Effects of tumour necrosis factor alpha (TNF α) on glucose transport and lipid metabolism of newly-differentiated human fat cells in cell culture

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Summary Tumour necrosis factor alpha (TNF α) has been found to cause a delipidation of fat cells and a decrease of the adipose tissue mass. In the present study, we tried to elucidate some of the mechanisms responsible for this phenomenon by investigating the action of TNFa on specific pathways which are involved in lipid storage. Cultured stromal cells from human adipose tissue were induced to differentiate into adipose cells by exposure to adipogenic factors and subsequently used for studying the effects of TNF α on fat cell metabolism. Presence of 5 nmol/l TNF α for 24 h resulted in a complete loss of the stimulatory effect of insulin on 2-deoxy-glucose transport. This inhibitory action was paralleled by a decrease of GLUT4 protein and mRNA levels. The amount of cellular GLUT4 protein was reduced by 49 ± 3 % after a 24-h exposure and by 82 ± 18 % after a 72-h exposure to 5 nmol/l TNFα. GLUT4 mRNA was almost undetectable after a 24-h incubation with

5 nmol/l TNF α In a similar time-dependent manner, TNF α dramatically reduced the lipoprotein lipase mRNA content of the cells. Furthermore, incubation of cultured human fat cells with TNF α resulted in a marked dose-dependent stimulation of lipolysis, assessed by glycerol release, by up to 400 % above controls, which became apparent after a 6-h exposure at the earliest. These data suggest that TNF α induces a catabolic state in human adipose tissue which includes a loss of the stimulatory effect of insulin on glucose transport. These multiple actions of TNF α may contribute to the loss of adipose tissue observed during cachexia in man. [Diabetologia (1995) 38: 764-771]

Key words Cachexia, glucose transport, human adipocytes, insulin resistance, lipolysis, tumour necrosis factor alpha.

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Abbreviations: TNFα, Tumour necrosis factor alpha; LPL, lipoprotein lipase; HEPES, N-2-Hydroxyethylpiperazine-N-2ethanesulfonic acid; MIX, 1-Methyl-3-isobutylxanthine; BSA, bovine serum albumin; GPDH, glycerophosphate dehydrogenase; PBS, phosphate buffered saline.

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Tumour necrosis factor alpha (TNF α), a 17-kD polypeptide synthesized mainly by activated macrophages but also by many other tissues, is known to exert pleiotropic metabolic effects in many species [1, 2]. Numerous in vivo and in vitro studies have tried to characterize the metabolic actions of TNFa. It was demonstrated in cell lines and animal experiments that this cytokine is able to decrease lipoprotein lipase activity [3-8], increase lipid mobilization from adipose tissue [4, 9–12] and stimulate de novo hepatic lipogenesis with the latter effect being probably dominant for the development of hypertriglyceridaemia [13, 14]. However, conflicting results have been reported from human studies. TNFa was not found to suppress lipoprotein lipase activity (LPL) activity in cultured human adipose cells [15], while LPL activity and synthesis were decreased when human adipose tissue fragments were cultured in the presence of TNF α [16]. We have recently reported that TNF α is able to suppress glycero-3-phosphate dehydrogenase activity, a lipogenic key enzyme, in cultured human fat cells in a time- and dose-dependent manner [17].

Other studies have shown that TNFa causes pronounced alterations in glucose metabolism that are similar to those observed in patients with chronic inflammatory diseases. These changes include an enhanced utilization of glucose mainly by muscle but also by other tissues, and an increased glucose production [18-20]. However, under hyperinsulinaemic clamp conditons the amount of glucose required to maintain euglycaemia was significantly reduced indicating a state of insulin resistance. The ability of insulin both to suppress hepatic glucose production and to stimulate peripheral glucose utilization was impaired in the presence of $TNF\alpha$ [21]. Recent studies in 3T3 L1 adipocytes suggest that TNF α -induced insulin resistance could be due to a transcriptional suppression of the GLUT4 gene resulting in a total depletion of insulin-sensitive glucose transporters [22]. A direct role of TNF α in the development of insulin resistance has been implicated by a recent study in genetically obese rodents that indicates an association between an increased expression of $TNF\alpha$ in adipose cells and insulin resistance [23]. These recent findings and our observation of a delipidation of human fat cells by TNFa under long-term exposure prompted us to study the effects of $TNF\alpha$ on metabolic pathways that are involved in the regulation of lipid storage in human adipose tissue. For this purpose, we used the model of newly-differentiated human fat cells which can be cultured for extended periods of time.

