

Insulin Signaling and Insulin Sensitizing in Muscle and Liver of Obese Monkeys: Peroxisome Proliferator-Activated Receptor Gamma Agonist Improves Defective Activation of Atypical Protein Kinase C

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Abstract

Obesity, the metabolic syndrome, and aging share several pathogenic features in both humans and non-human primates, including insulin resistance and inflammation. Since muscle and liver are considered key integrators of metabolism, we sought to determine in biopsies from lean and obese aging rhesus monkeys the nature of defects in insulin activation and, further, the potential for mitigation of such defects by an *in vivo* insulin sensitizer, rosiglitazone, and a thiazolidinedione activator of the peroxisome proliferator-activated receptor gamma. The peroxisome proliferator-activated receptor gamma agonist reduced hyperinsulinemia, improved insulin sensitivity, lowered plasma triglycerides and free fatty acids, and increased plasma adiponectin. In muscle of obese monkeys, previously shown to exhibit defective insulin signaling, the insulin sensitizer improved insulin activation of atypical protein kinase C (aPKC), the defective direct activation of aPKC by phosphatidylinositol (PI)-3,4,5-(PO₄)₃, and 5'-AMP-activated protein kinase and increased carnitine palmitoyltransferase-1 mRNA expression, but it did not improve insulin activation of insulin receptor substrate (IRS)-1-dependent PI 3-kinase (IRS-1/PI3K), protein kinase B, or glycogen synthase. We found that, although insulin signaling was impaired in muscle, insulin activation of IRS-1/PI3K, IRS-2/PI3K, protein kinase B, and aPKC was largely intact in liver and that rosiglitazone improved insulin signaling to aPKC in muscle by improving responsiveness to PI-3,4,5-(PO₄)₃. *Antioxid. Redox Signal.* 14, 207–219.

Introduction

THE PATHOGENESIS OF METABOLIC SYNDROME clearly involves alterations in the intracellular signaling cascades that are thought to underlie insulin resistance and to be specifically involved in diseases of aging (22). These diseases of aging are very similar in humans and in rhesus monkeys; the middle-aged onset of metabolic syndrome includes obesity, dyslipidemia, and impaired glucose tolerance, often progressing to overt type 2 diabetes mellitus (8, 9, 21–23, 32, 33, 67). Rhesus monkeys and humans have highly similar metabolism (17, 42, 46, 68) and genomic features (20, 50). Diabetes and insulin resistance in monkeys and other non-human primates are likely to have the same mechanistic causes as

underlie diabetes development in overweight middle-aged humans (11). Further, monkey insulin is identical to human insulin (51). Thus, this model is ideal for examining the nature of insulin signaling and insulin action, particularly in less accessible tissues that are difficult to obtain in humans under both basal and insulin-stimulated conditions, such as is true for liver. Insulin resistance seems to underlie the early stages in the development of the metabolic syndrome; and, thus, approaches to improving insulin action have been and remain key targets for potentially slowing or ultimately preventing type 2 diabetes (14, 18, 24).

Rhesus monkeys are also ideally suited to the examination of mechanisms of action of insulin-sensitizing agents, specifically where altered pathways may differ across different

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organs and at different stages of the disease progression. Such agents have been shown to have similar effects at the whole body level in humans and monkeys (16, 30, 52, 58). Prominent among the agents targeting insulin action are the peroxisome proliferator-activated receptor (PPAR) agonists and partial agonists including the thiazolidinediones (TZDs), such as rosiglitazone (RSGZ) and pioglitazone. Both are considered to be useful insulin-sensitizing agents for treating type 2 diabetes mellitus, and these, as well as other PPAR agonists, have been shown to be effective in humans and rhesus monkeys (10, 16, 30, 52, 56, 58, 86). The mechanisms by which they exert their insulin sensitizing effects are, however, only partially understood. The therapeutic effects of TZDs are thought to result from activation of PPAR γ receptors primarily located in adipose tissue (26), and possibly to a lesser degree, but of still uncertain significance, in muscle and liver. Nevertheless, TZDs and other PPAR agonists improve whole body insulin sensitivity and insulin signaling, not only in adipocytes [e.g., (28, 66)] but also in muscle of rats (27), rhesus monkeys (58), and humans (6, 34). In conjunction with improved insulin signaling, TZDs and other partial and full PPAR agonists improve insulin-stimulated glucose transport in isolated adipocytes (28) and muscle preparations (90). Moreover, insulin-stimulated glucose disposal rates increase in subjects with type 2 diabetes after TZD treatment [e.g., (6, 34, 58)], among many), reflecting improved muscle uptake and metabolism of glucose.

Most metabolic processes that are regulated by insulin in muscle, adipocytes, and liver appear to involve changes in the activity of protein phosphatases and of protein kinases, activities which may be altered by oxidative stress or intracellular redox imbalance. Insulin action at these organs involves the activation of insulin receptor substrate (IRS)-1-dependent and/or IRS-2-dependent phosphatidylinositol (PI) 3-kinase (PI3K), which, in turn, activate atypical protein kinase Cs (aPKCs) and protein kinase B ([PKB]/Akt). Both aPKC (2–5, 39, 69, 75) and PKB (1, 25, 38, 79, 85) appear to be required for insulin-stimulated glucose transport in muscle and adipocytes. Further, PKB, but not aPKC, is also required for insulin effects on glycogen synthesis in muscle, adipocytes, and liver and possibly other factors that limit hepatic glucose output, at least in rodents (12, 15, 65, 74) (Fig. 1). On the other hand, aPKCs appear to be important for mediating insulin effects on lipid synthesis in mouse liver (44, 78).

From studies of the knockout of genes encoding IRS-1 or IRS-2 in mice, multiple tissue-specific differences in mechanisms used by insulin to activate aPKCs and PKB have become apparent between muscle, liver, and adipose tissue. In muscle, the activations of both aPKCs (71) and PKB (71, 81) are dependent on IRS-1-dependent PI3K rather than IRS-2-dependent PI3K; in liver, however, PKB activation is dependent on both IRS-1-dependent (71, 81) and IRS-2-dependent (82) PI3K, whereas aPKC activation is dependent on IRS-2-dependent (72, 82) but not on IRS-1-dependent (71) PI3K; in adipocytes, aPKC activation is dependent on both IRS-1-dependent and IRS-2-dependent PI3K, whereas PKB activation is not compromised by deficiency of either IRS-1 or IRS-2 (49, 71).

