ORIGINAL ARTICLE

Protein ingestion does not impair exercise-induced AMPK signalling when in a glycogen-depleted state: implications for train-low compete-high

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Abstract The aim of the present study was to test the hypothesis that consuming protein does not attenuate AMPK signalling when exercise is commenced in a glycogendepleted state. After performing a glycogen-depleting protocol the evening before, the subsequent morning ten active men performed 45 min steady-state cycling at 50 % of peak power output (PPO) followed by an exercise capacity test (1-min intervals at 80 % PPO interspersed with 1-min periods at 40 % PPO). In a repeated measures design, subjects consumed 20 g of a casein hydrolysate solution (PRO) 45 min before exercise, 10 g during and a further 20 g immediately post-exercise, or an equivalent volume of a non-calorie taste matched placebo (PLA). Resting $(PRO = 134 \pm 29; PLA = 136 \pm 28 \text{ mmol kg}^{-1})$ and postexercise muscle glycogen (PRO = 43 ± 16 ; PLA = 47 ± 18 mmol kg⁻¹) was not different ($P > 0.05$) between trials nor was exercise capacity (PRO = 26 ± 9 ; PLA = 25 ± 1 10 min, $P > 0.05$). Phosphorylation of AMPK^{Thr172} increased threefold immediately post-exercise $(P < 0.05)$

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and PGC1-mRNA increased sixfold at 3 h post-exercise $(P<0.05)$, though there were no differences between conditions ($P > 0.05$). In contrast, there was a trend ($P = 0.08$) for a divergent response in $eEF2^{Thr56}$ phosphorylation such that 1.5 fold increases post- and 3 h post-exercise in PLA were blunted with PRO, thus indicative of greater eEF2 activation. We conclude that athletes who deliberately incorporate training phases with reduced muscle glycogen into their training programmes may consume protein before, during and after exercise without negating signalling through the AMPK cascade.

Keywords AMPK · PGC-1 α · Amino acids · Muscle glycogen

Introduction

At a molecular level, endurance training-induced increases in skeletal muscle mitochondrial biogenesis are thought to be due to the cumulative effects of transient increases in mRNA transcripts encoding mitochondrial proteins that follow each acute training session (Perry et al. [2010](#page-11-0)). Contractile-induced disruptions to cell homeostasis (e.g. increased AMP/ATP ratio, Ca^{2+} , reactive oxygen species, reduced substrate availability, etc.) result in the activation of a number of regulatory protein kinases which, in turn, phosphorylate downstream targets such as transcription factors or transcriptional co-activators (Hood [2009](#page-10-0)). Perhaps the most well-studied signalling kinase that is sensitive to muscle contraction is the adenosine monophosphate activated protein kinase (AMPK), given its role in regulating the proposed master regulator of mitochondrial biogenesis, peroxisome proliferator-activated γ receptor co-activator (PGC-1 α) (Irrcher et al. [2008;](#page-11-0) Jäger et al.

[2007\)](#page-11-0). Indeed, PGC-1 α expression is increased in human skeletal muscle in an exercise-intensity dependent manner that is associated with upstream signalling through AMPK (Egan et al. 2010). The importance of PGC-1 α in regulating mitochondrial content and function is evident from rodent studies demonstrating that overexpression increases oxidative enzyme activity (Lin et al. [2002\)](#page-11-0) and improves exercise capacity (Calvo et al. [2008](#page-10-0)).

In considering possible signals involved in regulating exercise-induced metabolic gene expression, there is growing evidence that reduction in carbohydrate (CHO) availability is a potent stimulus to enhance oxidative capacity which may be mediated, in part, through AMPK signalling. For example, AMPK displays higher activity at rest and following exercise when glycogen is low (Wojtaszewski et al. [2003;](#page-11-0) Yeo et al. [2010\)](#page-11-0) and ingestion of glucose during exercise also attenuates exercise-induced AMPK activation (Akerstrom et al. [2005](#page-10-0)). Furthermore, restricting CHO availability before (Pilegaard et al. [2002](#page-11-0)), during (Cluberton et al. [2005](#page-10-0)) and after exercise (Pilegaard et al. [2005\)](#page-11-0) augments the expression of many metabolic genes related to regulation of substrate utilisation. Data from these acute studies therefore suggested that commencing exercise with reduced muscle glycogen, restricting CHO provision during exercise and deliberately minimising post-exercise muscle glycogen re-synthesis may thereby provide an enhanced stimulus to up-regulate oxidative capacity of skeletal muscle. Accordingly, we (Morton et al. [2009\)](#page-11-0) and others (Hansen et al. [2005;](#page-10-0) Hulston et al. [2010;](#page-10-0) Yeo et al. [2008\)](#page-11-0) subsequently demonstrated that training in conditions of reduced endogenous and exogenous carbohydrate availability enhances traininginduced oxidative adaptations of skeletal muscle. This periodised approach to training and nutrition has since been termed as the train-low compete-high model surmising that selected training sessions be deliberately completed with reduced carbohydrate availability before, during and after exercise but yet competition always be performed with high carbohydrate availability (Burke [2010](#page-10-0)).

Despite the potential of the train-low approach to enhance skeletal muscle adaptations, there are a number of limitations associated with performing chronic exercise in the face of reduced carbohydrate availability. Indeed, reduced muscle glycogen levels compromises absolute training intensity (Hulston et al. [2010](#page-10-0); Yeo et al. [2008\)](#page-11-0) which if performed repeatedly, could lead to a de-training effect. In addition, contractile-induced protein degradation and synthesis are increased and decreased, respectively, to a greater extent when exercise is commenced with reduced glycogen availability compared with elevated glycogen stores (Blomstrand and Saltin [1999;](#page-10-0) Howarth et al. [2010](#page-10-0); Van Hall et al. [1999](#page-11-0)). In this way, chronic exposure to exercise training with low muscle glycogen availability could also lead to a progressive loss of lean mass thereby also negatively impacting upon physical performance. Indeed, we (Morton et al. [2010\)](#page-11-0) and others (Mettler et al. [2010](#page-11-0)) have observed that elite athletes can in fact lose skeletal muscle mass during times of heavy training especially when energy intake and CHO availability are considered sub-optimal.

