

Insulin-stimulated plasma membrane association and activation of Akt2, aPKC ζ and aPKC λ in high fat fed rodent skeletal muscle

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Several recent reports using cell lines have suggested that both Akt and atypical protein kinase C (aPKC) ζ/λ are translocated to the plasma membrane (PM) in response to insulin. However, it has yet to be determined in skeletal muscle whether: (1) insulin increases PM-associated Akt2, aPKC ζ and/or λ protein concentration, (2) the activity of these kinases is altered by insulin at the PM, and (3) high fat feeding alters the insulin-stimulated PM concentration and/or activity of Akt2 and aPKC ζ/λ . Sprague-Dawley rats were randomly assigned to either normal ($n = 16$) or high fat ($n = 16$) dietary groups. Following a 12 week dietary period, animals were subjected to hind limb perfusions in the presence ($n = 8$ per group) or absence ($n = 8$ per group) of insulin. In normal skeletal muscle, total PI3-kinase, Akt2 and aPKC ζ/λ activities were increased by insulin. PM-associated aPKC ζ and λ , and aPKC ζ/λ activity, but not Akt2 or Akt2 activity, were increased by insulin in normal muscle. High fat feeding did not alter total skeletal muscle Akt2, aPKC ζ or aPKC λ protein concentration. Insulin-stimulated total PI3-kinase, Akt2 and aPKC ζ/λ activities were reduced in the high fat fed animals. Insulin-stimulated PM aPKC ζ , aPKC λ , aPKC ζ/λ activity and GLUT4 protein concentration were also reduced in high fat fed animals. These findings suggest that in skeletal muscle, insulin stimulates translocation of aPKC ζ and λ , but not Akt2, to the PM. In addition, high fat feeding impairs insulin-stimulated activation of total aPKC ζ/λ and Akt2, as well as PM association and activation of aPKC ζ and λ .

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Insulin stimulation increases phosphoinositide 3-kinase (PI3-kinase) activity, resulting in phosphorylation and activation of Akt and atypical protein kinase C (aPKC) ζ/λ (Ducluzeau *et al.* 2002; Chen *et al.* 2003; Katome *et al.* 2003). Akt2 appears to have a primary role in the regulation of insulin-stimulated glucose transport (Cho *et al.* 2001a; Jiang *et al.* 2003) while Akt1 and Akt3 may contribute to the regulation of glucose transport in a secondary role (Walker *et al.* 1998; Cho *et al.* 2001b; Brozinick *et al.* 2003). However, insulin-stimulated activation of Akt2 does not appear to fully account for GLUT4 translocation (Cong *et al.* 1997; Kotani *et al.* 1998) as it has been observed that dominant-negative Akt expression reduces, but does not prevent, GLUT4 translocation (Cong *et al.* 1997; Bandyopadhyay *et al.* 1999b; Ducluzeau *et al.* 2002). Rather, it appears that aPKC ζ/λ functions in parallel with Akt to facilitate insulin-stimulated GLUT4 translocation (Bandyopadhyay *et al.* 1999a; Standaert *et al.* 1999).

In response to insulin stimulation, both Akt2 (Andjelkovic *et al.* 1997; Goransson *et al.* 1998; Chen

et al. 2003; Sasaoka *et al.* 2004) and aPKC ζ/λ (Standaert *et al.* 1999; Braiman *et al.* 2001) are translocated to the plasma membrane, where it is believed that they are activated (Galetic *et al.* 1999; Kanzaki *et al.* 2004). Insulin stimulation not only results in the translocation of aPKC ζ/λ to the plasma membrane in muscle cell cultures, but it is associated with GLUT4 vesicles (Braiman *et al.* 2001). It should be noted though that insulin-stimulated Akt and aPKC ζ/λ translocation to the plasma membrane has only been assessed in cell lines and adipocytes. However, it has been reported that exercise increases the activation and plasma membrane association of aPKC ζ/λ in human skeletal muscle (Nielsen *et al.* 2003; Perrini *et al.* 2004; Rose *et al.* 2004). Therefore, the initial aim of this investigation was to assess whether insulin increases the plasma membrane concentration and activation of Akt and/or aPKC ζ/λ in skeletal muscle.

It has been reported that a high-fat diet impairs carbohydrate metabolism in rodent skeletal muscle by reducing insulin-stimulated IRS-1-associated PI3-kinase,

Akt2, and aPKC ζ/λ activities (Tremblay *et al.* 2001; Singh *et al.* 2003; Krisan *et al.* 2004; Yaspelkis *et al.* 2004) and may account for a reduced plasma membrane GLUT4 protein concentration (Yaspelkis *et al.* 2001; Singh *et al.* 2003). Thus, the second aim of this investigation was to assess whether insulin-stimulated compartmentalization and/or activation of aPKC ζ/λ and Akt2 is altered by chronic high fat feeding and related to reduced GLUT4 translocation in skeletal muscle.

Methods

Experimental design

Thirty-two male Sprague-Dawley rats approximately 6 weeks old ranging in weight from 185 to 220 g (Harlan, San Diego, CA, USA) were placed randomly into one of two groups: normal diet ($n = 16$) or high-fat diet ($n = 16$). The normal diet (no. 112386, Dyets Inc., Bethlehem, PA, USA) consisted of 63% carbohydrates, 17% fat, and 15% protein. The high fat diet (no. 112387, Dyets Inc.) contained 26% carbohydrates, 59% fat, and 15% protein. The vitamin, mineral, and fibre content were the same for both groups. The animals were on their respective diets for 12 weeks and allowed to feed *ad libitum*. This high fat diet has previously been shown to induce skeletal muscle insulin resistance in male Sprague-Dawley rats (Singh *et al.* 2003; Krisan *et al.* 2004; Yaspelkis *et al.* 2004). Rats were housed two per cage in a temperature controlled room (21°C) with an artificial light cycle 12 : 12 light–dark cycle. After 12 weeks of feeding, animals were subdivided into four groups: normal diet insulin ($n = 8$), normal diet basal ($n = 8$), high fat diet insulin ($n = 8$), high fat diet basal ($n = 8$), and subjected to hind limb perfusion.