In rodent models, these insulin-induced activities have frequently been shown to differ between muscle and liver. For example, the activation of aPKC in muscle is consistently diminished in a variety of insulin-resistant conditions (6, 7, 27, 29, 35, 37, 76, 80), whereas aPKC activation in liver is main-

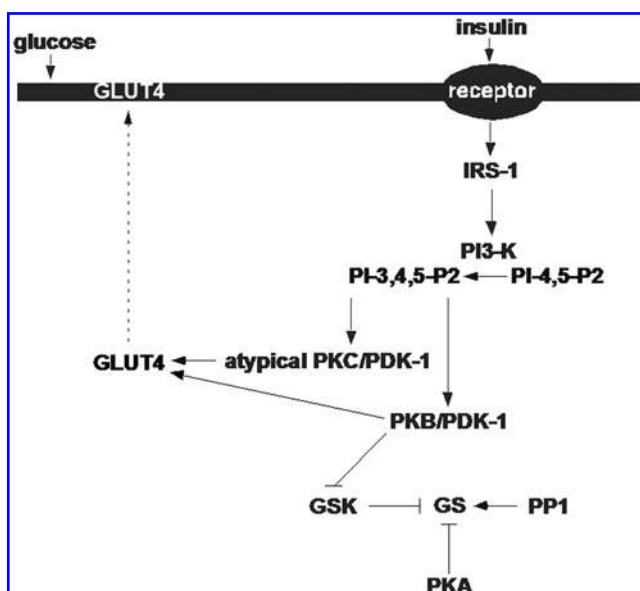


FIG. 1. Model of insulin signaling in skeletal muscle. IRS-1, insulin receptor substrate-1; PI 3-K, phosphatidylinositol 3-kinase; PDK-1, 3-phosphoinositide dependent protein kinase-1; PKB, protein kinase B; GSK, glycogen synthase kinase; GS, glycogen synthase; PP1, protein phosphatase-1; PKA, protein kinase A; GLUT 4, glucose transporter 4.

tained or increased (77). PKB activation in muscle is either maintained (6, 36, 76, 83) or diminished (40) and in liver either maintained or impaired (77) depending on the insulin-resistant state or animal model. Thus, there are divergent alterations in aPKC and PKB activation that seem to be, in part, due to alterations in IRS-1- and IRS-2-dependent PI3K. Accordingly, in high-fat-fed mice, an obesity-like model, insulin activation of IRS-1-dependent but not IRS-2-dependent PI3K is defective, thus explaining defects in aPKC and PKB activation in muscle; in contrast, in liver, insulin activation of both IRS-1- and IRS-2-dependent PI3K is intact, thus explaining the conserved activation of both aPKC and PKB (70, 71). In this high fat diet fed rodent model, there are defects in activation of aPKCs, with or without defects in PKB activation, in skeletal muscle, but apparently normal insulin signaling to IRS-1- and IRS-2-dependent PI3K, aPKCs, and PKB in liver (77). Further, in another insulin-resistant model, the GK-diabetic rat, insulin activation of IRS-1-dependent, but not IRS-2-dependent PI3K is defective in muscle, thus explaining the defect in aPKC activation (27, 72, 73, 77). Similarly, in liver of the GK rat, insulin activation of IRS-1-dependent PI3K is defective, thus explaining the defect in PKB activation, but, in contrast, insulin activation of IRS-2-dependent PI3K is intact, thus explaining conserved aPKC activation (73, 77). Of further note, in GK-diabetic rats, TZDs improve aPKC activation in muscle (27, 29, 34) but do not alter aPKC or PKB activation in liver (77).

In muscle and adipose tissue of type 2 diabetic rats (27–29) and humans (6, 27), the mechanism whereby TZDs improve insulin-stimulated aPKC activation in the absence of alterations in PI3K and PKB activation is enigmatic. However, the ability of aPKCs to directly respond to the lipid product of PI3K, namely, PI-3,4,5-(PO $_4$) $_3$ (PIP $_3$), is impaired in muscles of obese and type 2 diabetic rhesus monkeys (76) and humans

(6, 7) as well as in muscles of high-fat-fed rats and mice (27, 29) and diabetic mice (73). The possibility that TZDs may alter muscle aPKC responsiveness to PIP_3 , particularly in the absence of changes in IRS-1-dependent PI3K, has not been previously examined.

With regard to more naturally occurring forms of obesity, much less is known about alterations in insulin signaling in various tissues. Defective aPKC activation occurs in muscle of obese humans (7, 35) and monkeys (76) and in isolated adipocytes and myocytes of obese humans (70, 83). However, there is little information on insulin signaling in liver of spontaneously occurring forms of obesity, before the development of overt diabetes.

Here, we have examined insulin signaling in the liver of normal monkeys with or without insulin stimulation, for comparison to our previous study of insulin signaling in muscle. We also determined the effects of a TZD on mechanisms of insulin action in muscle of monkeys who spontaneously develop obesity while consuming standard chow *ad libitum*. Such obese monkeys have been previously shown to have defects in insulin action at skeletal muscle, specifically in the activation of IRS-1-dependent PI3K, PKB, aPKC (76), glycogen synthase (GS) (61), and protein phosphatase-1 (55) and in the inactivation of protein kinase A (53). We have previously found that in obese monkeys, insulin action on liver GS and glycogen phosphorylase is intact (57), although, as in muscle, liver triglyceride is higher in obese than in lean monkeys (64). It was, therefore, interesting to find that, whereas the insulin activation of IRS-1-dependent PI3K, PKB, aPKC and GS is impaired in muscle, the activation of all examined insulin signaling factors, namely, IRS-1-dependent PI3K, IRS-2-dependent PI3K, PKB, and aPKC, was not significantly compromised in liver. Due to the absence of demonstrably impaired signaling in liver, insulin signaling pathway alterations in response to whole body insulin sensitizing was pursued only in muscle. In muscle, the TZD, RSGZ, a PPAR γ agonist, selectively improved aPKC activation without altering defects in the activation of IRS-1-dependent PI3K, PKB, or GS in muscle. Interestingly, the improvement in muscle aPKC activation was, at least, partly due to improved responsiveness of aPKCs to PIP_3 .

Research Design, Materials, and Methods

Subjects

Fifteen adult male rhesus monkeys (*Macaca mulatta*) were studied. All were bred in the United States and were of Indian origin. The monkeys were individually housed, and consistent care was provided according to the Guide for the Care and Use of Laboratory Animals of the ILAR/NAS 1996, including attention to environmental enrichment. Food intake was daily monitored, and body weight was weekly or bi-weekly monitored throughout the study. Blood and tissue samples were always obtained after a consistent 16h overnight fast. Sedation for all blood draws and experimental procedures was accomplished with ketamine hydrochloride (10–15 mg/kg body weight) with supplemental ketamine, ~5–10 mg/kg, given as needed at 20–30 min intervals during the procedures. All protocols were approved by the Institutional Animal Care and Use Committee.

