

Hepatic Protein Phosphotyrosine Phosphatase

Dephosphorylation of Insulin and Epidermal Growth Factor Receptors in Normal and Alloxan Diabetic Rats

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Abstract

Polypeptide hormone signal transmission by receptor tyrosine kinases requires the rapid reversal of tyrosine phosphorylation by protein phosphotyrosine phosphatases (PPTases). We studied hepatic PPTases in the rat with emphasis on acute and chronic regulation by insulin. PPTase activity with artificial substrates ($[^{32}\text{P}]\text{Tyr}$ -reduced, carboxyamidomethylated, and maleylated lysozyme and $[^{32}\text{P}]\text{Tyr}$ -poly[glutamic acid:tyrosine] 4:1) was present in distinct membrane, cytoskeletal, and cytosolic fractions. These PPTase activities were unaffected by alloxan diabetes. Acute administration of insulin to normal animals also did not change PPTase activity in liver plasma membranes or endosomal membranes. Although alloxan diabetes did not affect PPTase activity measured with artificial substrates or with epidermal growth factor receptors, a decrease in insulin receptor dephosphorylation was noted. Dephosphorylation of hepatic receptors from normal and diabetic rats by membrane PPTase from control rats was similar. These results indicate that alloxan diabetes does not lead to a generalized effect on hepatic PPTase activity, although a substrate-specific decrease in activity with the insulin receptor may occur. (*J. Clin. Invest.* 1990. 85:1754–1760.) protein phosphotyrosine phosphatases • hormone receptors • diabetes

Introduction

The binding of epidermal growth factor (EGF)¹ (1) and insulin (2) to their respective receptors stimulates tyrosine kinase activity intrinsic to the receptors. This fact has led to the hypothesis that signal transmission for these and a number of other polypeptide hormones involves the phosphorylation of proteins on tyrosine. Thus far, receptor tyrosine kinases themselves represent the best characterized substrates for hormone-stimulated tyrosine phosphorylation (for reviews, see references 3 and 4), although a number of other potential substrates

have been identified. Regardless of the nature of the substrates for tyrosine phosphorylation, assignment of a regulatory function to this process requires its reversal by protein phosphatases. Several such phosphatases have been purified from diverse sources, including rabbit kidney (5), human placenta (6, 7), and rat (8) and bovine (9) brain.

Discovery of the insulin-activated receptor tyrosine kinase led a number of investigators to study alterations in this activity in clinical and experimental states in which sensitivity to insulin is diminished. Type I and type II diabetes mellitus as well as obesity in nondiabetic subjects are known to be associated with insulin resistance (10–12). Decreased insulin receptor tyrosine kinase activity has been reported in adipocytes (13, 14), erythrocytes (15), skeletal muscle (16) and liver (17) from non-insulin-dependent diabetic subjects, and adipocytes from nondiabetic, obese individuals (14). Correlative animal studies exist (18–20), although at least two studies have failed to demonstrate decreased hepatic insulin receptor kinase activity in experimental diabetes in rats (21, 22). Given the reciprocal action of tyrosine kinases and phosphatases, we hypothesized that an increase in protein phosphotyrosine phosphatase (PPTase) activity might contribute to the insulin resistance associated with decreased receptor-mediated tyrosine phosphorylation. Thus, we have undertaken the characterization of hepatic PPTase activity in the rat, with emphasis on the effects of insulin.

Methods

Animal studies. The acute effects of insulin on hepatic PPTase activities were studied by injecting female Sprague-Dawley rats (150–200 g; Charles River Ltd., St. Constant, Quebec, Canada) with 150 μg insulin/100 g body weight via the jugular vein after an overnight fast. Rats were killed before and 30 s, 2 min, and 10 min after injection. Plasma membrane and endosomal fractions were prepared from liver using sucrose density centrifugation (23).

Male Sprague-Dawley rats (Charles River Breeding Laboratories, Inc., Wilmington, MA) weighing 150–175 g were made diabetic by the intravenous administration of alloxan, 65 mg/kg, after an overnight fast. Animals were killed 4 d after alloxan administration by decapitation. Severity of diabetes was ascertained by measurements of plasma glucose (model 23A glucose analyzer; Yellow Springs Instrument Co., Yellow Springs, OH), immunoreactive insulin (using rat insulin standard; Amersham Corp., Arlington Heights, IL), and beta-hydroxybutyrate (24). A group of alloxan diabetic animals were treated with insulin (2 U recombinant human insulin [Humulin; Eli Lilly and Co., Indianapolis, IN] i.p. every 12 h) beginning 3 d after alloxan injection for 48 h before killing. Since alloxan diabetes was associated with a decreased rate of weight gain, three control animals were fasted for 48 h (ad lib water intake) before killing.