All experimental procedures were approved by the Institutional Animal Care and Use Committee at California State University, Northridge and conformed to the guidelines for the use of laboratory animals published by the United States Department of Health and Human Resources.

Hind limb perfusions

Animals were anaesthetized with an intraperitoneal injection of sodium pentobarbital (6.5 mg (100 g body wt)⁻¹) and surgically prepared for hind limb perfusion as previously described by Ruderman *et al.* (1971) and modified by Ivy *et al.* (1989). Following surgical preparation, cannulas were inserted into the abdominal aorta and vena cava, and the animals were killed via an intracardiac injection of pentobarbital as the hind limbs were washed out with 30 ml of Krebs–Henseleit buffer (KHB) (pH 7.55). Immediately, the cannulas were placed in line with a non-recirculating perfusion system and the hind limbs were allowed to stabilize during a 5 min washout period. The perfusate was continuously gassed

with a mixture of 95% O₂–5% CO₂ and warmed to 37°C. Perfusate flow rate was set at 7.5 ml min⁻¹ during the stabilization and subsequent perfusion during which rates of glucose transport were determined.

Perfusions were performed in the presence or absence of 500 μ U ml⁻¹ insulin. The basic perfusate medium consisted of 30% washed time-expired human erythrocytes (Ogden Medical Center, Ogden, UT, USA), KHB, 4% dialysed bovine serum albumin (Fisher Scientific, Fairlawn, NJ, USA) and 0.2 mM pyruvate. The hind limbs were washed out with perfusate containing 1 mM glucose for 5 min in preparation for the measurement of glucose transport. Glucose transport was measured over an 8-min period using an 8 mM concentration of non-metabolized glucose analogue 3-O-methylglucose (3-MG) (32 μ Ci 3-[³H]MG mm⁻¹, PerkinElmer Life Sciences, Boston, MA, USA) and 2 mM mannitol (60 μ Ci-[1-¹⁴C]mannitol mm⁻¹, PerkinElmer Life Sciences). Immediately after the transport period, portions of the red quadriceps (RQ) were excised from both hind limbs, blotted on gauze dampened with cold KHB, freeze clamped in liquid N₂ and stored at -80°C for later analysis.

3 MG transport

Rates of insulin-stimulated skeletal muscle 3-MG transport were calculated as previously described (Yaspelkis *et al.* 2001, 2004; Krisan *et al.* 2004).

Muscle homogenization for Western blotting

Portions were cut from the RQ, weighed frozen and homogenized in an ice-cold homogenization buffer (HB) 1 : 10 wt/vol containing 50.0 mM Hepes (pH 7.6), 150 mM NaCl, 200 mM sodium pyrophosphate, 20 mM β -glycerophosphate, 20 mM NaF, 2 mM orthovanadate, 20 mM EDTA, 10% IGEPAL, 10% glycerol, 20 mM phenylmethylsulfonylfluoride, 1 mM MgCl₂, 1 mM CaCl₂, 10 μ g ml⁻¹ leupeptin and 10 μ g ml⁻¹ aprotinin. The homogenate was then transferred to a microcentrifuge tube and centrifuged (19 600 g, 4°C) in a refrigerated microcentrifuge (Micromax RF, International Equipment Co., Needham Heights, MA, USA) for 15 min. The supernatant was collected, labelled as lysate and assayed for protein concentration using the Bradford method (Bradford, 1976) adapted for use with a Benchmark microplate reader (Bio-Rad, Richmond, CA, USA).

Plasma membrane fractionation

Plasma membrane fractions were prepared as we (Yaspelkis *et al.* 2001, 2002; Singh *et al.* 2003) and others (Ahmed *et al.* 1990; Turcotte *et al.* 1997, 1999) have previously described. Briefly, insulin-stimulated RQ samples were

Table 1. Membrane fractionation characterization

	Basal		Insulin	
	Normal Diet	High fat diet	Normal diet	High fat diet
5'-Nucleotidase activity (nmol min ⁻¹ mg protein ⁻¹)				
Plasma membrane	330.22 ± 57.3	405.72 ± 72.2	314.35 ± 66.7	354.22 ± 68.8
Crude homogenate	57.72 ± 11.0	50.33 ± 9.9	57.00 ± 11.4	48.36 ± 14.0
Fold Increase	7.2 ± 2.1	8.2 ± 2.8	8.9 ± 2.6	10.8 ± 2.4
Percentage yield	13.5 ± 5.2	16.6 ± 7.2	16.2 ± 7.4	22.4 ± 4.8

Values are means ± S.E.M.