In an attempt to overcome these limitations, one potential strategy may be to consume intact protein or essential amino acids, before, during and/or after exercise. For example, the provision of nutrients in the form of protein (as opposed to CHO) may still allow for the enhanced cell signalling pathways associated with training with reduced carbohydrate availability but yet would also provide an important substrate to promote skeletal muscle remodelling, and potentially minimise any progressive loss of lean mass during training. Indeed, increased protein availability reduces lean mass loss during heavy training periods when energy and CHO availability is restricted (Morton et al. [2010;](#page-11-0) Mettler et al. [2010](#page-11-0)). In this regard, accumulating data suggest that protein provision before (Coffey et al. [2011](#page-10-0)), during (Hulston et al. [2011](#page-11-0)) and after (Breen et al. [2011;](#page-10-0) Howarth et al. [2009](#page-10-0)) endurance or repeated sprint type protocols, enhances post-exercise skeletal muscle protein synthesis as well as activation of some of the signalling molecules regulating translation initiation and elongation. Aside from the additional energy content per se, there is also some evidence (albeit not consistent) that consumption of branched chain amino acids (BCAAs) during exercise improves exercise performance (Blomstrand et al. [1991](#page-10-0)) and reduces perceptions of effort during sub-maximal exercise that is commenced in a glycogen-depleted state (Blomstrand et al. [1997\)](#page-10-0), possibly through alleviation of central fatigue (Blomstrand [2006](#page-10-0)). Taken together, the above data suggest that protein ingestion before, during and after exercise performed with reduced CHO availability may therefore prove beneficial as it is likely to still allow for the enhanced cell signalling responses associated with reduced muscle glycogen but yet could also improve net protein balance, and potentially increase training capacity.

The aims of the present study were therefore to test the hypotheses that protein ingestion before, during and after exercise commenced in conditions of reduced muscle glycogen: (1) does not attenuate acute cell signalling responses associated with the regulation of mitochondrial biogenesis, (2) increases translational signalling of skeletal muscle protein synthesis and, (3) increases exercise capacity. In this way, we postulated that our experimental findings would have practical applications for those athletes who deliberately incorporate phases of train-low compete-high into their periodised training and nutritional programmes.

Materials and methods

Subjects

Ten recreationally active males (mean \pm SD: age, 22 \pm 2 years; body mass, 79 \pm 9 kg; height, 1.77 \pm 0.05 m; $\overline{VO}_{2\text{peak}}$, 53 \pm 6 ml kg⁻¹ min⁻¹; peak power output, 323 \pm 61 W), who cycled on average 1–6 h per week volunteered to take part in the study. The experimental procedures and potential risks associated with the study were explained and subjects gave written informed consent prior to participation. Subjects refrained from additional exercise outside of the study requirements as well as from alcohol and caffeine intake for at least 48 h prior to any of the testing sessions. None of the subjects had history of neurological disease or musculoskeletal abnormality and none were under pharmacological treatment during the course of the study. The study was approved by the Ethics Committee of Liverpool John Moores University.

Overview of experimental design

In a randomised, double-blind, cross-over and counterbalanced design and after having initially been assessed for maximal oxygen uptake, participants reported to the laboratory on the evening prior to the main experimental trial in order to perform a glycogen-depleting bout of intermittent exhaustive cycling. At the cessation of exercise, participants were deliberately provided with a low-carbohydrate meal $(<50 g)$ to be consumed within a 30 min period. The following morning, participants arrived at the laboratory in a fasted state and performed 45 min of steady-state cycling and a subsequent bout of intermittent exhaustive cycling as a measure of exercise capacity. Subjects ingested a protein (PRO) or placebo (PLA) beverage 45 min prior to exercise, at regular intervals during exercise and immediately post exercise. Muscle biopsies were sampled from the vastus lateralis immediately before and after exercise and also at 3 h post-exercise. Venous blood samples were also obtained at regular intervals during the experimental protocol. Participants returned to the laboratory 7–10 days after their initial trial so as to complete the second trial.

Assessment of maximal oxygen uptake

Peak oxygen uptake $(\dot{V}O_{2\text{peak}})$ and peak power output (PPO) were determined using an incremental cycle test to exhaustion on an electrically braked cycle ergometer (Daum Electronic Premium 8i, Furth, Germany). The test commenced with a 5 min warm-up at 50 W at a cadence corresponding to 60 revs min^{-1} followed by an increase in 30 W every 60 s until volitional exhaustion. Breathby-breath measurements were obtained throughout exercise using a CPX Ultima series online gas analysis system (Medgraphics, MN, US). $\dot{V}O_{2\text{peak}}$ was stated as being achieved by the following end-point criteria: (1) heart rate within 10 beats min^{-1} of age-predicted maximum, (2) respiratory exchange ratio >1.1 , and 3) plateau of oxygen consumption despite increased workload. Peak power output was defined as the highest power output that was able to be maintained for 60 s. The ergometer saddle and handle bar position were recorded for each participant during preliminary testing and replicated during the experimental trials.

