Increased \textit{in vivo} phosphorylation of insulin receptor at serine 994 in the liver of obese insulin-resistant Zucker rats

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Abstract

Serine phosphorylation of the insulin receptor (IR) has been proposed to exert an inhibitory influence on its tyrosine kinase activity. Previous works using site-directed mutagenesis suggested that serine 994 of the IR (IR Ser 994) might be part of an inhibitory domain of the receptor. In this study we examined whether this residue is subject to phosphorylation \textit{in vivo}. We used a site-specific antibody to determine the extent of phosphorylation of IR Ser 994 in insulin target tissues from two animal models of insulin resistance with different IR kinase (IRK) activity: obese (fa/fa) Zucker rats and transgenic mice overexpressing bovine growth hormone (PEPCK-bGH mice).

Phosphorylation at IR Ser 994 was markedly increased in liver of obese rats. This alteration appeared to be tissue-selective since no phosphorylation on Ser 994 was detected in IRs isolated from skeletal muscle of these animals. On the other hand, the phosphorylation level of IR Ser 994 was very low in liver of PEPCK-bGH mice and did not differ from that of the control group. We have also demonstrated that protein kinase (PK) C isoforms \(\alpha\) and \(\beta\) and \(\zeta\) are able to promote the \textit{in vitro} phosphorylation of the IR on Ser 994. Differential findings in these two models of insulin resistance might thus reflect increased PKC activity resulting from increased lipid availability in obese Zucker rats. Our results suggest that Ser 994 is a novel \textit{in vivo} IR phosphorylation site that might be involved in the regulation of the IRK in some states of insulin resistance.


Introduction

The insulin receptor (IR) is a tetrameric protein composed of two extracellular \(\alpha\)-subunits and two transmembrane \(\beta\)-subunits linked by disulfide bonds (White & Kahn 1994). Binding of insulin to the \(\alpha\)-subunits of the IR induces rapid phosphorylation of the IR \(\beta\)-subunit in tyrosine residues and activation of its intrinsic tyrosine kinase activity (White & Kahn 1994, Saltiel & Kahn 2001). In particular, the phosphorylation of three tyrosine residues in the kinase domain results in maximum tyrosine kinase activity (White & Kahn 1994, Saltiel & Kahn 2001), and enables the subsequent phosphorylation of several endogenous proteins. These events are essential for further signal transduction and consequently for insulin action (Saltiel & Kahn 2001). In addition to tyrosine, the IR undergoes serine and threonine (Ser/Thr) phosphorylation, which is detected in the basal state and in response to stimulation of cells by phorbol esters (Takayama \textit{et al}. 1984, Bollag \textit{et al}. 1986), cAMP analogues (Stadtmauer & Rosen 1986), and insulin itself (Kasuga \textit{et al}. 1982, Häring \textit{et al}. 1984, Pillay \textit{et al}. 1991). Although the functional significance of this phosphorylation is not completely understood, results from \textit{in vitro} studies indicate that it may attenuate signaling by decreasing insulin-stimulated tyrosine phosphorylation (Takayama \textit{et al}. 1984, Bollag \textit{et al}. 1986, Takayama \textit{et al}. 1988). This mechanism may also regulate insulin action \textit{in vivo}. Insulin resistance of polycystic ovary syndrome, gestational diabetes mellitus, and obesity appears to be associated with excessive Ser/Thr phosphorylation of the IR in skeletal muscle (Dunaif 1997, Itani \textit{et al}. 2000, Shao \textit{et al}. 2000). Increased Ser/Thr phosphorylation was demonstrated to be linked to reduced IR kinase (IRK) activity in liver of fasted rats (Karaski \textit{et al}. 1990) and in skeletal muscle of obese (fa/fa) Zucker rats (Zhou \textit{et al}. 1999). Among the serine kinases that might be involved in IR inhibition, protein kinase C (PKC) is one candidate with potential pathophysiological relevance. Increased PKC activity is associated with impaired insulin-stimulated IRK activity (Takayama \textit{et al}. 1988, Karaski \textit{et al}. 1990, Chin \textit{et al}. 1994, Itani \textit{et al}. 2000). Moreover,
the amount of membrane-associated PKC is increased in livers of obese insulin-resistant Zucker rats and in obese patients with non-insulin dependent diabetes mellitus, suggesting that excessive PKC activity may contribute to some states of insulin resistance (Considine et al. 1995, Qu et al. 1999).