Ten of these 15 monkeys were normal, studied to determine the normal action of insulin at the liver compared with

muscle. This group consisted of mature adults (~7 years old) that were lean (9.1 ± 0.5 kg body weight and $9.5 \pm 1.5\%$ body fat) and had normal fasting plasma glucose (FPG) (3.1 ± 0.1 mM) and normal insulin (300 ± 30 pM) levels. These monkeys also had normal chemistry and hematology values at the start of the study. Here we report the results of analysis of insulin signaling in the liver of these normal monkeys. We have previously reported the analysis of liver GS, glycogen phosphorylase (62), protein phosphatase-1, protein phosphatase-2C, protein kinase A, GS kinase-3, and PKC (54) regulation by insulin in these normal animals.

Five additional monkeys were both obese (body weights >12 kg) and insulin-resistant (glucose disposal rates under euglycemic clamp conditions of <8.0 mg/kg/FFM/min). These obese monkeys were prediabetic (FPG 3.3–7.0 mM) and were studied both before and at the end of treatment with the TZD, RSGZ (a PPAR γ agonist), (kindly supplied by Glaxo-SmithKline). RSGZ was administered orally at three doses (0.03, 0.1, 0.3 mg/kg body weight/day) for 4 weeks at each dose, a total of 12 weeks of treatment, preceded by vehicle and followed by washout periods. Blood samples were biweekly obtained, and tissue samples of muscle were obtained at the end of the 4-week period of administration of vehicle, and again at the end of the 12-week TZD treatment period. Blood and tissue samples were also obtained under maximally insulin-stimulated conditions during an euglycemic hyperinsulinemic clamp before and at the end of the treatment. Blood samples were analyzed for FPG, insulin, nonesterified fatty acids, adiponectin (human adiponectin RIA kit; Linco), triglycerides, very low-density lipoprotein (VLDL), and high-density lipoprotein lipid fractions, and routine clinical chemistry and hematology.

Procedures

Euglycemic hyperinsulinemic clamps to measure whole body insulin sensitivity were conducted after an overnight fast and included obtaining of skeletal muscle (vastus lateralis) and liver biopsies both basally and at maximal insulin stimulation at the end of the clamp period. Anesthesia was maintained with fentanyl citrate (0.01 mg/kg). Succinylcholine (1 mg/kg) was initially used followed by vecuronium bromide for muscle relaxation (0.1 mg/kg) and diazepam (2.5 mg/dose). The open biopsies of muscle and liver were obtained just before initiation of the insulin infusion and during steady-state insulin infusion (90–120 min after the onset of the insulin infusion), as previously described (61, 62). In normal monkeys, we determined the activities of multiple muscle and liver enzymes (see below) before (under basal conditions) and after maximal insulin stimulation and identified the differences in insulin signaling between these two target organs. In RSGZ-treated monkeys, the euglycemic hyperinsulinemic clamps were conducted just before the start of the initial dosing and at the end of the final highest dosing period after 12 weeks of RSGZ treatment. Maximal insulin-stimulated whole-body glucose disposal rate (M) was estimated using an insulin infusion rate of 400 mU/m²/min with maintenance of glucose at 4.7 pM using a variable rate infusion of 20% dextrose. M was corrected for fat-free mass determined by the tritiated water dilution method.

All tissues samples were rapidly frozen *in situ* (skeletal muscle) or immediately *ex situ* (liver) using aluminum clamps

cooled in liquid nitrogen. Tissue samples were lyophilized and stored in liquid nitrogen until enzyme assays were performed.

Enzyme and substrate assays

Muscle and liver tissue samples were homogenized in appropriate buffers (76). Atypical PKC activity was measured as described (76). In brief, aPKCs, ζ , λ , and ι , were immunoprecipitated from cell lysates with a rabbit polyclonal antiserum (Santa Cruz Biotechnologies, Inc.) that recognizes the nearly identical C-termini of PKC- ζ and PKC- λ/ι [The individual aPKCs apparently function interchangeably during insulin-stimulated glucose transport (2–5, 39, 75)]. Precipitates were collected on Sepharose-AG beads and incubated for 8 min at 30°C in 100 μ l buffer containing 50 mM Tris/HCl (pH 7.5), 100 μ M Na₃VO₄, 100 μ M Na₄P₂O₇, 1 mM NaF, 100 μ M PMSF, 4 μ g phosphatidylserine (Sigma), 50 μ M (γ -³²P) ATP (NEN Life Science Products), 5 mM MgCl₂, and, as substrate, 40 μ M serine analog of the PKC- ϵ pseudo-substrate (BioSource), a preferred substrate for aPKCs. In some assays, 10M PIP₃ (Matreya), a maximally effective concentration, was added to activate aPKCs (6, 27, 29, 76). After incubation, ³²P-labeled substrate was trapped on P-81 filter paper and counted.

PKB enzyme activity was measured using a kit obtained from Upstate Biotechnologies Inc. (UBI), as previously described (76). In brief, PKB α was immunoprecipitated with sheep polyclonal anti-PKB α antiserum (UBI), collected on Sepharose-AG beads, and assayed as per kit directions. PKB activation was also assessed by immunoblotting for phosphorylation of serine-473 (76, 77).

Immunoprecipitable IRS-1- and IRS-2-dependent PI3K activities were determined as previously described (76, 77) (rabbit polyclonal antisera for IRS-1 and IRS-2 were purchased from UBI). Autoradiographic results of chromatographically purified PI-3-³²PO₄, the lipid product of the PI3K assays, were quantified in a BioRad Phosphor-Imager, and finally expressed relative to the control samples developed on the same thin layer chromatography plate.

Immunoprecipitable 5'-AMP-activated protein kinase (AMPK) (combined 1 and 2) activity in muscle lysates was measured with the method of Wojtaszewski *et al.* (87) using rabbit polyclonal antiserum (Cell Signaling Technologies) and SAMS peptide (UBI) as substrate.

GS activity, glycogen, and glucose 6-phosphate (G6P) content were measured in lyophilized, micro dissected skeletal muscle as we have described (59, 63). We have previously reported these for liver.