Glycogen-depleting protocol

Participants arrived at the laboratory on the evening prior to each main experimental trial (\sim 1800 h) and performed an intermittent bout of cycling to volitional fatigue. After a 5 min warm up at 100 W, participants cycled for 2 min at 90 % PPO, followed immediately by a 2 min recovery period at 50 % PPO. They repeated this work to rest ratio until they could no longer complete 2 min cycling at 90 % PPO, determined as an inability to maintain a cadence of 60 revs min^{-1} . At this point, exercise intensity was lowered to 80 % PPO and when participants could no longer maintain this intensity, it was lowered to 70 % and finally, 60 % PPO with the same work to rest ratio. When the participants were unable to cycle for 2 min at 60 % PPO the exercise protocol was terminated. This intermittent pattern of exercise was employed so as to induce glycogen depletion in both type I and II fibres. Heart rate (HR) was measured continuously throughout exercise (Polar S610i, Kempele, Finland) and ratings of perceived exertion (RPE) were recorded at 2 min intervals during exercise (Borg [1970](#page-10-0)). The activity pattern and total time-to-exhaustion for the glycogen-depleting ride were recorded for each participant during their first trial and replicated during their second trial. Water intake was consumed ad libitum throughout exercise in the initial trial and was recorded and repeated for the subsequent trial. Participants were deliberately provided with a low carbohydrate meal consisting of 400 g of tomato soup (Heinz, UK: 950 kJ; 26.6 g CHO; 11.8 g fat; 3.6 g protein) within 30 min of completion of the depletion ride so as to induce minimal muscle glycogen re-synthesis prior to the main experimental trial on the subsequent morning. No further nutrient provision was provided after consumption of this meal.

Experimental trial

After an overnight fast, participants entered the laboratory on the subsequent morning (\sim 0630 h) and were fitted with a cannula into an antecubital vein. After obtaining a resting blood sample, the cannula was then flushed with \sim 2 ml of sterile saline (Kays Medical Supplies, Liverpool, UK) to keep the cannula patent and sterile, a procedure that was completed after each subsequent blood draw. Having obtained a resting blood sample, participants then consumed 11 of either the PRO (Science in Sport Ltd., Blackburn, UK) beverage-containing casein hydrolysate or a visually identical and taste-matched PLA (see Table 1 for drink compositions). Participants then rested in a supine position for a 45 min period during which further blood samples were obtained at 15 min intervals. After this period, a pre-exercise muscle biopsy was obtained from the vastus lateralis and participants subsequently cycled for 45 min at 50 % PPO. Venous blood samples were obtained at 15 min intervals during exercise and HR (Polar S610i, Kempele, Finland) and $VO₂$ (Medgraphics, MN, US) were measured continuously during exercise. Rates of carbohydrate and lipid oxidation $(g \text{ min}^{-1})$ during exercise were calculated according to the equations presented by Jeukendrup and Wallis [\(2005](#page-11-0)). RPE was also measured at 5 min intervals during exercise (Borg [1970\)](#page-10-0). Participants also ingested a further 500 ml of the PRO or PLA beverage during exercise, administered as four equal doses (125 ml)

Table 1 Nutrient compositions of the protein (PRO) and placebo (PLA) drinks

	PRO	PLA
Energy (kJ kcal ⁻¹ 1^{-1})	417/98	66/15
CHO $(g l^{-1})$	3.5	3.7
Fat $(g 1^{-1})$	0.3	0.1
Protein $(g l^{-1})$	20.3	θ
Alanine $(g l^{-1})$	0.95	Trace
Arginine (g 1^{-1})	1.01	Trace
Aspartic acid (g 1^{-1})	1.09	Trace
Cysteine $(g 1^{-1})$	0.48	Trace
Glutamic acid (g 1^{-1})	3.02	Trace
Glutamine (g 1^{-1})	$\overline{0}$	Trace
Glycine $(g l^{-1})$	1.58	Trace
Histidine (g 1^{-1})	0.39	Trace
Isoleucine(g 1^{-1})	1.50	Trace
Leucine $(g l^{-1})$	3	Trace
Lysine $(g l^{-1})$	1	Trace
Phenylalanine (g 1^{-1})	0.63	Trace
Proline $(g l^{-1})$	1.99	Trace
Serine $(g l^{-1})$	0.78	Trace
Threonine $(g l^{-1})$	0.57	Trace
Tryptophan $(g l^{-1})$	0.34	Trace
Tyrosine $(g l^{-1})$	0.63	Trace
Valine $(g l^{-1})$	1.49	Trace

immediately prior to exercise and after completion of 15, 30 and 45 min of exercise. Once the participants had completed the 45 min steady-state cycling protocol, they remained mounted on the ergometer and performed 5 min of active recovery (cycling at 50 W) before commencing the exercise capacity test. This test consisted of intermittent 1-min bout of cycling at intensities corresponding to 40 and 80 % of each subject's PPO until volitional fatigue. Upon exhaustion, participants dismounted the ergometer and immediately had a post-exercise blood sample and muscle biopsy taken, before consuming a final 1 l bolus of the PRO or PLA solution. In this way, participants ingested a total of 20 g protein prior to exercise, 10 g during exercise and a further 20 g immediately post-exercise. It is acknowledged that protein provision may be prescribed on the basis of g kg^{-1} body mass, however, on the basis of dose-response studies (albeit on resistance exercise, Moore et al. [2009](#page-11-0)) and also because 20 g post-exercise is sufficient to stimulate protein synthesis in the hours following steady-state endurance cycling (Breen et al. [2011](#page-10-0)), we considered an absolute amount of 20 g as a suitable initial dose for which to study. Furthermore, prescribing an absolute dose may also increase practical application for endurance athletes (who usually don't vary considerably in body mass) given that a 20 g bolus is also an easy to administer dose. Ingestion of 10 g of essential amino acids during exercise (60 min cycling at 70 % $\dot{V}O_{2\text{max}}$) has also been shown sufficient to stimulate post-exercise protein synthesis (Pasiakos et al. [2011\)](#page-11-0). As such, we considered our chosen dosing strategy as one that has both practical application and supporting rationale and that may be particularly pertinent to exercise in glycogen-depleted states given the increased demands upon amino acid metabolism in this instance. After the post-exercise biopsy, subjects remained seated in the laboratory (performing light activities such as reading or working on a computer) before having a final muscle biopsy sampled at 3 h post exercise. Laboratory conditions remained constant for both experimental trials (19–21 °C, 40–50 % humidity).

Dietary and physical activity control

Participants recorded their diet for 48 h prior to their initial glycogen-depleting exercise protocol and repeated this dietary intake prior to their second trial. Subsequent dietary analysis was performed using a computer software package (Microdiet, Downlee Systems, UK) and the average total daily energy intake was $2,334 \pm 528$ kcal $(52 \pm 13\%)$ CHO; 32 ± 12 % fat; 15 ± 5 % protein). Participants were instructed to refrain from ingesting alcohol and caffeine-containing food and drink for 48 h prior to each trial and refrain from physical activity during the 3–5 days preceding each trial.