Several serine residues in the IR β-subunit have been identified as phosphorylation sites. According to the numbering system of Ullrich et al. (1985), these include residues 955/56 (Feener et al. 1993, Liu F & Roth 1994a), 1023/25 (Liu F & Roth RA 1994b), 1293/94 (Lewis et al. 1990a, Tavare et al. 1991), 1309 (Al-Hasani et al. 1997), 1315 (Coghlan et al. 1994, Feener et al. 1994) and 1177/78/82 (Bossenmaier et al. 2000). In addition, Thr 1336 of the IR was identified as a major target for phosphorylation by PKC (Lewis et al. 1990b). However, no relationship between the level of phosphorylation of these sites and the activation state of the IRK has been established (Anderson & Olefsky 1991, Tavare et al. 1991, Coghlan et al. 1994, Kellner et al. 1995, Strack et al. 1997).

Insulin receptor Ser 994 has been identified as a potential inhibitory site of the IRK as determined by site-directed mutagenesis (Strack et al. 1997). This site appears to be important for mediating IRK inhibition by PKC (Strack et al. 2000).

Recently, we obtained a polyclonal antibody that specifically recognizes phospho–Ser 994 (Coba et al. 2003). By making use of this site-specific anti-phosphopeptide antibody, in the current work we have assessed the extent of phosphorylation of the IR on Ser 994 in insulin-target tissues from two animal models of insulin resistance: the Zucker fatty rat (fa/fa), a model of genetic obesity with severe hyperinsulinemia characterized by reduced insulin-stimulated IRK activity in liver and muscle (Hurrell et al. 1989, Kasiske et al. 1992, Zhou et al. 1999) and the PEPCK-bGH mouse, a model of growth hormone–induced insulin resistance characterized by increased basal levels of IRK activity in liver and muscle (Valera et al. 1993, Balbis et al. 1996, Dominici et al. 1998). As an indication of the activation status of the IRK, the in vivo level of IR tyrosine phosphorylation was monitored in parallel with the levels of phospho–Ser 994 and phospho–Thr 1336 in these models.

Materials and Methods

Materials

HEPES, Tris, phenylmethylsulfonyl fluoride (PMSF), aprotinin, leupeptin, ATP, Triton X-100, Tween 20, phosphatase acid Type IV– from potato, Protein A-Sepharose 6 MB, porcine insulin, bovine serum albumin, nitrocellulose membranes, and wheat germ agglutinin (WGA)-Sepharose were from Sigma Chemical Co. Affigel-10 and the reagents and apparatus for SDS-PAGE and immunoblotting were obtained from Bio-Rad. 125I–protein A was purchased from ICN Biomedicals.

Antibodies

The monoclonal anti-phosphotyrosine (anti–PY, PY99) and the polyclonal anti-insulin receptor β-subunit antibody (anti–IR, C–19) were purchased from Santa Cruz Biotechnology, Inc. Rabbit polyclonal antiserum raised against synthetic phosphopeptides containing Thr 1336 of the IR (anti–PT1336) was a generous gift of Dr MP Coghlan and was prepared as previously described (Coghlan et al. 1994). The polyclonal antibody against phospho–Ser 994 of the IR (anti–PS994) was obtained by immunization of the rabbits with the phosphorylated peptide LRELGQGSFGMVYE, as previously described (Coba et al. 2003).

Animals

Male obese Zucker rats (fa/fa) and their age–matched lean controls (+/?) were purchased from Charles River Laboratories and were used at the age of 4–5 months. Male transgenic mice overexpressing bovine growth hormone (bGH) were originally produced by a microinjection of the bGH gene fused to the control sequences of the rat phosphoenolpyruvate carboxykinase gene into the pronuclei of fertilized mouse eggs (Valera et al. 1993). The hemizygous transgenic mice used in the present study were derived from a founder male with ~25 copies of the hybrid gene and were produced by mating transgenic males with normal C57BL/6 X C3H F1 hybrid females. These matings produced approximately equal numbers of transgenic mice and normal animals that were used as controls. Transgenic mice had markedly accelerated postweaning growth, leading to a significant increase in body weight at the age of 3–5 months when these studies were conducted.

All animals were housed three to five per cage in a room with controlled light (12:12–h light–dark cycle) and temperature (22 ± 2 °C) with free access to water and standard chow. All animal studies were performed following the ‘Principles of laboratory animal care’ (NIH publication no. 85–23, revised 1985).

