Regulation of skeletal muscle glycogen phosphorylase and PDH during maximal intermittent exercise

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1Department of Medicine, McMaster University, Hamilton, Ontario L8N 3Z5; and 2Department of Human Biology and Nutritional Sciences, University of Guelph, Guelph, Ontario, Canada N1G 2W1

Parolin, Michelle L., Alan Chesley, Mark P. Matsos, Lawrence L. Spriet, Norman L. Jones, and George J. F. Heigenhauser. Regulation of skeletal muscle glycogen phosphorylase and PDH during maximal intermittent exercise. Am. J. Physiol. 277 (Endocrinol. Metab. 40): E890–E900, 1999.—The time course for the activation of glycogen phosphorylase (Phos) and pyruvate dehydrogenase (PDH) and their allosteric regulators was determined in human skeletal muscle during repeated bouts of maximal exercise. Six subjects completed three 30-s bouts of maximal isokinetic cycling separated by 4-min recovery periods. Muscle biopsies were taken at rest and at 6, 15, and 30 s of exercise during bouts 1 and 3. Phos was rapidly activated within the first 6 s of bout 1 from 12% at rest to 47% at 6 s. The activation of PDH increased from 14% at rest to 48% at 6 s and 95% at 15 s of bout 1. Phos reverted back to basal values at the end of the first bout, whereas PDH remained fully activated. In contrast, in the third bout, PDH was 42% at rest and was activated more rapidly and was nearly completely activated by 6 s, whereas Phos remained at basal levels (14–20%). Lactate accumulation was marked in the first bout and increased progressively from 2.7 to 76.1 mmol/kg dry wt with no further increase in bout 3. Glycogen utilization was also marked in the first bout and was negligible in bout 3. The rapid activation of Phos and slower activation of PDH in bout 1 was probably due to Ca2+ release from the sarcoplasmic reticulum. Lactate accumulation appeared to be due to an imbalance of the relative activities of Phos and PDH. The increase in H+ concentration may have served to reduce pyruvate production by inhibiting Phos transformation and may have simultaneously activated PDH in the third bout such that there was a better matching between pyruvate production and oxidation and minimal lactate accumulation. As each bout progressed and with successive bouts, there was a decreasing ability to stimulate substrate phosphorylation through phosphocreatine hydrolysis and glycolysis and a shift toward greater reliance on oxidative phosphorylation.

FOR MORE THAN A DECADE, WE HAVE BEEN EXAMINING THE REGULATION OF ENERGY PROVISION DURING REPEATED BOUTS OF MAXIMAL ISOKINETIC CYCLING (26, 35, 38, 45, 50). OTHER GROUPS HAVE ALSO PERFORMED SIMILAR STUDIES WITH THE USE OF DIFFERENT SPRINT MODELS (2, 6, 18, 20). DURING SPRINT EXERCISE, POWER OUTPUTS ARE ELICITED IN EXCESS OF TWO TO THREE TIMES THAT AT WHICH MAXIMAL OXYGEN CONSUMPTION (VO2max) IS ACHIEVED. TO MEET THE MAXIMAL DEMAND FOR ATP, THE MAJORITY OF ENERGY FOR SHORT-TERM MAXIMAL EXERCISE IS DERIVED VIA SUBSTRATE PHOSPHORYLATION FROM PHOSPHOCREATINE (PCr) HYDROLYSIS AND GLYCOLYSIS (26, 27, 33). HOWEVER, STUDIES USING INTERVAL EXERCISE CONSISTING OF TWO TO FOUR BOUTS OF 30 s OF MAXIMAL CYCLING SEPARATED BY 4 min OF REST HAVE DEMONSTRATED THAT THERE IS A PROGRESSIVE SHIFT TO OXIDATIVE METABOLISM IN LATER BOUTS (6, 38, 45, 50). IN ADDITION, SIGNIFICANT LACTATE ACCUMULATION OCCURRED IN THE FIRST BOUT COMPARED WITH THE THIRD AND FOURTH BOUTS, WHERE LACTATE ACCUMULATION WAS NEGligible (35, 38, 45). REDUCED FLUX THROUGH THE GLYCOLYTIC PATHWAY AND A GREATER DEGREE OF ACTIVATION OF PYRUVATE DEHYDROGENASE (PDH) BEFORE THE THIRD BOUT WAS ACCOMPANIED BY AN INCREASE IN OXYGEN UPTAKE (VO2), WHICH CONTRIBUTED TO REDUCED ACCUMULATION OF LACTATE AND INCREASED OXIDATION OF PYRUVATE (38). ASSOCIATED WITH A REDUCTION IN LACTATE ACCUMULATION WAS AN INCREASE IN OXIDATIVE PHOSPHORYLATION IN THE THIRD BOUT COMPARED WITH THE FIRST.


ALTHOUGH MANY STUDIES HAVE EXAMINED THE METABOLIC RESPONSES TO SPRINTING BEFORE AND AFTER 30-s BOUTS OF SPRINT EXERCISE (6, 18, 20, 26, 35, 38, 45), FEW HAVE EXAMINED THESE RESPONSES DURING THE EXERCISE BOUT. THEREFORE, TO EXAMINE THESE FACTORS MORE CLOSELY, THIS STUDY SPECIFICALLY EXAMINED THE TIME COURSE OF THE ACTIVATION OF PHOS AND PDH DURING THE FIRST AND THIRD BOUTS OF MAXIMAL INTERMITTENT CYCLING. MUSCLE BIOPSYs WERE OBTAINED AT REST AND AT 6, 15, AND 30 s OF THE FIRST...
and third bouts of exercise to construct a time course of Phos and PDH activation and to ascertain the regulatory factors responsible for Phos and PDH activity. This protocol also allowed us to calculate the relative contributions to ATP regeneration from substrate level phosphorylation by PCR hydrolysis and glycolysis and from oxidative phosphorylation within the two bouts of exercise.