Blood analysis

Blood samples were drawn from a superficial vein in the anticubital crease of the forearm using a cannula intravenous infusion set (NHS Supply Chain, Alfreton, UK). Samples were collected into vacutainers containing EDTA, lithium heparin or serum separation tubes (SST) and stored on ice or at room temperature (SST for approximately 1 h) until centrifugation at $1,500g$ for 15 min at 4 °C. Following centrifugation, aliquots of plasma or serum were stored at -80 °C for later analysis. Samples were analysed for plasma glucose, lactate, NEFA and glycerol concentration using commercially available kits (Randox Laboratories, Antrim, UK). Serum insulin concentration was measured using an automated ELISA insulin test kit (Insulin, Cobas, Roche Diagnostics, IN, USA).

Muscle biopsies

Muscle biopsies were obtained from separate incision sites (2–3 cm apart) from the lateral portion of the vastus lateralis muscle pre-, post- and 3 h after the exercise protocol using a Bard Monopty Disposable Core Biopsy Instrument (12 gauge \times 10 cm length, Bard Biopsy Systems, Tempe, AZ, USA). Samples were obtained (approximately 40–50 mg) under local anaesthesia (0.5 % marcaine) and immediately frozen in liquid nitrogen and stored at -80 °C for later analysis.

Muscle glycogen concentration

Muscle glycogen concentration was determined according to the method described by Van Loon et al. [\(2000](#page-11-0)). Approximately 3–5 mg of freeze dried sample was powdered, dissected free of all visible non-muscle tissue and subsequently hydrolyzed by incubation in 500 μ l of 1 M HCl for 3–4 h at 100 $^{\circ}$ C. After cooling to room temperature, samples were neutralized by the addition of $250 \mu l$ 0.12 mol 1^{-1} Tris/2.1 mol 1^{-1} KOH saturated with KCl. Following centrifugation, $150 \mu l$ of the supernatant was analysed in duplicate for glucose concentration according to the hexokinase method using a commercially available kit (GLUC-HK, Randox Laboratories, Antrim, UK). Glycogen concentration is expressed as mmol kg^{-1} dry weight (dw) and intra-assay coefficients of variation was \leq %.

Western blotting

Approximately 20 mg of frozen muscle was ground to powder and homogenised in 120μ l of ice cold lysis buffer [25 mM Tris/HCl (pH 7.4), 50 mM NaF, 100 mM NaCl, 5 mM EGTA, 1 mM EDTA, 10 mM Na-pyrophosphatase, 1 mM Na3VO4, 0.27 M sucrose, 1 % Triton X-100, 0.1 %

2-mercaptoethanol] and supplemented with a protease inhibitor tablet (Complete mini, Roche Applied Science, West Sussex, UK). Homogenates were centrifuged at 14,000g for 10 min at 4 \degree C and the supernatant was collected. The protein content of the supernatant was determined using a bicinchoninic acid assay (Sigma, UK). Each sample was diluted with an equal volume of $2 \times$ Laemmli buffer (National Diagnostics, USA) and boiled for 5 min at 100 \degree C. 50 µg of protein from each sample was then separated in Tris–glycine running buffer $(10 \times$ Tris/Glycine, Geneflow Ltd, Staffordshire, UK) using self-cast 4 % stacking and 10 % separating gels (National Diagnostics, USA). Gels were transferred semi-dry onto nitrocellulose membrane (Geneflow Ltd, Staffordshire, UK) for 2 h at 200 V and 45 mA per gel in transfer buffers (anode 1; 0.3 M Tris, 20 % methanol, pH 10.4; anode 2; 0.25 M Tris, 20 % methanol, pH 10.4; cathode; 0.4 M 6-amino hexanoic acid, 20 % methanol, pH 7.6). After transfer, membranes were blocked for 1 h at room temperature in Tris-buffered saline (TBST: 0.19 M Tris pH 7.6, 1.3 M NaCl, 0.1 %Tween-20) with 5 % non-fat milk. The membranes were then washed for 3×5 min in TBST before being incubated overnight at 4° C with antibodies for antiphospho AMPK^{Thr172}, eEF2^{Thr56} as well as total protein content of AMPK, GAPDH, eEF2 (Cell Signalling, UK), and PGC-1a (Calbiochem, Merck Chemicals, UK), all at concentrations of 1:1000 in $1 \times$ TBST. The next morning, membranes were washed for a further 3×5 min in TBST and subsequently incubated with anti-species horseradish peroxidise-conjugated secondary antibody (Bio-Rad, UK or Dako, UK) for 1 h at room temperature. After a further 3×5 min washes in TBST, membranes were exposed in a chemiluminescence liquid (SuperSignal, Thermo Fisher Scientific, Rockford, IL, USA) for 5-min. Membranes were visualised using a Bio-Rad Chemi-doc system, and band densities were determined using Quantity One imageanalysis software. Comparative samples from each subject for both exercise protocols were run on the same gel and all gels were run in duplicate to verify responses. It should be noted that all raw densitometry data were used for statistical analysis purposes so as to compare within-subject responses to both the PRO and PLA trials.