minced, diluted 1 : 7 in a 10 mM Tris–15% sucrose solution (pH 7.5) with 0.1 mM phenylmethylsulfonyl fluoride, 10 mM EGTA, 10 mg ml⁻¹ trypsin inhibitor and homogenized with a PT 2100 Polytron homogenizer (Kinemaica, Littau/Luzern, Switzerland). The homogenate was filtered and centrifuged at 100 000 g for 1 h at 4°C in a Sorvall TFT 50.38 fixed angle rotor (Kendro Laboratory Products, Newton, CT, USA). The pellet was resuspended in 6 ml of 1 mM Tris–15% sucrose buffer, and 50 µl of crude homogenate was collected and retained for later analysis. The remaining homogenate suspension was layered on continuous sucrose gradients (35% to 70%) and centrifuged at 120 000 g for 2 h at 4°C using a Sorvall Sure Spin 630 swinging bucket rotor (Kendro Laboratory Products). The plasma membrane layer was collected, washed in 10 mM Tris buffer, and centrifuged at 100 000 g for 1 h at 4°C using a Sorvall TFT 50.38 fixed angle rotor. The final plasma membrane pellet was resuspended in 200 µl of HB per gram of original tissue weight, frozen in liquid nitrogen and stored at –80°C for later analysis. The activity of the plasma membrane marker enzyme 5'-nucleotidase was assessed as previously described (Singh *et al.* 2003) in plasma membrane fractions and compared with the activity in crude homogenate fractions to insure that purified plasma membrane fractions were being used for analysis. Characterization of the membrane fractions obtained is presented in Table 1.

Western blotting

Lysate samples (140 µg protein for aPKC ζ and λ, 35 µg protein for Akt2, 70 µg of protein for GLUT4) and plasma membrane samples from the RQ (75 µg of protein for aPKC ζ and λ, 35 µg of protein for Akt2, 50 µg of protein for GLUT4) were added to Laemmli buffer (Laemmli, 1970). Sample proteins were subjected to SDS-PAGE run under reducing conditions on 7.5% (GLUT4) or 10% (Akt2, aPKC ζ and λ) resolving gels in a MiniProtean 3 dual-slab cell (Bio-Rad). Resolved proteins were transferred to polyvinylidene difluoride (PVDF) membranes using a semidry transfer unit (10 V for 30 min). Membranes were then blocked in 5% non-fat dry milk/Tris–Tween-buffered saline and incubated in affinity-purified rabbit polyclonal

anti-Akt2 (cat no. 07-372, Upstate Biotechnology (UBT) Charlottesville, VA, USA), anti-aPKC ζ/λ (sc-216, Santa Cruz Biotechnology (SCBT) Santa Cruz, CA, USA) or GLUT4 (donated by Dr Samuel W. Cushman, National Institute of Diabetes and Digestive and Kidney Disease, Bethesda, MD, USA) followed by incubation with goat anti-rabbit IgG conjugated horseradish peroxidase (sc-2004, SCBT). Antibody binding was visualized by enhanced chemiluminescence (ECL) in accordance with the manufacturer's instructions (West Femto, Pierce Chemical Company, Rockford, IL, USA). Images were captured using a ChemiDoc system (Bio-Rad) equipped with a CCD camera and saved to a Macintosh G4 computer. Protein bands were quantified as a percentage of a muscle sample standard run on each gel using Quantity One analysis software (Bio-Rad).

IRS-1-associated PI3-kinase activity

IRS-1-associated PI3-kinase activity was determined as described elsewhere (Singh *et al.* 2003). Approximately 150 µg of RQ protein lysate was immunoprecipitated with 4 µg of anti-IRS-1 antibody (cat. no. 06-248, UBT), and HB for 2 h at 4°C. Following 1 h of separation the TLC plate was dried, exposed to a storage phosphor screen (Eastman Kodak Company, Rochester, NY, USA) and scanned with a phosphor imager (Personal Molecular Imager FX System, Bio-Rad). The image was imported into a Macintosh G4 computer and quantified using Quantity One analysis software (Bio-Rad). Kinase activity was calculated as a percentage of an insulin-stimulated muscle standard run on each TLC plate.

Akt2 kinase activity

Two hundred and fifty micrograms of either RQ lysate or plasma membrane protein were combined with 4 µg of anti-Akt2 (UBT) and incubated overnight at 4°C. One hundred microlitres of a slurry containing Pro-A beads was added to each immunoprecipitate and incubated with rotation at 4°C for 1.5 h. After incubation, the samples were centrifuged (18 300 g, 4°C) for 10 min, and the immunocomplex was washed with the same protocol as described for PI3-kinase activity (Singh *et al.* 2003).

After the wash protocol, the samples were centrifuged (18 300 g, 4°C) for 10 min and the supernatant was removed. Ten microlitres of assay dilution buffer (cat. no. 20-108, UBT) was added to the immunocomplex in addition to PKA inhibitor peptide (cat. no. 12-151, UBT) and 10 μCi [γ - ^{32}P]ATP (PerkinElmer Life Sciences). Kinase reactions were initiated by addition of the Crosstide substrate oligopeptide (cat. no. 12-331, UBT), and warmed to 37°C with constant mixing for 10 min. Reactions were halted by addition of Laemmli buffer (1:1). For analysis, 15 μl of the sample/Laemmli buffer were loaded onto a 20% Tris–tricine polyacrylamide gel in duplicate and electrophoresed at 100 V for 130 min using a MiniProtean 3 electrophoresis system (Bio-Rad). After electrophoresis, gels were wrapped in plastic wrap and exposed to a phosphor screen for 8 h. Images were captured and quantified as described above.

