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# Isolation by preparative free-flow electrophoresis and aqueous twophase partition from rat adipocytes of an insulin-responsive small vesicle fraction with glucose transport activity

Dorothy M. Morré<sup>a,\*</sup>, David W. Sammons<sup>b</sup>, Jonathan Yim<sup>a,c</sup>, Maria Bruno<sup>a,c</sup>, Timothy Snyder<sup>a</sup>, Timothy Reust<sup>a</sup>, Lidia Maianu<sup>d</sup>, W. Timothy Garvey<sup>d</sup>, D. James Morré<sup>c</sup>

<sup>a</sup>Department of Foods and Nutrition, Purdue University, West Lafayette, IN 47907, USA

<sup>b</sup>Department of Veterinary Science, University of Arizona, Tucson, AZ 85721, USA

<sup>c</sup>Department of Medicinal Chemistry and Pharmacognosy, Purdue University, West Lafayette, IN 47907, USA

<sup>d</sup>Department of Medicine, Medical University of South Carolina and the Charleston Veterans Affairs Medical Center, Charleston, SC 29425, USA

#### Abstract

Preparative free-flow electrophoresis and aqueous two-phase polymer partition were used to obtain a plasma membrane-enriched fraction of adipocytes isolated from epididymal fat pads of the rat together with a fraction enriched in small vesicles with plasma membrane characteristics (thick membranes, clear dark-light-dark pattern). The electrophoretic mobility of the small vesicles was much less than that of the plasma membrane consistent with an inside-out orientation whereby charged molecules normally directed to the cell surface were on the inside. When plasma membranes and the small vesicle fraction were isolated from fat cells treated or not treated with  $100 \, \mu \text{U/ml}$  insulin and the resident proteins of the two fractions analyzed by SDS-PAGE, the two fractions exhibited characteristic responses involving specific protein bands. Insulin treatment for 2 min resulted in the loss of a 90 kDa band from the plasma membrane. At the same time, a ca. 55-kDa peptide band that was enhanced in the plasma membrane was lost from the small vesicle fraction. The latter corresponded on Western blots to the GLUT-4 glucose transporter. Thus, we suggest that the small vesicle fraction with characteristics of inside-out plasma membrane vesicles may represent the internal vesicular pool of plasma membrane subject to modulation by treatment of adipocytes with insulin.

Keywords: Free-flow electrophoresis; Partitioning; Vesicles, insulin-responsive; Glucose transport

# 1. Introduction

Our laboratory has used preparative free-flow electrophoresis [1,2] and aqueous two-phase polymer partition [3,4] to obtain a plasma membrane-enriched fraction from adipocytes of epididymal fat pads of

the rat together with a fraction enriched in small vesicles with plasma membrane characteristics (thick membranes, clear dark-light-dark pattern) [4,5]. The electrophoretic mobility of the small vesicles was much less than that of the plasma membrane consistent with an inside-out orientation whereby charged molecules normally directed to the cell surface were on the inside. These fractions have been

<sup>\*</sup>Corresponding author.

utilized in the present report to investigate the cascade of membrane-associated events that occur in response to insulin treatment of isolated adipocytes. These include modulation of the 90-kDa peptide reported previously by Schoenle et al. [6] to be lost from the plasma membrane of adipocytes upon insulin treatment and a redistribution of the GLUT-4 glucose transporter consistent with translocation from a small vesicle fraction to the plasma membrane.

The insulin action sequence begins with ligand binding to cell-surface receptors which are heterodimers comprised of 135 kDa alpha subunits and 95 kDa beta subunits. Binding of insulin to alpha subunits initiates autophosphorylation of tyrosine residues in the beta subunits which in turn augments receptor tyrosine phosphorylase activity for other intracellular substrates [7,8]. Using isolated rat adipocytes, we found increased numbers of cell surface invaginations and loss of vesicular profiles associated with the plasma membrane during the first 3 min of insulin administration and return to initial values by 5 min [9]. The findings are expected to have significance for the elucidation of early events responsible for the down regulation of insulin receptors [10-12] at the cell surface.