Real-time RT-PCR

Total RNA was isolated from muscle biopsies (20 mg) using Trizol reagent (Life Technologies/Invitrogen, USA), according to the manufacturer's protocol. For further purification of the RNA, Qiagen All Prep DNA/RNA Micro Kit (Qiagen, UK) was used using manufacturer's recommended guidelines. The final volume of the purified RNA samples was 14μ l. RNA quality and quantity were determined using Implen Nanophotometer (Implen,

Munchen, Germany) and the RNA was stored at -80 °C. The cDNA was synthesised using random hexamers (Applied Biosystems, Foster City, USA) and Superscript III enzyme (Life Technologies/Invitrogen, USA), using manufacturer's protocol. Gene-specific expression data were obtained using custom-designed primers, supplied by Eurofins (MWG Eurofins, Germany). For PGC-1a, forward primer 5'-TGAGAGGGCCAAGCAAAG-3' and reverse primer 5'-ATAAATCACACGGCGCTCTT-3' were used, and for GAPDH, forward primer 5'-AGCCACATCGCT CAGACAC-3' and reverse primer 5'-GCCCAATACGAC $CAAATCC-3'$ were applied. Each cDNA sample was analysed in triplicate with negative controls using AB 7500 Real-Time Quantitative PCR instrument (Applied Biosystems) and Agilent Brilliant II qPCR Master Mix with Low ROX (Agilent Technologies, USA). One microliter of cDNA together with 200 nM of final concentration of primers was used for each reaction with total volume of 20 µl. The following cycling parameters were used: 50° C for 2 min, initial denaturation at 95 °C for 10 min, followed by 40 cycles of denaturation at 95 \degree C for 15 s and annealing/elongation at 60 $^{\circ}$ C for 1 min. Data were collected and analysed using proprietary software for the instrument (SDS version 1.43, Applied Biosystems, Foster City, USA). Changes in mRNA content were calculated according to the $2^{-\Delta\Delta Ct}$ method where GAPDH was used as the housekeeping gene.

Statistical analysis

Statistical analysis was conducted using the Statistical Package for Social Sciences software programme (version 15). Data were analysed using a two way repeated measures general linear model where the within factors were time and exercise condition (i.e. PLA versus PRO). Where there was a significant interaction effect or main effect of time, Bonferroni post hoc tests were used to locate specific differences. Student's t test for paired samples was also used to compare average physiological responses during the glycogen-depletion protocol as well as exercise capacity and physiological variables at the point of exhaustion. All data in text, figures and tables are presented as means (SD) with P values ≤ 0.05 indicating statistical significance.

Results

Physiological responses to the glycogen-depleting exercise protocol

Exercise time to exhaustion for the glycogen-depletion protocol for both the placebo and protein condition was 64.1 ± 6.4 min. Mean HR and RPE for the placebo and

protein conditions during exercise were 160 ± 15 versus 163 ± 14 b min⁻¹ and 14.7 ± 1.2 versus 15 ± 1.4 , respectively. There was no significant difference in either average heart rate $(P = 0.106)$ or RPE $(P = 0.175)$ between trials.

Physiological responses to the steady-state exercise and exercise capacity test

Muscle glycogen

The glycogen depletion and nutritional protocol was successful in ensuring that participants commenced the steadystate exercise protocol with reduced glycogen availability (placebo and protein conditions = 136 ± 28 and 134 ± 29 mmol kg dw^{-1} , respectively). Exercise decreased muscle glycogen similarly in both trials $(P < 0.001)$ to levels \leq 50 mmol kg dw⁻¹ and there' was also no apparent muscle glycogen re-synthesis at 3 h post-exercise ($P = 0.328$) in either condition (see Fig. 1).

Substrate oxidation

CHO oxidation significantly decreased during steady-state exercise $(P < 0.001)$ though there was no difference $(P = 0.12)$ $(P = 0.12)$ between trials (see Table 2). Similarly, lipid oxidation increased ($P < 0.001$) during exercise though no difference $(P = 0.23)$ was apparent between protein and placebo ingestion (see Table [2](#page-6-0)). Both \overline{VO}_2 and RPE increased (both $P < 0.001$) during steady-state cycling though the magnitude of increase was not different $(P = 0.89$ and 0.92, respectively) between conditions (see Table [2](#page-6-0)).

Fig. 1 Muscle glycogen concentration immediately pre-, post- and at 3 h post-exercise. Asterisk denotes significant difference from preexercise values, $P < 0.05$. Data are presented as mean \pm SD $(n = 10)$

Table 2 Substrate oxidation $\dot{V}O_2$ and RPE during the 45 min steadystate cycling protocol

	Time (min)					
	$10 - 15$	$25 - 30$	$40 - 45$			
CHO oxidation $(g \text{ min}^{-1})$						
Protein	1.86(0.28)	$1.68~(0.34)^a$	1.52 $(0.36)^{a, b}$			
Placebo	1.70(0.32)	1.43 $(0.35)^{a}$	1.25 $(0.38)^{a, b}$			
	Lipid oxidation $(g \text{ min}^{-1})$					
Protein	0.44(0.21)	$0.59(0.24)^{a}$	0.67 $(0.25)^{a, b}$			
Placebo	0.53(0.28)	$0.69(0.30)^a$	0.79 $(0.34)^{a, b}$			
\dot{VO}_2 (% $\dot{VO}_{2\text{peak}}$)						
Protein	46 (10)	48 $(9)^a$	49 $(9)^a$			
Placebo	47 (9)	49 $(9)^a$	50 $(9)^a$			
RPE (AU)						
Protein	12(2)	$13(2)^{a}$	$14 (1)^{a, b}$			
Placebo	12(1)	$13(1)^{a}$	$15 (1)^{a, b}$			

Data are presented as mean \pm SD ($n = 10$)

^a Significant difference from 10 to 15, $P < 0.05$

^b Significant difference from 25 to 30, $P < 0.05$

Plasma metabolites

Plasma glucose, lactate, NEFA, glycerol and serum insulin concentrations are presented in Table 3. Plasma glucose

concentration decreased throughout the experimental protocol ($P < 0.001$) in both conditions with no effect of protein ingestion ($P = 0.98$). Exercise-induced increases in plasma lactate ($P < 0.001$) during the protocol were also not different between conditions $(P = 0.18)$. Plasma NEFA concentration increased during exercise under both placebo and protein ingestion ($P = 0.001$) with a tendency for significant differences between trials $(P = 0.075)$. Similarly, plasma glycerol concentration increased during exercise ($P < 0.001$) and also displayed significant effects for condition ($P = 0.03$) and interaction ($P = 0.003$). In accordance with the provision of amino acids in the protein trial, serum insulin concentration displayed a significant main effect for time ($P = 0.001$), condition ($P = 0.01$) and interaction ($P = 0.002$).

