

Pharmacological activation of the pyruvate dehydrogenase complex reduces statin-mediated upregulation of FOXO gene targets and protects against statin myopathy in rodents

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Key points

- Statin myopathy impairs phosphatidylinositol 3-kinase/Akt signalling and activates forkhead box protein O (FOXO) transcription factors *in vivo* in rodent skeletal muscle. This is associated with upregulation of downstream gene targets known to increase proteasomal and lysosomal-mediated protein breakdown, oxidative stress and inflammation, and inhibit muscle carbohydrate (CHO) oxidation.
- We hypothesised that forcibly increasing muscle CHO oxidation *in vivo*, using the pyruvate dehydrogenase complex activator, dichloroacetate (DCA), would blunt statin-mediated increases in mRNA expression of these FOXO gene targets, thereby reducing statin myopathy.
- Chronic administration of DCA with simvastatin dampened statin-mediated increases in muscle atrophy F-box (MAFbx), cathepsin-L and pyruvate dehydrogenase kinase-4 mRNA in a dose-dependent manner, which was corroborated by protein activity and expression measurements, and blunted statin myopathy.
- These results provide convincing evidence that pharmacologically increasing muscle CHO oxidation reduces simvastatin-induced myopathy by dampening the upregulation of genes known to increase proteasomal and lysosomal protein breakdown and inhibit CHO oxidation.

Abstract We previously reported that statin myopathy is associated with impaired carbohydrate (CHO) oxidation in fast-twitch rodent skeletal muscle, which we hypothesised occurred as a result of forkhead box protein O1 (FOXO1) mediated upregulation of pyruvate dehydrogenase kinase-4 (PDK4) gene transcription. Upregulation of FOXO gene targets known to regulate proteasomal and lysosomal muscle protein breakdown was also evident. We hypothesised that increasing CHO oxidation *in vivo*, using the pyruvate dehydrogenase complex (PDC) activator, dichloroacetate (DCA), would blunt activation of FOXO gene targets and reduce statin myopathy. Female Wistar Hanover rats were dosed daily for 12 days (oral gavage) with either vehicle (control, 0.5% w/v hydroxypropyl-methylcellulose 0.1% w/v polysorbate-80; $n = 9$), 88 mg kg⁻¹ day⁻¹ simvastatin ($n = 8$), 88 mg kg⁻¹ day⁻¹ simvastatin + 30 mg kg⁻¹ day⁻¹ DCA ($n = 9$) or 88 mg kg⁻¹ day⁻¹ simvastatin + 40 mg kg⁻¹ day⁻¹ DCA ($n = 9$). Compared with control, simvastatin reduced body mass gain and food intake, increased muscle fibre necrosis, plasma creatine kinase levels, muscle PDK4, muscle atrophy F-box (MAFbx) and cathepsin-L mRNA expression, increased PDK4

protein expression, and proteasome and cathepsin-L activity, and reduced muscle PDC activity. Simvastatin with DCA maintained body mass gain and food intake, abrogated the myopathy, decreased muscle PDK4 mRNA and protein, MAFbx and cathepsin-L mRNA, increased activity of PDC and reduced proteasome activity compared with simvastatin. PDC activation abolished statin myopathy in rodent skeletal muscle, which occurred at least in part via inhibition of FOXO-mediated transcription of genes regulating muscle CHO utilisation and protein breakdown.

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Abbreviations BCAA, branched-chain amino acid; CHO, carbohydrate; CK, creatine kinase; DCA, dichloroacetate; FOXO, forkhead box protein O; HMBS, hydroxymethylbilane synthase; LC3, microtubule-associated protein 1, light chain 3; MAFbx, muscle atrophy F-box; MuRF-1, muscle RING finger-1; PDC, pyruvate dehydrogenase complex; PDK, pyruvate dehydrogenase kinase; PI3k, phosphatidylinositol 3-kinase.

Introduction

Statins inhibit synthesis of mevalonate by inhibiting 3-hydroxy-3-methyl-glutaryl coenzyme A reductase, the rate-limiting step in hepatic cholesterol and isoprenoid biosynthesis. Statins are used clinically for cholesterol reduction in hypercholesterolaemia, and their therapy has been associated with a 30% reduction in cardiovascular events in patients with vascular disease (Vaughan & Gotto, 2004). Statins have also been found to be protective against heart failure in patients where cholesterol levels are not elevated (Ridker *et al.* 2008). Furthermore, because of reported pleiotropic effects, statin use has expanded to the treatment of other conditions, including ventricular arrhythmias, idiopathic dilated cardiomyopathy, cancer, osteoporosis and diabetes (Dirks & Jones, 2006).

Statins are generally well tolerated, but can have severe myopathic effects, albeit relatively infrequently (Thompson *et al.* 2003). Statin-related myopathy has varying degrees of severity, ranging from muscle myositis and myalgia (muscle aches or weaknesses with and without increased serum creatine kinase (CK) concentration, respectively) and, in the severest case, rhabdomyolysis (>10 times the upper limit of normal serum CK). Randomised trials suggest rhabdomyolysis is a rare event, and even myalgia and myositis, although more common, are not seen as being highly prevalent (Thompson *et al.* 2003). Nevertheless, in general practices the incidence of adverse muscular events associated with the most commonly prescribed statin, simvastatin, is high, particularly at higher doses (>40 mg; 18%; Bruckert *et al.* 2005). Furthermore, recent meta-data analyses have associated intensive high-dose statin therapy with increased risk of new-onset diabetes compared with moderate-dose therapy (Sattar *et al.* 2010; Preiss *et al.* 2011). Given the recommendation by the National Institute of Clinical Excellence (NICE) to prescribe simvastatin over other statins (NICE, 2006), and to prescribe high-dose statin therapy (80 mg day⁻¹) to older

patients (Deedwania *et al.* 2007) when in fact older age per se has been reported to be a major myopathy risk factor (Nichols & Koro, 2007), adverse muscular events could potentially further increase with increased use. Current treatment for statin myopathy is the discontinuation of statin use, in which symptoms are most often reversible (Dirks & Jones, 2006), and therefore it is important to identify the mechanism of statin-induced myopathy.