Propagation of signal transduction by ligand-activated insulin receptors results in stimulation of glucose transport into adipose and muscle cells. The increase in transport activity is due to translocation of an intracellular pool of facilitative glucose transporter proteins to the cell surface [13,14]. This process involves exocytosis of transporter-containing vesicles, together with other insulin-responsive vesicles containing alternative receptors [15], which fuse with plasma membrane allowing an exofacial orientation. The predominant insulin-responsive glucose transporter isoform has been designated GLUT 4. While the involvement of GLUT 4 vesicles has been established [16,17], the actual location of the intracellular pool and organelle source of translocating GLUT 4 has not been identified with certainty. This is because conventional methods for membrane subfractionation have not yielded a pure intracellular fraction containing GLUT 4, and morphological localization using immunocytochemistry has not been definitive. The current study utilizes new approaches, namely free-flow electrophoresis and aqueous two-phase polymer partitioning, to study insulin's effects on vesicular traffic in isolated adipocytes.

#### 2. Materials and methods

#### 2.1. Preparation of adipocytes

Intact insulin-sensitive adipocytes were isolated from rat epididymal fat pads according to the basic procedure of Rodbell [18]. The preparations of collagenase and albumin used for digestion and incubation of fat cells were carefully monitored. Albumin was trypsinized to remove insulin. Male Wistar rats (fed standard laboratory chow), weighing approximately 200 g, were killed by decapitation and the epididymal fat pads removed. Fat cells were isolated by collagenase digestion [18] in Krebs-Ringer bicarbonate buffer (pH 7.4) containing 0.6% calcium, 3% bovine serum albumin, 0.2% glucose and 7.0 mg collagenase for 30 min at 37°C. The isolated cells were washed in 10 volumes of homogenization medium containing 10 mM Tris-HCl (pH 7.4 at  $4^{\circ}$ C), 0.25 M sucrose and 0.1 mM phenylmethyl-sulfonyl fluoride (PMSF) and centrifuged at 400 g for 1 min. Cell intactness was verified by light microscopic examinations. Insulin from porcine was a gift of Eli Lilly and Company, and was added to the cells and membranes at a near optimum final concentration of 100 µU/ml.

# 2.2. Electron microscopy

Reproducible fixations of high quality have been difficult to achieve with adipocytes and special fixation protocols have appeared [19]. Our protocol is based on addition of a small quantity of concentrated glutaraldehyde at ambient temperature directly to the culture medium followed by rapid mixing to a final concentration of 0.2%. After an additional 5 min, the culture medium containing fixative was removed and replaced by 2% glutaraldehyde prepared in 0.1 M phosphate buffer, pH 7.2, beginning at ambient temperature and then continuing at 4°C for 2 to 8 h. Postfixation was in 1% osmium tetroxide with dehydration using an acetone series and embedment in Epon. The sections were post-

stained with alkaline lead citrate to enhance membrane contrast and examined and photographed using a Philips EM200 electron microscope.

# 2.3. Morphometry

Well-fixed cells with high membrane contrast were photographed and the micrographs printed at a final magnification of about 35 000. All vesicular profiles were analyzed as to number, diameter, morphological characteristics and distance from the plasma membrane as a function of time after insulin addition. Surface evaginations were analyzed in proportion to total membrane surface, as were locations of subsurface vesicles. The findings were analyzed for statistical significance and to ensure that sample sizes were sufficient to provide reliability.

# 2.4. Isolation of plasma membranes

Isolated fat cells were incubated in buffer in the presence or absence of insulin, resuspended in 3.5 volumes of the homogenization medium, transferred to a glass homogenizer fitted with a Teflon pestle and homogenized with ten up-and-down strokes by means of a variable-speed motor driven homogenizer at 2600 rpm. The homogenates were centrifuged at 35 000 g for 15 min to obtain a 'heavy' microsomal pellet. The pellet was resuspended in 0.5 ml phosphate buffer, pH 7.4, and used for two-phase separation of plasma membranes. The 4-g two-phase system contained equal portions (6.6% w/w) of dextran T500 (Pharmacia, Uppsala, Sweden) and poly(ethylene glycol) 3350 (Fisher, Chicago, IL, USA) plus 0.1 M sucrose, 5 mM potassium phosphate, pH 7.2, and 0.5 g of resuspended microsomes (0.5 mg protein). The contents of the two-phase systems were mixed thoroughly by 30 to 40 inversions of the tubes at 4°C and centrifuged at 750 g for 5 min to separate the phases. The upper phase was collected, diluted three- to five-fold with 1 mM sodium bicarbonate and collected by centrifugation at 20 000 g as the source of plasma membranes.