Materials and methods

Chemicals. Human transferrin, triiodothyronine (T_3) , pantothenate, and bovine serum albumin (BSA) were purchased from Sigma (Munich, Germany). 1-methyl-3-isobutylxanthine was from Serva (Heidelberg, Germany). Collagenase CLS type 1 was obtained from Worthington (Freehold, N.J., USA). Semisynthetic human insulin and cortisol were kindly donated by Hoechst (Frankfurt, Germany). Fetal bovine serum and media were from Biochrom (Berlin, Germany). Recombinant human TNF α , antibiotics, trypsin-EDTA solution, glycerol kinase, glycerol 3-phosphate dehydrogenase, triosephosphate isomerase, flavin mononucleotide reductase and luciferase were obtained from Boehringer (Mannheim, Germany). Sterile plastic ware for tissue culture was purchased from Flow Laboratories (Irvine, UK).

2-Deoxy-D-(1-³H)glucose (15 Ci/mmol), (α -³²P)dCTP (3000 Ci/mmol) and ¹²⁵I-labelled protein A (30 mCi/mg) were from Amersham (Braunschweig, Germany). (γ -³²P)ATP (6000 Ci/mmol) was from New England Nuclear (Dreieich, Germany). Reagents for SDS-PAGE were supplied by Pharmacia (Freiburg, Germany) and Sigma. Polyclonal GLUT1and GLUT4-antisera were a product of Calbiochem (Bad Soden, Germany). All other chemicals were of the highest purity grade available and were purchased from Sigma and from Merck (Darmstadt, Germany).

Cell preparation and culture. Adipose tissue samples (20 to 80 g) were obtained from the mammary adipose of young normal-weight women (body mass index < 26 kg/m², age between 18 and 37 years) undergoing surgical mammary reduction. All subjects were otherwise healthy, and had no evidence of diabetes mellitus according to routine laboratory tests. The procedure for obtaining adipose tissue has been approved by the Ethical Committee of the University of Düsseldorf.

Stromal cells from human adipose tissue samples were isolated as described recently [24]. Briefly, the samples were carefully dissected from other tissues, all visible fibrous material and blood vessels were discarded. The remaining adipose tissue was minced into pieces of approximately 10 mg weight and digested in 10 mmol/l phosphate buffered saline (PBS) containing 1.5 mg/ml collagenase and 20 mg/ml bovine serum albumin, pH 7.4. To obtain full disaggregation the cells were isolated in a two-step procedure. After short centrifugation at $200 \times g$, the floating fat cells and the incubation solution were removed by aspiration. The sedimented cells were resuspended and incubated with an erythrocyte-lysing buffer consisting of 154 mmol/l NH₄Cl, 5.7 mmol/l K₂HPO₄, and 0.1 mmol/l EDTA for 10 min to remove contaminating erythrocytes. The dispersed material was filtered through a nylon mesh with a pore size of 150 µm. After additional washing and centrifugation steps, the sedimented cells were resuspended in Dulbecco's Modified Eagle's/Ham's F-12 medium (v/v, 1:1) supplemented with 10% fetal bovine serum and inoculated into 35mm or 100-mm dishes at a density of 30,000 to 50,000 cells per cm². After a 16-h incubation period, cultures were repeatedly washed with PBS to remove non-adhering material and re-fed with a serum-free medium consisting of Dulbecco's Modified Eagle's/Ham's F-12 medium (v/v, 1:1) supplemented with 15 mmol/l NaHCO₃, 15 mmol/l Hepes, 33 µmol/l biotin, 17 µmol/l pantothenate, 10 mg/l human transferrin, 100 U/ml penicillin, and 0.1 g/l streptomycin. To induce and maintain adipose differentiation, 100 nmol/l cortisol, 66 nmol/l insulin, 0.2 nmol/l T₃ and, for the first 3 days, 0.25 mmol/l MIX were added. The medium was changed every 2-3 days.