Measurement of glucose, insulin and triglycerides concentrations

Fasting serum insulin concentration was determined using a solid phase radioimmunoassay kit from Diagnostic Products Inc. Measurements were performed in duplicate and variations between duplicate samples of less than 5% were considered acceptable. The sensitivity of the assays was 3 µIU/ml. Serum glucose was measured with...
the glucose oxidase procedure (Trender; Sigma). Serum triglycerides were estimated using a colorimetric assay [triglyceride (INT), procedure 336; Sigma).

**Experimental design and preparation of tissue extracts**

Animals were fasted overnight and 15 min before the experiment they were killed by the intraperitoneal administration of 100 mg of sodium pentobarbital per kg body weight. After anesthesia was induced, the portal vein was exposed and 10 IU insulin per kg body weight in normal saline (0·9% NaCl) were injected in a final volume of 0·1 ml. Additional animals were injected with diluent to obtain data under basal conditions. The liver and the rectus abdominus muscle were removed 50 s and 2 min after injection respectively, frozen under dry ice and kept at −70 °C until further use. Tissues were minced coarsely and homogenized with a polytron homogenizer in 10 volumes of ice-cold lysis buffer containing 1% Triton X-100, 50 mM HEPES (pH 7·4), 100 mM Na3P2O7, 100 mM NaF, 10 mM EDTA, 10 mM Na3VO4, 2 mM PMSF and 0·5% aprotinin. Tissue extracts were centrifuged at 100 000 × g for 1 h at 4 °C to remove insoluble material; protein concentration was measured using the Bradford method (Bradford 1976).

**Immunoprecipitation**

Supernatants containing equal amounts of protein (8–16 mg) were incubated overnight at 4 °C with anti-IR (4 µg/ml). Immune complexes were collected by incubation with protein A-Sepharose 6 MB for 2 h at 4 °C, washed three times with 50 mM Tris buffer (pH 7·4) containing 1% Triton X-100 and 1 mM Na3VO4, boiled in Laemmli sample buffer, and stored at −70 °C until needed for electrophoresis.

**Immunoblotting**

Immunoprecipitated proteins were resolved by SDS-polyacrylamide gel electrophoresis (7·5% acrylamide) in a Bio-Rad miniature slab gel apparatus. Electrophoretic transfer of proteins from the gel to nitrocellulose membranes was performed for 1 h at 100 V (constant) using the Bio-Rad miniature transfer apparatus in 25 mM Tris, 192 mM glycine and 20% (v/v) methanol. To reduce non-specific binding, the nitrocellulose membranes were incubated for 2 h with T-TBS buffer (50 mM Tris–HCl, pH 7·6, 150 mM NaCl, and 0·02% Tween 20) containing 3% BSA. Blots were then incubated overnight with anti-PY (1 µg/ml) to detect tyrosine phosphorylation of IR, with anti-PT1336 (1/1000) to detect IR phosphorylation on threonine 1336 or with anti-PS994 (1/250) to detect IR phosphorylation on serine 994. The membranes were subjected to four 5 min washes in T-TBS buffer and were then incubated with 125I-protein A in T-TBS containing 3% BSA for 1 h at room temperature and then washed again for 60 min as described above. 125I-protein A bound to antibodies was detected by autoradiography and band intensities were quantified by densitometry.

**Insulin receptor purification**

Preparation of a microsomal fraction containing liver plasma membranes was performed using a previously described procedure (Dominici et al. 1998) with slight modifications. The preparation buffer was 50 mM Tris–HCl (pH 7·6), 0·25 M sucrose, 1 mM PMSF and 0·5% aprotinin, plus 30 mM Na4P2O7, 1 mM Na3VO4 and 10 mM NaF as phosphatase inhibitors. The resulting membrane pellet was resuspended in solubilization buffer (50 mM Tris–HCl pH 7·6, 1% Triton X-100, 1 mM PMSF, 0·5% aprotinin, 30 mM Na4P2O7, 1 mM Na3VO4 and 10 mM NaF), incubated with gentle agitation for 1 h at 4 °C, and centrifuged for 90 min at 100 000 × g. The supernatant (solubilized receptor) was loaded onto a column containing WGA–Sepharose and mixed end over end for 16 h at 4 °C for affinity chromatography purification. The column was washed with 30 volumes of buffer (50 mM HEPES, pH 7·6, 0·15 M NaCl, 0·1% Triton X-100, 30 mM Na4P2O7, 1 mM Na3VO4 and 10 mM NaF). Bound receptors were eluted with buffer containing 50 mM HEPES, (pH 7·6), 0·15 M NaCl, 0·1% Triton X-100 and 0·3 M N-acetyl glucosamine. The phosphatase inhibitors were omitted in the preparation when the purification of the IR was to be followed by treatment with acid phosphatase.