Both upper and lower phases were analyzed for the plasma membrane marker, K<sup>+</sup>-stimulated p-nitrophenylphosphatase (EC 3.1.3.1), the mitochondrial marker succinate-INT reductase (EC 2.3.99.1) and protein (with BSA as standard) [3]. As the polymer concentration of the two phases was increased, mitochondria and mitochondrial enzyme activities were partitioned to the lower phase.  $K^+$ -stimulated p-nitrophenylphosphatase specific activity increased to an optimum at about 6.6% of each polymer. The recovery of plasma membrane in the upper phase obtained by aqueous two-phase partition was about 2% of the total protein.

## 2.5. Loss of 90-kDa protein

Two-dimensional gel electrophoresis used the ISO-DALT system (Ref. [6] and references cited therein). The proteins on the polyacrylamide gels were stained with the silver-based color development system using Gelcode (Upjohn, Kalamazoo, MI, USA). Molecular mass was determined by applying molecular mass standards (10–200 kDa) designed specifically for use with the color silver stains (Health Products) to each of the second dimension gels.

#### 2.6. Free-flow electrophoresis

For isolation of the small vesicle fraction, freeflow electrophoresis was used. In this technique, the mixture of components to be separated was introduced as a fine jet into a separation buffer moving perpendicular to the field lines of an electric field. Membranes bearing different electrical charge densities migrated different distances across the separation chamber and thus were resolved. The electrophoresis medium (chamber buffer) contained 10 mM triethanolamine, 10 mM acetic acid, 5 mM glucose, 0.25 M sucrose and 0.5 mM MgCl<sub>2</sub>, final pH 6.5 (1 M NaOH). The electrode buffer contained 100 mM triethanolamine and 100 mM acetic acid, pH 6.5. Electrophoresis conditions were 167 mA (constant current), 131±10% V/cm, buffer flow 2.75 ml per fraction per h, sample injection 3.5 ml/h, temperature 6°C. The electrophoresis equipment was a VAP-21 continuous free-flow electrophoresis unit (Bender and Hobein, Munich, Germany).

## 2.7. In vitro glucose uptake

For the in vitro glucose uptake experiments, purified plasma membrane vesicles, purified by

aqueous two-phase partition, or the free-flow electrophoretic fractions were absorbed onto 0.5×0.5 cm nitrocellulose strips. The fractions were resuspended at a final concentration of 1 to 2 mg of protein per milliliter of saline and incubated 6 strips/300 to 500  $\mu$ l for 30 min, with continuous shaking. The strips, each loaded with about 30 µg of plasma membrane protein, were rinsed through four changes of 4-(2-hydroxyethyl)-1-piperazineethanesulphonic (Hepes)-Mg(OAc)2-KCl, and edge blotted to remove excess solution. The strips were added to Eppendorf cups containing 0.1  $\mu$ Ci [methyl-<sup>14</sup>C]-3-O-methylglucose (30-60 mCi/mmol) (Sigma, St. Louis, MO, USA), 15 µM ATP plus an ATP regenerating system [20] with or without 1 mM glucose and in the presence or absence of  $0.2 \mu M$ cytochalasin B in a final volume of 150 µl. At appropriate times of incubation, strips were removed. washed through four changes of Hepes-Mg(OAc),-KCl, 0.5 s per change, and edge blotted. Radioactivity was determined by scintillation methods.

# 3. Results

Rat adipocytes consist of a thin band of cytoplasm surrounding a large central fat core (Fig. 1A). Upon lysis, the membrane-rich periphery and the central fat droplet were separated at the initial centrifugation step.

The morphological appearance of the plasma membrane fractions prepared by aqueous two-phase partition from adipocytes demonstrated the predominant plasma membrane composition of the fraction (Fig. 1B). The plasma membranes were isolated as smooth vesicles having thick membranes. Contaminants were predominantly mitochondria. Other internal membranes, chiefly mitochondria and endoplasmic reticulum, entered the lower phase of the separation (Fig. 1C).