aPKC ζ/λ activity

Five hundred micrograms of either RQ lysate or plasma membrane proteins were added to 4 μg of anti-aPKC ζ/λ (sc-216, SCBT) and incubated overnight at 4°C. One hundred microlitres of a slurry containing Protein A–Sepharose (Pro-A) beads were combined with each of the samples and incubated with rotation at 4°C for 1.5 h. After incubation, the samples were centrifuged (18 300 g, 4°C) for 10 min and the immunocomplex was washed by using the same protocol as described for PI3-kinase activity (Singh *et al.* 2003). After the wash protocol, samples were centrifuged (18 300 g, 4°C) for 10 min, and the supernatant was discarded. The Pro-A beads were then resuspended in 75 μl of kinase buffer. The kinase reaction was started by the combining 5 μCi [γ - ^{32}P]ATP (PerkinElmer Life Sciences), 40 μM ATP and 5 μl myelin basic protein (M8-184, Sigma-Aldrich, St Louis, MO, USA). The samples were vortexed periodically while incubating for 12 min at 37°C. Reactions were stopped by addition of Laemmli buffer (1:1). For analysis 15 μl of the sample/Laemmli buffer and subjected to SDS-PAGE run under reducing conditions on a 20% Tris–tricine resolving gel. After electrophoresis the gels were wrapped in plastic wrap and exposed to a phosphor screen for 8 h. Images were captured and quantified as described above.

Statistical analysis

A one-way analysis of variance (ANOVA) was used on all variables to determine whether significant differences exist between groups. When a significant *F*-ratio was obtained, a Tukey HSD *post hoc* test was used to identify statistically significant differences ($P < 0.05$) among the means. Statistical analyses were performed using JMP software (SAS Institute Inc., Cary, NC, USA) and all values were expressed as means \pm s.e.m.

Results

Body and epididymal fat pad mass

At the onset of the dietary period, the body mass of normal diet basal (202.1 ± 6.1 g), normal diet insulin (198.1 ± 3.0 g), high fat diet basal (204.2 ± 3.3 g), and high fat diet insulin (203.0 ± 3.5 g) animals were similar. Although the body mass of all animals increased throughout the study, no significant differences in body mass were observed among normal diet basal (472.8 ± 10.3 g), normal diet insulin (477.3 ± 12.3 g), high fat diet basal (488.1 ± 11.3 g), and high fat diet insulin (491.7 ± 11.9 g) animals at the end of the 12 week dietary period. At the end of the dietary period, the epididymal fat pad mass of the high fat diet basal (11.6 ± 1.0 g), and high fat diet insulin (12.3 ± 1.1 g) animals exhibited a trend ($P = 0.09$) to be heavier than the normal diet basal (10.0 ± 1.2 g) and normal diet insulin (9.0 ± 0.5 g) animals.

Glucose transport

Rates of 3-MG transport were similar between the normal diet (3.92 ± 0.17 $\mu\text{mol h}^{-1} \text{g}^{-1}$) and high fat fed (3.40 ± 0.24 $\mu\text{mol h}^{-1} \text{g}^{-1}$) animals in the absence of insulin. In the presence of insulin, rates of 3-MG transport were increased above basal levels. However, rates of insulin-stimulated 3-MG transport in the high fat fed animals (5.07 ± 0.28 $\mu\text{mol h}^{-1} \text{g}^{-1}$) were significantly less than ($P < 0.05$) that of the normal diet animals (7.04 ± 0.40 $\mu\text{mol h}^{-1} \text{g}^{-1}$).

IRS-1-associated PI3-kinase activity

In the absence of insulin, IRS-1-associated PI3-kinase activity was similar between the normal diet ($20.87 \pm 3.9\%$ standard) and high fat fed ($16.88 \pm 2.7\%$ standard) animals. While insulin increased PI3-kinase activity above basal levels in both the normal diet and high fat diet groups, insulin-stimulated PI3-kinase activity was greater in the normal diet animals ($70.85 \pm 4.3\%$ standard) compared to high fat diet animals ($39.72 \pm 2.1\%$ standard).

Akt2 protein concentration

Total Akt2 protein concentration was not different in the absence (141.3 ± 19.1 versus $154.45 \pm 12.1\%$ standard) or presence (147.30 ± 4.8 versus $151.82 \pm 10.3\%$ standard) of insulin between the normal diet and high fat diet groups, respectively. In addition, plasma membrane Akt2 protein concentration (Fig. 1A) was not different among the normal diet and high fat diet groups either in the absence or presence of insulin.

Akt2 activity

In the absence of insulin, total Akt2 activity was not different between the normal diet and high fat diet groups (Fig. 2A). In the presence of insulin, total Akt2 activity was increased above basal levels for both the normal diet and high fat diet animals. However, total insulin-stimulated Akt2 activity of the high fat diet animals was less than that of the normal diet animals. Although the high fat diet and insulin affected total Akt2 activity, plasma membrane Akt2

activity (Fig. 2B) was not different among the experimental groups.

aPKC ζ and λ protein concentration

Total aPKC ζ protein concentration was similar in the normal diet and high fat diet groups in the absence (50.36 ± 9.5 versus $55.33 \pm 12.1\%$ standard) or presence of insulin (56.35 ± 12.2 versus $43.76 \pm 6.1\%$ standard),