**Treatment of purified IRs with acid phosphatase**

Acid phosphatase type IV from potato (8 mg) in 50 mM HEPES buffer, (pH 6·0), was incubated overnight at 4 °C with 2 ml of Affi-gel 10 (Bio-Rad). The protein–coupled resin was then washed with 100 ml ice cold water, incubated for 90 min at 4 °C with 1 M ethanolamine (pH 8·0) to block unreacted amino groups and then washed and equilibrated with 50 mM HEPES (pH 5·0). Affinity-purified IRs from liver of obese Zucker (fa/fa) rats were incubated for 4 h at 25 °C with 500 units of the immobilized enzyme in 50 mM HEPES, (pH 5·0). Dephosphorylated samples were centrifuged, and 500 µl of supernatants were immunoprecipitated with anti-IR in 50 mM HEPES buffer, (pH 7·6), 1% Triton X-100, 2 mM PMSF and 1 µg/ml aprotinin. Control samples were prepared by treating IRs that had been isolated in conditions that preserved basal phosphorylation, with immobilized acid phosphatase that had been formerly inactivated by boiling for 1 h. The resulting samples were then subjected to immunoprecipitation with anti-IR in buffer 50 mM HEPES, (pH 7·6), 1% Triton X-100,
100 mM Na\textsubscript{4}P\textsubscript{2}O\textsubscript{7}, 100 mM NaF, 10 mM EDTA, 10 mM Na\textsubscript{3}VO\textsubscript{4}, 2 mM PMSF and 10 µg/ml aprotinin. Immune complexes were collected with protein A-Sepharose 6 MB, eluted by boiling with Laemmli buffer for 5 min and kept at −70 °C until electrophoresis.

In vitro phosphorylation of purified IRs by PKC isoforms
IRs isolated from liver of lean rats were subjected to dephosphorylation by incubation with immobilized acid phosphatase as described above and then incubated with 0·2 units of different PKC isoforms (α, βI, βII, δ, ε, γ, ζ and η) in independent experiments. The incubation was performed in the absence or presence of 7 µg/ml phosphatidylserine, 2 µM diolein and 1·5 mM CaCl\textsubscript{2} in a reaction mixture containing 1 mM EGTA, 10 mM MgCl\textsubscript{2}, 3 mM MnCl\textsubscript{2}, 15 mM Tris, (pH 7·4) and 50 µM ATP. After 3 h at 22 °C, the reaction was terminated by the addition of 900 µl 50 mM Tris buffer, (pH 7·4), 1% Triton X-100, 2 mM PMSF, 1 µg/ml aprotinin, 10 mM EDTA, 100 mM Na\textsubscript{2}P\textsubscript{2}O\textsubscript{7}, 100 mM NaF and Na\textsubscript{3}VO\textsubscript{4} and the IR was immunoprecipitated by incubation with anti-IR and protein A-Sepharose as described above.

Statistical analysis
Results are presented as means ± s.e.m. Experiments were performed by analyzing all groups of animals in parallel. Data were analyzed with ANOVA followed by the Tukey-Kramer test. Student’s t-test was used when the values of two groups were analyzed. P values less than 0·05 were considered significant.

Results
Animal characteristics
Data describing the metabolic parameters of the two animals model used are presented in Tables 1 and 2. A state of severe insulin resistance was evident in obese Zucker rats. Fasting glucose levels were increased by 2·6-fold in the obese rats (P<0·001), whereas insulin levels were increased by 12·7-fold (P<0·001) relative to lean controls (Table 1). In addition, when compared with the lean control animals, obese rats displayed a large increase in serum triglyceride concentrations (24·1-fold increase; (Table 1). PEPCK-bGH transgenic mice displayed a state of insulin resistance, shown by the concomitant presence of normal fasting glucose levels and substantially elevated plasma insulin concentration (8·5-fold above control values; P<0·001; Table 2).
Figure 1 Phosphorylation status of the IR in liver of lean and obese Zucker rats. Animals were injected with saline (-) or 10 IU insulin (+) per kg/body weight. Equal amounts of solubilized liver proteins obtained as described in Materials and Methods were immunoprecipitated with an anti-IR antibody (anti-IR), separated by SDS-PAGE and immunoblotted either with an anti-phosphotyrosine antibody (anti-PY) (A, upper panel), with an antibody that recognizes the IR phosphorylated on Ser 994 (anti-PS994) (C, upper panel) or with an antibody that recognizes the IR phosphorylated on Thr1336 (anti-PT1336) (E, upper panel). Membranes were stripped and reblotted with anti-IR (A, C, and E, lower panels); IRβ, IR β-subunit. B, D and F, data quantification by densitometric analysis. Values are expressed as relative to control assigning a value of 1 to the mean of saline-injected normal rats. Each bar represents mean ± S.E.M. of the indicated number of independent experiments. * P<0.05 vs the corresponding insulin-stimulated lean value. ** P<0.001 vs saline-injected lean.
while in skeletal muscle of obese rats insulin-stimulated IR tyrosine phosphorylation was reduced by 56% (n=5; P<0.05) relative to the insulin-stimulated lean value (Fig. 2A and B). The extent of phosphorylation of the IR on Ser 994 was analyzed in rectus abdominus muscle from obese and lean rats. Basal IR Ser 994 phosphorylation could not be detected in any of the samples analyzed (n=8; Fig. 2C). In an attempt to increase the sensitivity of the
assay, the amount of protein used for the immuno-
precipitation step was increased from 8 to 16 mg). In
addition, a different detection method was tried. Enhanced
chemiluminescence was used instead of 125I-protein A. In
spite of these modification, Ser 994 phosphorylation was
not detected in skeletal muscle (data not shown).