Unfractionated membrane preparations were made from individual, fresh livers by the method of Williams et al. (25) and diluted with 50 mM Hepes, pH 7.5, to a concentration of 10 mg protein/ml. Protein

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1. Abbreviations used in this paper: EGF, epidermal growth factor; polyEY, poly[glutamic acid:tyrosine] 4:1; PPTase, protein phosphotyrosine phosphatase; RCAM-lysozyme, reduced, carboxyamidomethylated and maleylated lysozyme.

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determinations were by the bicinchoninic acid method (BCA; Pierce Chemical Co., Rockford, IL) using BSA as standard.

The supernatants from membrane preparations were retained for measurement of soluble cytosolic activity. Detergent extracts of membrane preparations, referred to as the membrane fraction, were made by diluting 1 vol of the unfract ionated membrane suspension with 1 vol 50 mM Hepes, pH 7.4, and 2% (wt/vol) Triton X-100. Insoluble material was collected by centrifugation at 40,000 *g* for 20 min and extracted with 0.5 vol 25 mM Hepes, pH 7.4, and 0.6 M KCl. This second extract, clarified by centrifugation at 40,000 *g* for 20 min, is referred to as the cytoskeletal fraction. Alloxan diabetes was not associated with altered membrane, cytoskeletal, or cytosolic protein contents.

PPTase assays. PPTase activity was measured as the release of ³²P from substrates phosphorylated on tyrosine (5). Assays were carried out in the presence of 1 mM EDTA and 15 mM 2-mercaptoethanol. Reduced, carboxyamidomethylated and maleylated lysozyme (RCAM-lysozyme) and poly[glutamic acid:tyrosine] 4:1 (polyEY) were labeled on tyrosine using partially purified human placental insulin receptor tyrosine kinase as described by Tonks et al. (6). Dephosphorylation of [³²P]Tyr-polyEY and [³²P]Tyr-RCAM-lysozyme was measured at a final substrate concentration of 1 μM. PPTase activities are reported as nanomoles or picomoles of ³²P released per minute.

Receptor preparation, binding, and autophosphorylation. Rat liver insulin and EGF receptors were prepared for use as phosphatase substrates as follows. Rat liver membranes were extracted with 1% Triton X-100. The extract (32–91 mg protein in 2.5–4.0 ml) was applied to a 10-ml column of wheat germ lectin-Sepharose (Pharmacia Fine Chemicals, Piscataway, NJ) and unbound proteins were applied a second time. The column was washed with 250 ml of 40 mM imidazole-HCl, 0.5 M NaCl, 10% glycerol, and 0.05% Triton X-100, pH 7.2. Receptors were then eluted with the same buffer containing 0.3 M *N*-acetyl glucosamine. Fractions that bound ¹²⁵I-insulin were pooled. This same pool contained EGF receptors.

For determination of the specific binding of insulin in lectin-purified preparations and in solubilized plasma membrane and endosomal fractions, 50 μl receptor was incubated with ¹²⁵I-human insulin (40,000 cpm, 100 μl; New England Nuclear, Boston, MA) for 1 h at 22°C. Bound insulin was separated from unbound insulin with the addition of 50 μl 2% bovine gamma globulin and 1.5 ml 15% polyethylene glycol followed by centrifugation (26).

Receptor autophosphorylation was accomplished by incubating receptor preparations (40 μl) with hormones (20 μl, 3 × 10⁻⁷ M insulin and EGF, final concentrations) overnight at 4°C. Receptor activation was followed by the addition of 15 μl containing 6.75 μCi [gamma-³²P]ATP, 50 μM ATP, and 25 mM MnCl₂. Incubation at 4°C proceeded for 30 min and was carried out in the absence of phosphatase inhibitors. Receptor phosphorylation was terminated by the addition of 2-mercaptoethanol (15 mM, final concentration) and EDTA (6 mM, final concentration) in 15 μl.

Receptor dephosphorylation was initiated by the addition of 8 μl PPTase. An additional 2-μl volume was reserved for the addition of phosphatase inhibitors or buffer as control. The reaction proceeded at 37°C and was terminated by the addition of 5× gel electrophoresis sample buffer containing 50 mM ATP, 10% dodecyl sulfate, and 0.5 M dithiothreitol, followed by a 10-min incubation in a boiling water bath. Analysis was by electrophoresis in 7.5% polyacrylamide gels followed by exposure of the dried gel to Kodak XAR-5 film at -70°C in the presence of intensifying screens. Radiolabeled bands were quantified with a scanning densitometer (model 300S; Hoefer Scientific Instruments, San Francisco, CA) connected to a Hewlett-Packard integrator (model 3390A). Densitometric scanning was reported as the mean of triplicate determinations.

