Insulin binds to and promotes the phosphorylation of a M_r 210 000 component of its receptor in detergent extracts of rat liver microsomes

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Insulin in the presence of Mn^2 and $[\gamma^{32}P]ATP$ promoted the phosphorylation of two proteins of M_r 95 000 and M_r 210 000 in detergent extracts of rat liver microsomes. The M_r 210 000 protein was identified as a component of the insulin receptor by immunoprecipitation. It also bound [^{125}I]insulin specifically, was phosphorylated largely on a tyrosine residue and could not be cleaved to smaller subunits under extreme reducing conditions. The M_r 210 000 protein appears to be a component of a sub-population of liver membrane insulin receptors in which insulin-binding and insulin-stimulated tyrosine kinase phosphorylation site(s) reside in a single polypeptide chain.

Insulin binding
Autophosphorylation

Tyrosine kinase Proreceptor Insulin receptor Liver membrane

1. INTRODUCTION

Recent investigations have established that insulin, when added to intact cells, or detergent extracts of membranes, can promote the phosphorylation of the M_r 95 000 (3) subunit of its cellular receptor [1,2]. In addition, insulin added to partially purified insulin receptor preparations activates a tyrosine-specific protein kinase [3], (submitted). The insulin receptor may itself be the tyrosine kinase, in a manner analogous to the cellular receptor for epidermal growth factor (EGF) [4]. Direct evidence favoring this formulation is the demonstration of an ATP-binding site on the M_r 95 000 subunit of the insulin receptor [5,6]. Since insulin appears to bind exclusively to the M_r 130 000 (α) subunit of its receptor, it has been assumed that the M_r 130 000 subunit is the binding subunit and the M_r 95 000 subunit is the effector subunit of a disulfide-linked oligomer.

During studies in which we evaluated insulinstimulated receptor autophosphorylation in detergent extracts of rat hepatic microsomes, we repeatedly noted insulin-stimulated phosphorylation of a minor phosphoprotein of $M_{\rm r}$ 210000 (submitted). Here, we present evidence that this $M_{\rm r}$ 210000 subunit is a component of at least a fraction of the insulin receptors in these membrane extracts, and that it binds insulin in a displaceable manner. These findings suggest that in a proportion of insulin receptors in this tissue, insulinbinding and tyrosine kinase activities may reside on a single polypeptide chain.

2. MATERIALS AND METHODS

Microsomal extracts were prepared from fed, fasted or fasted/refed male Sprague-Dawley rat (200-300 g) livers by a method described in detail elsewhere (submitted). Briefly, livers were minced, homogenized with a Teflon/glass homogenizer in an ice-cold buffer (10 mM sodium phosphate (pH 7.5), 0.25 M sucrose, 10% (v/v) glycerol, 5 mM EDTA, 10 mM benzamidine-HCl, 10 mM ε-aminoisocaproic acid, 0.1% (w/v) bacitracin, 10³ KIU/ml aprotinin, 2 μM pepstatin and 2 μM

leupeptin), and centrifuged at $12\,000 \times g$ for 30 min. The resulting supernatant was centrifuged at $100\,000 \times g$ for 60 min, and the microsomal pellet was resuspended in 50 mM Tris-HCl (pH 7.4), 0.25 M sucrose, at about 10 mg protein/ml, and stored at -70° C. Membranes were thawed in the presence of (final concentration) 1% (w/v) Triton X-100, 2 μ M leupeptin, 2 μ M pepstatin, and 10^{3} KIU/ml aprotinin, extracted at 0° C for 40 min, and the supernatant of a $100\,000 \times g$ for 60 min centrifugation was recovered. In a standard assay, the supernatants (about 10 mg protein/ml) were exposed to insulin ($1\,\mu$ M) or control conditions for

10 min at room temperature, followed by chilling to 0°C. This extract was added to a reaction mixture consisting of (final concentration) 7.5 mM MnCl₂, 50 mM Tris-HCl (pH 7.5), 16.7 mM NaF, 0.05 mM ZnSO₄ and 200 μ M [γ -³²P]ATP (1500 cpm/pmol). After 5 min at 0°C, the reaction was stopped by the addition of 1% (w/v) SDS and dithiothreitol (DTT) (40 mM), or as noted in the figure legends. The samples were boiled for 5 min prior to detergent gel electrophoresis [7].