METHODS

Subjects

Six male subjects (mean \(\pm\) SE, age 28.5 \(\pm\) 1.8 yr, height 178.8 \(\pm\) 2.1 cm, and weight 85.8 \(\pm\) 3.1 kg) participated in this study. Informed consent was obtained from each subject after a verbal and written explanation of the experimental protocol and its attendant risks. The study was approved by the McMaster University Ethics Committee.

Experimental Protocol

The exercise protocol for this study consisted of three 30-s bouts of maximal isokinetic cycling at 100 rpm separated by 4 min of rest. The isokinetic cycle ergometer used in this study has been described previously (34). Briefly, the cycle was fitted with a 3-hp direct current motor that set the upper limit of crank velocity at 100 rpm. The forces exerted by the subjects were recorded by strain gauges attached to the pedal cranks. Electrical signals were transferred to a computer, and work was calculated for each pedal stroke as the product of impulse and angular velocity. The total work for each 30-s period was calculated as the sum of 50 pedal strokes for two legs. The total work and power output accomplished during each time interval (i.e., 0–6, 6–15, and 15–30 s) were calculated from the sum of the pedal strokes for two legs over each interval.

Biopsies of the vastus lateralis were taken at rest and at 6, 15, and 30 s of bouts 1 and 3. Because it was not possible to obtain all of the biopsies on the same day, the sampling was distributed over three trials each separated by 1 wk (Fig. 1). The study was designed such that 1) the predetermined exercise intervals (i.e., 6, 15, or 30 s) were completed uninterrupted from rest before each biopsy was taken and 2) the resting and 30-s biopsies were obtained on the same day for bouts 1 and 3, respectively, so that muscle glycogenolysis could be determined within each bout from the difference between resting glycogen content and glycogen content at 30 s of exercise. During trial 1, two biopsies were taken at rest, one at 30 s of bout 1, and another at 6 s of bout 3. During trial 2, one biopsy was taken at 15 s of bout 1, two biopsies were taken at rest before bout 3, and one biopsy was taken at 30 s of bout 3. During trial 3, biopsies were taken at 6 s of bout 1 and 15 s of bout 3. A total of 10 biopsies were taken. The order of the trials was randomized for each subject and took place at the same time each day. Subjects were asked to consume a high-carbohydrate diet 48 h before trial 1 and to replicate this diet before the remaining two trials. The subjects were instructed to abstain from the consumption of caffeine and to refrain from intense exercise for 48 h before each trial.

Muscle Sampling

Before each trial, one thigh was prepared for needle biopsies of the vastus lateralis. Two or four incisions were made through the skin to the deep fascia under local anesthesia (2% lidocaine without epinephrine). Alternate legs were used on each visit so that one leg received four biopsies and the contralateral leg received six biopsies in total. Biopsies of the vastus lateralis were obtained as described by Bergström (5). All muscle biopsies were taken while the subject was on the bike and were immediately plunged in liquid N\(_2\). At the appropriate time, the exercise was interrupted and the clock was stopped. After the biopsy, cycling was resumed for the time remaining in the bout so that subjects completed a total of 30 s of maximal cycling for each bout. The time delay between interrupting and resuming cycling was 6–9 s. The time elapsed between the start of the biopsy to complete freezing was \(\sim\) 10 s. One of the resting biopsies before bouts 1 and 3 was frozen in liquid N\(_2\) after a 30-s delay to obtain resting measurements of the active form of Phos (Phos\(_a\); see Ref. 41). The biopsies were removed from the needle while frozen and were stored in liquid N\(_2\).

Analyses

A 5- to 15-mg piece of muscle was chipped from each biopsy under liquid N\(_2\) and was dissected free of blood and connective tissue for the determination of the active form of PDH (PDH\(_a\)). On two larger biopsies, another piece of wet muscle was used for the determination of total PDH activity (PDH\(_t\)).

Fig. 1. Experimental design. Muscle biopsies were taken at the times indicated by the arrows.
PDH\(_a\) and PDH\(_t\) were measured as described by ConstantinTeodosiu et al. (13) and modified by Putman et al. (39). The remaining muscle was freeze-dried, dissected free of blood, connective tissue, and fat, and stored dry at \(-70^\circ\text{C}\) for subsequent analysis.

One aliquot of powdered muscle was used for the determination of Phos activity, as described by Young et al. (52). Briefly, 3–4 mg of powdered muscle were homogenized at \(-25^\circ\text{C}\) in 200 µl of buffer containing 100 mM Tris, 60% glycerol, 50 mM KF, and 10 mM EDTA (pH 7.5). Homogenates were then diluted with 800 µl of the above buffer without glycerol and were homogenized further at 0°C. Total (a + b) Phos activity (in the presence of 3 mM AMP) and Phos V at 30°C. Maximal velocity (activity in the absence of AMP) were measured by following the production of glucose 1-phosphate spectrophotometrically at 30°C. Maximal velocity (V\(_\text{max}\)) and the mole fraction of Phos a and a + b was calculated from the measured activities as described by Chasiotis et al. (11).

A second aliquot of powdered muscle was assayed enzymatically for glycogen, as described by Harris et al. (21). The remaining dry powdered muscle was extracted in a solution of 0.5 M perchloric acid (PCA), and 1 mM EDTA and was neutralized with 2.2 M KHCO\(_3\). The neutralized PCA extracts were assayed for ATP, PCr, creatine, pyruvate, lactate, glucose, glucose 6-phosphate (G-6-P), fructose 6-phosphate (F-6-P), and glyceral 3-phosphate (Gly-3-P) as described by Bergmeyer (4) and acetyl-CoA, the reduced form of coenzyme A (CoASH), and acetyl carnitine as described by Cedergard et al. (8). Muscle metabolites and enzyme activities were corrected to the highest total creatine content for each subject to correct for nonmuscle contamination.