Using an animal model of statin-induced myopathy (Westwood *et al.* 2005), we have previously shown simvastatin administration impairs phosphatidylinositol 3-kinase (PI3k)/Akt signalling in muscle, resulting in the dephosphorylation (and activation of) the forkhead box protein O (FOXO) (1 and 3) transcription factors, and the induction of ubiquitin and lysosomal proteolysis through upregulation of the FOXO downstream target genes muscle atrophy F-box (MAFbx), muscle RING finger-1 (MuRF-1) and cathepsin-L mRNA. Furthermore, these changes preceded a statin-induced decline in muscle mass reported as the protein:DNA ratio (Mallinson *et al.* 2009). Carbohydrate (CHO) oxidation was also impaired in simvastatin-treated animals, reflected by a decline in muscle glycogen utilisation and a marked increase in the FOXO downstream target pyruvate dehydrogenase kinase-4 (PDK4) mRNA expression above control (Mallinson *et al.* 2009). Combined muscle protein loss and impairment of muscle glycogen oxidation, in parallel with blunted Akt signalling and activated FOXO and its target genes, have also been reported with fibrate therapy (Motojima & Seto, 2003; Constantin *et al.* 2007), in an *in vivo* model of endotoxaemia (Crossland *et al.* 2008) and in critical care patients (Constantin *et al.* 2011). Additionally, exposure of human rhabdomyosarcoma cells to statins has been shown to increase protein degradation by induction of autophagy (Araki & Motojima, 2008). A hallmark of muscle autophagy during *in vivo* catabolic conditions, which include starvation (Lecker *et al.* 2004; Mammucari *et al.* 2007), denervation (Mammucari *et al.*

2007), cancer cachexia and diabetes (Lecker *et al.* 2004), is upregulation of the microtubule-associated protein 1, light chain 3 (LC3) gene. However, to date, no study has examined the role of autophagy in an *in vivo* model of statin-induced myopathy.

Upregulation of PDK2 and PDK4, the main isoforms identified in skeletal muscle (Bowker-Kinley *et al.* 1998), phosphorylate and inactivate muscle pyruvate dehydrogenase complex (PDC; Wieland, 1983), thereby limiting the rate of CHO oxidation. PDC is a multi-enzyme complex located on the inner mitochondrial membrane that catalyses the irreversible conversion of pyruvate to acetyl-CoA (Denton *et al.* 1972). Dichloroacetate (DCA) is a halogenated carboxylic acid that has been shown to increase the activity of the PDC (Whitehouse & Randle, 1973; Whitehouse *et al.* 1974) by competitively inhibiting PDK2 and PDK4 (Bowker-Kinley *et al.* 1998). Several studies have demonstrated DCA increases CHO oxidation and reduces muscle lactate accumulation, through increased PDC activation, in rat (Clark *et al.* 1987), canine (Timmons *et al.* 1996, 1998c; Grassi *et al.* 2002) and human skeletal muscle (Timmons *et al.* 1998a; Constantin-Teodosiu *et al.* 1999, 2012). In keeping with this, DCA has also been shown to decrease blood glucose concentration in healthy humans (Wells *et al.* 1980; Jahoor *et al.* 1989) and patients with diabetes (Stacpoole *et al.* 1978), probably due to increased muscle and liver oxidative glucose disposal (Shangraw *et al.* 1989; Constantin-Teodosiu *et al.* 1999).

Interestingly, DCA administration has also been linked to the regulation of muscle protein turnover. Goodman *et al.* reported DCA attenuated muscle branched-chain amino acid (BCAA) accumulation in the perfused rat hind limb in the fed state, which may reflect a diminished net muscle proteolysis. Furthermore, DCA administration has been used to increase muscle glutamine concentrations in an attempt to increase muscle protein synthesis in burn injury (Ferrando *et al.* 1998), sepsis (Vary *et al.* 1988) and diabetes (Goodman *et al.* 1978), although the acute elevation did not affect muscle protein metabolism.

In the present study we aimed to test the hypothesis that forcibly increasing muscle CHO oxidation by increasing PDC activity and flux, using DCA, would suppress statin-mediated upregulation of FOXO target genes, thereby blunting muscle necrosis and the rise in plasma CK levels in an *in vivo* rodent model of statin myopathy. To do this we measured muscle PDC activity and flux through the PDC reaction, mRNA and protein expression levels of muscle PDK4, mRNA expression levels of the other FOXO gene targets MAFbx and cathepsin-L, and also muscle proteasome and cathepsin-L activity, and related these events to functional end-points, namely body mass and food intake, and the incidence of muscle necrosis and plasma CK concentration. Using this approach we hoped to dampen the myopathic effects associated

with simvastatin, thereby providing novel and important insight into a pharmacological approach directed towards the treatment of statin-induced myopathy.

Methods

Ethical approval

This study was planned in accordance with the standards of animal care and ethics described in 'Guidance on the Operations of the Animal (Scientific Procedures) Act 1986' issued by the UK Home Office, and was conducted so that any clinical expression of toxicity remained within a moderate severity limit as described in guidelines agreed with the UK Home Office Inspector.

Animals and treatments

Simvastatin lactone (purity 98.4%) was obtained from Wuhan S&M Biochemie (Wuhan Hubei, China). The statin was formulated for dosing as a suspension in water containing 0.5% w/v hydroxypropyl methylcellulose and 0.1% w/v polysorbate 80. Sodium DCA was obtained from Sigma-Aldrich (St Louis, MO, USA). Female Wistar Hanover rats, substrain Crl:WI (Glx/BRL/Han) BR (Charles River, UK), aged 8 weeks, were multiple-housed appropriate to each study and were acclimatised for 6 days. The animal rooms were illuminated in a 12 h light/dark cycle, and temperature and humidity were controlled (limits $21 \pm 2^\circ\text{C}$ and $55 \pm 15\%$ relative humidity). Pelleted RM1(E) SQC rodent diet and drinking water were freely available. Animals were dosed with 88 mg kg^{-1} body weight per day ($\text{mg kg}^{-1} \text{ day}^{-1}$) of simvastatin for 12 days (simvastatin, $n = 8$), $88 \text{ mg kg}^{-1} \text{ day}^{-1}$ simvastatin and $30 \text{ mg kg}^{-1} \text{ day}^{-1}$ DCA for 12 days (simvastatin + DCA 30 mg, $n = 9$), $88 \text{ mg kg}^{-1} \text{ day}^{-1}$ simvastatin and $40 \text{ mg kg}^{-1} \text{ day}^{-1}$ DCA for 12 days (simvastatin + DCA 40 mg, $n = 9$) or vehicle (0.5% w/v hydroxypropyl methylcellulose and 0.1% w/v polysorbate 80) for 12 days (control, $n = 9$) by oral gavage. The animals in the simvastatin + DCA groups were primed with $50 \text{ mg kg}^{-1} \text{ day}^{-1}$ DCA for 5 days prior to the administration of simvastatin, and DCA was given for a further 24 h after the simvastatin administration ceased at 12 days to ensure PDC was fully activated (Fig. 1).

Statin administration is known to predominantly affect type IIB fibres (Westwood *et al.* 2005, 2008), and therefore biceps femoris muscle, composed of approximately 70% type IIB fibres (Armstrong & Phelps, 1984), was investigated in this study. Muscles were harvested under terminal anaesthesia with halothane, immediately snap-frozen and stored in liquid nitrogen.