The isolated adipocytes were insulin responsive. Morphologically, the cells responded to insulin by the appearance of numerous fusion profiles with the plasma membrane resembling pinosomes (Fig. 2). The numbers of the pinosome profiles were quantitated and found to be maximal after about 2 min of insulin addition (Fig. 3). At the same time, the microvesicle population of the internal cytoplasm

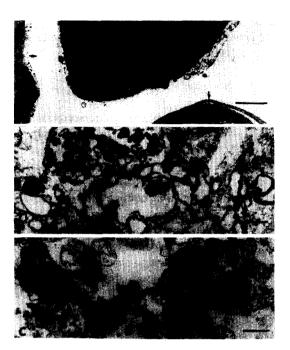


Fig. 1. Electron micrographs of isolated fat cells (A) and membrane fractions by aqueous two-phase partition (B and C). (A) Isolated fat cell from an epididymal fat pad of the rat. The electron micrograph shows the thin band of cytoplasm (arrows) surrounding the large central fat store. The membrane-rich periphery and the central fat droplet are separated in the initial centrifugation step. Scale bar=5  $\mu$ m. (B) Top phase (plasma membrane-enriched) after aqueous two-phase partitioning. Electron microscopy shows the top phase material enriched in smooth vesicles with morphological characteristics typical of plasma membrane vesicles from other cells and tissues. Scale bar=0.5  $\mu$ m. (C) Lower phase (mitochondria-enriched). The lower phase of the two-phase separation contains contaminating membranes, chiefly mitochondria (m) as well as rough-surfaced endoplasmic reticulum (er). Scale bar=0.5  $\mu$ m.

decreased in response to insulin treatment (Fig. 4). These results suggested that at least some of the fusion profiles of Fig. 2 were due to cytoplasmic vesicles fusing with the plasma membrane. As with the appearance of pinosomes, the loss of cytoplasmic microvesicles was maximal after about 2 min after insulin addition and tended to return to initial values within 30 min after insulin administration.

If the plasma membranes of the adipocytes were coated with cationized ferritin prior to insulin addition, some microvesicles carrying cationized ferritin were observed within the cytoplasm (Fig. 5) but not

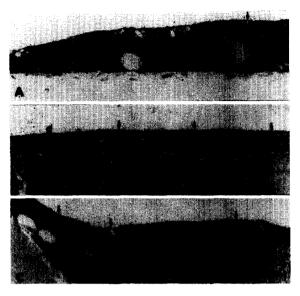


Fig. 2. Isolated fat cells treated for 1 min in the absence (A) or presence (B and C) of insulin (100  $\mu$ U/ml). Both cell populations showed abundant pinosomes (invaginations of the external membrane surface) (arrows) but the pinosomes were more abundant with the cells treated for brief (30 s to 2 min) intervals with insulin. Bar=0.5  $\mu$ m.

in sufficient numbers to account for all of the apparent pinosomes suggested by observations of Fig. 2 and Fig. 3. These findings, as well, were consistent with enhanced microvesicle traffic to and from the plasma membrane with a net decrease in the level of cytoplasmic microvesicles.

Additionally, when analyzed by two-dimensional gel electrophoresis and silver staining, a 90-kDa protein similar to that reported previously by Schoenle et al. [6] was lost from the plasma membrane in response to insulin treatment (Fig. 6). In these experiments, the plasma membranes were isolated by aqueous two-phase partition as illustrated in Fig. 1. The 90-kDa protein was restricted to the plasma membrane-enriched fraction of the upper phase. Lower phase membranes of either untreated fat cells or of fat cells incubated with 100  $\mu$ U/ml of insulin for 1 min did not contain evidence of the 90-kDa protein.