Calculations

Intracellular H\(^+\) concentration ([H\(^+\)]) was calculated from muscle lactate and pyruvate concentrations according to Sahlin et al. (44) and validated by Spriet et al. (45) for the type of exercise used in the present study. The concentrations of free ADP and free AMP were calculated from the near-equilibrium reactions of creatine kinase and adenylate kinase, respectively (15); however, these calculations do not provide information as to the exact location and local concentrations of these metabolites within the cell.

The concentration of free Pi was calculated as the difference between the concentrations of resting and exercise PCr, less the accumulation of G-6-P, F-6-P, and Gly-3-P plus the assumed resting concentration of 10.8 mmol/kg dry wt (15). The concentrations of the monoprotonated (HPO\(_2\)\(^-\)) and diprotonated (H\(_2\)PO\(_4\)\(^-\)) forms of Pi were calculated from the total concentration of Pi, and the calculated pH, using a pK\(_a\) of 6.78.

The rate of pyruvogenesis, expressed as millimole glucosyl units per kilogram dry weight per second, was estimated according to Spriet et al. (47) from the accumulation of glycogenic intermediates plus the flux of pyruvate through PDH\(_a\) during each time interval. Because previous studies have shown that flux through PDH is equivalent to its level of activity (19, 22, 38), PDH\(_a\) was converted to units of millimole per kilogram dry weight per second, assuming a wet-to-dry muscle ratio of 4:1 at rest and 4.5:1 during exercise (37).

\[
glycogenolysis = (\Delta[G-6-P] + [F-6-P]) + (\Delta[Gly-3-P] + [lactate] + [pyruvate]) / 2 \div \text{time} + \text{PDH}_a / 2
\]

where Δ indicates change and brackets indicate concentration.

The rate of pyruvate production, expressed as millimole pyruvate units per kilogram dry weight per second, was calculated from the accumulation of pyruvate and lactate plus the flux of pyruvate through PDH\(_a\) during each time interval. From a previous study using the same protocol (31), lactate efflux from the muscle was found to be very small and was thus not included in the calculation of pyruvate production:

\[
\text{pyruvate production} = \Delta[[\text{lactate}] + [\text{pyruvate}]] / \text{time} + \text{PDH}_a
\]

The rate of pyruvate oxidation was estimated from the flux through PDH\(_a\), which was converted to units of millimole per kilogram dry weight per second. The rate of lactate accumulation was calculated from the differences in concentration between two time points, divided by the elapsed time.

The rate of ATP turnover from PCr was calculated from the breakdown of PCr, whereas the rate of ATP turnover from glycolysis was calculated from the accumulation of lactate and the flux of pyruvate through PDH\(_a\). The rate of ATP turnover from oxidative phosphorylation was calculated from total acetyl-CoA production as the area under the PDH\(_a\) curves for each bout; 1 mmol of acetyl-CoA from glycolysis was equal to 15 mmol of ATP. The contribution of fat fuels (25) and exogenous glucose (29) was assumed to be negligible.

Statistical Analysis

Data were analyzed by a two-way ANOVA with repeated measures over time. When a significant F ratio was found, the Newman-Keuls post hoc test was used to compare the means. Net changes of metabolites during bouts 1 and 3 were compared using a paired dependent-samples t-test. Results were considered significant at P < 0.05.

RESULTS

Total Work and Average Power Output

Total work performed during the 30-s bouts of isokinetic cycling was 18.7 ± 0.8 and 13.8 ± 1.0 kJ in bouts 1 and 3, respectively, and was significantly greater in bout 1. The total power output generated during each bout decreased significantly from 622 ± 27 W during
the first bout to 459 ± 32 W in the third bout. The average power declined progressively during each bout, and this decline was more pronounced in bout 1 (Fig. 2).

Phos

The maximal total Phos activities were similar between bouts and across time (Table 1). In contrast, the calculated maximal Phos_a activity increased from rest during bout 1, returned to basal levels at the end of bout 1, and remained unchanged throughout bout 3 (Table 1). Consequently, during bout 1, the percentage of Phos in the a form increased rapidly from a resting value of 11.8 ± 2.7 to 46.8 ± 5.3% at 6 s, remained elevated at 15 s, and reverted back to basal levels at 30 s (Fig. 3). In bout 3, the resting Phos_a mole fraction was similar to the resting value before bout 1 and remained unchanged throughout exercise (Fig. 3).

PDH

Before bout 1, resting PDH_a was 0.53 ± 0.02 mmol·min⁻¹·kg wet wt⁻¹ and increased in a linear fashion to 1.81 ± 0.16 mmol·min⁻¹·kg wet wt⁻¹ at 6 s and to 3.56 ± 0.50 mmol·min⁻¹·kg wet wt⁻¹ at 15 s (Table 1). Because PDH was 3.74 ± 0.59 mmol·min⁻¹·kg wet wt⁻¹, this represented complete activation of PDH (95.2 ± 13.2%), which was maintained for the remainder of bout 1 (Fig. 3). Resting PDH_a was approximately threefold higher before bout 3 vs. bout 1. In bout 3, PDH was nearly completely activated to PDH_a by 6 s and was fully activated for the remainder of exercise (Table 1 and Fig. 3).