Quantification of muscle necrosis

Muscle necrosis was quantified in biceps femoris sections using the method previously described (Westwood *et al.* 2005). The magnitude of necrosis was graded as follows: minimal, up to 10 muscle fibres affected in whole section; mild, up to 20% of fibres in section affected; moderate, up to 50% of fibres in section affected; and severe, more than 50% of fibres in section affected (Westwood *et al.* 2005). Detailed histological images depicting muscle necrosis in the model used in the present study were presented in Westwood *et al.* (2005). Plasma CK concentration (IU l^{-1}) was also determined using a spectrophotometric assay (Szasz *et al.* 1976).

Muscle cathepsin-L, MAFbx, LC3 and PDK4 mRNA expression

RNA was extracted from approximately 30 mg snap-frozen tissue using RNA Plus (Qbiogene, Uxbridge, UK). First strand cDNA was synthesised from 1 μg total RNA, using Powerscript reverse transcriptase (BD Biosciences, Oxford, UK) and random primers (Promega, Southampton, UK). A reaction was also carried out with the reverse transcriptase omitted to assess genomic DNA contamination.

All reactions were performed in the ABI Prism 7000 Sequence Detection System (Applied Biosystems, Foster City, CA, USA). Each well contained 2 μl cDNA, 12.5 μl Universal Taqman 2 \times PCR master mix (Eurogentec, Liege, Belgium), 18 μM primer, 5 μM probe and 9.25 μl

RNase-free water to make a volume of 25 μl . Each sample was measured in duplicate. The housekeeping gene used was hydroxymethylbilane synthase (HMBS), which was selected because it was not affected by the statin treatment (data not shown). The thermal cycling conditions used were 2 min at 50°C, 10 min at 95°C, followed by 40 cycles at 95°C for 15 s and 60°C for 1 min. Genes analysed were cathepsin-L (Rn01442127), MAFbx (Rn00591730), LC3 (Rn00563181) and PDK4 (Rn00585577; Applied Biosystems). Relative quantification of the genes was determined from linear interpolation of standard curves constructed using pooled cDNA synthesised from extracted RNA from control and treated tissue. The relative values for the gene of interest were normalised with the relative values of HMBS (Rn00565886). The control group was used as calibrator with a value of 1.

Muscle PDK4 protein expression

Protein was extracted from approximately 30 mg snap-frozen tissue using a method modified from Blough *et al.* (1999). Samples were homogenised in 50 mM Tris buffer pH 7.5 with the addition of protease and phosphatase inhibitors (Sigma-Aldrich, UK), centrifuged and the supernatant containing the cytosolic protein fraction was collected. Protein content was measured using the Bradford assay. Protein samples were run on a 4–12% Bis-tris acrylamide gel (Invitrogen, Paisley, UK) at 200 V for 1 h and transferred onto a polyvinylidene difluoride membrane for 1 h 45 min. Protein transfer was checked with Ponceau S staining before being blocked in 5% BSA-TBS-Tween. Membranes were probed with PDK4 primary antibody (1:500; Abcam, Cambridge, UK) overnight at 4°C. Membranes were washed with TBS-Tween and incubated with the secondary antibody HRP-linked anti-rabbit IgG (GE Healthcare, Amersham, UK; 1:2000) in 1% BSA-TBS-Tween. Membranes were incubated with ECL chemiluminescence detection reagent (Pierce, Cramlington, UK) and exposed to X-ray film. Blots were scanned using a Duoscan T1200, Agfa, Mortsel, Belgium. Bands were identified with Genetools from Syngene (Cambridge, UK), and the density volume was adjusted by subtracting the local background and then normalised with actin (1:1000; Sigma-Aldrich, UK).

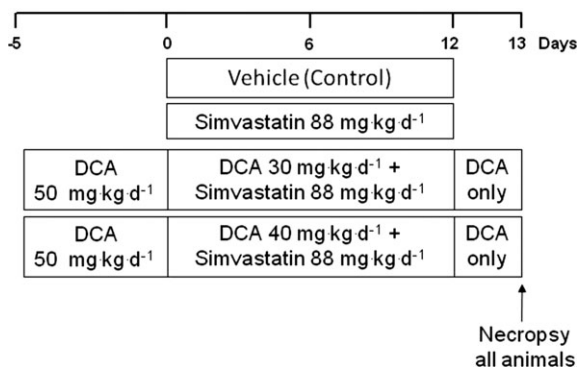


Figure 1. Protocol of simvastatin and dichloroacetate (DCA) dosing during the study

Vehicle (0.5% w/v hydroxypropyl methylcellulose and 0.1% w/v polysorbate 80, control) and simvastatin (88 mg kg⁻¹ day⁻¹) were given daily for 12 days. In the DCA groups, animals were primed with DCA (50 mg kg⁻¹ day⁻¹) for 5 days before dosing with either DCA 30 mg kg⁻¹ day⁻¹ + simvastatin 88 mg kg⁻¹ day⁻¹ or DCA 40 mg kg⁻¹ day⁻¹ + simvastatin 88 mg kg⁻¹ day⁻¹ for 12 days. DCA dosing continued for a further 24 h after the simvastatin administration ceased at 12 days. All animals were necropsied on day 13.

Muscle PDC activity

A small portion of frozen 'wet' muscle was used to determine PDC activity as previously described (Constantin-Teodosiu *et al.* 1991). Briefly, the activity of PDC in its dephosphorylated active form (PDCa) was assayed in a buffer containing NaF and DCA, and was expressed as a rate of acetyl-CoA formation ($\text{mmol min}^{-1} \text{kg}^{-1}$ wet muscle) at 37°C.

Muscle acetylcarnitine content

After removal of visible blood and connective tissue, freeze-dried muscle samples were powdered and extracted with 0.5 mol l^{-1} perchloric acid (containing 1 mmol l^{-1} EDTA), and neutralised with 2.2 mmol l^{-1} KHCO_3 . Acetylcarnitine was measured as an index of PDC flux by a radioisotope enzymatic assay as previously described (Cederblad *et al.* 1990), as an indicator of the magnitude of flux through the PDC reaction (Constantin-Teodosiu *et al.* 1993).

Muscle 20S proteasome and cathepsin-L activity

Frozen muscle (15–25 mg) was homogenised, and the soluble muscle extract used to measure the chymotrypsin-like activity of the 20S proteasome using the fluorogenic substrate N-Suc-Leu-Leu-Val-Tyr-7-amido-4-methyl-coumarin (Sigma-Aldrich), according to Dawson *et al.* (1995). The same homogenate was used to determine cathepsin-L activity using the substrate Z-Phe-Arg-7-amido-4-methyl-coumarin (Sigma-Aldrich), according to Bergmeyer (1983).

Statistics

Data are expressed as mean \pm SEM. Treatment effects were investigated using one-way ANOVA. When a significant *F*-ratio was found, a least significance difference *post hoc* test was performed to locate specific differences. A significance level of $P < 0.05$ was used.