Exercise capacity test and physiological responses

Following the completion of the 45 min steady-state cycling protocol, there was no difference in time to exhaustion ($P = 0.61$) between trials during the subsequent exercise capacity test (24.7 \pm 10.2 and 25.6 \pm 8.7 min for placebo and protein, respectively, see Fig. [2](#page-7-0)). Heart rate $(160 \pm 16$ and 163 ± 15 beats min⁻¹, $P = 0.30$), RPE $(16 \pm 1$ and 16 ± 1 , $P = 0.76$) and plasma glucose $(4.6 \pm 0.97 \text{ and } 4.6 \pm 0.72 \text{ mmol } 1^{-1}, P = 0.88)$, lactate

Table 3 Plasma glucose, lactate, non-esterified fatty acid (NEFA), glycerol and serum insulin concentrations during the experimental protocols

Time (min)								
	-45	-30	-15	θ	$+15$	$+30$	$+45$	
Glucose (mmol 1^{-1})								
Protein	4.8(0.5)	5.1(0.6)	5.0(0.8)	4.9(0.8)	4.4 $(0.4)^a$	4.5 $(0.5)^a$	4.4 $(0.7)^a$	
Placebo	4.8(0.4)	50(0.5)	5.2(0.6)	50(0.4)	4.5 $(0.6)^a$	4.4 $(0.7)^a$	4.3 $(0.8)^a$	
Lactate (mmol 1^{-1})								
Protein	1.1(0.2)	1.1(0.3)	1.1(0.2)	1.2(0.3)	1.8(0.5)	1.7 $(0.3)^b$	1.6 $(0.3)^b$	
Placebo	1.1(0.3)	1.0(0.2)	0.9(0.4)	1.1(0.2)	1.5(0.3)	1.5 $(0.4)^{b}$	$1.5(0.3)^{b}$	
NEFA (mmol 1^{-1})								
Protein	0.66(0.2)	0.51(0.2)	0.48(0.2)	0.46(0.2)	0.45(0.2)	$0.68(0.3)^d$	$0.85(0.4)^c$	
Placebo	0.91(0.4)	0.74(0.2)	0.61(0.1)	0.63(0.2)	0.71(0.3)	$0.94~(0.4)^d$	$1.06~(0.3)^c$	
Glycerol (μ mol 1^{-1})								
Protein* [*]	68 (22)	46(19)	34(11)	33(17)	99 $(46)^{e, s}$	$220(64)^{e, f, s}$	285 $(74)^{e, f, g, \$}$	
Placebo	85 (37)	55 (21)	56 (27)	47 (13)	164 $(50)^{e, s}$	285 $(90)^{e, f, s}$	350 $(100)^{e, f, g, s}$	
Insulin (pmol 1^{-1})								
Protein* \cdot #	34(25)	114 $(87)^{5}$	64 $(28)^{\$}$	$28(20)$ ^h	15 $(11)^{h}$, $\frac{1}{s}$	$10(8)^{h,i, 8}$	$10(9)^{h, i}$	
Placebo	23(14)	33 $(21)^{s}$	31 $(17)^{$	$26 (16)^h$	$8(5)^{h. s}$	7 $(5)^{h, i, s}$	$6(4)^{h, i}$	

Data are presented as mean \pm SD ($n = 10$)

* Significant main effect of condition, $P < 0.05$. $*$ Significant interaction effect, $P < 0.05$. $*$ Significant different from respective time-points between trials, $P < 0.05$

^a Significant difference from -30 , $P < 0.05$. ^b Significant difference from -30 and -15 , $P < 0.05$. ^c Significant difference from -15 , 0 and $+15$, $P < 0.05$. d Significant difference from $+15$, $P < 0.05$. e Significant difference from -45 , -30 , -15 and 0, $P < 0.05$. f Significant difference from $+15$, $P < 0.05$. ^g Significant difference from $+30$, $P < 0.05$. ^h Significant difference from -15 , $P < 0.05$. ⁱ Significant difference from 0, $P < 0.05$

Fig. 2 Exercise capacity during the placebo and protein trials. Data are presented as mean \pm SD ($n = 10$). *Individual lines* also represent capacity of individual participants during each trial

 $(4.5 \pm 1.95 \text{ and } 5.4 \pm 2.55 \text{ mmol l}^{-1}, P = 0.60)$, glycerol $(421.4 \pm 115.4 \text{ and } 390.8 \pm 109.8 \text{ mmol } 1^{-1}, P = 0.30)$ and NEFA (1.35 ± 0.59) and 1.36 ± 0.56 mmol 1^{-1} , $P = 1.0$) were also not different at the point of exhaustion between the placebo and protein trials, respectively.

Skeletal muscle cell signalling responses

Phosphorylation of $AMPK^{Thr172}$ increased by approximately threefold post-exercise $(P = 0.03)$ with no difference $(P = 0.545)$ between conditions and returned to basal levels at 3 h post-exercise (see Fig. 3). There was a statistical trend ($P = 0.08$) for greater activation of eEF2^{Thr} following the ingestion of protein, compared to the placebo

Fig. 3 Phosphorylation of AMPK^{Thr172} expressed relative to total AMPK immediately pre-, post- and at 3 h post-exercise in the placebo and protein trials. Asterisk denotes significant difference from preexercise, $P < 0.05$. Data are presented as mean \pm SD (n = 10)

Fig. 4 Phosphorylation of eEF2^{Thr56} expressed relative to total eEF2 immediately pre-, post- and at 3 h post-exercise in the placebo and protein trials. Data are presented as mean \pm SD ($n = 10$)

solution, as illustrated by the greater dephosphorylation of $eEF2^{Thr56}$ in the protein condition (see Fig. 4).

$PGC-1\alpha$ mRNA and protein content

Muscle PGC-1 α mRNA content increased sixfold at 3 h following exercise ($P = 0.001$) with no difference between conditions ($P = 0.77$; Fig. 5). Total protein content of $PGC-1\alpha$ did not change at any time-point post-exercise $(P = 0.81)$ in either trial (see Fig. [6](#page-8-0)).

