group differences [baseline vs post]. Statistical significance was accepted at P < 0.05.

Result

Maximal intensity interval exercise (MIE) was set at 245 \pm 36 watts corresponding to 100% VO₂max, High intensity interval exercise (HIE) at 179 \pm 23 watts corresponding to 74 \pm 3% VO₂max and moderate intensity at anaerobic threshold interval exercise (AIE) at 113 \pm 14 watts corresponding to 48 \pm 7% VO₂max. Exercise duration was 60 s for MIE, 82 \pm 4s for HIE, and 132 \pm 19s for AIE to the same amount of exercise and rest of the time for 5 min were stop exercise and repeated 20 times.

The results are shown in Figure 1. Three hours after the exercise, the volume of PGC-1 α mRNA increased significantly in all

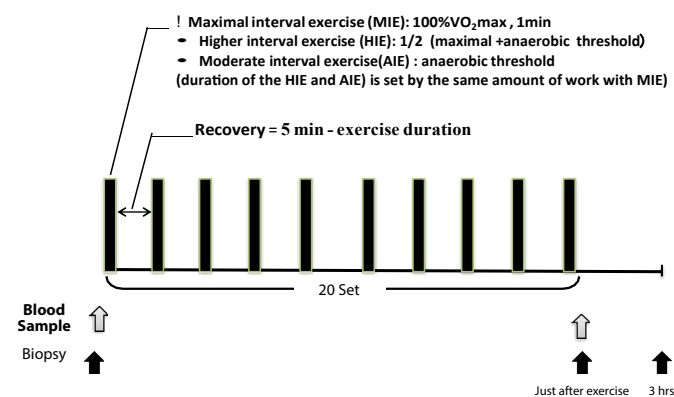
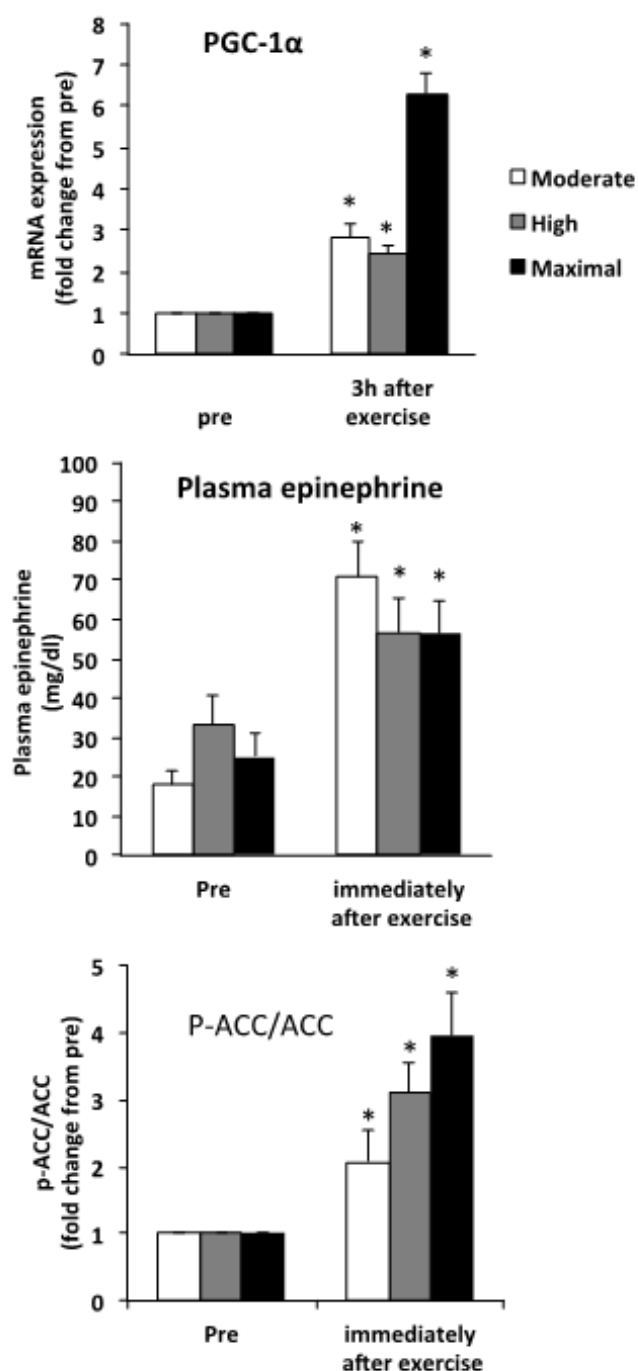