Results

DCA prevents the reduced food intake and body mass loss associated with high-dose simvastatin administration

Average daily food intake declined from day 5 in the simvastatin and simvastatin + DCA 30 mg groups ($P < 0.05$ compared with control; Fig. 2A). By the end of the study, food consumption was similar in the control and simvastatin + DCA 30 mg groups, but approximately 25% lower in the simvastatin group ($P < 0.001$, compared with control). There was a decline in the average body mass of both the simvastatin and simvastatin + DCA 30 mg groups from day 7 ($P < 0.05$ compared with control; Fig. 2B). In both control and simvastatin + DCA 40 mg groups body mass had increased by 10% at the end of the study ($P < 0.001$, $P < 0.05$, respectively, compared with simvastatin).

DCA administration protects against simvastatin-induced myopathy

Simvastatin increased mean plasma CK levels 95-fold above control ($P < 0.05$; Fig. 3). There was a trend ($P = 0.10$) for plasma CK to be reduced in simvastatin + DCA 30 mg, whilst the administration of simvastatin + DCA 40 mg completely abolished this increase in plasma CK with simvastatin ($P < 0.05$ vs. simvastatin). In line with these observations, histological analysis revealed evidence of muscle damage (necrosis) in

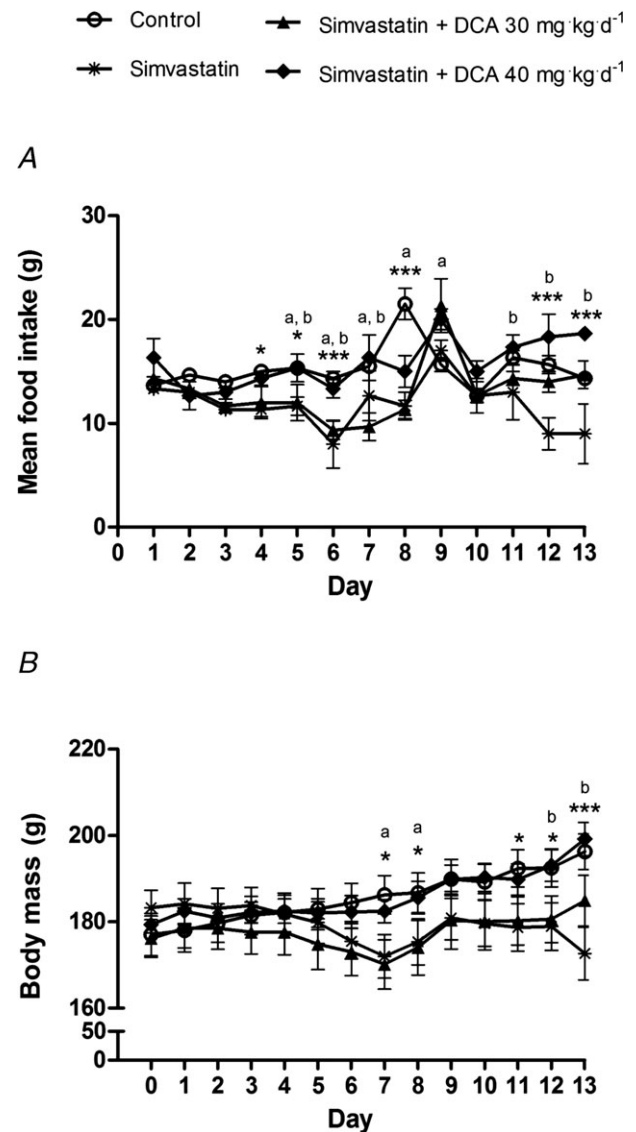


Figure 2. Daily food intake (A, g) and body mass (B, g) of control, simvastatin- and simvastatin plus dichloroacetate (DCA)-treated animals

Values expressed as mean \pm SEM. * $P < 0.05$, *** $P < 0.001$ when control compared with simvastatin, ^a $P < 0.05$ when control compared with simvastatin + DCA 30 mg, ^b $P < 0.05$ when simvastatin compared with simvastatin + DCA 40 mg.

simvastatin compared with control (Table 1). The severity grade and frequency of muscle necrosis was unchanged from simvastatin to simvastatin + DCA 30 mg. However, the severity of muscle necrosis was similar to control in simvastatin + DCA 40 mg.

DCA reverses the simvastatin-induced increase in muscle PDK4 mRNA and protein expression, and inhibition of muscle PDC activity and acetylcarnitine accumulation

Muscle PDK4 mRNA expression in the simvastatin-treated group was elevated threefold above control ($P < 0.001$; Fig. 4A). In keeping with this, PDK4 protein expression was also increased 0.5-fold above control ($P < 0.05$; Fig. 4B). Compared with simvastatin, mRNA and protein expression levels were dampened in simvastatin + DCA 30 mg ($P < 0.001$, $P < 0.05$, respectively) and simvastatin + DCA 40 mg ($P < 0.001$, $P < 0.05$, respectively), and were no different from expression levels seen in control. Muscle PDC activity was reduced in simvastatin compared with control ($P < 0.05$; Fig. 4C). However, muscle PDC activity was unchanged from control in simvastatin + DCA 30 mg and simvastatin + DCA 40 mg. There was a trend for muscle acetylcarnitine content to decrease in simvastatin compared with control ($P = 0.10$; Fig. 4D). Muscle acetylcarnitine content increased above control and simvastatin alone with simvastatin + DCA 30 mg ($P < 0.05$, $P < 0.05$, respectively) and simvastatin + DCA 40 mg ($P < 0.01$, $P < 0.01$, respectively; Fig. 4D), demonstrating PDC flux was increased.

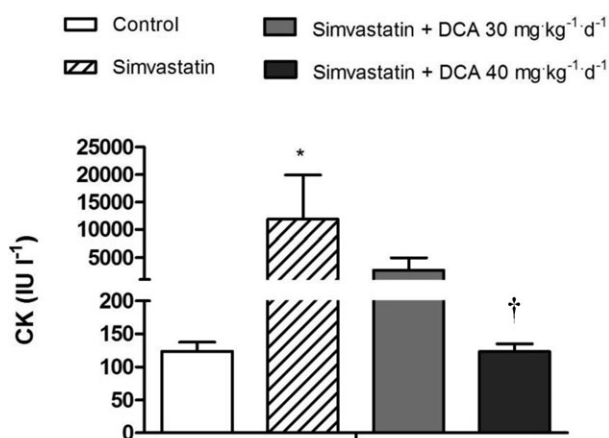


Figure 3. Plasma creatine kinase (CK) concentration in control, simvastatin- and simvastatin plus dichloroacetate (DCA)-treated animals

Values expressed as mean \pm SEM. * $P < 0.05$ when compared with control, and † $P < 0.05$ when compared with simvastatin.