Fig. 5 PGC-1a mRNA content expressed relative to GAPDH immediately pre-, post- and at 3 h post-exercise in the placebo and protein trials. Asterisk denotes significant difference from preexercise, $P < 0.05$. Data are presented as mean \pm SD (n = 10)

Fig. 6 Total PGC-1a protein content expressed relative to GAPDH immediately pre-, post- and at 3 h post-exercise in the placebo and protein trials. Data are presented as mean \pm SD ($n = 10$)

Discussion

Although recent data demonstrate that training with reduced CHO availability enhances oxidative adaptations of skeletal muscle (Hansen et al. [2005](#page-10-0); Hulston et al. [2010](#page-10-0); Morton et al. [2009](#page-11-0); Yeo et al. [2008\)](#page-11-0), incorporating this approach to training may have detrimental effects if performed long-term such as reduced absolute training intensity (Hulston et al. [2010](#page-10-0); Yeo et al. [2008](#page-11-0)) and potentially loss of lean mass (Morton et al. [2010\)](#page-11-0). With this in mind, we postulated that provision of nutrients in the form of protein (as opposed to CHO) may still allow for the activation of acute cell signalling pathways associated with regulating the enhanced training response in CHO restricted states but yet also provide important substrate to promote skeletal muscle remodelling. To this end, we employed an experimental design in which subjects performed acute exercise in a glycogen-depleted state but with consumption of protein or placebo before, during and after exercise. Given that subjects replicated the precise pattern of intensities and duration of the glycogen-depletion protocol between trials (thus resulting in similar heart rates and RPE) and that pre-exercise glycogen availability prior to the main experimental trial was almost identical, our data indeed suggest that subjects commenced the experimental trial in a similar metabolic state and with similar CHO availability. Importantly, we provide novel data by demonstrating that protein ingestion before and during glycogen-depleted exercise does not impair the magnitude of AMPK^{Thr172} phosphorylation or PGC-1 α gene expression. Additionally, we observed that protein feeding before, during and after exercise induced a trend (which we consider indicative of physiological significance) for greater dephosphorylation and hence, activation of eEF2. These data were therefore considered to have important practical implications in that athletes who deliberately incorporate the train-low compete-high model into their periodised training programmes may consume protein in close proximity to the exercise stimulus without apparently negating AMPK mediated signalling.

A contractile-induced increase in AMPK activation is a well-documented component of a signalling cascade thought to regulate the early transcriptional responses to exercise (Hood [2009](#page-10-0)). Given that glycogen-binding domains have been observed on AMPK subunits (McBride et al. [2009\)](#page-11-0), commencing exercise with reduced muscle glycogen levels increases the activity of AMPK α 2 (Wojtaszewski et al. 2003) and AMPK^{Thr172} phosphorylation in human skeletal muscle (Yeo et al. [2008\)](#page-11-0). Augmented signalling through AMPK in a glycogen-depleted state has therefore been suggested as a contributory mechanism underpinning the enhanced oxidative adaptations associated with training with reduced carbohydrate availability (Hansen et al. [2005](#page-10-0); Hulston et al. [2010](#page-10-0); Morton et al. [2009](#page-11-0); Yeo et al. [2008\)](#page-11-0). Although the present study design did not incorporate a high glycogen trial, the threefold increases in $AMPK^{Thr172}$ phosphorylation immediately post-exercise is consistent with the magnitude of increase observed by other authors when exercise has been commenced in a glycogen-depleted state compared with high glycogen availability (Yeo et al. [2010](#page-11-0)). Importantly, consuming 20 g of protein 45 min before exercise and a further 10 g during exhaustive exercise does not cause any impairment in AMPK signalling, thus demonstrating that protein ingestion still allows for the increased magnitude of AMPK phosphorylation associated with reduced glycogen availability.

In accordance with the role of AMPK as an upstream signalling kinase regulating PGC-1 α expression (Jäger et al. [2007](#page-11-0)), we observed a similar sixfold increase in PGC-1 α mRNA between trials at 3 h post-exercise with no concomitant changes in PGC-1 α protein. This time-course of PGC-1 α expression is consistent with data from our laboratory (Bartlett et al. [2012](#page-10-0)) and others (Little et al. [2010,](#page-11-0) [2011](#page-11-0)) and indeed, it is now well documented that increased $PGC-1\alpha$ protein is typically only observed in the days following an acute exercise bout (Little et al. [2010,](#page-11-0) [2011](#page-11-0)) or with repeated days and weeks of exercise (Perry et al. [2010](#page-11-0)). In this regard, future studies should include additional sampling points in an extended recovery period (i.e. >24 h post-exercise) and/or with repeated bouts of glycogendepleted exercise (i.e. exercise training) so as to ascertain the role of increased protein availability in stimulating upregulation of oxidative proteins within a more physiologically relevant time-course. We also acknowledge that measurement of nuclear and mitochondrial translocation of $PGC-1\alpha$

as well as acetylation status would have also been advantageous to have been measured here (Philp et al. [2011\)](#page-11-0) though the limited tissue obtained by our chosen biopsy technique did not permit this analysis.