Muscle Metabolites

Muscle glycogen utilization was 89.2 ± 31.3 mmol/kg dry wt during bout 1 but was negligible in bout 3 (4.2 ± 28.5 mmol/kg dry wt; Fig. 4). The rate of muscle glycogenolysis during the initial 6 s of bout 1 was 4.04 mmol glucosyl units·kg dry wt⁻¹·s⁻¹, remained elevated from 6 to 15 s, and dropped to 0.58 mmol glucosyl units·kg dry wt⁻¹·s⁻¹ in the final 15 s (Fig. 5). Glycogenolytic rates were depressed throughout bout 3 to levels similar to those in the last 15 s of bout 1.

The concentrations of G-6-P, F-6-P, and Gly-3-P increased significantly from rest to 15 s of exercise in bout 1 and remained elevated (Table 2). In contrast, the concentrations of G-6-P, F-6-P, and Gly-3-P did not increase above resting levels during bout 3. The concentration of free glucose did not change within either bout but was significantly greater in bout 3 (Table 2).

During bout 1, the concentrations of lactate, pyruvate, and H⁺ increased progressively from rest until 15 s and remained elevated at 30 s (Fig. 4 and Table 2). Before bout 3, the concentrations of lactate, pyruvate, and H⁺ were similar to those found at the end of bout 1 and did not increase further during bout 3.

The rates of total pyruvate production and oxidation and lactate accumulation are shown in Fig. 6. The rate of pyruvate production was highest in the first 15 s of bout 1 but was reduced to <1/3 to 1/3 in the last 15 s of bout 1 and throughout bout 3. In contrast, the rate of pyruvate oxidation increased more than threefold dur-

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Table 1. Muscle Phos activities and PDH_a at rest and during maximal intermittent isokinetic cycling

<table>
<thead>
<tr>
<th></th>
<th>V_max for Phos a + b</th>
<th>V_max for Phos a</th>
<th>PDH_a</th>
</tr>
</thead>
<tbody>
<tr>
<td>Rest</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Bout 1</td>
<td>1.97 ± 0.22</td>
<td>0.24 ± 0.08</td>
<td>0.53 ± 0.02</td>
</tr>
<tr>
<td>Bout 3</td>
<td>2.56 ± 0.43</td>
<td>0.35 ± 0.13</td>
<td>1.56 ± 0.25†</td>
</tr>
</tbody>
</table>

|       | 15 s                |                 |       |
| Bout 1| 2.73 ± 0.30         | 1.28 ± 0.22*    | 1.81 ± 0.16* |
| Bout 3| 3.08 ± 0.55         | 0.39 ± 0.05†    | 3.08 ± 0.55† |

|       | 30 s                |                 |       |
| Bout 1| 2.38 ± 0.25         | 1.09 ± 0.17*    | 3.56 ± 0.50‡ |
| Bout 3| 2.60 ± 0.33         | 0.54 ± 0.10†    | 3.31 ± 0.40* |

Data are means ± SE. V_max for phosphorylase (Phos) a + b, calculated maximal total Phos activity; V_max for Phos a, calculated maximal Phos activity in mmol·kg dry wt⁻¹·s⁻¹. Active form of pyruvate dehydrogenase (PDH_a) is expressed in mmol·kg wet wt⁻¹·min⁻¹. *Different from rest of same bout. †Difference between bouts 1 and 3. ‡Different from previous time point in same bout.
High energy phosphates. The concentration of PCr decreased from a resting value of 88.2 ± 4.7 to 7.6 ± 3.2 mmol/kg dry wt at 30 s of exercise in bout 1 (Table 3). PCr resynthesis was incomplete before bout 3 and was 79% of the amount available before bout 1. Total PCr hydrolysis was greater in bout 1 compared with bout 3 (80.7 vs. 59.9 mmol/kg dry wt).

The concentration of ATP was unchanged during exercise and between bouts (Table 3). The calculated concentrations of free ADP and AMP increased in a time-dependent fashion and were significantly greater at 30 s of exercise in bout 1 compared with bout 3 (Table 3).

The concentration of total free P_i increased rapidly during the first 6 s of exercise and was significantly elevated throughout bout 3 compared with bout 1, due to increased binding to accumulated hexose monophosphates in bout 1 (Table 3). Due to a greater change in [H^+] in the first bout compared with the third, differences were shown between the two bouts when total free P_i was partitioned into its two molecular species. The diprotonated form (H_2PO_4^-) followed a similar pattern of change as total P_i, whereas the monoprotonated form (HPO_4^{2-}) increased as a function of time only with no differences between bouts 1 and 3 (Table 3).

Acetyl group accumulation. The concentrations of total and free CoA remained constant throughout both bouts of cycling (Table 2). The concentration of acetyl-CoA increased from resting values during each bout and was significantly elevated throughout bout 3 compared with bout 1 (Table 2). The concentration of acetyl carnitine remained constant throughout each exercise bout but was greater in bout 3 compared with bout 1 (Table 2).

DISCUSSION
In the present study, we examined the regulation of metabolic changes in muscle during the first and last of three bouts of maximal exercise for 30 s separated by 4 min of rest. During bout 1, the challenge of the extreme ATP demand was met through substrate phosphorylation by rapid PCr breakdown and the rapid onset of glycogenolysis. Glycogenolysis and pyruvate production were maximal during the first 15 s and far exceeded the rate of oxidation through PDH, resulting in significant lactate accumulation. However, during the last 15 s, Phos had returned to resting levels with low rates of glycogenolysis and pyruvate production, whereas PDH was fully active by 15 s and pyruvate oxidation was more closely matched to its rate of production, resulting in minimal further lactate accumulation. In bout 3, Phos b-to-a transformation was inhibited such that glycogenolysis and pyruvate production were low. PDH activity was elevated before bout 3, and full activation was rapidly achieved during the third bout such that pyruvate oxidation was more closely matched to pyruvate production with minimal lactate accumulation. As each bout progressed and with
Intermittent isokinetic cycling