Figure 1. Experimental protocol.



* Significant ($P < 0.05$) main effects of time were found.

Figure 2. Summary data (mean \pm SE) from the moderate, higher, and maximal conditions, showing the magnitude of the change in muscle PGC-1 α mRNA abundance, plasma epinephrine and muscle ACC-p level. Total RNA was isolated from muscle biopsies pre-exercise, , and 3 h post exercise plasma epinephrine and ACC-p/ACC from the biopsies immediately after exercise.

conditions 4.9 fold for MIE, 2.5 fold for HIE and 2.8 fold for AIE, but no significant difference in conditions. Plasma epinephrine levels increased significantly, and the increase was similar in all conditions. ACC increased significantly in all conditions 3.9 fold for MIE, 3.1 fold for HIE and 2.1 fold for AIE, but no significant difference in conditions.

Discussion

A number of studies have reported increased PGC-1 α mRNA expression as a result of endurance training for 40 to 90 min [3]. However, no previous studies have shown an increase in PGC-1 α with easily performed moderate exercise interrupted by short breaks. The current study revealed an increase in PGC-1 α following moderate interval exercise similar to previous reports of increased PGC-1 α following HIE [9-11] and very MIE [12,13]. Gibala *et al.* [5] instructed subjects to cycle for four 30 sec “all out” exercise bouts interspersed with 4 min rest periods. That study found a relative increase in PGC-1 α mRNA expression that was relatively smaller than the increase we observed in the current study following AIE [6].

Interestingly, even AIE training ($48 \pm 7\%$ $\text{VO}_{2\text{max}}$) was observed to induce a similar increase in PGC-1 α mRNA compared with HIE ($74 \pm 3\%$ $\text{VO}_{2\text{max}}$). Previous studies have also reported an attenuated response in post-training exercise bouts at a similar absolute training intensity [10, 14]. Discrepancies among previous studies may be related to the amount of exercise. The difference between the current study and the previous studies was in the duration: In previous studies it remained the same pre and post training, however in the current study it different (i.e. longer for lower intensity). β_2 -adrennergic receptor activation is a potential mechanism underlying the PGC-1 α mRNA increase we observed. In mouse studies, even very slow treadmill running was found to induce a 2-fold increase in PGC-1 [15], which was inhibited by a β_2 -AR antagonist [16]. It has been well established that plasma catecholamine levels are dependent on the intensity and duration of exercise [17,18] We used moderate-intensity exercise at the anaerobic threshold, which is an indirect method for determining LT, because the intensity corresponded to the catecholamine threshold [19]. In a previous study, we found that the exercise below LT, which is below catecholamine threshold, could not induce an increase in PGC-1 α even though the total amount was the same as exercise above LT [20]. As expected, we observed a significant increase in plasma epinephrine following moderate intensity interval exercise, similar to that induced by higher intensity interval exercise. This is likely because the duration of every bout of exercise was longer in the moderate intensity interval exercise, so the total amount of exercise was the same. AMPK also stimulates PGC-1 α expression. Despite the easily performed AIT, AMPK activity may be increased as ACC phosphorylation, sensitive indicator of skeletal muscle AMPK activity [21, 22], increased.

In conclusion, the present results provide evidence that even moderate intensity interval exercise, which can be easily performed by every body including cardiac patient and elderly people, is sufficient to activate the genes that contribute to endurance training adaptation such as mitochondrial biogenesis.

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