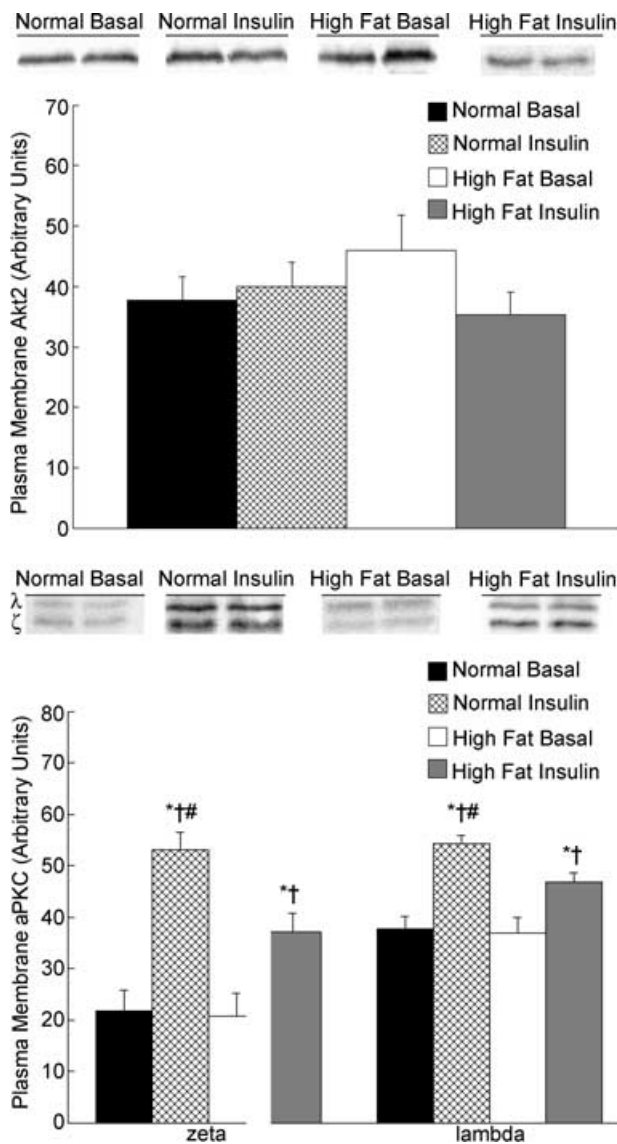


Figure 1. Red quadriceps total plasma membrane Akt2 (A) and plasma membrane aPKC ζ and λ (B) protein concentration from normal diet basal, normal diet insulin stimulated, high fat diet basal and high fat diet insulin stimulated animals

*Significantly different from normal diet basal ($P < 0.05$).
 †Significantly different from high fat diet basal ($P < 0.05$).
 #Significantly different from high fat diet insulin stimulated ($P < 0.05$).
 Values are means \pm s.e.m.

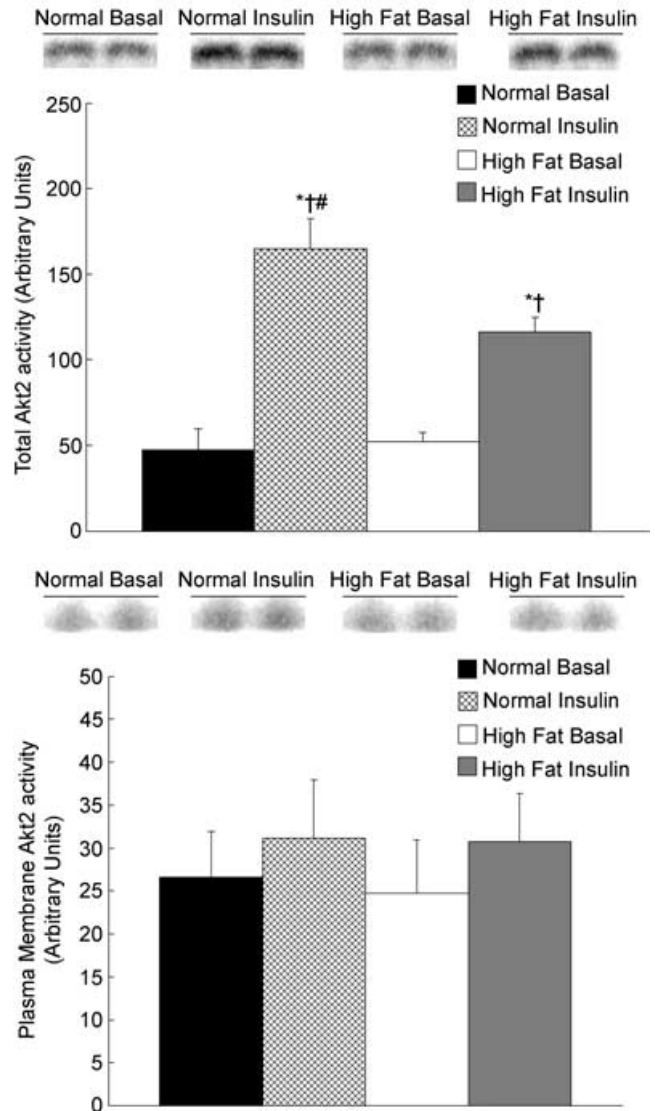


Figure 2. Red quadriceps total Akt2 protein activity (A) and plasma membrane Akt2 activity (B) from normal diet basal, normal diet insulin stimulated, high fat diet basal and high fat diet insulin stimulated animals

*Significantly different from normal diet basal ($P < 0.05$).
 †Significantly different from high fat diet basal ($P < 0.05$).
 #Significantly different from high fat diet insulin stimulated ($P < 0.05$).
 Values are means \pm s.e.m.

respectively. Similarly, total PKC λ protein concentration was not different in the normal diet and high fat diet groups in the absence (84.44 ± 15.0 versus $74.66 \pm 13.3\%$ standard) or presence (89.67 ± 21.3 versus $86.78 \pm 17.3\%$ standard) of insulin, respectively. In the absence of insulin, plasma membrane-associated aPKC ζ and λ (Fig. 1B) protein concentration were similar in the normal diet and high fat diet animals. In the presence of insulin, plasma membrane-associated aPKC ζ and λ were increased above basal levels in both dietary groups. However, insulin-stimulated plasma

membrane-associated aPKC ζ and λ were reduced in the high fat diet animals compared to the normal diet animals.