Immunoprecipitation with antiserum directed against the human insulin receptor was carried out as in [2]. To cross-link [125] linsulin to its receptor,

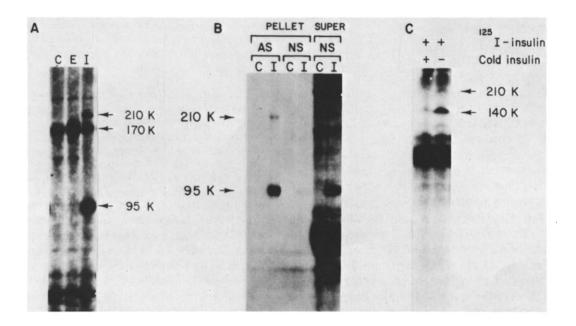


Fig. 1. Insulin-stimulated phosphorylation and immunoprecipitation of and [125 I]insulin binding to the M_r 210000 protein. (A) Microsomal extracts (10 mg protein/ml) from starved rat livers were exposed to control conditions (C), EGF (10 M; E) or insulin (10 M; I) for 10 min at room temperature, then phosphorylated under standard conditions as described in the text. The arrows point to the M_r 95 000 β subunit of the insulin receptor (95 k), the M_r 170 000 EGF-receptor (170 k), and the M_r 210 000 protein (210 k). (B) Phosphorylated extracts of rat liver microsomes, exposed to control conditions (C) or insulin (I), were immunoprecipitated (PELLET) with normal human serum (NS) or anti-insulin receptor antiserum (AS) as described in the text; in addition, the supernatant of the immunoprecipitation reaction in the presence of normal serum (SUPER) is shown. The arrows point to the M_r 95 000 (95 k) and the M_r 210 000 (210 k) subunits of the insulin receptor. (C) [125 I]Insulin (10 M) was incubated with rat liver microsomal membrane extracts (10 mg/ml) for 30 min at room temperature in the presence (+) or absence (-) of excess non-radioactive insulin (10 M) (cold insulin), and then cross-linked to the insulin receptor with disuccinimidyl suberate. Proteins of M_r 210 000 (210 K) and M_r 140 000 (140 K) were specifically labelled with [125 I]insulin. The minor band of M_r ~ 115 000 which appears to be specifically labelled was shown in other studies to be a proteolytic fragment of the M_r 140 000 protein. A large amount of non-specific labelling was also noted on unidentified proteins of M_r of about 60 000.

the membrane extract was allowed to bind $[^{125}I]$ -insulin (10^{-10} M; spec. act. about $100 \,\mu\text{Ci/\mu g}$) for 30 min at room temperature in the presence or absence of 10^{-6} M non-radioactive insulin. Disuccinimidyl suberate was then added (final conc. 1 mM), and cross-linking was allowed to proceed for 5 min at room temperature.

The identification of ³²P-linked amino acid residues was carried out as in [2].

3. RESULTS

Insulin promoted the phosphorylation of two proteins in detergent extracts of rat liver microsomes, a major phosphoprotein of M_r 95 000 and a minor phosphoprotein of M_r 210000 (fig. 1A). The latter protein was clearly distinct from the M_r 170 000 protein whose phosphorylation was stimulated by EGF in these extracts, which is presumably the EGF receptor. The insulinstimulated increment in 32P incorporated into the M_r 210 000 protein represented about 14% of total insulin-stimulated protein phosphorylation in extracts of livers from fed, starved or fasted/refed rats. Insulin-stimulated protein phosphorylation per unit protein was greatest in liver extracts prepared from 48-h starved animals, and these extracts were used in all subsequent experiments.

Antiserum directed at the human insulin receptor specifically immunoprecipitated both the insulin-stimulated $M_{\rm r}$ 210 000 and $M_{\rm r}$ 95 000 phosphoproteins, indicating that both were components of the insulin receptor (fig. 1B). In contrast, the $M_{\rm r}$ 210 000 phosphoprotein was not detected in immunoprecipitates of placental insulin receptors (not shown). [125 I]Insulin was specifically cross-linked to peptides of $M_{\rm r}$ 210 000 and $M_{\rm r}$ 140 000 (fig. 1C).