Results

Characterization of rat liver particulate PPTase activities. Preliminary studies showed that ~ 60–70% of total hepatic

PPTase activity measured with [³²P]Tyr-RCAM-lysozyme was soluble. Most of the particulate activity (80%) could be extracted with 1% Triton X-100. The remaining particulate activity (20%) was soluble in 0.6 M KCl. Further studies showed that extraction with Triton X-100 a second time did not diminish the amount of activity that could be obtained by extraction of the detergent-insoluble proteins with 0.6 M KCl. Furthermore, reversing the order of extraction (KCl followed by detergent) did not significantly alter the yield of activity in the two fractions. Thus, we concluded that particulate PPTase was localized to distinct membrane (soluble in Triton X-100) and cytoskeletal (soluble in 0.6 M KCl) fractions. Initial studies were aimed at characterizing activities in these two fractions as well as the soluble fraction to determine whether these activities should be attributed to more than one enzyme.

Membrane and cytoskeleton PPTase measured with [³²P]Tyr-RCAM-lysozyme displayed neutral pH optima (not shown). They displayed similar sensitivity to the PPTase inhibitors vanadate and zinc (IC₅₀ ~ 250 μM, data not shown). Cytosolic, membrane, and cytoskeleton PPTase measured with [³²P]Tyr-RCAM-lysozyme were similar with regard to inhibition by polyEY, dodecyl sulfate, or heat inactivation at 57°C (Fig. 1).

Because PPTase measurements made with [³²P]Tyr-polyEY consistently resulted in higher specific activities than those done with [³²P]Tyr-RCAM-lysozyme (see below), we investigated the possibility of a second type of PPTase specifically active against [³²P]Tyr-polyEY. Membrane PPTase activities measured with both substrates were indistinguishable based on inhibition by vanadate, denaturation by dodecyl sulfate, and inactivation at 45°C (Fig. 2). Likewise, no differences were observed using cytoskeletal PPTase (not shown). We concluded that these substrates were reactive with one PPTase, or with catalytic proteins with nearly identical properties.

During our initial studies we observed that soluble PPTase was liberated from frozen membrane preparations upon thawing. We therefore performed an experiment to ascertain if soluble PPTase could be released from membranes by proteolysis. Particulate membranes were treated with trypsin (Fig. 3) and the soluble proteins recovered after centrifugation were assayed for PPTase activity. Trypsin

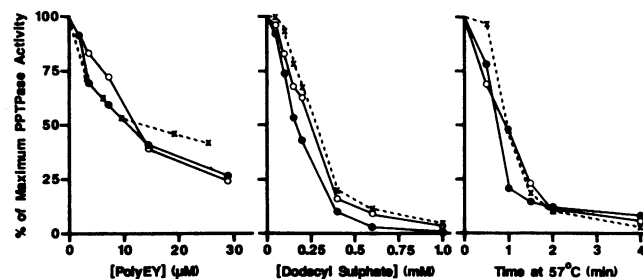


Figure 1. Comparison of particulate and soluble PPTase activities. Activity was measured in membrane (●), cytoskeleton (○), and cytosolic (×) fractions using [³²P]Tyr-RCAM-lysozyme. Heat inactivation at 57°C was measured by adding enzyme to prewarmed tubes. Heating was stopped at the designated times by diluting an aliquot of the enzyme in ice-cold assay buffer (1:50 for the membrane fraction, 1:20 for the cytoskeleton fraction, and 1:10 for the cytosolic fraction).

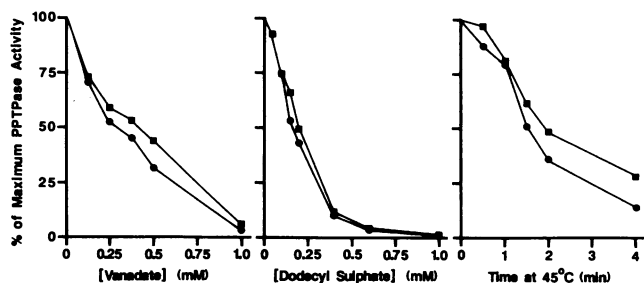


Figure 2. Comparison of membrane PPTase activities measured with [32 P]Tyr-polyEY (●) and [32 P]Tyr-RCAM-lysozyme (■). Heat inactivation at 45°C was determined as in Fig. 1.

treatment of membranes resulted in time-dependent release of a soluble PPTase. Conversely, exposure of soluble PPTase to trypsin led to loss of activity (Fig. 3). The results shown in the figure were replicated in a second, independent experiment.