Lipid was isolated from skeletal muscle using a modified method of Folch. Twenty milligram of lyophilized, micro dissected sample was homogenized in 4 ml chloroform:methanol (2:1) and rotated overnight in a glass vial. The following day, 2 ml of 0.6% NaCl was added to the vial, vortexed, and centrifuged for 10 min @ 2000 \times g (4°C). The lower phase was removed and placed into a clean glass vial. After complete evaporation, 250 μ L of 100% ethanol was added to the dried sample. Triglyceride content was determined by using an enzymatic kit (Glycerol; r-biopharm) in conjunction with the enzymes esterase and lipase (Sigma). Triglyceride content was expressed as nmol/mg dry tissue weight.

In all assays, comparisons between baseline and RSGZ-treated samples from the same monkey, or between samples

from normal lean controls and obese monkeys, were made with samples that were simultaneously assayed.

Carnitine palmitoyltransferase-1 mRNA expression in skeletal muscle with and without insulin stimulation before and during TZD treatment

Total RNA was extracted from 20 mg lyophilized muscle obtained before and during the euglycemic hyperinsulinemic clamp during vehicle administration and at the end of the 12 weeks of RSGZ administration and processed as described (56). Muscle carnitine palmitoyltransferase-1 (CPT-1) mRNA expression was determined by real-time reverse transcription-polymerase chain reaction expressed relative to the house-keeping ribosomal gene 36B4 and based on concentrations from standard curves. Taqman probes (Assays on Demand) for this gene were purchased from Applied Biosystems (CPT-1 assay ID Hs00992651 g1) and used in the LightCycler (Roche Diagnostics) with the LightCycler FastStart DNA Master Hybridization Probe Kit. All samples were run in triplicate.

Western analyses

Lysate proteins were resolved by sodium dodecyl sulfate-polyacrylamide gel electrophoresis, transferred to nitrocellulose membranes, and immunoblotted as described (76, 77). Antibodies used for blotting included rabbit polyclonal anti-C-terminal PKC- $\zeta/\lambda/\iota$ antiserum (Santa Cruz Biotechnologies, Inc.) (PKCs ζ and λ/ι have nearly identical C-termini that are recognized by this antiserum); sheep polyclonal anti-PKB α antiserum (UBI); rabbit polyclonal anti-IRS-1 and IRS-2 antisera (UBI); and rabbit polyclonal anti-phospho-ser-473-PKB antiserum (Cell Signaling). Immunoblots were quantified by measurement of chemiluminescence in a BioRad Molecular Analyst Chemiluminescence/Phosphorescence Imaging System or by laser scanning densitometry.

Statistical methods

Data are expressed as mean \pm standard error of the mean. Means were compared by Student's *t*-test for paired or unpaired samples. Pearson's correlation coefficient was used to test for significant linear relationships between variables.

Results

Assessment of regulation of insulin signaling in liver of lean and obese insulin-resistant monkeys

In the present study, we compared insulin signaling in the liver of obese monkeys with that of lean control monkeys and compared hepatic insulin signaling with our previous study of muscle insulin signaling in lean and obese monkeys (76). Unexpectedly, in view of the significant differences in insulin signaling at muscle in lean *versus* obese monkeys, in liver there were no obesity associated changes. Comparing basal activity with activity during the maximal hyperinsulinemic stimulation of a euglycemic clamp, the activities of several kinases involved in insulin signaling were equally increased in activity in lean and obese monkeys. As shown in Figure 2, there were no apparent differences in the activation by insulin of either IRS-2-dependent PI3K (Fig. 2A) or IRS-1-dependent PI3K (Fig. 2B). Thus, it appears that the initial steps of insulin signaling are not compromised in liver of obese monkeys compared with lean animals.

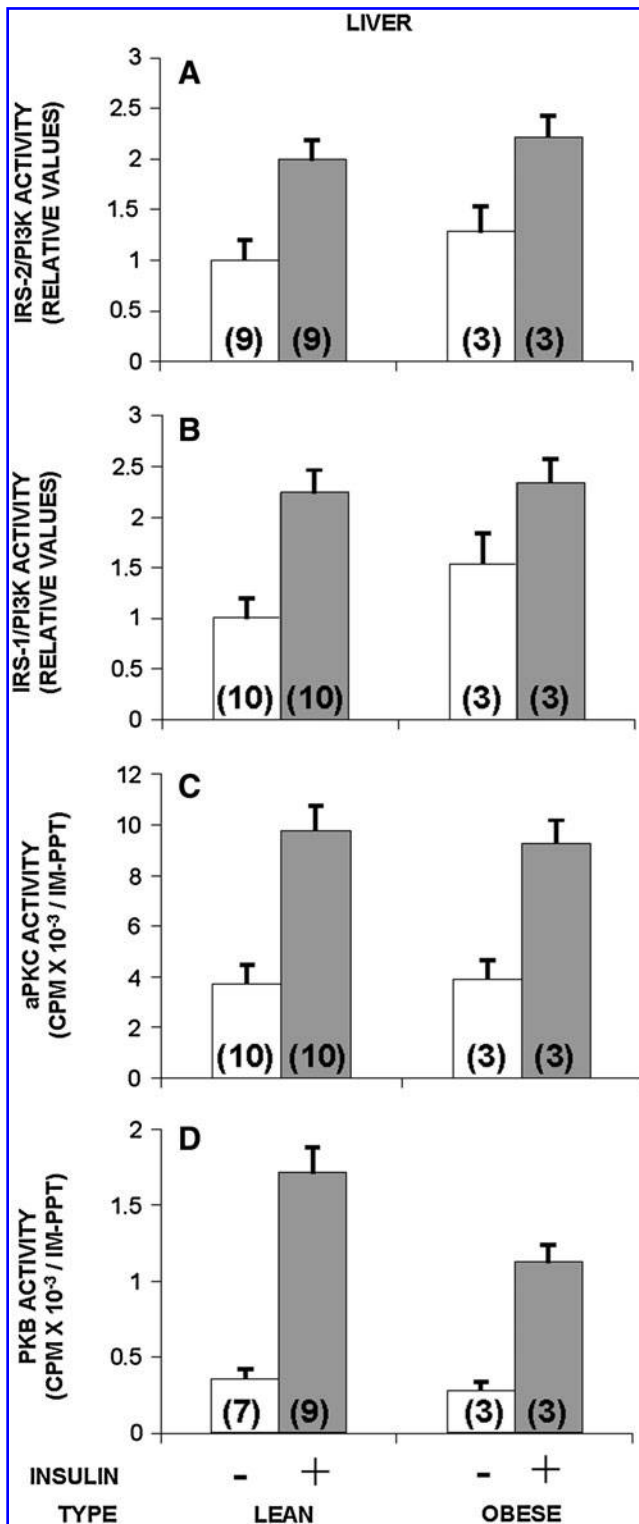


FIG. 2. Comparison of insulin signaling to IRS-2-dependent PI3K (A), IRS-1-dependent PI3K (B), aPKCs (C), and PKB (D), and in liver of lean versus obese monkeys before (-) and during (+) the insulin stimulation of a euglycemic hyperinsulinemic clamp. Basal values are shown by clear bars, and insulin-stimulated values are shown by shaded bars. The number of determinations is shown in parentheses. In the liver, there were no significant differences between the effects of insulin in lean versus obese (insulin-resistant) monkeys (all p 's > 0.05). aPKC, atypical protein kinase C.