Table 2. Muscle metabolite content in the vastus lateralis at rest and during maximal intermittent isokinetic cycling

<table>
<thead>
<tr>
<th>Measure</th>
<th>Rest 6s</th>
<th>15 s</th>
<th>30 s</th>
<th>Rest 6s</th>
<th>15 s</th>
<th>30 s</th>
</tr>
</thead>
<tbody>
<tr>
<td>Glucose, mmol/kg dry wt</td>
<td>1.3±0.2</td>
<td>4.0±0.2</td>
<td>2.0±0.3</td>
<td>4.3±1.6</td>
<td>12.8±3.4†</td>
<td>12.9±2.8†</td>
</tr>
<tr>
<td>G-6-P, mmol/kg dry wt</td>
<td>0.9±0.2</td>
<td>9.0±2.1*</td>
<td>18.4±2.6*</td>
<td>16.8±1.6*</td>
<td>8.5±2.5†</td>
<td>7.0±1.4</td>
</tr>
<tr>
<td>F-6-P, mmol/kg dry wt</td>
<td>0.2±0.1</td>
<td>1.5±0.4*</td>
<td>3.4±0.5*</td>
<td>3.9±1.3*</td>
<td>1.6±0.4†</td>
<td>1.8±0.5</td>
</tr>
<tr>
<td>Gly-3-P, mmol/kg dry wt</td>
<td>0.5±0.1</td>
<td>4.0±0.7*</td>
<td>7.4±0.3*</td>
<td>6.5±1.4*</td>
<td>11.1±1.7†</td>
<td>10.8±1.4†</td>
</tr>
<tr>
<td>H+, mM</td>
<td>623±0.3</td>
<td>110.3±5.4</td>
<td>1574.7±9.8*</td>
<td>182.9±12.4*</td>
<td>2164±48.0†</td>
<td>2545±45.8†</td>
</tr>
<tr>
<td>Acetyl-CoA, µmol/kg dry wt</td>
<td>8.0±1.4</td>
<td>9.6±0.7</td>
<td>11.0±1.1*</td>
<td>14.4±4.3†</td>
<td>14.3±1.5†</td>
<td>18.9±1.1†</td>
</tr>
<tr>
<td>Free CoASH, µmol/kg dry wt</td>
<td>584±9.6</td>
<td>67.2±13.2</td>
<td>66.6±7.6</td>
<td>60.9±21.6</td>
<td>64.0±12.0</td>
<td>62.8±8.8</td>
</tr>
<tr>
<td>Total CoASH, µmol/kg dry wt</td>
<td>664±9.9</td>
<td>76.8±13.6</td>
<td>77.7±8.1</td>
<td>75.2±23.4</td>
<td>78.3±13.2</td>
<td>81.7±9.8</td>
</tr>
<tr>
<td>Acetylcarnitine, mmol/kg dry wt</td>
<td>4.6±1.6</td>
<td>2.0±0.4</td>
<td>3.3±1.0</td>
<td>5.4±18</td>
<td>11.8±0.9†</td>
<td>11.4±2.0†</td>
</tr>
<tr>
<td>Glucose, mmol/lactate units·kg dry wt</td>
<td>1.3*</td>
<td>1.6</td>
<td>1.8</td>
<td>1.4*</td>
<td>1.7*</td>
<td>1.6*</td>
</tr>
<tr>
<td>Fructose 6-phosphate, mmol/kg dry wt</td>
<td>0.4†</td>
<td>2.0</td>
<td>2.8†</td>
<td>0.9†</td>
<td>0.9†</td>
<td>1.4†</td>
</tr>
<tr>
<td>Glycerol 3-phosphate, mmol/kg dry wt</td>
<td>1.4†</td>
<td>11.1</td>
<td>12.3</td>
<td>11.7†</td>
<td>11.7†</td>
<td>11.8†</td>
</tr>
<tr>
<td>Phospho-enolpyruvate, mM</td>
<td>62.3±0.3</td>
<td>1103.3±54</td>
<td>1574.7±98*</td>
<td>182.9±12.4*</td>
<td>2164±48.0†</td>
<td>2545±45.8†</td>
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<tr>
<td>Pyruvate oxidation, mM</td>
<td>8.0±1.4</td>
<td>9.6±0.7</td>
<td>11.0±1.1*</td>
<td>14.4±4.3†</td>
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<td>Phosphating enzyme activity, U/mg dry wt</td>
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<td>75.2±23.4</td>
<td>78.3±13.2</td>
<td>81.7±9.8</td>
</tr>
<tr>
<td>Phosphorylation kinase activity, U/mg dry wt</td>
<td>4.6±1.6</td>
<td>2.0±0.4</td>
<td>3.3±1.0</td>
<td>5.4±18</td>
<td>11.8±0.9†</td>
<td>11.4±2.0†</td>
</tr>
</tbody>
</table>

Data are means ± SE. G-6-P, glucose 6-phosphate; F-6-P, fructose 6-phosphate; Gly-3-P, glycerol 3-phosphate. *Different from rest of same bout. †Different from previous time point in same bout.

Each progressive bout, there was decreased substrate phosphorylation by PCR hydrolysis and glycolysis and a greater reliance on oxidative phosphorylation.