IR levels in muscle were similar in lean and obese rats
(n=8; Fig 2C). Similar results were found in samples from
tensor fasia latae and gastrocnemius muscle obtained from
both groups of animals (data not shown). The extent of
IR Thr 1336 phosphorylation in skeletal muscle under
basal conditions was very low in both groups of animals
(Fig. 2D and E). In skeletal muscle of lean rats, insulin
increased IR Thr 1336 phosphorylation by 4.9-fold, while
in skeletal muscle from obese rats insulin increased the
phosphorylation of this site by only 2.3-fold (n=3;
P<0.05; Fig. 2D and E).

IR Tyr Phosphorylation in Liver of Bovine Growth Hormone Transgenic Mice
Basal Tyr phosphorylation of the IR increased 4.5-fold
in liver of PEPCK-bGH transgenic mice (n=3; P<0.05),
relative to control mice (Fig. 3A and B). Insulin increased
IR Tyr phosphorylation by approximately 7-fold in liver of
normal mice (Fig. 3A and B). In liver of PEPCK-bGH transgenic mice, insulin-stimulated levels of
IR Tyr phosphorylation were similar to those
detected in insulin-treated control animals. (Fig. 3A and
B). A very low anti-PS994 immunoreactivity was detected
in IRs isolated from liver of PEPCK-bGH transgenic mice
or the corresponding normal control mice with (n=3;
Fig. 3C). Quantification by densitometric analysis of
the signal obtained was not possible. In vivo insulin
administration had no detectable effect on IR Ser 994
phosphorylation in liver from any of the mice analyzed
(Fig. 3C). In contrast to the large increase in basal IR
Tyr phosphorylation exhibited by transgenic mice, basal IR
Thr 1336 phosphorylation levels in liver of PEPCK-bGH mice were low and similar to those
detected in liver of normal control mice (n=3; Fig. 3D
and E). In liver of both groups of animals, insulin increased
IR Thr 1336 phosphorylation to a similar extent (Fig. 3D
and E).

Characterization of the Phospho-Ser 994 Antibody
IRs purified from livers of obese Zucker rats were
dephosphorylated by treatment with immobilized acid
phosphatase and subjected to immunoblot analysis with
anti-PS994. This treatment completely abolished the
immunoreactivity of the anti-PS994 antibody (Fig. 4,
lane 1), while a strong signal was detected in control IR
samples that had been treated with inactive immobilized
acid phosphatase (Fig. 4, lane 2). Additional confirmation of
the specificity of the anti-PS994 antibody was obtained
when aliquots of extracts prepared from liver of obese
Zucker rats were immunoprecipitated with anti-IR and
immunoblotted with anti-PS994 in the presence of the
phosphorylated peptide used as immunogen. The presence
of the phosphorylated peptide containing PS994 in the
incubation media prevented the immunoblotting of
anti-PS994 (Fig. 4, lanes 3 and 4). In contrast, when
the incubation with anti-PS994 was performed in the
presence of either the same peptide sequence unphos-
phorylated or with a serine-phosphorylated non-
related peptide sequence (VKTVNE(PS)ASLRERI) the
immunoreactivity of anti-PS994 towards IR isolated from
liver of obese Zucker rats remains unaltered (Fig. 4, lanes
5-8).