The time course of insulin-stimulated phosphorylation of both the $M_{\rm r}$ 210 000 and $M_{\rm r}$ 95 000 proteins was identical when $[\gamma^{-32}{\rm P}]$ ATP was added after pre-binding insulin (fig. 2a). This time course identity is against the possibility that the $M_{\rm r}$ 210 000 protein serves as a substrate for a protein kinase associated exclusively with the $M_{\rm r}$ 95 000 subunit. When the $M_{\rm r}$ 210 000 phosphoprotein was cut from the gel, extracted, hydrolyzed in 6 N HCl, and subjected to thin-layer electrophoresis in two dimensions, the major increment in phosphorylation promoted by insulin was on a tyrosine residue,

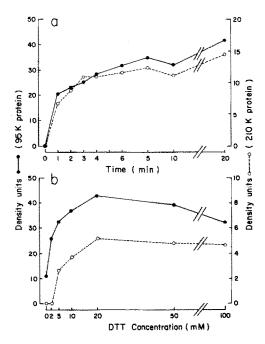


Fig. 2. Time course (a) and effect of DTT concentration (b) on the appearance of the M_r 95 000 and M_r 210 000 insulin-stimulated phosphoproteins. (a) microsomal extracts were exposed to insulin (10-6 M) at room temperature for 10 min, followed by cooling to 0°C and addition of 7.5 mM MnCl₂ and $[\gamma^{-32}P]ATP$. Aliquots of the reaction mixture were removed at the specified intervals and subjected to electrophoresis. The densities of the M_r 95000 (\bullet and M_r 210000 (O-O) [32P]proteins at each time point are shown. In (b) the effect of increasing concentrations of DTT on the appearance of the M_r 95 000 (\bullet and the M_r 210 000 (0-0) [32P]proteins at their appropriate positions in the autoradiograph are shown. After phosphorylation under standard conditions after exposure to insulin, aliquots of the reaction mixture were mixed with (final concentration) 1% (w/v) SDS and the concentrations of DTT noted in the abscissa, and then boiled for 5 min. See the text for further details.

although traces of phosphoserine were noted in both control and insulin-stimulated samples (not shown).

To establish that the M_r 210000 [32 P]peptide could not be further cleaved by disulfide reduction, the reaction was stopped by the addition of 1% SDS and 40 mM DTT or 1% SDS, 5% β -mercaptoethanol and 100 mM DTT [8], followed by boiling in both cases. The more stringent

disulfide-reducing conditions did not alter the relative proportions of the insulin-stimulated phosphoproteins ($M_{\rm r}$ 210 000 and $M_{\rm r}$ 95 000) in the gel (not shown). Furthermore, the relative proportions of the $M_{\rm r}$ 210 000 and $M_{\rm r}$ 95 000 phosphoproteins were unaffected over a wide range of DTT concentrations (fig. 2b), a result which would not be expected if the $M_{\rm r}$ 210 000 protein were merely an incompletely reduced heterodimer of the α - and β -subunits.

4. DISCUSSION

This study demonstrates that detergent extracts of rat liver microsomal membranes contain a M_r 210 000 protein whose tyrosine-specific phosphorylation is stimulated by the addition of insulin directly to the extract This $M_{\rm r}$ 210 000 protein is identified as a component of the insulin receptor by its ability to be specifically cross-linked to [125] Ilinsulin as well as by its immunoprecipitation by antisera directed at the insulin receptor. The M_r 210 000 protein does not appear to be a disulfide-linked heterodimer of the M_r 130 000 (α) and M_r 95 000 (3) subunits of the insulin receptor. Nevertheless, there do appear to be structural homologies between the M_r 210 000 and M_r 95 000 [³²P]proteins based on partial [³²P]peptide maps (unpublished) and based on peptide maps using other labelling procedures [8].

These findings should be compared to those in IM9 lymphocytes [8] and 3T3-L1 cells [9]. Both groups identified a $M_{\rm r}$ 210000 protein immunoprecipitated by antisera to the insulin receptor, and suggested that it may represent a biosynthetic precursor of both the $M_{\rm r}$ 130000 and $M_{\rm r}$ 95000 subunit. They also provided evidence that at least a fraction of the $M_{\rm r}$ 210000 proteins were at the cell surface, as determined by surface labelling and limited proteolysis in living cells.

We are uncertain as to whether the hepatic M_r 210 000 insulin receptor subunit observed here is a receptor precursor, or separate functional pool of

mature receptor subunits. In any case, the hepatic $M_{\rm r}$ 210 000 subunit appears to be part of a subpopulation of insulin receptors in which insulinbinding and autophosphorylation activities reside on a single peptide chain, and whose role in insulin action remains to be defined.

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