Contribution of Substrate Phosphorylation and Oxidative Phosphorylation to ATP Production

In response to the high demand for ATP turnover during the initial 6 s of maximal exercise, substrate level phosphorylation by rapid PCR hydrolysis provided an immediate source of ATP at a rate of 7.0 mmol·kg dry wt⁻¹·s⁻¹ (Fig. 7). The activation of Phos and glycogenolysis allowed substrate level phosphorylation by glycolysis at a rate of 6.2 mmol·kg dry wt⁻¹·s⁻¹, whereas the rate of ATP turnover by oxidative phosphorylation was 1.32 mmol·kg dry wt⁻¹·s⁻¹ in the first 6 s of exercise. Thus a total maximal ATP turnover rate of 14.5 mmol·kg dry wt⁻¹·s⁻¹ was obtained in the first 6 s of bout 1 and was in agreement with values obtained in other studies using similarly strenuous protocols (24). In the last 15 s of bout 1, when the PCR stores had been depleted and glycogenolysis was inhibited, substrate level phosphorylation accounted for a decreasing proportion of the total ATP turnover rate. In bout 3, PCR was still an important source of ATP regeneration, but, due to the inhibition of Phos, substrate level phosphorylation by glycolysis contributed very little. As each bout progressed and with each progressive bout, the decreasing ability to maintain substrate phosphorylation by PCR hydrolysis and glycolysis was supplemented by oxidative phosphorylation. As each bout progressed, and the average power output was reduced, the ATP turnover rate was reduced accordingly (Figs. 2 and 7).

Regulation of Phos

Glycogen supplies the majority of the energy required for rapid ATP provision during short-term maximal exercise (7, 18, 26, 35, 46). During the initial 6 s of the first bout, the glycolytic rate was estimated to be 4.04 mmol glucosyl units·kg dry wt⁻¹·s⁻¹, which remained elevated during the next 9 s, and declined to

![Graph A](https://via.placeholder.com/150)

![Graph B](https://via.placeholder.com/150)

Fig. 6. Comparison of the rates of pyruvate production, pyruvate oxidation, and lactate accumulation during each time interval in the first (A) and third (B) bouts.
The breakdown of glycogen is regulated by the rate-limiting enzyme Phos. Phos is regulated by reversible enzymatic phosphorylation and exists in two interconvertible forms, a less active b form and a more active a form. Phos a is transformed to its a form by Phos kinase, whereas dephosphorylation back to the b form is catalyzed by Phos phosphatase. The transformation of Phos is regulated at the hormonal level by cAMP and at the contractile level by Ca\(^2+\) release from the sarcoplasmic reticulum at the onset of contraction.

Posttransformational regulation of Phos activity and glycogenolysis occurs via the availability of substrate (P\(_i\) and glycogen) and the presence of positive (free AMP and IMP) and negative (G-6-P) allosteric modulators. The Michaelis constant (K\(_m\)) values of Phos a and a + b for P\(_i\) are 26.2 and 6.8 mM, respectively (11). Furthermore, the presence of 0.01 mM free AMP reduces the K\(_m\) of Phos a for P\(_i\) to 11.8 mM (42). Phos b is also active in the presence of high levels of free AMP (>0.02 mM) and IMP (1). In response to the high demand for ATP turnover during the initial 6 s of maximal exercise, the rapid hydrolysis of PCr resulted in significant accumulation of free P\(_i\), which falls within the range of the K\(_m\) for Phos.

The high glycogenolytic rate attained during the first 6 s of bout 1 and maintained during the next 9 s would require a Phos activity of 4.04 mmol glycosyl units·kg dry wt\(^{-1}\)·s\(^{-1}\). The calculated V\(_{max}\) of Phos a at 6 s was not sufficient to accommodate such a high glycogenolytic rate. This suggests that complete activation of Phos a + b would be required to achieve this flux. Due to the rapid reversal of Phos a to b (41), the 10-s delay from muscle sampling to complete freezing in our study may have led to an underestimation of Phos a activity and hence Phos activation.

Previous studies using submaximal exercise intensities ranging from 30 to 90% V\(_{O_2\ max}\) have shown that activation of Phos is in excess of the estimated flux through the enzyme and that glycogenolytic flux through

### Table 3. Muscle high energy phosphate content in the vastus lateralis at rest and during maximal intermittent isokinetic cycling

<table>
<thead>
<tr>
<th>Measure</th>
<th>Rest</th>
<th>6 s</th>
<th>15 s</th>
<th>30 s</th>
<th>Rest</th>
<th>6 s</th>
<th>15 s</th>
<th>30 s</th>
</tr>
</thead>
<tbody>
<tr>
<td>ATP, mmol/kg dry wt</td>
<td>20.8 ± 1.6</td>
<td>22.4 ± 0.5</td>
<td>20.4 ± 2.0</td>
<td>16.2 ± 4.9</td>
<td>17.6 ± 1.3</td>
<td>16.4 ± 2.0</td>
<td>18.1 ± 0.9</td>
<td>17.8 ± 1.4</td>
</tr>
<tr>
<td>Free ADP, µmol/kg dry wt</td>
<td>101.0 ± 8.7</td>
<td>208.8 ± 21.6</td>
<td>274.8 ± 17.3*</td>
<td>651.2 ± 13.8*</td>
<td>492.6 ± 6.4</td>
<td>168.9 ± 38.5</td>
<td>295.4 ± 58.0*</td>
<td>455.4 ± 95.9*† †</td>
</tr>
<tr>
<td>Free AMP, µmol/kg dry wt</td>
<td>0.5 ± 0.1</td>
<td>1.9 ± 0.4</td>
<td>3.8 ± 0.6</td>
<td>20.0 ± 0.3†</td>
<td>0.1 ± 0.0</td>
<td>1.8 ± 0.6</td>
<td>5.1 ± 1.6</td>
<td>15.6 ± 4.1† †</td>
</tr>
<tr>
<td>PCr, mmol/kg dry wt</td>
<td>88.2 ± 4.0*</td>
<td>46.2 ± 4.0*</td>
<td>28.2 ± 2.6†</td>
<td>7.6 ± 3.2*</td>
<td>69.7 ± 4.3†</td>
<td>25.7 ± 3.9†</td>
<td>14.9 ± 4.5*</td>
<td>9.8 ± 1.8*</td>
</tr>
<tr>
<td>Total P(_i), mmol/kg dry wt</td>
<td>10.8 ± 3.4*</td>
<td>36.7 ± 3.4*</td>
<td>45.5 ± 9.4*</td>
<td>47.8 ± 6.5*</td>
<td>18.8 ± 10.4</td>
<td>55.7 ± 6.5*</td>
<td>62.1 ± 7.4†</td>
<td>63.7 ± 8.6†</td>
</tr>
<tr>
<td>HPO(_4)(^2-), mmol/kg dry wt</td>
<td>7.8 ± 0.0</td>
<td>22.1 ± 1.6*</td>
<td>24.1 ± 5.3*</td>
<td>22.1 ± 2.9*</td>
<td>7.7 ± 4.2</td>
<td>20.8 ± 2.0*</td>
<td>25.4 ± 2.9*</td>
<td>24.3 ± 3.5*</td>
</tr>
<tr>
<td>H(_2)PO(_4)(^-), mmol/kg dry wt</td>
<td>3.0 ± 0.0</td>
<td>14.6 ± 1.8*</td>
<td>21.4 ± 4.2*</td>
<td>25.7 ± 3.6*</td>
<td>11.2 ± 6.5†</td>
<td>35.0 ± 5.5†</td>
<td>36.7 ± 4.6*</td>
<td>39.4 ± 5.8*†</td>
</tr>
</tbody>
</table>