Table 1. Incidence and severity of muscle fibre necrosis for control, simvastatin- and simvastatin plus DCA-treated animals

	Animal number	CT	Sol	BF	Gas
Group 1	1				
Vehicle (control)	2	*			
	3		*		
	4				
	5				
	6		*	*	
	7				
	8				
	9		*		
	10	***		**	***
Simvastatin, 88 mg kg ⁻¹ day ⁻¹	11	**		**	***
	12	**		**	**
	13	*		*	**
	14				
	15				*
	16				
	17				
	18				
Group 3 Simvastatin + sodium DCA, 88/30 mg kg ⁻¹ day ⁻¹	19	**	—	**	**
	20	*		**	**
	21	*		*	*
	22			**	
	23		*	*	
	24			*	
	25		*		
	26				
	27				
Group 4 Simvastatin + sodium DCA, 88/40 mg kg ⁻¹ day ⁻¹	28				
	29			*	
	30				
	31		*		
	32		*		
	33	*			
	34			*	
	35		*		
	36				*

BF, biceps femoris; CT, cranial tibial; DCA, dichloroacetate; Gas, gastrocnemius; Sol, soleus. —: muscle not sampled; no symbol: muscle necrosis not present. *Minimal necrosis: up to 10 muscle fibres affected in whole section; **mild necrosis: up to 20% of fibres in section affected; ***moderate necrosis: up to 50% of fibres in section affected.

The upregulation of transcriptional markers of ubiquitin proteasome, lysosomal and autophagy-mediated protein breakdown with simvastatin is attenuated with DCA administration

MAFbx mRNA expression was sixfold greater in simvastatin compared with control ($P < 0.001$; Fig. 5A).

Simvastatin + DCA 30 mg administration resulted in a fourfold decrease in MAFbx mRNA expression compared with simvastatin ($P < 0.05$), and simvastatin + DCA 40 mg returned MAFbx mRNA levels to that of the control ($P < 0.001$, compared with simvastatin). Muscle proteasome activity was increased above control with simvastatin administration ($P < 0.05$; Fig. 6A). There was a strong trend ($P = 0.08$) for simvastatin + DCA 40 mg to reduce proteasome activity levels to that of the control. Simvastatin alone increased cathepsin-L mRNA expression fivefold above control ($P < 0.001$; Fig. 5B), and simvastatin + DCA 40 mg reduced expression to that seen in control ($P < 0.001$). In keeping with this, cathepsin-L activity was increased in simvastatin ($P < 0.05$; Fig. 6B), and both simvastatin + 30 mg DCA and simvastatin + 40 mg DCA brought cathepsin-L activity levels to that of control. LC3 mRNA expression

was upregulated sixfold above control in simvastatin ($P < 0.001$; Fig. 5C), and both simvastatin + DCA 30 mg ($P < 0.001$) and simvastatin + DCA 40 mg ($P < 0.001$) significantly decreased LC3 mRNA expression when compared with simvastatin alone.

Discussion

The present study provides novel and impactful insight into the inhibition of statin-mediated myopathy by demonstrating that forcibly increasing muscle CHO oxidation with DCA can prevent the myopathic effects seen with simvastatin in an animal model of statin-induced myopathy. Specifically, DCA reversed the statin-mediated inhibition of muscle PDC activity and flux, abolished statin-mediated upregulation of FOXO target genes

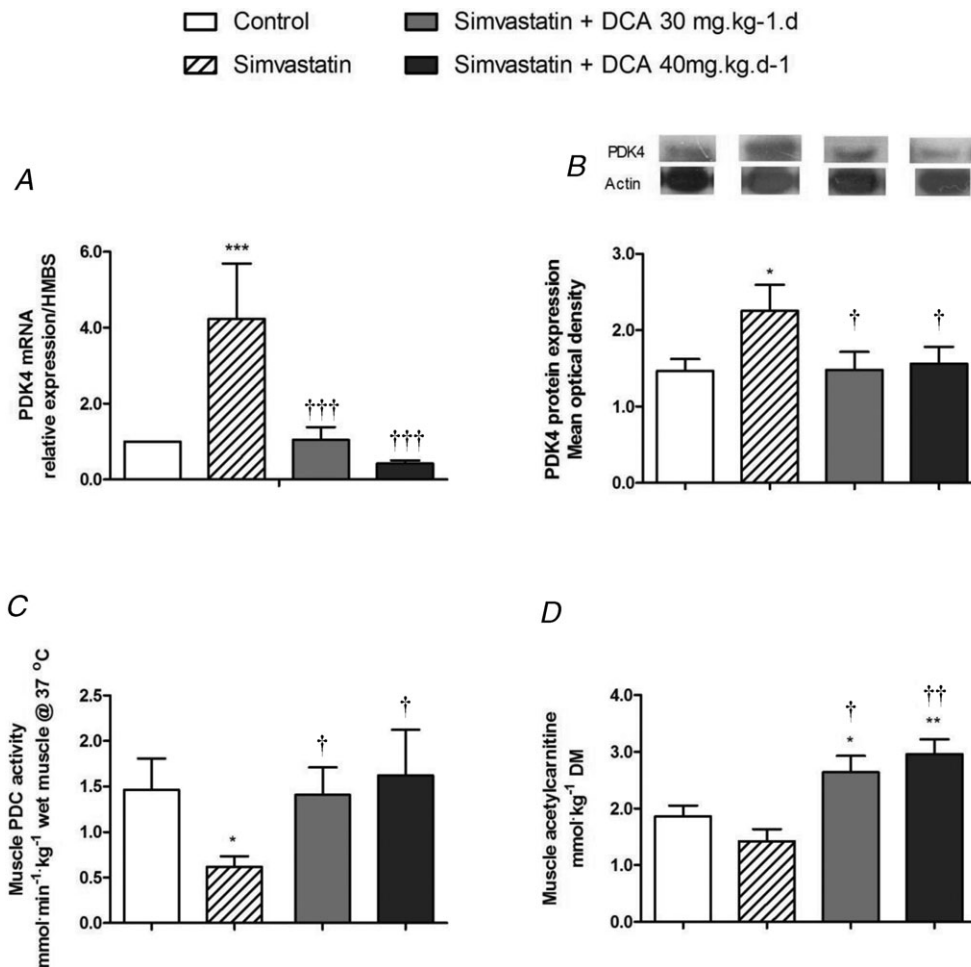


Figure 4. Muscle pyruvate dehydrogenase kinase-4 (PDK4) mRNA and protein expression, pyruvate dehydrogenase complex (PDC) activity and acetylcarnitine content after simvastatin and dichloroacetate (DCA) administration

A, PDK4 mRNA expression, values expressed as mean fold changes relative to control. B, PDK4 protein expression, values normalised to actin and expressed as mean \pm SEM. C, muscle PDC activity, values expressed as mean \pm SEM. D, muscle acetylcarnitine content, values expressed as mean \pm SEM. * $P < 0.05$ and ** $P < 0.01$ when compared with control, † $P < 0.05$ and †† $P < 0.01$ when compared with simvastatin.