As a result of the morphological studies, an attempt was made to isolate a subcellular fraction containing the microvesicles depleted from the cyto-

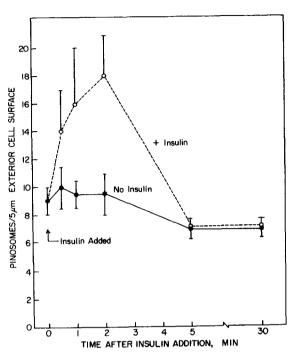


Fig. 3. Pinosomes per 5  $\mu$ m of exterior cell surface of isolated fat cells were increased two-fold during the first 2 min after addition of 100  $\mu$ U/ml insulin. Pinosome numbers then returned to control values by 5 min.

plasm in response to insulin treatment. Free-flow electrophoretic profiles of homogenates of cells treated for 5 min with or without  $100~\mu\text{U}$  of insulin/ml exhibited evidence for such a fraction (Fig. 7). The electrophoretic profiles exhibited membrane material distributed over about 30 electrophoretic fractions. The most electrophoretically mobile material was designated zone 1 while material collecting between fractions 33 and 39 were combined to give zone 2. The least electrophoretically mobile material was designated zone 6 and fractions 46 to 51 were combined as zone 5. The mid portion of the separation between fractions 40 and 45 were divided equally into zones 3 and 4.

The electron microscopic appearance of the pooled fractions is illustrated in Fig. 8. Zone 1 consisted of filaments. Zone 2 was fragmented mitochondria. Zone 3 was enriched in vesicles resembling plasma membranes whereas the bulk of the mitochondria and endoplasmic reticulum were found in zone 4.

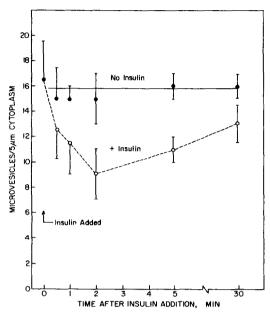


Fig. 4. Microvesicles per 5  $\mu$ m of cytoplasm of isolated fat cells were decreased about two-fold during the first 2 min after addition of 100  $\mu$ U/ml insulin. Compare with Fig. 3.

Zone 5 consisted of small vesicles resembling the microvesicles of the adipocyte cytoplasm whereas zone 6 consisted of a mixture of these small vesicles and filaments.

As evidenced from the electrophoretic profiles, zone 5, the fraction enriched in the microvesicles, decreased in response to insulin treatment (Fig. 7). This profile was very reproducible and was a consistent feature of electrophoretic separations comparing homogenates of adipocytes treated with or without insulin.

Analysis by SDS-PAGE of the electrophoretic fractions revealed an important difference in the membranes from adipocytes treated with or without insulin that was characteristic of zone 5, the putative microvesicle fraction. The membrane fraction from cells treated with insulin was depleted in a ca. 55-kDa protein band (Fig. 9). Each of the major electrophoretic fractions was compositionally distinct in terms of the pattern of protein bands but, except for the apparent loss of the 90 kDa protein from the plasma membrane-enriched fraction (Fig. 6), the loss of the ca. 55-kDa protein from the microvesicle



Fig. 5. The insulin-treated cells show greater internalization of a surface marker. After 5 min of treatment with  $100 \mu U/ml$  insulin of fat cells to which cationized ferritin was bound to the plasma membrane, insufficient cationized ferritin (arrows) was internalized to account for the abundance of pinosomes at 2 min. Scale bar=0.5  $\mu m$ .

fraction was the major protein alteration seen in response to insulin treatment.

The identity of the 55-kDa protein is suggested from Western blot analysis to be the GLUT-4 glucose transporter (Fig. 10).

The adipocytes isolated and treated with insulin exhibited increased glucose transport as determined by the uptake of methyl-D-glucose (data not shown). However, it was of interest to determine if the small vesicle fraction of these same adipocytes would exhibit glucose transport activity and if this transport activity could be lost upon insulin treatment.

In order to test this idea, an in vitro assay to measure glucose transport was developed in which the plasma membrane vesicles isolated by aqueous two-phase partition were immobilized on nitrocellulose, exposed to [methyl-<sup>14</sup>C]-D-glucose and then washed rapidly through four changes of buffer for

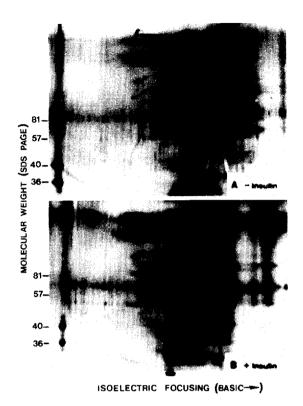


Fig. 6. Upper phase membranes after aqueous two-phase partition of fat cells treated with and without insulin. Two-dimensional gel electrophoretograms of proteins stained with a silver-based detection system [6] from upper phase membranes of (A) untreated fat cells showing presence of 90-kDa protein (arrow) and (B) fat cells incubated with 100  $\mu$ U/ml of insulin for 1 min.