aPKC ζ/λ activity

Total aPKC ζ/λ activity under basal conditions was similar between the normal diet and high fat diet animals. Insulin stimulation increased total aPKC ζ/λ kinase activity above basal levels in both dietary groups (Fig. 3A). However, insulin-stimulated total aPKC ζ/λ kinase activity from the high fat diet animals was significantly lower when compared with the normal diet animals. Similar observations were made for aPKC ζ/λ activity in the plasma membrane (Fig. 3B). Specifically, in the absence of insulin no differences existed between normal diet and high fat diet animals; insulin increased plasma membrane aPKC ζ/λ activity above basal levels in both dietary groups; and insulin-stimulated aPKC ζ/λ activity was lower in the high fat diet animals when compared to normal diet animals (Fig. 3B).

GLUT4 protein concentration

Total GLUT4 protein concentration was similar in the normal diet basal and normal diet insulin-stimulated animals, and was significantly greater than that of both the high fat diet basal and high fat diet insulin-stimulated animals (Fig. 4A). In the absence of insulin, plasma membrane GLUT4 protein concentration was similar in the normal diet and high fat diet animals (Fig. 4B). Insulin increased the plasma membrane GLUT4 protein concentration above basal levels in both dietary groups. However, insulin-stimulated plasma membrane GLUT4 protein concentration was significantly lower in the high fat diet animals compared to the normal diet animals.

Discussion

In normal rodent skeletal muscle we observed that insulin stimulation increased rates of skeletal muscle glucose transport above basal levels and appeared to be a result of the plasma membrane GLUT4 protein concentration being increased, which is consistent with previous investigations that report a relationship existing between GLUT4 and rates of 3-MG transport (Friedman *et al.* 1990; Banks *et al.* 1992). We also observed that insulin increased the activities of PI3-kinase, Akt2 and aPKC ζ/λ above basal levels in the skeletal muscle lysate from the normal diet animals, which is also consistent with a number of previous investigations (Krisan *et al.* 2004; Yaspelkis *et al.* 2004). Having established that insulin activated components of the insulin signalling cascade, we turned our focus towards identifying whether insulin

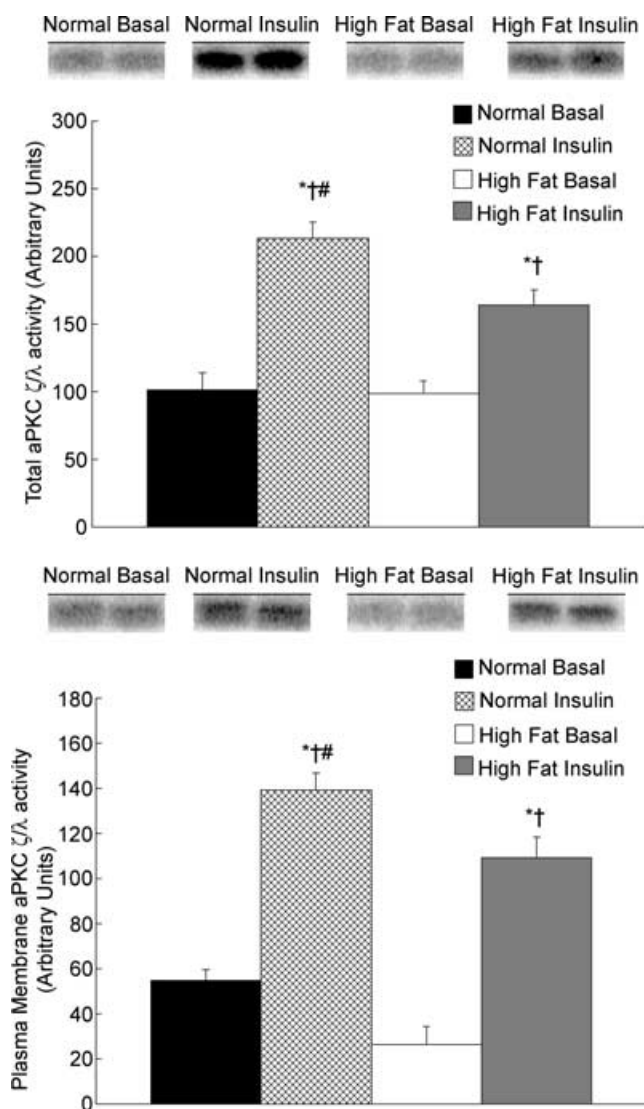


Figure 3. Red quadriceps total aPKC ζ/λ protein activity (A) and plasma membrane total aPKC ζ/λ activity (B) from normal diet basal, normal diet insulin stimulated, high fat diet basal and high fat diet insulin stimulated animals

*Significantly different from normal diet basal ($P < 0.05$).

†Significantly different from high fat diet basal ($P < 0.05$).

#Significantly different from high fat diet insulin stimulated ($P < 0.05$).

Values are means \pm S.E.M.

altered the plasma membrane content and/or activities of Akt2 and aPKC ζ/λ in the skeletal muscle of normal rodents.