MAFbx and cathepsin-L, decreased PDK4 mRNA and protein expression, and normalised statin-mediated increases in muscle cathepsin-L and proteasome activity, which collectively was associated with a blunting of muscle

necrosis and myopathy and the maintenance of body mass and food intake. To our knowledge, this is the first study to date to identify such a promising pharmacological approach in the prevention of statin-induced myopathy.

Our previous work (Mallinson *et al.* 2009) showed one of the main effects of simvastatin on muscle metabolism is the inhibition of CHO oxidation, specifically by upregulation of PDK4 mRNA, via blunted Akt signalling and activation of FOXO transcription factors. Indeed, a similar switch in muscle fuel metabolism is prominent in other animal models of muscle atrophy, namely sepsis (Crossland *et al.* 2008), peroxisome proliferator-activated receptor delta agonism (Constantin *et al.* 2007; Constantin-Teodosiu *et al.* 2009) and in critically ill patients (Constantin *et al.* 2011), where in all scenarios PDK4 mRNA and protein expression are upregulated and PDC activity decreased. In the present study, as hypothesized, simvastatin administration increased muscle PDK4 mRNA and protein expression (Fig. 4A and B), and inhibited muscle PDC activity (Fig. 4C) and flux (Fig. 4D). In keeping with this, work by Motojima and Seto showed increased PDK4 mRNA expression in muscle of statin-treated rodents (Motojima & Seto, 2003). However, the present study is the first to our knowledge to demonstrate that simvastatin-mediated upregulation of PDK4 protein inhibits PDC activity and flux, clearly identifying the metabolic impact of statin-mediated PDK4 upregulation. Importantly, we also showed simultaneous treatment with DCA and simvastatin completely reversed simvastatin-mediated upregulation of PDK4 mRNA and protein expression (Fig. 4A and B), and inhibition of muscle PDC activity (Fig. 4C) and flux (Fig. 4D) to that seen in control. We feel that the effect of DCA on PDK4 mRNA and protein expression was unlikely to be an immediate or direct effect, but rather occurred secondary to chronically increased muscle glucose uptake and CHO oxidation, Akt phosphorylation and downstream inhibition of FOXO transcription factors. Indeed, as outlined above, DCA is known to decrease blood glucose concentration (Stacpoole *et al.* 1997), probably by increasing muscle glucose uptake and CHO oxidation *in vivo*. Furthermore, PDK4 expression is known to be controlled through activation of FOXO transcription factors (Furuyama *et al.* 2003), which in turn are activated via decreased phosphorylation of Akt (Zhao *et al.* 2007). In keeping with the acute effect of DCA on PDC activation and flux (rather than PDK4 mRNA and protein expression levels), studies have also shown DCA can bind to and inhibit PDK2 and PDK4 activity *in vitro* (Li *et al.* 2009), and inhibit PDK activity in mitochondrial extracts of rat skeletal muscle (Vary & Hazen, 1999).

In addition to having myopathic effects, statins are known to cause muscle weakness and premature fatigue in patients (Phillips *et al.* 2002), and based on the findings of the present study, it is plausible that statin-mediated

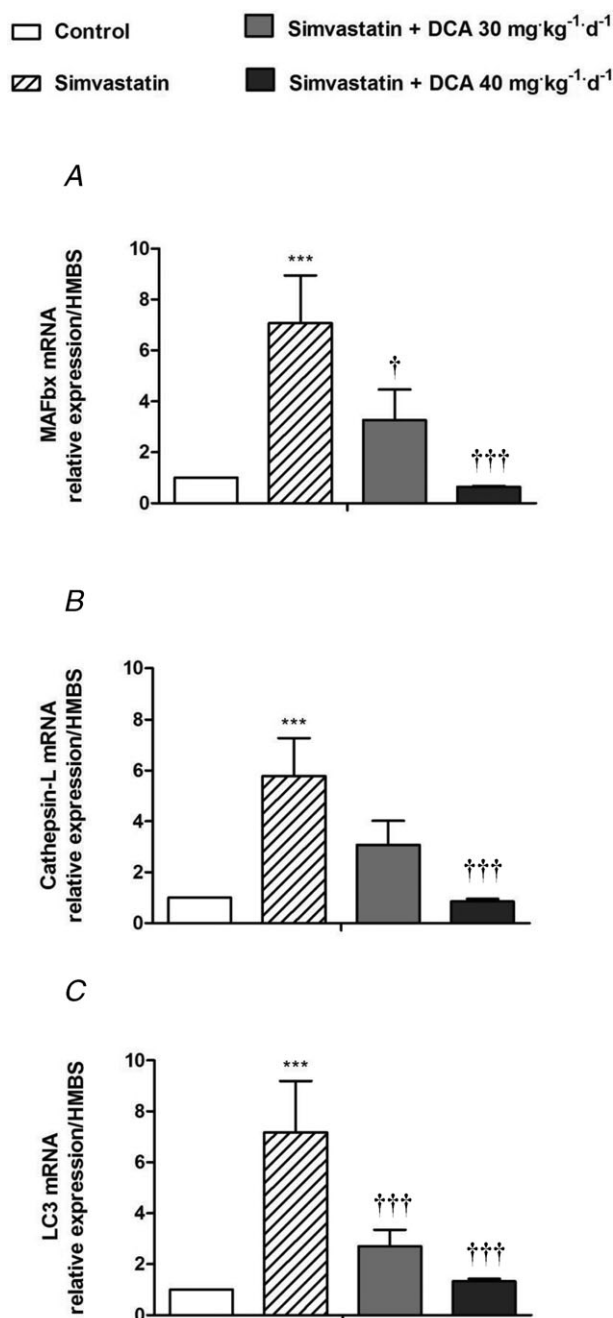


Figure 5. mRNA expression of markers of proteasome-, lysosomal- and autophagy-mediated proteolysis (fold changes relative to control) in biceps femoris muscle of animals treated with simvastatin or simvastatin plus dichloroacetate (DCA)

A, muscle atrophy F-box (MAFbx) expression. B, cathepsin-L expression. C, LC3 expression. Values expressed as mean fold changes relative to control \pm SEM. *** $P < 0.001$ when compared with control (which is set at 1), † $P < 0.05$ and ††† $P < 0.001$ when compared with simvastatin.

inhibition of PDC and CHO oxidation may contribute to these side-effects. Indeed, inhibition of muscle PDC has been shown to impair muscle CHO oxidation and contractile function (Constantin-Teodosiu *et al.* 2009). Furthermore, PDK inhibition in instances of statin-mediated muscle weakness and premature fatigue may prove to be a viable strategy in reducing these symptoms. It is also interesting to note there is increasing evidence of a positive association between high-dose statin administration and new-onset type II diabetes (Sattar *et al.* 2010; Preiss *et al.* 2011), which could be, at least partly, explained by statin-mediated upregulation of muscle PDK4 and PDC inhibition; both of which have been advocated as being causative factors in skeletal muscle insulin resistance (Wu *et al.* 1999). Pertinent to the latter contention, work from our laboratory has recently shown that pharmacological activation of PDC through DCA administration can abrogate the high-fat diet-mediated inhibition of CHO oxidation during exercise in humans (Constantin-Teodosiu *et al.* 2012).