Acute effects of insulin administration on PPTase activities. The intravenous injection of insulin into rats (Fig. 4) resulted in an increase in insulin binding in hepatic endosomal fractions within 2 min and a concomitant decrease in plasma membrane insulin binding. In contrast, PPTase activity did not change significantly (as determined by analysis of variance) in either plasma membrane or endosomal fractions after insulin administration. Of note, PPTase specific activities were similar in the two membrane fractions.

Effects of alloxan diabetes on PPTase activities. The injection of alloxan resulted in hyperglycemia (597 ± 89 mg/dl [mean \pm SD] vs. 189 ± 29 in control animals) and relative insulinopenia (median, 0.2 ng/ml [range, 0.1–0.9] vs. 2.8 [1.8–5.1] in control animals). Diabetic animals had elevated plasma beta-hydroxybutyrate concentrations (0.69 mM [0.57–4.37]) compared with control animals (0.22 mM [0.18–0.25]); levels were insufficient to produce acidosis. The overall well-being of diabetic animals was confirmed by their ability to gain weight after alloxan administration (7.8 ± 4.0 vs. 11.8 ± 1.2 g/d in control animals). Insulin treatment of alloxan diabetes resulted in restoration of normal plasma glucose (100 ± 17 mg/dl) and beta-hydroxybutyrate (0.22 mM [0.10–0.33]) concentrations.

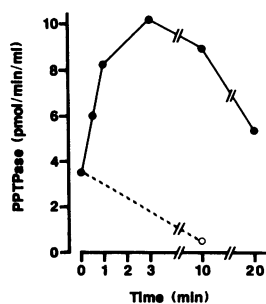


Figure 3. Generation of soluble PPTase activity by treatment of particulate membranes with trypsin. A membrane preparation at 1 mg/ml protein content was incubated with trypsin at a final concentration of 4 μ g/ml. Proteolysis was stopped at the designated times by transferring 80 μ l of the mixture to a tube on ice containing 0.4 μ g lima bean trypsin inhibitor. The membranes were collected by centrifugation in a microfuge for 10 min at 4°C. The

supernatant was assayed directly for PPTase activity using [32 P]Tyr-RCAM-lysozyme (●). Treatment of the membrane supernatant from $t = 0$ with trypsin for 10 min resulted in loss, rather than generation, of activity (○). Particulate membranes recovered after trypsin treatment were also assayed for PPTase. Activity was totally depleted by 5 min.

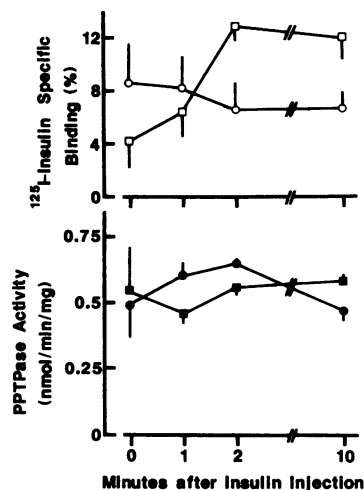


Figure 4. Acute effects of insulin administration on hepatic membrane-associated PPTase activity. Rat liver was obtained at the designated times after the intravenous administration of insulin. Plasma membranes (circles) and endosomal membranes (squares) were prepared for the measurement of insulin binding (open symbols) and PPTase activity (closed symbols). Insulin binding was corrected for protein content to allow comparison between preparations. PPTase activity was measured using [32 P]Tyr-RCAM-lysozyme. Measurements represent mean \pm SE for three experiments.

Alloxan diabetes did not alter soluble, membrane, or cytoskeletal PPTase activity measured with [32 P]Tyr-RCAM-lysozyme or [32 P]Tyr-polyEY (Fig. 5). Significantly higher specific activities were obtained in all fractions using the latter substrate. To study the significance of the decreased rate of weight gain in diabetic animals, we measured PPTase activity in three control rats fasted for 48 h before killing. Fasting did not alter PPTase specific activity measured with both substrates in any of the three liver fractions (Fig. 5).

To further investigate a possible effect of alloxan diabetes on PPTase activities, kinetic analyses were performed using [32 P]Tyr-RCAM-lysozyme. Double-reciprocal analysis of substrate concentration versus rate of hydrolysis (not shown) demonstrated that the K_m for substrate was unchanged in alloxan diabetes for membrane (382 ± 46 nM [mean \pm SD] vs. 423 ± 75 nM in controls) and cytoskeleton (702 ± 358 nM vs. 629 ± 337 nM in controls). In all cases these analyses were linear, indicating reactions with a single substrate affinity.