Data are means ± SE. PCr, phosphocreatine. *Different from rest of same bout. † Different between bouts 1 and 3. †† Different from previous time point in same bout.
Phos is mediated primarily through posttransformational regulation (substrate and allosteric; see Refs. 12, 16, 22). In the present study, the estimated glycolytic flux was equal to the total Phos activity, and the peak glycolytic rates were ~20-fold higher than rates of 0.17–0.25 mmol glucosyl units·kg dry wt·min⁻¹ observed at 80–90% Vo₂max (12, 16, 22). The power outputs generated during the first 15 s of exercise in bout 1 were 3.5-fold greater than that obtained in the study by Howlett et al. (22) in which the average output was 229 ± 19 W at 90% Vo₂max. Thus it appears that, during the first 15 s of maximal exercise, total Phos activation is required to maintain glycolytic flux. This involves upregulation of Phos activity by both transformational and posttransformational mechanisms.

Compared with the first 15 s of bout 1, glycolysis was reduced by 86–95% in the latter one-half of bout 1 and throughout bout 3. This large reduction in glycolysis coincided with Phos a to b reversion and the inability to reactivate Phos transformation in bout 3. In addition, the predicted flux through Phos a during the last 15 s of bout 1 and throughout bout 3 corresponded more closely to the observed glycolytic fluxes. At the end of the first bout, Vₘₐₓ for Phos a corresponded more closely to the glycolytic rate estimated over the last 15 s of bout 1 (Table 1 and Fig. 5). Similarly, during the third bout, the estimated glycolytic rates were in the range of the calculated Vₘₐₓ for Phos a during each time interval (Fig. 5 and Table 1). In contrast, the maximal Phos a + b activity remained stable during the entire exercise protocol (Table 1). These observations imply that Phos b was minimally activated during this time despite elevations in free AMP. It is conceivable that the accumulation of G-6-P, a known allosteric inhibitor of Phos b (17), could have overridden positive modulation by free AMP.

The reduction in Phos transformation observed at the end of the first bout and throughout bout 3 suggests that the mechanism of activation by Ca²⁺ was being overridden. Krebs et al. (30) have shown in vitro that elevated [H⁺] inhibited Phos kinase, thus reducing the transformation of Phos to its a form. During the last 15 s of the first bout, [H⁺] increased significantly to 182.9 ± 12.4 nM (pH 6.74) and up to 254.5 ± 45.8 nM (pH 6.59) at 6 s in the third bout (Table 2). Chasiotis et al. (10) also observed in human skeletal muscle that released from the sarcoplasmic reticulum at the onset of exercise was probably the primary stimulus for the activation of PDH. The Ca²⁺ released activates PDH phosphatase (23) and may simultaneously inactivate PDH kinase (40), thus increasing the proportion of PDH. Increases in Ca²⁺ concentration in rat heart mitochondria have previously been shown in vitro to result in 60% activation of PDH (14), which is similar to what we observed in the first 6 s of contraction. The remaining 40% of the activation of PDH may have been due to other allosteric activators of PDH. Although the effects of Ca²⁺ on PDH activation were studied in cardiac muscle (14), it is generally believed that Ca²⁺ is also important in PDH activation in skeletal muscle (51).

Pyruvate is a known activator of PDH and inactivates PDH kinase (23). Pyruvate increased linearly during the first 15 s, served as substrate for the reaction, and may also have functioned as an allosteric regulator that stimulated activation of PDH. The levels of pyruvate appear to be within the estimated range of the inhibitory constant of PDH kinase for pyruvate of 0.5–2.0 mM (51). Increases in [H⁺] followed the same time course as pyruvate and may have played a role in PDH activation. Again, the only available evidence has been demonstrated in cardiac muscle. A study in per-
fused rat heart reported that acidosis increased PDH activation (36). Hucho et al. (23) also reported pH optima of 7.0–7.2 for bovine kidney and rat heart PDH kinases and 6.7–7.1 for the PDH phosphatases, suggesting that an increase in \([\text{H}^+\]) would favor the activity of the phosphatase over the kinase and increase the transformation of PDH to its active form. Although acetyl-CoA is known to inhibit activation of PDH by inhibition of PDH phosphatase and activation of PDH kinase, significant increases in acetyl-CoA in the last 15 s of exercise did not appear to exert an inhibitory effect over the activation of PDH. It appears that the increases in \(\text{Ca}^{2+}\) with contraction and increases in [pyruvate] and \([\text{H}^+\]) override the acetyl-CoA inhibition.