There is now a growing body of literature examining the regulation of skeletal muscle protein synthesis in the early recovery period (i.e. ≤ 4 h) following acute endurance type exercise as opposed to the more frequently studied resistance protocols. It is generally thought muscle protein synthesis is suppressed during exercise, possibly due to Ca^{2+} induced activation of eEF2 kinase (Rose et al. [2009\)](#page-11-0) such that eEF2 becomes phosphorylated (Rose et al. [2005\)](#page-11-0) and hence deactivated during exercise thereby suppressing translational elongation. In contrast, endurance exercise elevates muscle protein synthesis in the recovery period even when the subjects remain in a fasted state (Mascher et al. [2011](#page-11-0)), though it is well documented that exercise and post-exercise feeding act synergistically to elevate synthesis rates and ensure that net protein balance becomes positive. Indeed, Hulston et al. [\(2011\)](#page-11-0) observed that co-ingestion of protein with carbohydrate during 3 h cycling exercise and throughout recovery increased net leg protein balance during exercise and recovery as well as enhancing post-exercise mixed muscle protein synthesis when compared with carbohydrate per se. Interestingly, Breen et al. [\(2011](#page-10-0)) also observed that the enhanced post-exercise protein synthesis (in a 4 h recovery period) induced by coingestion of protein with carbohydrate immediately postexercise is specifically due to increased rates of myofibrillar and not mitochondrial protein synthesis. The latter data are of particular importance for the present design given that our participants commenced exercise in a glycogen-depleted state, a condition which exacerbates contractile-induced increases and decreases in muscle protein degradation and synthesis, respectively (Howarth et al. [2010\)](#page-10-0). The repeated absence of protein provision in the early recovery period following glycogen-depleting exercise may therefore lead to a progressive loss of myofibrillar protein, muscle mass and hence, an impaired capacity to generate force.

Although the present study is limited in that we did not directly measure muscle protein turnover via stable isotope methods, we chose to measure the phosphorylation status of eEF2 given its role as a molecular regulator of elongation. Making inferences on protein turnover based on snapshots of signalling molecules is of course not without limitations. However, it is important to note that Breen et al. [\(2011](#page-10-0)) observed that the enhanced myofibrillar protein synthesis induced by post-exercise protein feeding was associated with reduced phosphorylation (i.e. hence increased activation) of $eEF2^{Thr56}$ at 4 h post-exercise, therefore supporting the role of eEF2 as a key regulator of muscle protein synthesis. Our data are in agreement with these authors as we also observed that protein feeding apparently reduces phosphorylation of $eEF2^{Thr56}$ at 3 h post-exercise (we acknowledge that our chosen sample size may have been underpowered to achieve full statistical significance). Furthermore, we also provide novel data by demonstrating that protein feeding before and during glycogen-depleting exercise appears to reduce $eE F2^{Thr56}$ phosphorylation immediately post-exercise, thereby suggestive of improved metabolic conditions to attenuate the contraction-induced phosphorylation of $eEF2^{Thr56}$ (Rose et al. [2005](#page-11-0)) and hence reduce the suppression in muscle protein synthesis rates which occur during exercise (Rose et al. [2009](#page-11-0)). Future studies utilising stable isotope methodology are now required to fully test this hypothesis. Unfortunately, the limited tissue obtained by our chosen biopsy technique also did not permit us to measure other important signalling nodes regulating protein synthesis (e.g. mTOR, p70S6 kinase etc.) or indeed, markers of muscle protein degradation such as the atrogenes.

Given that training power outputs are reduced in a glycogen-depleted state compared with normal glycogen availability (Hulston et al. [2010;](#page-10-0) Yeo et al. [2008\)](#page-11-0), we also tested the hypothesis that protein ingestion before and during glycogen-depleting exercise would be able to restore exercise capacity whilst simultaneously allowing for the enhanced signalling responses associated with low glycogen stores. Aside from additional energy per se, this hypothesis was based on a small body of literature demonstrating that branched chain amino acid supplementation (BCAAs) reduces central fatigue and enhances performance, as evidenced by decreased ratings of perceived exertion (RPE) (Blomstrand et al. [1997](#page-10-0); Greer et al. [2007\)](#page-10-0) and improved time to exhaustion and time-trial perfor-mance (Blomstrand et al. [1991;](#page-10-0) Crowe et al. [2006](#page-10-0)). Ergogenic mechanisms of action in these studies have been postulated to be due to increased plasma concentrations of BCAA relative to free tryptophan (f-Trp) such that f-Trp entry to the brain is reduced thereby suppressing the synthesis of the neurotransmitter 5-hydroxytryptamine (5-HT) and ultimately reducing central fatigue (Blomstrand [2006](#page-10-0); Newsholme and Blomstrand [2006](#page-11-0)). Although we did not measure plasma amino acid concentrations, Watson et al. [\(2004](#page-11-0)) observed BCAA ingestion neither reduce RPE nor improve exercise capacity at 50 % $\dot{V}O_{2\text{max}}$ when commenced in a glycogen-depleted state, despite a fourfold reduction in the plasma concentration of f-TRP to BCAA. When taken together, both the present data and that of the aforementioned researchers suggest that essential amino acid provision offers no advantage to exercise capacity or subjective feelings of perceived exertion when exercise is commenced in a glycogen-depleted state.

A limitation of our chosen experimental design was that post-exercise biopsies were sampled at the point of exhaustion and not at matched time-points across both the PLA and PRO trials. Indeed, although we observed no significant differences in mean exercise capacity between trials (25 ± 10) and 26 ± 9 min for PLA and PRO, respectively, $P = 0.61$, it is important to consider individual differences in exercise capacity (range: -9 to 7 min difference between trials) and as such, differences in total exercise duration may have contributed to subtle differences in cell signalling responses. Nevertheless, because one of our experimental aims was to assess the effects of protein ingestion on exercise capacity, we felt that disrupting the subject from their natural flow of exercise (as well as including a brief rest period in order to perform a time-matched biopsy) followed by subsequent exercise until exhaustion may not have allowed us to appropriately address this experimental aim. Future studies may therefore benefit from adopting a time-matched biopsy approach.

In summary, we provide novel data by demonstrating that protein ingestion before, during and after exercise does not attenuate AMPK signalling in human skeletal muscle when exercise is commenced in a glycogen-depleted state. As such, we therefore consider the present data to be of practical implications for those athletes who deliberately incorporate the train-low compete-high approach into their training programmes as protein does not attenuate activation of a cell signalling cascade thought to contribute to the enhanced training response observed when restricting CHO before, during and after training. Further acute studies utilising a greater portfolio of measures of protein synthesis and degradation are now warranted as well as longitudinal studies investigating the effects of protein ingestion on training adaptations (and estimates of lean muscle mass) when in glycogen-depleted states.

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