Studies on PPTase activities with insulin and EGF receptors as substrates. The aforementioned studies were performed

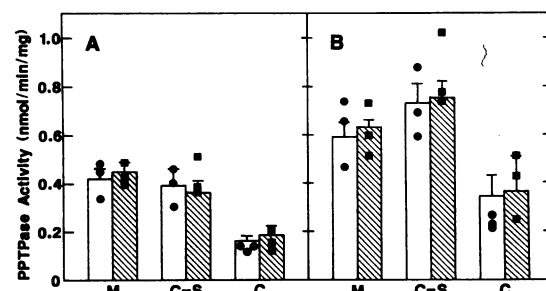


Figure 5. Hepatic PPTase activity in control and alloxan-diabetic animals. Activity was determined in membrane (M), cytoskeleton (C-S), and cytosolic (C) fractions from control animals ($n = 6$, open bars), alloxan-diabetic animals ($n = 6$, hatched bars), insulin-treated alloxan-diabetic animals (circles), and animals fasted for 48 h (squares). PPTase activity was measured using both [32 P]Tyr-lysozyme (A) and [32 P]Tyr-RCAM-polyEY (B). Error bars represent 1 SD.

using artificial substrates for PPTase measurements. Such artificial substrates can be prepared in sufficiently high concentrations to allow determination of reaction rates. Nonetheless, interpretation of such measurements is limited by the nonphysiologic nature of these substrates. We therefore sought to perform correlative studies using autophosphorylated rat liver insulin and EGF receptors as physiologic substrates. Autophosphorylated receptors were identified as proteins with $M_r = 170,000$ (EGF receptor) and 95,000 (beta subunit of the insulin receptor), which were phosphorylated in response to hormones (see Methods). Insulin receptor phosphorylation was stimulated 15 to 20-fold by insulin, whereas EGF receptor phosphorylation was stimulated approximately 5-fold.

Our initial studies demonstrated that PPTase activity against autophosphorylated receptors had characteristics in common with activity measured with artificial substrates. Both had an absolute requirement for reducing agent (15 mM 2-mercaptoethanol); in its absence, no activity could be detected. Both activities were preserved in the presence of EDTA and were fully inhibited by micromolar concentrations of vanadate.

To correlate PPTase activities measured with [32 P]Tyr-RCAM-lysozyme and receptors, hepatic PPTase was subjected to gel filtration chromatography (Fig. 6). A liver homogenate was extracted with 1% Triton X-100. The soluble proteins were concentrated ~ 10 -fold by vacuum dialysis and 1 ml was applied to a 1.5×33 -cm column of Ultrogel AcA 44

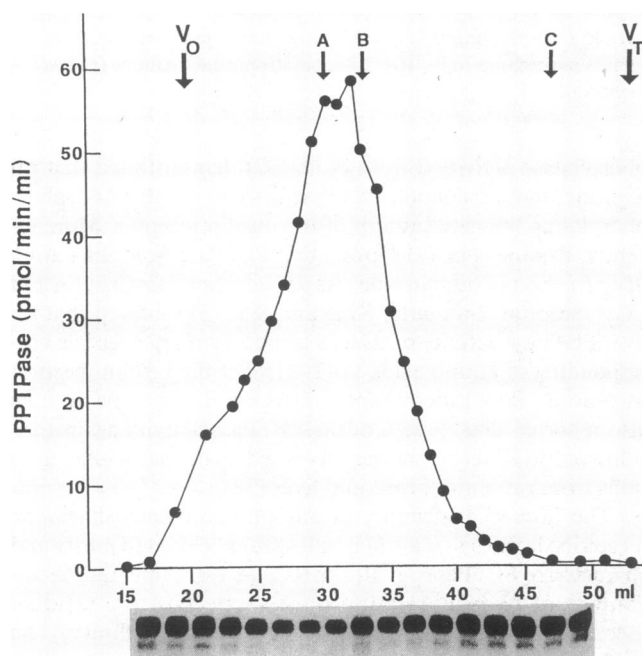


Figure 6. Gel filtration chromatography of membrane-associated PPTase. A concentrated liver extract was prepared and applied to a column of Ultrogel AcA 44 as described in Results. PPTase activity, measured with [32 P]Tyr-RCAM-lysozyme, is shown in the graph. The autoradiogram at the bottom of the figure shows the ability of corresponding fractions to dephosphorylate an autophosphorylated EGF receptor preparation from a normal rat. Arrows at the top of the figure denote the positions of the following: V_0 , void volume; A, BSA ($M_r = 67,000$); B, ovalbumin ($M_r = 43,000$); C, chymotrypsinogen A ($M_r = 25,000$); and V_T , total column volume.