0.5 s each. A control incubation contained glucose so that only glucose-inhibited uptake was measured. As shown in Fig. 11, these vesicles showed a rapid glucose-dependent uptake of methyl-D-glucose that was inhibited by cytochalasin B. This glucose-inhibited methyl-D-glucose uptake prevented by cytochalasin B was greater for plasma membrane vesicles isolated from adipocytes treated with insulin than from plasma membrane vesicles isolated from adipocytes treated in the absence of insulin (Fig. 11).

When applied to the analysis of the cell-fractions prepared by preparative free-flow electrophoresis, zone 3, enriched in plasma membrane, again exhibited greater uptake of glucose-inhibited methyl-pglucose from insulin-treated cells than from plasma membranes prepared from cells not treated with

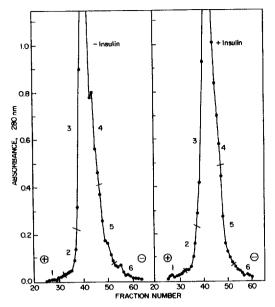


Fig. 7. Free-flow electrophoretic profiles comparing homogenates from fat cells treated for 5 min with (+) and without (-) 100  $\mu$ U/ml insulin. The numbers (in order of decreasing electrophoretic mobility) refer to the pooled fractions analyzed.

insulin (Fig. 12A). In contrast, in fractions of zone 3 which were enriched in microvesicles, uptake of glucose-inhibited methyl-p-glucose uptake was greater for fractions prepared from homogenates obtained from cells untreated with insulin than for fractions prepared from homogenates obtained from cells treated with insulin (Fig. 12B). This uptake was inhibited by cytochalasin B for both the plasma membrane-enriched (Fig. 12A) and the microvesicle-enriched (Fig. 12B) fractions.

### 4. Discussion

The responses of fat cells to insulin involve potentially important membrane alterations. There is now considerable evidence to support the concept that insulin stimulates glucose transport by including a rapid and reversible translocation of glucose transporters from an intracellular membrane pool to the plasma membrane [21–24]. However, little is known about the cellular mechanism of insulin action or the process or biochemical events whereby glucose

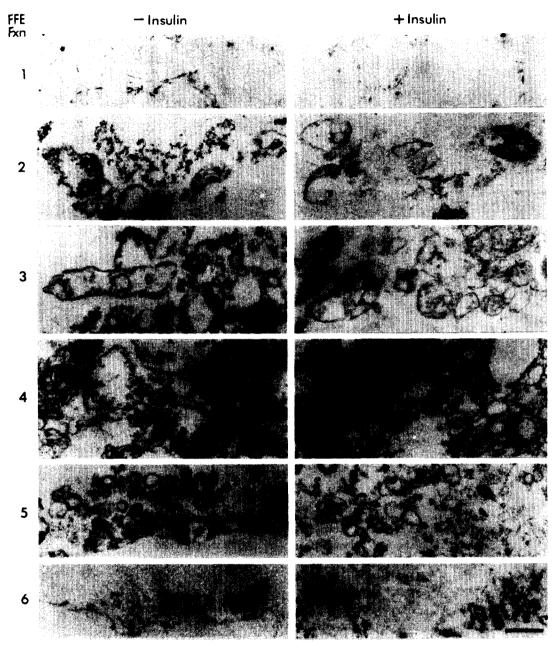


Fig. 8. Electron micrographs of the pooled electrophoretic fractions comparing cells treated with and without insulin. The numbers refer to pooled fractions indicated on the electrophoretic separations. Scale bar= $0.5 \mu m$ .

transporter levels at the cell surface may be modulated. Work of Schoenle et al. [6] used a sensitive silver staining method to visualize membrane proteins in two-dimensional gel electrophoretograms in conjunction with computerized scanning and quantification. A 90-kDa protein was lost from the plasma membranes upon incubation of fat cells with insulin and calcium. This protein has not been characterized

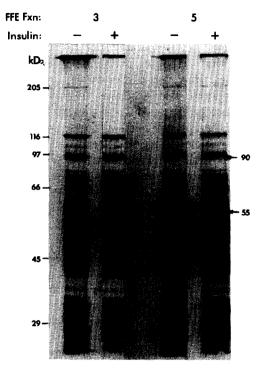


Fig. 9. SDS-PAGE gels of the pooled electrophoretic fractions comparing cells treated with and without insulin. The lane numbers refer to the pooled fractions of the electrophoretic separations of Fig. 7 and Fig. 8.