As with IRS-2-dependent PI3K, which presumably functions upstream of aPKC in liver (71, 82), the activation of aPKC by insulin was essentially the same in liver of lean and obese monkeys (Fig. 2C).

The enzymatic activation of PKB by insulin, which appears to be dependent on both IRS-1- and IRS-2-dependent PI3K in liver (71, 81, 82), tended downward, but not significantly, in the liver of obese monkeys, relative to that seen in lean monkeys (Fig. 2D).

Effects of PPAR γ agonist RSGZ on *in vivo* metabolic parameters

Treatment of obese insulin-resistant prediabetic monkeys with the TZD insulin sensitizer, RSGZ, over a 12 week escalating dose regimen reaching a maximal dose of 0.3 mg/kg in the final 4 weeks, resulted in an average body weight gain of 5%. Amount of weight gain widely ranged from 0% to 8% over 3 months (mean + 4.5%; p = n.s.; Fig. 3A), including a gain of 1% to 3% during 1 month at the highest dose (\sim threefold human recommended dose levels; [p = n.s.]). Fasting plasma insulin levels were consistently reduced in all monkeys by an average of 16% (p = 0.02) in response to RSGZ treatment (Fig. 3B). Whole body insulin sensitivity increased by 17%, with four of five monkeys showing an improvement in insulin sensitivity under RSGZ treatment (Fig. 3C). The decline in fasting insulin was significantly negatively correlated with the improvement in insulin sensitivity. Plasma triglyceride levels significantly decreased (p = 0.008), with those having the highest initial triglyceride levels showing the greatest reduction in response to RSGZ (Fig. 3D); VLDL triglycerides declined at an average of 0.34 ± 0.12 mM (p < 0.05) (not shown). Fasting plasma nonesterified fatty acids decreased from elevated levels (>400 mEq/L) in three monkeys but were unchanged in the two monkeys in whom initial pretreatment levels were normal (Fig. 3E). Plasma adiponectin level increases were highly consistent across monkeys and were highly significant (p = 0.002, Fig. 3F). FPG levels were unchanged in the four monkeys that were normoglycemic at study initiation (64 vs. 70 mg/dl start to end of study) and slightly increased in the early diabetic monkey (not shown).

Effects of RSGZ on insulin signaling and insulin action in skeletal muscle

During euglycemic hyperinsulinemic clamp studies to examine whole body insulin sensitivity, we previously identified defects in the *in vivo* insulin activation in muscle of IRS-1-dependent PI3K, PKB, aPKC (76), and GS (61) (vastus lateralis) in obese monkeys. In the present study, we found that relative to pretreatment values, administration of RSGZ had no effect on the basal or insulin-stimulated activities of IRS-1/PI3K (Fig. 4A) in muscle. In keeping with the failure to observe alterations in IRS-1/PI3K, RSGZ did not affect the enzymatic activation of PKB by insulin in muscle of obese monkeys (Fig. 4B). In addition, RSGZ failed to alter the phosphorylation of serine-473 in PKB (Fig. 5).

By contrast, RSGZ treatment significantly improved the activation of aPKCs by insulin (p < 0.05, Fig. 4C), although, as seen in Figure 5, RSGZ did not alter protein expression of aPKCs in muscle.

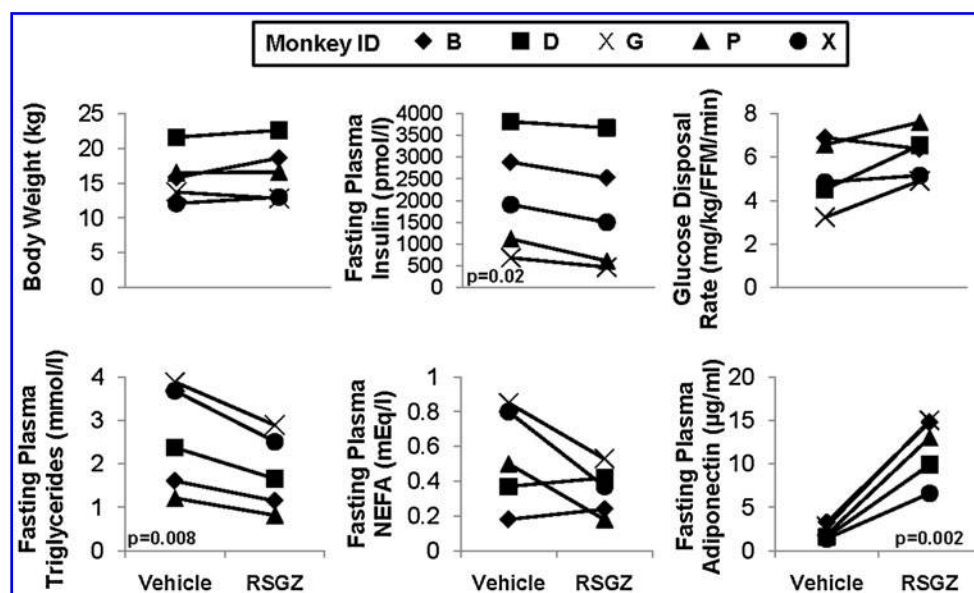


FIG. 3. Alterations in body weight (*upper left*), fasting plasma levels of insulin (*upper middle*), and insulin-mediated glucose disposal rate (whole body insulin sensitivity) (*upper right*), and of fasting triglycerides (*lower left*), NEFA (*lower middle*), and adiponectin levels (*lower right*) after RSGZ treatment of obese insulin-resistant monkeys. RSGZ, rosiglitazone; NEFA, nonesterified fatty acid.

In this study, we did not examine IRS-2-dependent PI3K activation, as other studies have suggested that in muscle the activation of aPKC and PKB are dependent on IRS-1 rather than on IRS-2 (71, 77) (also found in unpublished observations in IRS 2-knockout mice).