equilibrated in 20 mM Hepes, pH 7.4, 10% glycerol, 50 mM NaCl, and 30 mM 2-mercaptoethanol. PPTase activity measured with [32 P]Tyr-RCAM-lysozyme eluted as a broad peak with maximal activity centered at $M_r \sim 55,000$. The fractions containing PPTase activity with [32 P]Tyr-RCAM-lysozyme were also active against autophosphorylated EGF receptors (Fig. 6). These same fractions had activity with insulin receptors (not shown). No [32 P]Tyr-RCAM-lysozyme, EGF receptor, or insulin receptor dephosphorylation was seen in fractions coincident with the elution of chymotrypsinogen A ($M_r = 25,000$). Other studies have found PPTases of this size with activity against synthetic peptides corresponding to portions of the insulin receptor (see below).

Membrane extracts from control and alloxan diabetic rats contained PPTase activity capable of dephosphorylating hepatic insulin and EGF receptors from normal rats (Fig. 7). Using membrane PPTase from normal animals, the extent of insulin receptor dephosphorylation was consistently greater than EGF receptor dephosphorylation (Fig. 7, *left*). In contrast, membrane PPTase from diabetic animals dephosphorylated both receptors similarly (Fig. 7, *right*). EGF receptor dephosphorylation was similar in the two groups, thereby indicating a decrease in the extent of insulin receptor dephosphorylation with PPTase from diabetic animals. Similar results were obtained in replicate studies using three additional control and diabetic membrane extracts (not shown).

In addition, sensitivity of receptor dephosphorylation to polyEY was studied. As shown in Fig. 7, EGF receptor dephosphorylation in all cases was at least partially inhibited by the addition of 100 μ M polyEY. In contrast, insulin receptor dephosphorylation by normal or diabetic extract PPTase was not sensitive to inhibition by polyEY. Dose response of the inhibition of EGF receptor dephosphorylation (not shown) indicated half maximal inhibition at ~ 10 –30 μ M polyEY.

To complement these studies on alterations in receptor PPTases in alloxan diabetes, we examined the possibility that alloxan diabetes might alter receptor dephosphorylation via effects on the receptors themselves. Studies similar to those shown in Fig. 7 were carried out using PPTase extracted from control membranes to dephosphorylate insulin and EGF receptors from two control and two alloxan diabetic animals. Results (not shown) demonstrated that alloxan diabetes did not alter the susceptibility of EGF or insulin receptors to dephosphorylation by control PPTase. Finally, EGF and insulin receptors from control and alloxan diabetic animals were dephosphorylated similarly by PPTase from a diabetic animal.

Discussion

Until recently the only physiologic substrates for the insulin and EGF receptor kinases that had defined roles were the receptors themselves. Currently, a number of potential substrates for receptor-associated tyrosine kinases have been described, including several proteins without assigned functions (27–31), lipocortin (32), a protein kinase oncogene product (33), and an EGF-activated phospholipase (34). Studies on PPTases have been limited by the availability of physiologic tyrosine kinase substrates. PPTases that can dephosphorylate insulin and/or EGF receptors have been purified from human placenta (7, 35). In these studies, monitoring of activity during