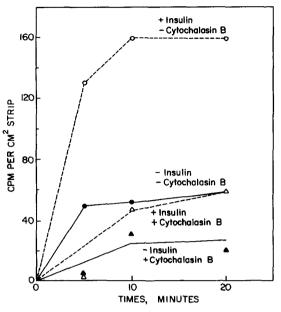


Fig. 11. In vitro glucose uptake by plasma membrane vesicles prepared by aqueous two-phase partition. To correct for unspecific uptake or trapping of methyl-p-glucose, uptake in the presence of excess glucose was measured. In the assay, vesicles were attached to strips of nitrocellulose. The strips were then incubated in the radiolabeled methyl-p-glucose for varying periods of time and rapidly rinsed through four changes of buffer. Two sets of strips were incubated in parallel. To one set was added excess unlabeled glucose. The unlabeled glucose was omitted from the other set. Results are the difference of uptake in the absence of glucose minus that in the presence of glucose.

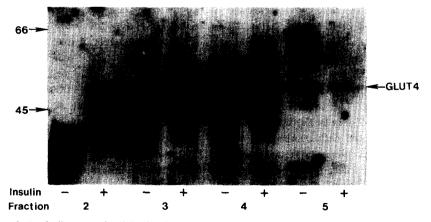


Fig. 10. Western blot analysis of adipocyte subcellular fractions using a monoclonal antibody to the GLUT-4 glucose transporter. Adipocytes were treated for 5 min with (+) or without (-) 100  $\mu$ U/ml of insulin. Membranes were isolated and stored frozen in 0.1 mM PMSF to retard proteolysis prior to analysis. The numbers refer to the pooled fractions analyzed as for Fig. 7. Each lane was loaded with 15  $\mu$ g protein. After 4 days of exposure of the blot to X-ray film.

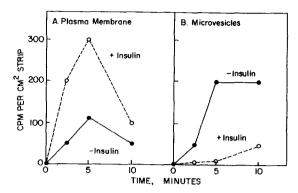


Fig. 12. In vitro glucose uptake comparing vesicles of electrophoretic zone 3 (plasma membrane) (A) and electrophoretic zone 5 (microvesicles) (B). Conditions were as in Fig. 11.

further and its characteristics on two-dimensional electrophoresis remain as the sole basis for its identification.

Fat cell plasma membranes usually have been purified by differential and density-gradient centrifugation [25,26]. However, such techniques are based on using 16–24 rats for each preparation. An alternative procedure, such as partitioning in aqueous polymer two-phase systems might be useful for purification of plasma membranes from fat cells isolated from as few as 1 or 2 rats. Use of two-phase partition in dextran-polyethylene glycol is well established in the purification and characterization of plant membranes [27] and, to a degree, has been used for animal membranes, particularly plasma membranes of liver [28]. In our study, this technique has been utilized to prepare highly purified plasma membranes from isolated fat cells.

Sufficient plasma membranes were obtained from fat pads of 1–2 rats for both enzymatic assays and analysis by two-dimensional gel electrophoresis with detection by silver staining. The insulin-responsive 90 kDa protein was detected only in the upper phase membranes of cells not treated with insulin and was not detected in either lower phase untreated membranes or in both upper and lower phase membranes of cells treated with insulin. The findings confirm the previous observations of Schoenle et al. [6] and establish the utility of aqueous two-phase partition as a method for plasma membranes of fat cells.

Micropinocytic activity and membrane flux involving microvesicles has long been considered as an aspect of insulin action in the cell. However, there have been few studies to quantitate the changes.