Three isoforms of Akt have been identified in rodent skeletal muscle and it has been shown that the activation of Akt2 (Cho *et al.* 2001a), but not Akt1 (Cho *et al.* 2001b), is involved in insulin-stimulated glucose transport. It has also been reported in 3T3-L1 adipocytes that if Akt2 is deleted then GLUT4 translocation is reduced (Bae *et al.* 2003), and in response to insulin stimulation Akt2 becomes associated with the plasma membrane (Calera *et al.* 1998; Mitsuuchi *et al.* 1998; Bae *et al.* 2003). Thus, we chose to narrow our focus to Akt2 for this investigation. In response to insulin we did not observe plasma membrane Akt2 protein concentration or plasma membrane Akt2 activity to be increased above basal levels in normal rodent skeletal muscle. We are unaware of any other investigations that have evaluated the effects of insulin on plasma membrane-associated Akt2 protein concentration or activity in skeletal muscle. In contrast to our findings it has been observed that insulin increases plasma membrane-associated Akt in human ovarian carcinoma cells (Mitsuuchi *et al.* 1998), human embryonic kidney cells (Andjelkovic *et al.* 1997), rat adipocytes (Goransson *et al.* 1998; Bae *et al.* 2003; Chen *et al.* 2003) and 3T3-L1 adipocytes (Hill *et al.* 1999; Ducluzeau *et al.* 2002; Sasaoka *et al.* 2004). The differences existing among tissues are not readily reconcilable, but a plausible explanation may be that skeletal muscle has the capacity for glycogen synthesis whereas the previously investigated cell lines do not store glycogen. Specifically, insulin-stimulated activation of Akt causes serine phosphorylation of glycogen synthase kinase 3β (GSK3 β), which causes deactivation of GSK3 and in turn allows for the activation of glycogen synthase and the subsequent storage of glycogen (Frame & Cohen, 2001). As glycogen storage occurs primarily in the sarcoplasm of skeletal muscle it may be of greater metabolic relevance that Akt does not increase its concentration and activity in the plasma membrane of skeletal muscle, unlike cell lines or other tissues. However, this does not minimize the fact that activation of Akt2 is clearly necessary in order for glucose transport to occur in skeletal muscle (Cho *et al.* 2001a).

In contrast to Akt2, we observed that insulin increased aPKC ζ protein concentration, aPKC λ protein concentration and aPKC ζ/λ activity in plasma membrane prepared from normal rodent skeletal muscle. This finding is consistent with previous reports that have shown insulin increases plasma membrane associated aPKC ζ/λ in myotubes (Braiman *et al.* 2001), rat adipocytes (Standaert *et al.* 1999), and 3T3-L1 adipocytes (Bandyopadhyay *et al.* 1997; Standaert *et al.* 1999; Kanzaki *et al.* 2004). The physiological significance of plasma membrane aPKC ζ and λ protein concentration, and

aPKC ζ/λ activity being increased by insulin has not been fully elucidated, but it may contribute to facilitating GLUT4 translocation as it has been reported that aPKC ζ/λ directly interacts with GLUT4 containing vesicles (Standaert *et al.* 1999; Braiman *et al.* 2001; Kanzaki *et al.* 2004).

Our next line of inquiry was to evaluate high fat feeding effects on skeletal muscle Akt2 and aPKC ζ/λ protein concentration, plasma membrane association and activation. Confirming that the high fat diet impaired insulin-stimulated skeletal muscle carbohydrate

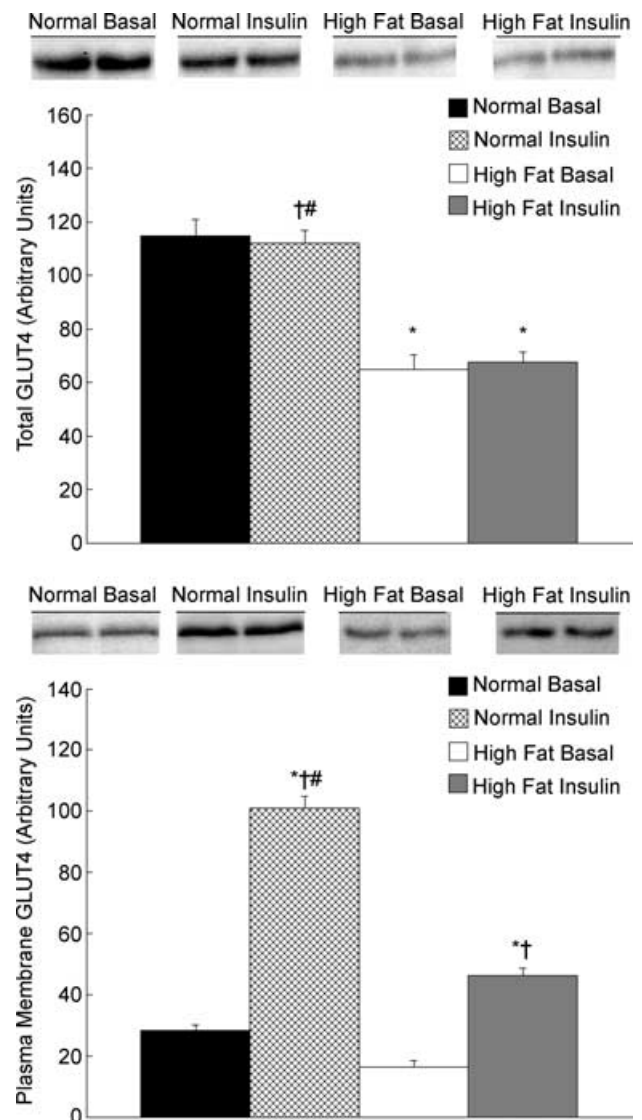


Figure 4. Red quadriceps total GLUT4 protein concentration (A) and plasma membrane GLUT4 protein concentration (B) from normal diet basal, normal diet insulin stimulated, high fat diet basal and high fat diet insulin stimulated animals

*Significantly different from normal diet basal ($P < 0.05$).