Phosphorylation of IR Ser 994 by PKC Isoforms
To determine if Ser 994 of the IR is a target for
phosphorylation by PKC, WGA-purified IRs from lean
Zucker rats liver were subjected to dephosphorylation
with immobilized acid phosphatase and incubated with
PKC isoforms belonging to the three subclasses of PKC.
Phospho-Ser 994 levels were then analyzed by subjecting
the samples to immunoprecipitation with anti-IR fol-
lowed by immunoblotting with anti-PS994. Out of eight
PKC isoforms tested, only three were able to promote
phosphorylation of the IR on Ser 994. They included
PKCa and PKCβII (Fig. 5, left and centre) belonging to
the classical or conventional class and PKCζ from the
atypical class (Fig. 5, right). Incubation of IRs with PKC
isoforms belonging to the novel group (δ, ε and η) failed
induce phosphorylation of Ser 994 as detected by
immunoblotting with anti-PS994 (Fig. 5, right). The
highest level of IR phosphorylation on Ser 994 was attained after incubation with PKCζ (Fig. 5).
To confirm
that the negative result obtained by incubation with isoforms BI, δ, ε, γ and η was the consequence of a lack
of induction of phosphorylation of the IR on Ser 994
instead of a defective activation of these enzymes, their
activity was evaluated by incubation with a specific
peptide substrate. All the isoforms of PKC were able to
phosphorylate the substrate used (data not shown).

Discussion
Among the factors affecting the insulin-signaling pathway,
phosphorylation of the IR and the IR substrates (IRSs) on
serine residues, emerges as a potential cause of attenuation
of the IRK leading to decreased insulin-stimulated tyro-
sine phosphorylation (Saltiel & Kahn 2001). Although not
completely characterized, this inhibitory phosphorylation
may provide a negative feedback to insulin signaling and
serve as a mechanism for crosstalk from other pathways
(Saltiel & Kahn 2001). Among the serine kinases impli-
cated in this process, PKC is one candidate with potential
**Figure 3** Phosphorylation status of the IR in liver of normal and PEPCK-bGH transgenic mice. Animals were injected with saline (-) or 10 IU insulin (+) per kg/body weight. Equal amounts of solubilized liver proteins obtained as described in Materials and Methods were immunoprecipitated with an anti-IR antibody (anti-IR), separated by SDS-PAGE and immunoblotted either with an anti-phosphotyrosine antibody (anti-PY) (A, upper panel), with anti-PS994 (C, upper panel) or with anti-PT1336 (D, upper panel). Membranes were stripped and reblotted with anti-IR (A, C, and D, lower panels); IRβ, IR β-subunit. B and E, data quantification by densitometric analysis. Values are expressed as relative to control assigning a value of 1 to the mean of saline-injected normal mice. Each bar represents mean ± S.E.M. of the indicated number of independent experiments. * P<0.05 vs saline-injected normal mice.
This blot is representative of four different experiments. SDS-PAGE and immunoblotted with anti-PS994; IR samples were immunoprecipitated with anti-IR, separated by prepared by incubation in the absence of PKC (-). The resulting described in Materials and Methods. Control samples were phosphatase and incubated with different PKC isoforms as WGA-purified IRs from lean Zucker rats livers were (lanes 7 and 8); IR serine-phosphorylated peptide sequence. (VKTVNE(SP)ASLRERI), unphosphorylated (lanes 5 and 6) or with a non-related they were raised against (samples were run in duplicate and are presented in lanes 3 and 4) or the same peptide sequence unphosphorylated (lanes 5 and 6) or with a non-related serine-phosphorylated peptide sequence. (VKTVNE(SP)ASLRERI), (lanes 7 and 8); IRβ, IR β-subunit.

Figure 4 Characterization of the specificity of the PS994 antibody. WGA-purified IRs from obese Zucker rats liver were treated with either active immobilized acid phosphatase (lane 1) or with the immobilized enzyme that had been previously inactivated by boiling for 1 h (lane 2). After treatment, samples were immunoprecipitated with anti-IR and immunoblotted with anti-PS994. Solubilized extracts from livers of obese Zucker rats were subjected to SDS-PAGE and immunoblotted with anti-PS994 in the presence (50 μg/ml) of the phosphorylated peptide which they were raised against (samples were run in duplicate and are presented in lanes 3 and 4) or the same peptide sequence unphosphorylated (lanes 5 and 6) or with a non-related serine-phosphorylated peptide sequence. (VKTVNE(SP)ASLRERI), (lanes 7 and 8); IRβ, IR β-subunit.