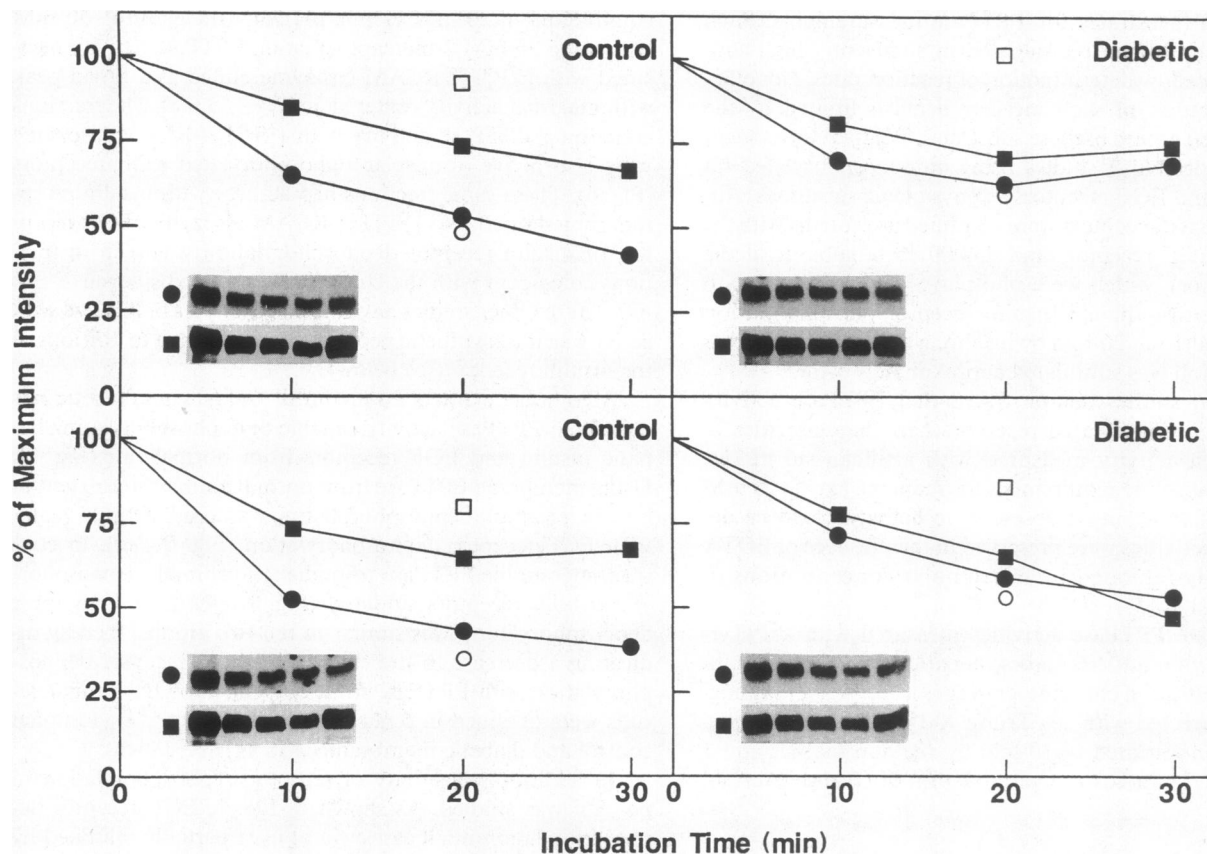


Figure 7. Dephosphorylation of hepatic insulin receptor beta subunit (circles) and EGF receptors (squares) by liver membrane extracts from control and alloxan-diabetic rats. Lanes (from left to right) in the autoradiograms correspond to 0-, 10-, 20-, and 30-min incubations (closed symbols) and 20-min incubation in the presence of 100 μ M polyEY (open symbols).

purification using artificial substrates yielded enzymes active against the physiologic receptor substrates.

In these studies we have attempted to characterize particulate and soluble hepatic PPTases using artificial substrates. Our results are most consistent with a single class of PPTase that dephosphorylates both [32 P]Tyr-RCAM-lysozyme and [32 P]Tyr-polyEY. This activity is characterized by well-recognized functional criteria; i.e., neutral pH optimum and sensitivity to inhibition by zinc, vanadate, and polyEY. Linear double-reciprocal analyses of enzyme kinetics and inhibition/inactivation data are most consistent with a single catalytic moiety.

Hepatic PPTase is distributed in distinct membrane and cytoskeletal fractions as well as the soluble fraction. Our data indicate that the latter may be partly derived from particulate fractions by limited proteolysis. Furthermore, membrane-associated PPTase was distributed in both plasma membrane and endosomal fractions, consistent with accessibility to phosphorylated receptors during the process of ligand-stimulated receptor internalization.

We have concentrated our efforts on the measurement of particulate PPTase activities for several reasons. First is the observation that soluble PPTase activity can be generated by proteolysis of rabbit kidney membranes (36), a finding that we have confirmed in rat liver. Second, the presence of PPTase activity in the membrane and cytoskeleton might result in colocalization of tyrosine kinases, their substrates, and protein

phosphatase activity. Strout et al. (37) demonstrated that rat liver membranes contain PPTase activity capable of dephosphorylating the beta subunit of the insulin receptor. More recently, Roome et al. (7) showed that PPTase from the particulate fraction of human placenta dephosphorylates insulin and EGF receptors. Similarly, King and Sale (38) showed that rat liver PPTase activity measured using a synthetic peptide corresponding to amino acids 1142-1153 of the insulin receptor was mostly (80%) membrane associated. These same authors also reported that \sim 75% of the PPTase activity against insulin and EGF receptors was recovered in the particulate fractions from rat brain, heart, and liver (38).