As in a previous study (38), before the initiation of exercise in the third bout, the activity of PDH_a was significantly elevated above resting values before bout 1. The increased activation was probably due to elevated levels of pyruvate and \([\text{H}^+\]), which continued to stimulate activation to PDH_a overridding the elevated levels of the negative allosteric modulator acetyl-CoA. Before the onset of contraction in the third bout, the mole fraction of PDH_a was 41.8 ± 6.8%, and with the onset of contraction PDH became totally activated within 6 s, likely due to the release of \(\text{Ca}^{2+}\) during contraction. Therefore, compared with the first bout, the higher levels of PDH_a in bout 3 would provide increased levels of acetyl-CoA for increased oxidative phosphorylation during the initial 6 s.

Timmons et al. (48, 49) have demonstrated that, at the onset of submaximal exercise, oxidative phosphorylation was limited by the availability of acetyl-CoA for the TCA cycle. They showed that dichloroacetate administration before exercise increased PDH_a and acetyl carnitine levels in human skeletal muscle and reduced PCr hydrolysis, glycogenolysis, and lactate production during the rest to exercise transition. In the present study, there was a tendency for acetyl carnitine to decrease (P = 0.11) at the onset of exercise in the first bout, and the major contribution of energy provision was from substrate level phosphorylation (Fig. 7), which suggests that the availability of acetyl groups may be limiting to oxidative phosphorylation. Before the third bout of exercise, we observed a similar degree of activation of PDH and accumulation of acetyl groups as in the studies of Timmons et al. The increased availability of the acetyl groups did not prevent the large and rapid rate of PCr hydrolysis in the first 6 s but did result in a shift to oxidative phosphorylation with minimal utilization of glycogenolysis, resulting in lactate accumulation. In a previous study, we observed a 10% increase in \(V_\text{O}_2\) in the third bout compared with bout 1 (38), which was mirrored by an overall increase of 10% in the rate of ATP provision from oxidative phosphorylation in the present study. Other investigators have observed an 18% increase in \(V_\text{O}_2\) in the second bout relative to the first bout (6) and an increase in the transient rise in \(V_\text{O}_2\) observed in a second exercise bout when it was preceded by high-intensity exercise (32).

### Lactate Production

During intense exercise, Phos provides substrate to the glycolytic pathway. Glycolysis in turn provides substrate for the TCA cycle in the form of pyruvate via decarboxylation by the rate-limiting enzyme PDH. Thus the accumulation of pyruvate is determined by its rate of production by glycolysis and its rate of oxidation by PDH. Glycolysis produced pyruvate at a rate of 4.31, 3.89, and 1.36 mmol·kg dry wt\(^{-1}\)·s\(^{-1}\) from 0 to 6, 6 to 15, and 15 to 30 s in bout 1, respectively (Fig. 6). The corresponding rates at which PDH_a could oxidize pyruvate were 0.09, 0.20, and 0.27 mmol·kg dry wt\(^{-1}\)·s\(^{-1}\) (Fig. 6). Thus the glycolytic rate was 48- and 19-fold greater than PDH_a during the first 15 s of bout 1 but only 5-fold greater in the last 15 s of bout 1. The large disparity between the rates of glycolytic pyruvate production and oxidation resulted in the net accumulation of pyruvate and lactate in the first 15 s of exercise (Fig. 6). As bout 1 progressed, there was a better match between pyruvate production and oxidation due to decreased glycolysis and increased PDH activation, which was evidenced by the reduced accumulation of pyruvate and thus lactate toward the end of the first bout (Fig. 6). Throughout the third bout, the rates of pyruvate production were considerably reduced to <0.67 mmol·kg dry wt\(^{-1}\)·s\(^{-1}\), whereas the rates of pyruvate oxidation through PDH were maintained at 0.17–0.25 mmol·kg dry wt\(^{-1}\)·s\(^{-1}\), representing less than a fourfold difference. The better matching between the rates of pyruvate production and pyruvate oxidation in the third bout resulted in no further accumulation of pyruvate or lactate.

During the initial seconds of maximal exercise, large rates of glycogenolysis and glycolysis are required to provide a rapid and immediate source of ATP, and Ca\(^{2+}\) appears to be the primary mechanism responsible for Phos and PDH activation. The rapid hydrolysis of ATP and PCr resulted in an accumulation of free P, and AMP that increased Phos activity and substrate for glycolysis. The increased glycolytic rate translated into enhanced pyruvate production, which may have exerted a positive allosteric effect on the activation of PDH. The increase in \([\text{H}^+\]) toward the end of the first bout may have inhibited Phos to a transformation and glycogenolysis, while simultaneously activating PDH. The elevated \([\text{H}^+\]) at the beginning of and throughout the third bout may have completely inhibited the transformation of Phos and glycogenolysis flux and may have simultaneously maintained a high level of PDH activity. This mechanism of regulation may have served to spare the glycogen stores, inhibit further accumulation of lactate, and limit further increases in \([\text{H}^+\]). Thus the increase in \([\text{H}^+\]) that accompanies intense exercise may be the primary mechanism responsible for limiting further increases in \([\text{H}^+\]) by inhibiting Phos and the accumulation of lactate while increasing PDH activity and the oxidation of lactate. The result was a tighter match between flux through Phos and PDH and a shift toward greater reliance on oxidative phosphorylation and less...
on substrate phosphorylation by PCr hydrolysis and glycolysis.

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