In view of the failure to observe an increase in insulin-stimulated IRS-1-dependent PI3K activation following RSGZ treatment, we examined the ability of muscle aPKC to respond to PIP₃, the lipid product of PI3K. As seen in Figure 4D (shaded bars), aPKCs, immunoprecipitated from muscle of monkeys treated with RSGZ, responded significantly more effectively to PIP₃ than aPKCs immunoprecipitated from muscle obtained prior to such treatment ($p < 0.05$). (Note that PIP₃ was added only to aPKCs immunoprecipitated from muscle under basal conditions, not stimulated by insulin.) It, therefore, seems reasonable to suggest that this improvement in aPKC responsiveness to PIP₃ importantly contributed to the increase in aPKC activation observed in response to insulin administration during the hyperinsulinemic clamp procedure (Fig. 4C).

In contrast to aPKC, but in keeping with a failure to see alterations in IRS-1-dependent PI3K and PKB, in these insulin-resistant monkeys, neither did RSGZ, compared with vehicle, increase the already impaired activation of GS by insulin (Maximal insulin-stimulated GS fractional activity during vehicle: $16.4\% \pm 3.7\%$ vs. during RSGZ treatment: $17.5\% \pm 3.2\%$) nor did treatment affect GS independent activity or total GS activity. There was also no significant effect of RSGZ on muscle G6P or glycogen content. Muscle triglyceride content was slightly, but not significantly, lower during RSGZ treatment (vehicle; 73 ± 21 vs. RSGZ 50 ± 14 nmol/mg dry weight, n.s.). Nevertheless, the monkeys with the highest basal skeletal muscle triglyceride content before RSGZ treatment had the greatest decrease in basal triglyceride content after RSGZ treatment ($r = -0.85$, $p < 0.05$).

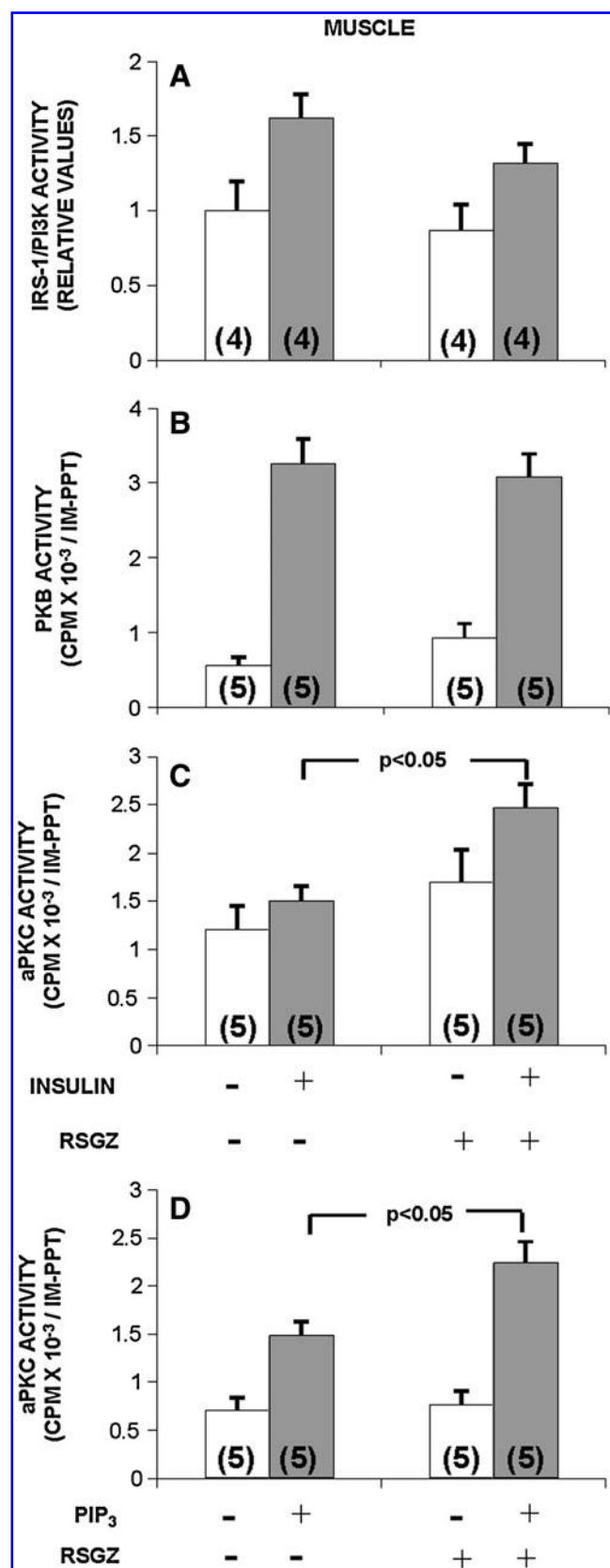
Effects of RSGZ on AMPK activity and CPT-1 gene expression in muscle

The failure to see an effect of RSGZ on IRS-1-dependent PI3K, and, on the other hand, the ability to see an effect of RSGZ on aPKC responsiveness to PIP₃, prompted us to examine whether factors that may influence the intracellular metabolism of lipids (that conceivably may down-regulate aPKC activity in conditions of obesity) might be involved in RSGZ actions in muscle. One such factor is AMPK, which has been shown to be activated both by RSGZ itself (19) and by adiponectin (88). Similar to the robust effect of RSGZ treatment on plasma adiponectin, RSGZ treatment provoked significant increases in basal fasting AMPK activity in muscle of obese monkeys (Fig. 6).

Another important factor that affects the intramyocellular metabolism of lipids, specifically long-chain acyl CoAs (LCA-CoAs), is CPT-1, the gene that controls fatty acid mitochondrial β -oxidation (48). Similar to the response of aPKC to *in vivo* insulin after RSGZ treatment (Fig. 4), CPT-1 mRNA expression in muscle during the maximal insulin stimulation of an euglycemic hyperinsulinemic clamp was significantly higher after RSGZ treatment compared with vehicle (Fig. 7).