Pathophysiological relevance, since numerous studies have linked excessive PKC activity to diminished insulin sensitivity, especially to that occurring together with increased lipid availability (Considine et al. 1995, Qu et al. 1999, Itani et al. 2000). Such conditions can increase the amount of lipid-signaling molecules derived from free fatty acids that can activate isoforms of the PKC family, such as diacylglycerol or ceramide (Schnittz-Peffer 2000).

The features of the obese Zucker (fa/fa) rat include insulin resistance, hyperinsulinemia, glucose intolerance, dyslipidemia and impaired insulin signaling in liver and skeletal muscle (Hurrell et al. 1989, Kasiske et al. 1992, Zhou et al. 1999), together with increased PKC activity in insulin target tissues (Considine et al. 1995, Qu et al. 1999), suggesting that this could be a suitable model to analyze the relationship between IR serine phosphorylation and impaired insulin signaling.

In the present study we have analyzed the status of phosphorylation of Ser 994, a putative inhibitory site of the IR, in liver and skeletal muscle of obese Zucker rats. Phosphorylation of Ser 994 was determined using a recently generated anti-phosphopeptide antibody that specifically recognizes IR phosphorylated at this site (Coba et al. 2003). In liver of both lean rats and normal mice, basal IR Ser 994 phosphorylation levels were very low and did not change significantly after in vivo insulin administration, suggesting that phosphorylation at this site may not be modulated by insulin. In contrast, basal phosphorylation levels of IR Ser 994 were found to be markedly elevated in liver of obese Zucker rats. Previous work from Strack et al. demonstrated that insulin-stimulated autophosphorylation was increased in mutant human IRSs in which serine was exchanged to alanine at position 994, suggesting that this residue plays an inhibitory role on the regulation of the IR kinase (Strack et al. 1997). Based on that result we postulate that excessive phosphorylation of the IR on Ser 994 could have a role in the impairment of the IRK displayed by obese Zucker rats (Karasik et al. 1990). Insulin administration resulted in a significant decrease in the phosphorylation levels of IR Ser 994 in liver of obese rats suggesting that activation of the IRK by insulin in vivo might be associated with or might require dephosphorylation of this site.

Phosphorylation levels of Ser 994 were also analyzed in liver of PEPCK-bGH mice, a model of insulin resistance with increased basal IR tyrosine kinase activity (Dominici et al. 1998). In contrast to results obtained in Zucker rats, phosphorylation levels of the IR at Ser 994 were almost not detectable in liver of PEPCK-bGH mice and did not differ from those found in control animals. In a previous study, IRs were purified from liver of PEPCK-bGH mice and treated with immobilized alkaline phosphatase to remove phosphate groups from Ser/Thr residues (Dominici et al. 1998). The in vitro autophosphorylation of the IR was unchanged after this treatment, suggesting that Ser/Thr phosphorylation of the IR β-subunit was apparently not involved in the regulation of the IRK in these mice (Dominici et al. 1998). Consequently, the current results agree with our previous report in PEPCK-bGH mice (Dominici et al. 1998).

Insulin-induced phosphorylation of the IR at Thr 1336 has been previously demonstrated in cells in culture (Tavare et al. 1991). However, this is the first report to show that IR Thr 1336 is phosphorylated in vivo in an insulin target tissue and that exogenous insulin administration results in an increase of its phosphorylation level. The role of the insulin-stimulated phosphorylation of the IR at Thr 1336 is not clear. Previous in vitro studies demonstrated that this residue of the IR does not participate in the activation or regulation of the IRK (Anderson & Olefsky 1991, Tavare et al. 1991, Coghlan et al. 1994, Kellner et al. 1995). Possibly, this site could have a regulatory role in the interaction between the IR and intracellular substrates. In liver of obese Zucker rats the phosphorylation level of the IR on Thr 1336 did not correspond with that of Ser 994 and follows the same pattern of phosphorylation observed to tyrosine.

Figure 5 Phosphorylation of IR-Ser 994 by various PKC isoforms. WGA-purified IRs from lean Zucker rats livers were dephosphorylated by incubation with immobilized acid phosphatase and incubated with different PKC isoforms as described in Materials and Methods. Control samples were prepared by incubation in the absence of PKC (-). The resulting samples were immunoprecipitated with anti-IR, separated by SDS-PAGE and immunoblotted with anti-PS994; IRβ, IR β-subunit. This blot is representative of four different experiments.