The limited availability of autophosphorylated hormone receptors dictates that artificial substrates be used to purify and characterize PPTases. [32 P]Tyr-RCAM-lysozyme has the advantage of phosphorylation on a single tyrosine residue, thereby avoiding multiple sites with varying affinity. The [32 P]Tyr-polyEY substrate is unusual in that hepatic particulate PPTase activity is inhibited by micromolar concentrations of polyEY. In fact, we used this substrate in an attempt to measure a polyEY-insensitive enzyme. Tonks et al. (35) characterized two protein tyrosine phosphatases from human placenta that were most readily distinguished by sensitivity to polyEY. PPTase 1A and 1B were inhibited 50% by \sim 4 μ M and 50 nM polyEY, respectively. The sensitivity of unpurified rat liver particulate and soluble PPTase activities measured with [32 P]Tyr-RCAM-lysozyme substrate (50% inhibition with

~ 10 μ M polyEY) is sufficient to characterize it as polyEY sensitive, but not to permit comparison with the purified placental enzymes.

Several studies have been published recently that use synthetic peptides phosphorylated on tyrosine to monitor PPTase activity. The primary structure for these peptides has been derived from the structure of autophosphorylated tyrosine kinases (either the insulin receptor [38, 39] or the oncogene pp60src [40]). Sparks and Brautigan (41) reported that cytosolic rabbit kidney PPTases could dephosphorylate serum albumin, casein, and myosin light chains, while two peptide substrates ([Val⁵]angiotensin and the pp60src-derived peptide noted above) were not substrates for this enzyme.

Examination of the recent report by Myerovitch et al. (39) indicates further the need to be cautious in interpreting PPTase measurements using artificial substrates. These authors measured PPTase activity in normal and diabetic rats with a peptide corresponding to residues 1142–1153 of the insulin receptor. Their results showed zinc-sensitive activities in rat liver membrane preparations of ~ 2–3 pmol/min per mg protein (compared with ~ 400 pmol/min per mg measured in the present study with [³²P]Tyr-RCAM-lysozyme). While no change in particulate PPTase activity was seen with streptozotocin diabetes, a reproducible increase of 30–80% in soluble activity was seen. In addition, it was demonstrated that this soluble activity resided in an enzyme with an M_r of 20,000–25,000 based on gel filtration chromatography. However, our own gel filtration analysis of PPTase activity against autophosphorylated hormone receptors indicates no activity in the range of $M_r = 25,000$.

In an attempt to correlate PPTase activity with artificial substrates and results using physiologic substrates (insulin and EGF receptors) we found that activity against the two receptor substrates differed in sensitivity to inhibition by polyEY and effects of alloxan diabetes. This result might be interpreted as indicating that distinct enzymes dephosphorylate the two receptor substrates. It also raises the possibility that different enzymes are measured using the artificial versus physiologic substrates. Since initiating these studies we have undertaken purification of rat hepatic PPTase using [³²P]Tyr-RCAM-lysozyme to monitor activity (manuscript in preparation). Affinity chromatography on polylysine agarose yields two peaks of activity. Both are fully inhibited by micromolar concentrations of polyEY. However, both PPTases are capable of dephosphorylating insulin and EGF receptors and only activity against the latter is inhibited by polyEY. Furthermore, the enzymes maintain an apparent M_r of 50,000–60,000 throughout purification. This finding supports the identity of these enzymes with the major insulin and EGF receptor PPTases in rat liver membranes. The results of the studies presented herein are therefore interpreted as indicating that (a) inhibition by polyEY is substrate dependent, and (b) PPTases with M_r 's of 50,000–60,000 are capable of dephosphorylating both the artificial and physiologic substrates used in the present study.

An additional finding still demands explanation, however. The difference in the effects of alloxan diabetes on dephosphorylation of insulin versus EGF receptors must be interpreted as indicating distinct PPTases that are substrate specific and independently regulated. This situation is reminiscent of the relationship between glycogen synthase

phosphatase and phosphorylase phosphatase. These glycogen-associated phosphatases are both forms of type 1 protein phosphatase (42). However, alloxan diabetes in the rat results in a marked decrease in synthase phosphatase activity with no effect on phosphorylase phosphatase activity (43, 44). It is unknown whether substrate specificity for type 1 protein phosphatase is dependent on associated regulatory proteins or posttranslational modification of the catalytic subunits. Similarly, the relationship between PPTases that dephosphorylate insulin and EGF receptors will have to await isolation and characterization of catalytic and regulatory subunits.

The present studies also should be interpreted in relation to our original hypothesis: an increase in PPTase activity will contribute to the insulin resistance seen in states associated with decreased insulin receptor kinase activity. Our data indicate that, in the specific case of experimental diabetes in the rat, no generalized alteration in PPTase activity occurs. Indeed, studies with autophosphorylated receptors indicate that a decrease in insulin receptor dephosphorylation may occur. Most importantly, the constitutive expression of hepatic PPTase activity after acute insulin administration and the distribution of PPTase activity in both cell surface and intracellular membranes suggest a primary regulatory role for receptor-associated tyrosine kinases.

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