Discussion

Insulin resistance and obesity, accompanied by dyslipidemia, have been well documented in rhesus monkeys with metabolic syndrome (8, 9, 17, 22, 23, 32, 33, 41, 43, 67, 68, 84). Calorie restriction and calorie restriction mimetic agents have been demonstrated in rhesus monkeys to reverse or prevent these metabolic disorders (13, 21, 24, 42, 45, 46, 50, 60, 64). From previous findings in obese insulin-resistant monkeys (55, 57, 61, 76) and the present study, it seems clear that, whereas insulin signaling to IRS-1-dependent PI3K, PKB, aPKC, and GS is compromised in muscle of obese monkeys



relative to lean healthy control monkeys (63, 76) and humans (7, 19, 35, 36, 40, 83, 87, 88), such signaling to each of these factors, and to IRS-2-dependent PI3K, is largely intact in liver of obese monkeys (57, 62). This pattern of compromised insulin signaling in muscle, coupled with conserved insulin signaling in liver, presently observed in obese insulin-resistant monkeys, is similar to the signaling pattern observed in muscle and liver of high-fat-fed mice (72, 77). These findings of differential apparent effects of insulin insensitivity in muscle and liver are in accord with our previous identification of insulin resistance (as measured by the euglycemic hyperinsulinemic clamp) as an early event in the progression from normal through sequential phases to overt diabetes (9), occurring in obese normoglycemic, normal glucose tolerant monkeys. Since glucose uptake rate under insulin stimulation is thought to principally reflect uptake by muscle, muscle insulin resistance is an early event, developing in the earliest phases of this diabetes trajectory (9, 53, 55, 61). By contrast, failure of insulin to suppress hepatic glucose production is a very late event in this progression, occurring in direct relationship to developing hyperglycemia (9, 62). Thus, this pattern would be in full accord with the absence of defects in insulin signaling at the liver in obese prediabetic monkeys (9). Thus, the development of impaired insulin action at the liver appears to be a late event in the progression to overt diabetes.

Taken together, the present and previous findings (9, 44, 57, 61, 62, 64, 71–73, 77, 78, 82) suggest that insulin signaling mechanisms in liver are initially largely intact in simple obesity and deteriorate only with the development of overt diabetes, selectively impairing IRS-1-dependent PI3K and PKB but sparing IRS-2-dependent PI3K and aPKCs (71, 77). The initial conservation of hepatic PKB activation contributes to the maintenance of relatively normal glucose tolerance in obesity, as PKB is important in regulating glycogen synthesis (65, 71), gluconeogenesis (15), and glucose release (74). The conservation of hepatic aPKC activation in both primate obesity (present results) and diabetic rodent models (72, 77), on the other hand, contributes to the maintenance of insulin-dependent synthesis and release of VLDL-triglycerides, as lipid synthesis effects of insulin in liver are thought to be largely mediated *via* aPKCs (44) and presumably IRS-2-dependent PI3K, which functions upstream of aPKCs in hepatocytes (82). Indeed, the development of hyperinsulinemia together with defects in insulin signaling in skeletal muscle, coupled with intact insulin action on IRS-2-dependent PI3K and aPKCs in liver, would be expected to lead to increased hepatic synthesis and release of VLDL-triglycerides, as seen in obese monkeys (17).

FIG. 4. Effects of RSGZ on basal and insulin-stimulated (A–C) or PIP₃-stimulated (D) activation of IRS-1-dependent PI3K (A), PKB (B), and aPKCs (C and D) in muscle of obese monkeys. Euglycemic clamps with muscle biopsies were performed before and at the end of 3 months of RSGZ treatment. Basal values are shown by clear bars, and insulin-stimulated values are shown by shaded bars. Note that PIP₃ was added only to assays of aPKCs immunoprecipitated from basal (noninsulin-treated) muscle. The number of determinations is shown in parentheses. RSGZ significantly increased aPKC activity ($p < 0.05$) and in basal (noninsulin-stimulated) samples RSGZ enhanced the response to PIP₃ ($p < 0.05$). PIP₃, PI-3,4,5-(PO₄)₃; RSGZ, rosiglitazone.

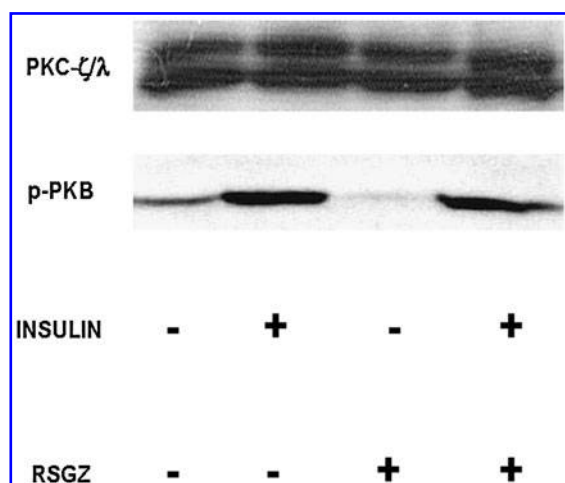


FIG. 5. Failure of RSGZ to alter aPKC levels or insulin effects on serine-473 phosphorylation of PKB in muscle of obese monkeys. Shown here are representative blots from two monkeys.

The reason insulin signaling is impaired in muscle, while normal in liver of both obese monkeys and high-fat-fed mice, is not entirely clear. Recently, we have found in high-fat-fed mice that basal activities of both conventional and novel diacylglycerol-sensitive PKCs are increased in muscle but not in liver, despite marked hyperlipidemia and hepatosteatosis (unpublished observations). Accordingly, the presence of insulin signaling defects in muscle and their absence in liver may be related to activities of these conventional/novel PKCs in muscle. Studies of conventional and novel PKC activities in monkey tissues across the entire progression from normal lean to overtly diabetic status are needed to test this hypothesis.

Effects of TZD treatment on insulin action

In the present study, we did not evaluate insulin signaling in liver before and after RSGZ treatment, as there were no

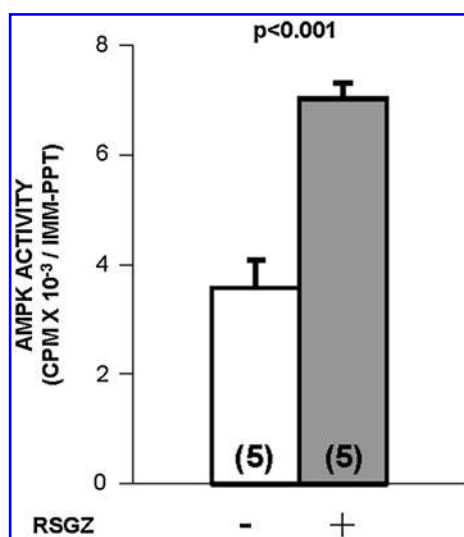


FIG. 6. Treatment of obese monkeys with RSGZ significantly increased AMPK activity in muscle ($p < 0.001$). The number of determinations is shown in parentheses. AMPK, 5'-AMP-activated protein kinase.