Another key event possibly involved in the development of statin-induced myopathy is the upregulation of proteasomal and lysosomal proteolysis, specifically by decreased Akt/FOXO signaling, and increased transcription of MAFbx, MuRF-1 and cathepsin-L, which occurs before evidence of extensive muscle damage becomes apparent (Mallinson *et al.* 2009). We therefore aimed to determine whether forcibly increasing muscle CHO oxidation using DCA would suppress statin-mediated upregulation of FOXO downstream target genes, thereby blunting muscle proteolysis and simvastatin-induced muscle myopathy. In keeping

with this, the highest dose of DCA administered ($40 \text{ mg kg}^{-1} \text{ day}^{-1}$) blunted the development of muscle myopathy, and blocked the reduction in food intake and body mass loss observed with simvastatin (Table 1; Fig. 2). High-dose statin administration has been associated with body mass loss and decreased food intake in this model (Westwood *et al.* 2005). Furthermore, these events were paralleled by a marked 95-fold decline in plasma CK concentration, an accepted marker of muscle damage, such that levels returned to that seen in control (Fig. 3). This $40 \text{ mg kg}^{-1} \text{ day}^{-1}$ dose of DCA also abolished the simvastatin-induced increase in MAFbx (Fig. 5A) and cathepsin-L (Fig. 5B) mRNA, and attenuated proteasome and cathepsin-L activity (Fig. 6). Of note, the dampening of the mRNA expression levels with DCA occurred in a dose-dependent manner (Fig. 5). Collectively, these data provide convincing evidence that increasing CHO oxidation with DCA reduces simvastatin-induced necrosis by simultaneously inhibiting proteasomal- and lysosomal-mediated protein breakdown. *In vitro* studies have shown that upregulation of MAFbx and cathepsin-L mRNA expression appears to occur via Akt signaling-mediated changes in FOXO transcription factors (Sandri *et al.* 2004; Stitt *et al.* 2004). Indeed, we have previously shown that statin administration decreases muscle Akt phosphorylation and increases expression of FOXO target genes similar to the present *in vivo* model of statin myopathy (Mallinson *et al.* 2009). In line with the present reported effect of DCA on PDK4 mRNA and protein expression, it is suggested that a chronic DCA-mediated increase in CHO oxidation was associated with increased Akt phosphorylation, leading to

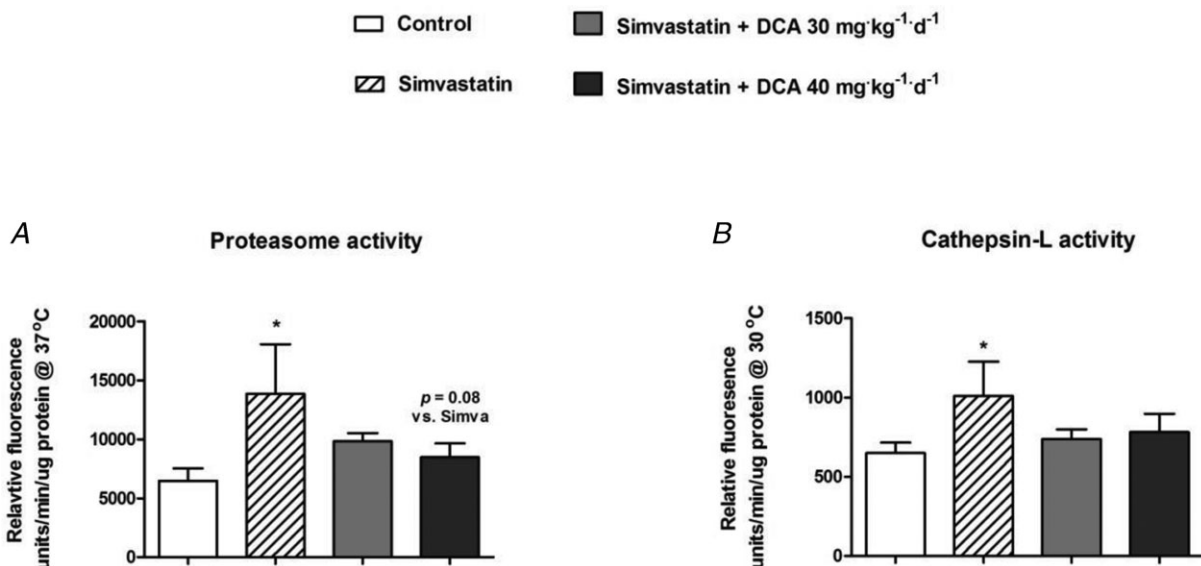


Figure 6. Proteasome (A) and cathepsin-L (B) activity in biceps femoris muscle of animals treated with simvastatin or simvastatin plus dichloroacetate (DCA)

Values expressed as mean \pm SEM. * $P < 0.05$ compared with control.

inactivation of FOXO transcription factors, and thereby the decreased expression of MAFbx and cathepsin-L mRNA and associated protein activity observed. However, more work is needed to clarify the exact mechanism by which forcibly increasing CHO oxidation with DCA prevents statin-induced upregulation of proteasomal and lysosomal protein breakdown. Previous studies have suggested that DCA treatment can blunt muscle catabolism, as evidenced by reduced levels of BCAA and phenylalanine in rodent muscle during sepsis (Vary *et al.* 1988), and reduced necrosis in rabbit tibialis muscle during acute limb ischaemia (Platz *et al.* 2007). Furthermore, Goodman *et al.* reported attenuated BCAA release from the perfused hind limb of fed healthy animals following DCA administration, suggesting DCA may either decrease protein breakdown and/or increase protein synthesis (Goodman *et al.* 1978). Because intramuscular glutamine concentrations have been correlated to muscle protein synthesis, a possible mechanism by which DCA exerts its effect on muscle protein metabolism is through its ability to increase skeletal muscle-free glutamine levels (Ferrando *et al.* 1998). Alternatively, increased glutamine availability has been shown to increase signalling through the PI3k/Akt pathway in HepG2 cells (van Meijl *et al.* 2010) and in muscle of rats with diet-induced obesity (Prada *et al.* 2007). Furthermore, new findings indicate that glutamine availability also relates strongly to the induction of apoptosis, working both as a nutrient and as a signalling molecule, acting directly or indirectly against the pathways leading to programmed cell death (Mates *et al.* 2006). However, it is not possible to distinguish from the present findings the exact mechanisms involved in the blunting of simvastatin-induced muscle myopathy by DCA.