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phosphorylation, suggesting that the mechanism of impairment of the IRK in liver of obese Zucker rats does not involve alterations in the phosphorylation of the IR at Thr 1336. Strikingly, the concordance between the phosphorylation of the IR on tyrosine residues and on Thr 1336 that was observed in liver of obese Zucker rats was not observed in liver of PEPCK-bGH mice, where basal IR tyrosine phosphorylation levels were high but those of Thr 1336 were low and similar to those detected in liver of control mice. This result suggests that phosphorylation of the IR at Thr residues could be differentially regulated from phosphorylation at tyrosine residues.

During the last years the involvement of IRSs Ser/Thr phosphorylation in the development of insulin resistance has received increased experimental support. A relationship between a decrease in the insulin-mediated signaling pathway and an increase in the phosphorylation of the IRSs on Ser residues was consistently reported (Sykotis et al. 2001, Gao et al. 2002, Qiao et al. 2002). Several agents including phorbol esters and free fatty acids are able to induce this phosphorylation (Johnston et al. 2003). Importantly, hyperinsulinemia and TNF-α, two inducers of insulin resistance trigger increased Ser/Thr phosphorylation of IRS-1 and IRS-2 (Johnston et al. 2003).

Tumor necrosis factor α activates PKCζ and its downstream effector IkappaB kinase beta (IKKβ) (Lallena et al. 1999) a potent inducer of insulin resistance (Kim et al. 2001, Yuan et al. 2001). Protein kinase Cζ activation leads to a negative control of the insulin-signaling pathway mediated by Ser/Thr phosphorylation of IRS-1 (Liu et al. 2001). Moreover, activation of IKKβ by TNF-α attenuates insulin signaling and inhibits insulin-mediated IRS-1 tyrosine phosphorylation, whereas IKKβ inhibition by high doses of salicylates prevents Ser/Thr phosphorylation of the IRS proteins (Yuan et al. 2001).

The same inhibitory mechanism that controls the insulin-signaling pathway at the level of the IRS proteins could be involved in the regulation of the IRK. The effects on IRS proteins mediated by salicylates are in part secondary to enhanced IRK activity (Yuan et al. 2001), suggesting that the IKKβ pathway can negatively regulate both the activity of the IR and the engagement of the IRS proteins in productive insulin signaling. Though there is no direct evidence of a correlation between decreased Ser/Thr phosphorylation of the IRSs and enhanced IRK, the change in the levels of IR Ser/Thr phosphorylation could have a role in the regulation of the IRK.

Among the serine kinases that might be involved in the phosphorylation of the IR on serine residues, PKC is one candidate with potential pathophysiological relevance. In PKC-mediated insulin-resistant states such as obesity or conditions of elevated lipid availability, both a tissue and isoform-selective PKC activation has been detected (Avignon et al. 1996, Griffin et al. 1999, Cortright et al. 2000). PKC activity is increased in tissues from obese Zucker rats. In liver this increase appears to be the result of an increase in the membrane content of PKC isoforms α, β, ε and ζ, (Considine et al. 1995, Qu et al. 1999), while in skeletal muscle this increase has been ascribed to an elevation in the membrane content of PKC isoforms θ (Cortright et al. 2000) and ε (Avignon et al. 1996). In the current study, we have demonstrated that IR Ser 994 is an in vitro phosphorylation target for PKC. Moreover, by incubating purified IRs with different PKC isoforms, we analyzed the PKC isoform-specificity towards IR Ser 994. Of all the isoforms analyzed, the highest phosphorylation level was attained after incubation with PKCζ, PKCα and PKC βI were also able to promote phosphorylation at this site but to a lesser extent. In contrast, none of the novel PKC tested (δ, ε, η) were able to promote phosphorylation of the IR at Ser 994. This tissue and isoform-selective PKC activation might explain the different pattern of phosphorylation observed on Ser 994 in liver and skeletal muscle of obese Zucker rats. Moreover, this suggests that, in muscle of obese Zucker rats, other IR residues could be implicated in the regulation of the IRK.

However, the participation of other serine kinases beside PKC, such as protein kinase A, cannot be ruled out since the latter has been shown to participate in the decreased autophosphorylation that is exhibited by IR from fibroblasts of patients with PCOS (Li et al. 2002). In addition, the Ser/Thr kinases IKKβ and JNK are implicated in the phosphorylation of the IRS proteins and they could also be involved in the phosphorylation of the IR on Ser/Thr residues (Johnston et al. 2003).

In conclusion, we have demonstrated that Ser 994 and Thr 1336 are in vivo phosphorylation sites of the IR. In addition we have detected a large increase in the phosphorylation of IR Ser 994 in liver of obese Zucker rats under basal conditions that could be involved in the attenuation of the IR kinase displayed by these animals.

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