†Significantly different from high fat diet basal ($P < 0.05$).

#Significantly different from high fat diet insulin stimulated ($P < 0.05$).

Values are means \pm S.E.M.

metabolism we found that rates of hind limb 3-MG transport, PI3-kinase activity and plasma membrane GLUT4 protein concentration were significantly reduced in the high fat fed animals compared to the normal diet animals. These observations are in agreement with a number of investigations that have also shown a high fat diet to decrease insulin-stimulated 3-MG transport (Kraegen *et al.* 1986; Barnard *et al.* 1998; Hansen *et al.* 1998; Wilkes *et al.* 1998; Buettner *et al.* 2000; Halseth *et al.* 2000; Yaspelkis *et al.* 2001; Singh *et al.* 2003; Krisan *et al.* 2004), PI3-kinase activity (Tremblay *et al.* 2001; Singh *et al.* 2003; Krisan *et al.* 2004; Yaspelkis *et al.* 2004) and plasma membrane GLUT4 protein concentration (Hansen *et al.* 1998; Tremblay *et al.* 2001; Yaspelkis *et al.* 2001; Singh *et al.* 2003; Yaspelkis *et al.* 2004) in rodent skeletal muscle. Additionally, we found that total skeletal muscle insulin-stimulated Akt2 and aPKC ζ/λ activities were reduced in the high fat fed animals even though total skeletal muscle Akt2, aPKC ζ and λ protein concentrations were unaltered. These observations are not entirely unexpected since they are consistent with those of several recent investigations that have also reported Akt2 and aPKC ζ/λ activities to be reduced in the absence of alterations in total protein concentration and phosphorylation (Tremblay *et al.* 2001; Kanoh *et al.* 2003; Krisan *et al.* 2004). However, to the best of our knowledge, it has not previously been addressed whether insulin-stimulated Akt2, aPKC ζ and λ protein concentration and/or activation of these kinases are altered in plasma membrane prepared from high fat fed rodent skeletal muscle.

Similar to what we observed in the plasma membranes of the normal diet animals, we did not find that insulin increased either the plasma membrane associated Akt2 protein concentration or activity above basal levels. Nevertheless, despite a lack of effect at the plasma membrane, the high fat diet did alter total Akt2 kinase activity. While the mechanism by which high fat feeding results in a decrease total Akt2 activity has not been fully resolved, it is likely to be due to insulin-stimulated PI3-kinase activity being reduced.

With respect to the plasma membrane protein concentration and activation of aPKC ζ and λ protein concentration in the high fat fed skeletal muscle, we found that they were increased above basal levels in response to insulin. However, the insulin-stimulated increase in plasma membrane associated aPKC ζ and λ , and aPKC ζ/λ activities in the skeletal muscle obtained from high fat fed animals was less than that of the normal diet animals. While this observation has not been previously reported in rodent skeletal muscle, it is likely that the reduced plasma membrane protein concentration and activation of aPKC ζ and λ is a direct result of insulin-stimulated PI3-kinase activity being reduced in the high fat fed animals. It is also possible that alterations in the CAP/Cbl insulin signalling

cascade (novel insulin signalling cascade) may contribute to the concentration and activation of aPKC ζ/λ being reduced in the plasma membranes obtained from the high fat fed rodent skeletal muscle. Kanzaki *et al.* (2004) have recently reported that insulin stimulates the recruitment of aPKC ζ/λ to the plasma membrane of 3T3L1 adipocytes through the activation of the CAP/Cbl insulin signalling cascade rather than through the PI3-kinase-dependent (classical) insulin signalling cascade, and that the CAP/Cbl cascade converges with the PI3-kinase-dependent pathway via the interaction of TC10 with aPKC ζ/λ at the plasma membrane. Of interest, we have recently observed that insulin-stimulated activation of the CAP/Cbl insulin signalling cascade in skeletal muscle is improved by aerobic exercise training (Bernard *et al.* 2005). However, additional investigation of the CAP/Cbl insulin signalling cascade is warranted to elucidate the existence and functional significance of the interaction of TC10 with aPKC ζ/λ , and if these contribute to GLUT4 translocation in skeletal muscle.

In summary, we observed in normal skeletal muscle that insulin increased total PI3-kinase, Akt2 and aPKC ζ/λ activities above basal levels. In response to insulin, plasma membrane-associated aPKC ζ , aPKC λ and GLUT4, but not Akt2, were increased, and that only aPKC ζ/λ activity was increased at the plasma membrane. High fat feeding did not alter total Akt2 or aPKC ζ and λ protein concentrations. However, insulin-stimulated total PI3-kinase, Akt2 and aPKC ζ/λ activities were reduced in the skeletal muscle of the high fat fed animals. Insulin-stimulated plasma membrane-associated aPKC ζ and λ protein concentration, aPKC ζ/λ activity and GLUT4 protein concentration were also reduced in the high fat fed animals. These findings suggest that in skeletal muscle, insulin stimulates translocation of aPKC ζ and λ to the plasma membrane, whereas Akt2 is undetected at the plasma membrane. In addition, high fat feeding impairs insulin-stimulated translocation and activation of aPKC ζ/λ , which may contribute to decreasing insulin-stimulated plasma membrane GLUT4 protein concentration and rates of insulin-stimulated glucose transport in the skeletal muscle of the high fat fed animals.

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