Statins have also been shown in human rhabdomyosarcoma cells to mediate protein degradation by the induction of autophagy (Araki & Motojima, 2008). However, to date, no study has examined the role of autophagy in an *in vivo* model of statin-induced myopathy. Autophagy is a cellular process where intracellular components are degraded within lysosomes (Meijer & Codogno, 2004). Consistent with many *in vivo* models of muscle mass loss, including starvation (Lecker *et al.* 2004; Mammucari *et al.* 2007), denervation (Mammucari *et al.* 2007), cancer cachexia and diabetes (Lecker *et al.* 2004), is the upregulation of the gene LC3. Our data show, for the first time, that simvastatin administration increases the expression of LC3 mRNA in biceps femoris muscle, which suggests autophagy is involved in the muscle myopathy seen with statins. Furthermore, co-administration of simvastatin with DCA inhibited this increase in LC3 and returned mRNA expression levels to that seen in control, and in a dose-dependent manner (Fig. 5C). Furthermore, this was paralleled by decreased muscle necrosis (Table 1) and plasma CK (Fig. 3). Interestingly, autophagy has been

shown to be induced by activated FOXO3a (Mammucari *et al.* 2007), and it may therefore be suggested that DCA administration inhibited induction of autophagy in statin-induced myopathy in a similar manner to the FOXO1-mediated inhibition of MAFbx and cathepsin-L mRNA expression.

Given PDC is key to the regulation of mitochondrial CHO oxidation, it is interesting to note that in line with our observations reported here mitochondrial dysfunction is evident in models of statin myopathy (Bouitbir *et al.* 2011; Mullen *et al.* 2011; Galtier *et al.* 2012; Kwak *et al.* 2012). Specifically, skeletal muscle tissue from patients receiving statin therapy shows impaired mitochondrial respiration restricted to mainly complex I of the respiratory chain (Sirvent *et al.* 2012). Such mitochondrial dysfunction is proposed to increase mitochondrial NADH (Sharma *et al.* 2011), thereby increasing the mitochondrial NADH/NAD⁺ ratio, which in turn could activate PDK (Bowker-Kinley *et al.* 1998) or directly inhibit flux through the PDC reaction (Constantin-Teodosiu *et al.* 1993). Furthermore, statin-induced mitochondrial dysfunction could lead to increased oxidative stress (Bouitbir *et al.* 2011; Kwak *et al.* 2012) and apoptosis (Kwak *et al.* 2012), which could be at least partly responsible for the statin-related muscle necrosis seen in statin myopathy.

The results of this study suggest the use of DCA could offer potential to prevent statin myopathy in the clinic, particularly as the highest dose administered (40 mg kg⁻¹ day⁻¹) is less than that previously administered to humans. DCA has been used clinically for the treatment of several conditions, including lactic acidosis (100 mg kg⁻¹ day⁻¹; Stacpoole *et al.* 1988), diabetes mellitus (50 mg kg⁻¹ day⁻¹; Stacpoole *et al.* 1978), hyperlipoproteinaemia (50 mg kg⁻¹ day⁻¹; Stacpoole *et al.* 1978), mitochondrial encephalomyopathies (2.4 g day⁻¹; Curto *et al.* 2006) and to offset muscle fatigue associated with anaerobic ATP production during ischaemic muscle contraction (50 mg kg⁻¹; Timmons *et al.* 1996, 1998a, 1998b). Chronic use (months to a year) of DCA is, however, associated with adverse side-effects, which has detracted from its widespread use (Kurlemann *et al.* 1995; Stacpoole *et al.* 1998; Spruijt *et al.* 2001). Nonetheless, it has been administered to adults at a dose of 50 mg kg⁻¹ day⁻¹ for up to 5 years to treat mitochondrial encephalomyopathy, which resulted in only mild side-effects that did not warrant discontinuation (Mori *et al.* 2004). A lower dose of 25 mg kg⁻¹ day⁻¹ was administered to children for up to 6 months in a controlled trial for treatment of lactic acidosis with no signs of any side-effects (Stacpoole *et al.* 2006). Clearly dose-response studies would be needed to determine whether a tolerated dose of DCA (or any other potential PDK inhibitor) could prevent statin myopathy in humans, and any clinical evaluation

of DCA as a preventative treatment for statin myopathy would need careful consideration of its side-effect profile.

In summary, we have presented novel insight into the inhibition of statin myopathy by pharmacological means by demonstrating that increasing muscle CHO oxidation with DCA can reverse the myopathic effects seen with simvastatin in an animal model of statin-induced myopathy. Specifically, DCA increased muscle PDC activity and flux, abolished statin-mediated upregulation of FOXO target genes MAFbx and cathepsin-L, decreased PDK4 mRNA and protein expression, and decreased cathepsin-L and proteasome activity, which occurred concomitantly with the maintenance of food intake and body mass, and the blunting of muscle necrosis and myopathy. The finding that simvastatin activates the transcription and translation of PDK4 and thereby inhibits PDC-mediated CHO oxidation may account for the recent association between statin use and development of diabetes. Further studies are warranted to identify the precise mechanisms by which increased oxidative use of CHO prevents the myopathic effects of simvastatin administration.

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Author contributions

J.E.M.: data analysis and interpretations, drafting and revising the manuscript. D.C.-T.: conception and experimental design, data interpretation, and revising the manuscript. P.D.G.: experimental design and revising the manuscript. E.A.M.: performed *in vivo* experiments and revised the manuscript. W.J.D.: performed *in vivo* experiments and revising the manuscript. F.R.W.: performed histological analysis and revised the manuscript. J.E.S.: conception and experimental design, data interpretation, and revising the manuscript. P.L.G.: conception and experimental design, data interpretation and major revising. All authors approved the final version of the manuscript.

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Translational perspective

Statins are used widely to reduce hypercholesterolaemia in people with a heightened cardiovascular risk. Nevertheless, although rhabdomyolysis is rare, myositis and myalgia commonly accompany statin use, with a major symptom being muscle weakness and fatigue. Furthermore, increasing evidence suggests long-term, high-dose statin use is associated with an increased incidence of new-onset diabetes. The molecular mechanisms by which statins bring about these adverse effects are not fully resolved, but we have previously shown upregulation of downstream gene targets of the transcription factor FOXO, and evidence of increased proteasomal- and lysosomal-mediated muscle protein breakdown and inhibition of muscle CHO oxidation in an animal model of statin myopathy. The present study demonstrates that increasing CHO oxidation *in vivo*, using the PDK inhibitor, DCA, blunted activation of FOXO gene targets regulating muscle CHO utilisation and protein breakdown, which inhibited statin myopathy. This is the first demonstration that pharmacological intervention can prevent statin myopathy. The impairment of muscle CHO oxidation is a feature of many forms of muscle atrophy, and therefore the use of a pharmacological intervention to forcibly drive muscle CHO use could be of therapeutic benefit. However, further studies are warranted to assess the impact of long-term DCA use, or any other PDK inhibitor, and to determine the efficacy and minimal dose required to elicit similar responses in patients experiencing statin myopathy symptoms.