Our findings show what appear to be budding of vesicles in response to insulin treatment of adipocytes. However, a microvesicle population of the cytoplasm also decreases in parallel to the appearance of the budding profiles suggesting that some of the apparent budding profiles may actually be microvesicles fusing with the plasma membrane.

One possibility for the function of the fusing microvesicles would be to carry glucose transporters to the plasma membrane in response to the insulin stimulus [21,22,29]. We provide additional evidence for this function from studies where a microvesicle fraction has been isolated by preparative free-flow electrophoresis. In approximately twenty repetitions of the electrophoretic separation, the profile from homogenates prepared from cells treated with or without insulin shows the same characteristic appearance. A shoulder on the separation containing vesicles of lower electrophoretic mobility than the bulk of the plasma membrane and of lower electrophoretic mobility than that of either mitochondria or bulk endoplasmic reticulum was decreased in amount with insulin treatment. The electrophoretic mobility of this vesicle fraction would be that comparable to or less than that of a plasma membrane vesicle with an inside-out or cytoplasmic side-out orientation [3]. This fraction which has a morphology when examined by electron microscopy of a fraction enriched in microvesicles is enriched in the glucose-transporter protein GLUT-4. The glucose transporter protein GLUT-4 is depleted from the fraction when prepared from homogenates of insulin-treated cells with a net redistribution to the fractions enriched in right sideout plasma membrane vesicles.

As a final confirmation of the glucose transport activity of the fraction, an in vitro assay system was devised to measure uptake of radiolabeled methyl-D-glucose by isolated plasma membrane vesicles. This glucose-inhibited uptake of methyl-D-glucose was sensitive to inhibition by cytochalasin B and provides a measure of glucose transport amenable to analysis of the internal membrane fraction. When applied to the analysis of the fractions from preparative free-flow electrophoresis, the combined fractions of zone 3 when prepared from homogenates of insulin-treated cells were stimulated in their ability to

take up methyl-D-glucose identifying them as being of plasma membrane origin. Their methyl-D-glucose uptake characteristics and inhibition by glucose and by cytochalasin B were similar to those of plasma membrane vesicles obtained by aqueous two-phase partition. In contrast, combined fractions of zone 5 from the electrophoretic separations showed a reduced ability to transport glucose when isolated from homogenates of cells treated with insulin. These results further indicate that the microvesicle fraction contains the internal vesicular pool of plasma membrane subject to modulation by treatment of adipocytes with insulin and responsible for transport of glucose transporters to the cell surface.

After insulin stimulation, GLUT-4 unambiguously becomes the major glucose transporter of the plasma membrane [29–31]. In the rat adipocyte plasma membrane, insulin treatment results in a fifteen- to twenty-fold increase in GLUT-4 in the plasma membrane [30]. Using impermeant photoprobes it has been shown that GLUT-4 recycles between its intracellular compartment and the plasma membrane [32,33].

The identity of the intracellular pool of GLUT-4 remains controversial. The *trans* Golgi has been suggested [34,35]. Smith et al. [36] detected GLUT-4 not only in *trans* Golgi but also concentrated heavily in subplasma membrane vesicles using immuno-gold electron microscopy.

Taken together our findings agree with those of Smith et al. [36] to suggest that the intracellular pool for insulin responsive glucose transporters, represents at least in part a plasma membrane-like compartment perhaps not unlike an early endosome (see Garvey [29] for references of localization of GLUT-4 to early endosomes, coated pits and coated vesicles). The putative internal compartment of microvesicles that we describe, if a plasma membrane derivative, would be inside-out relative to the cell surface, as determined by its electrophoretic mobility [3] and behavior in aqueous two-phase partitioning. The latter technique is especially sensitive to surface properties. Right side-out vesicles enter the upper phase whereas inside-out vesicles are recovered with the lower phase [37].

While it is clear that diabetic conditions resulting from cellular insulin resistance may result from multiple defects [29], among these there may be abnormalities in cellular traffic or targeting. Some may relegate the GLUT-4 to a membrane compartment from which insulin is unable to recruit transporters to the cell surface [38]. Therefore, the availability of new subcellular fractionation technology that permits facile resolution of intracellular and plasma membrane glucose transport compartments is expected to aid future studies of direct relevance to the mechanisms of certain types of insulin resistance.

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