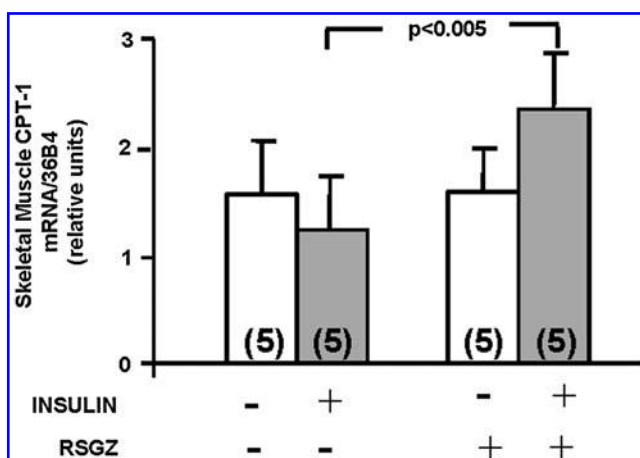


FIG. 7. Effect of RSGZ to increase CPT-1 mRNA gene expression during the euglycemic hyperinsulinemic clamp in muscle of obese monkeys. The number of determinations is shown in parentheses. CPT-1, carnitine palmitoyltransferase-1.

significant defects in insulin signaling in liver of obese monkeys. Specifically, in previous studies of liver of GK diabetic rats, RSGZ did not improve defects in IRS-1-dependent PI3K or PKB activation (77); also, in a preliminary study involving liver biopsies in insulin-resistant obese monkeys during another TZD treatment (R-102380; Sankyo, Inc.), there was additional evidence of no alteration in hepatic insulin signaling to IRS-1-dependent PI3K, IRS-2-dependent PI3K, PKB, aPKC, and GS, despite significant whole body insulin resistance (unpublished observations).

The mechanism for improved aPKC activation in muscle after RSGZ treatment is uncertain. Surprisingly, we found no evidence for improvement in the activation of IRS-1-dependent PI3K or PKB. On the other hand, muscle aPKC responsiveness to PIP_3 was increased after RSGZ treatment, and this most likely contributed to improved aPKC activation during insulin stimulation.

RSGZ did not increase GS activity, glycogen, or G6P content in muscle during the euglycemic hyperinsulinemic clamp assessing whole body insulin resistance in these obese monkeys. In contrast, a different TZD, R-102380, increased insulin activation of muscle GS and increased G6P content during a clamp in insulin-resistant monkeys (58). This difference may reflect the relative strength of the insulin sensitizing effects of RSGZ (mild) versus R-102380 (profound). Similarly, a PPAR α agonist, K-111, increased GS activity and glycogen content during a clamp in obese insulin-resistant monkeys (56). Whether the PPAR agonists R-102380 or K-111 increase insulin activation of IRS-1-dependent PI3K, PKC and/or PKB in muscle of obese monkeys has not yet been determined.

Despite improvement in insulin sensitivity at the whole body level, RSGZ did not fully repair the defect in muscle aPKC activation by insulin in obese monkeys. The fold increase in insulin-stimulated aPKC activity was relatively small in RSGZ-treated muscle of obese monkeys (about 50%), as compared with that seen in normal muscle, that increased ~two to fourfold (76). A more complete repair of the defect in aPKC activation by insulin would obviously require an improvement in IRS-1-dependent PI3K as well as an increase in responsiveness of aPKC to PIP_3 . Whether PIP_3 could fully

normalize aPKC activation in RSGZ-treated muscle was not directly tested.

The mechanism whereby aPKC responsiveness to PIP_3 was improved by RSGZ treatment is uncertain, but several factors should be considered. First, in rats high-fat feeding impairs aPKC responsiveness to PIP_3 (27, 29), and it is reasonable to propose that RSGZ may improve the lipid environment within muscle cells and thereby restore aPKC activity or activation by PIP_3 . In this regard, TZDs activate AMPK (19), which increases fatty acid oxidation and diminishes fatty acid synthesis. TZDs also increase adiponectin secretion from adipose tissue, and this factor similarly activates AMPK and increases fatty acid oxidation in muscle (88). Second, decreases in plasma levels of free fatty acids and triglycerides may improve the lipid environment within muscle cells. Third, TZD-induced increases in CPT-1 gene expression in muscle may increase β -oxidation of fatty acids and thereby diminish lipids. In this regard, although intramyocellular triglycerides were not affected by RSGZ treatment in diabetic humans (47), or in the obese monkeys in the present study, pioglitazone decreased triglyceride and LCA-CoA levels in basal muscle samples and further decreased LCA-CoA content during hyperinsulinemic clamps in high-fat-fed rats (89). In the present study, mean triglyceride content for the group was not significantly affected by RSGZ treatment; however, the monkeys with the highest basal triglyceride before RSGZ treatment had the greatest improvement in triglyceride content after RSGZ treatment.

In conjunction with alterations in insulin signaling to aPKC in muscle, RSGZ improved circulating VLDL-triglyceride levels of these obese monkeys. Presumably, this plasma lipid-lowering effect of RSGZ reflected increased lipid synthesis in adipose or other tissues, due to improved glucose uptake (to provide glycerol-3- PO_4) and activation of lipid-clearing enzymes, and/or decreased hepatic lipid synthesis. Given that insulin signaling to IRS-2-dependent PI3K and aPKCs, and thus lipid synthesis, in liver would be expected to be increased in insulin-resistant states, such as in obese hyperinsulinemic monkeys, any decrease in hepatic lipid synthesis would most likely be due to enhanced insulin action on glucose transport in muscle, leading to diminished insulin resistance in association with decreased ambient circulating insulin levels. Further studies are needed to test these hypotheses.

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Author Disclosure Statement

None of the authors have any competing financial interest, as confirmed by all authors.

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Abbreviations Used

AMPK = 5'-AMP-activated protein kinase
aPKC = atypical protein kinase C
CPT-1 = carnitine palmitoyltransferase-1
FPG = fasting plasma glucose
G6P = glucose 6-phosphate
GS = glycogen synthase
GSK-3 = glycogen synthase kinase-3
IRS-1/PI3K = insulin receptor substrate-1-dependent
phosphatidylinositol 3-kinase

LCA-CoA = long-chain acyl CoA
NEFA = nonesterified fatty acid
PIP₃ = PI-3,4,5-(PO₄)₃
PKA = protein kinase A
PKB = protein kinase B
PP1 = protein phosphatase-1
PPAR γ = peroxisome proliferators-activated
receptor gamma
RSGZ = rosiglitazone
TZD = thiazolidinedione
VLDL = very low-density lipoprotein

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