Glucose transport assay. 2-Deoxy-D-glucose uptake was determined as a measure of the glucose transport system. Assays were performed in newly differentiated human fat cells cultured in 35-mm dishes. Three days prior to the assay, both insulin and glucose concentrations of the medium were reduced to 20 pmol/l and 5 mmol/l, respectively, and Hepes was completely removed from the medium. Twenty-four hours before the assay, cells were washed and re-fed with an insulin-free medium. For assessment of the stimulatory effect of insulin on glucose uptake, some dishes were preincubated with 100 nmol/l insulin for 15 min immediately before the assay. (³H)-labelled 2deoxy-D-glucose (1 µCi/dish, concentration 4.5 µmol/l) was added to the medium which already contained 5 mmol/l glucose and uptake was measured at 37 °C for 20 min. Glucose transport was terminated by transferring the dishes to an ice-bath. Cells were repeatedly washed using ice-cold PBS and incubated for 20 min with 0.1 % SDS. The radioactivity of the cell material was counted in a liquid scintillation counter (Beckman, Munich, Germany). Values were corrected for the unspecific uptake which was assessed by incubating cells with labelled 2deoxy-D-glucose for 20 min at 4°C. A similar unspecific rate of 2-deoxy-D-glucose uptake was obtained, when cells were incubated with 25 µmol/l cytochalasin B for 20 min (data not shown).



Fig. 1(A, B). Photomicrographs of primary cultures of stromal cells from human adipose tissue 16 h after inoculation (A) and 16 days after exposure to an adipogenic serum-free medium (B) as described in Methods. Magnification $125 \times$

Preparation of crude membrane fraction. Cells were twice washed in TES buffer (225 mmol/l saccharose, 1 mmol/l EDTA, 20 mmol/l Tris, 2.5 µg/ml leupeptin, 2.5 µg/ml pepstatin, 2.5 µg/ml aprotinin and 0.2 mmol/l phenylmethylsulfonyl fluoride, pH 7.4). Then cells were scraped from the dishes using a rubber policeman, immediately frozen in liquid nitrogen und stored at -70 °C. For analysis, the cell material of one sample was homogenized in 10 ml TES buffer using a teflonglass-homogeniser at 4°C. The first centrifugation was carried out at $1000 \times g$ for 10 min., the supernatant was centrifuged again at $100,000 \times g$ for 90 min. The membrane pellet was resuspended in TES buffer. After a new homogenisation step using a glass-glass-homogenizer the material was stored in aliquots at - 70 °C. The protein content of the samples was determined according to a modification of the Biorad protein assay (Biorad, Munich, Germany).

Western blotting. Prior to Western blotting cells were treated as described for the glucose transport experiments except that 100-mm dishes were used. Membrane protein samples (30 to 50 μ g) were subjected to SDS-PAGE and transferred to nitrocellulose filters in a semidry blotting apparatus. Filters were blocked for 10 min in Tris-buffered saline (20 mmol/l Tris/HCl, 137 mmol/l NaCl, 1 CaCl₂, pH 7.6) containing 0.05 % Tween-20 and 5 % BSA. Then, filters were incubated for 16 h at 4 °C with a 1 : 500 dilution of a polyclonal GLUT4 antiserum. After extensive washing with Tris-buffered saline containing 0.05 % Tween-20 and 0.5 % BSA, filters were incubated for 2 h with ¹²⁵I-protein A (0.3 μ Ci/ml) at room tempera-

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ture. Filters were again extensively washed, airdried and exposed to Hyperfilm-MP films using intensifying screens. Autoradiographs were quantified by laser scanning densitometry (LKB, Gräfelfing, Germany).

Northern blotting. Total RNA was isolated from the cell material using the guanidinium thiocyanate/phenol/chloroform method [25], separated by 2% agarose gel electrophoresis and transferred to nylon membranes as described previously [26]. Densitometry of ethidium bromide-stained 18 S rRNA was used to normalize the amount of RNA loaded per lane. Blots were hybridized either with the oligonucleotide GLUT4 probe which was 5'-labelled to a specific radioactivity of 1.11×10^8 dpm/µg by using (³²P)-ATP and T4-kinase or to a LPL cDNA probe (kindly provided by Dr. A. Singh, Genen-tech, San Francisco, Calif., USA) multiprime labelled to a specific activity of 2.2×10^8 dpm/µg by using (³²P)-CTP. After washing under high stringency conditions and air-drying, the blots were exposed to MP Hyperfilms at - 70 °C using intensifying screens. The autoradiographs were quantified by a FU-JIX BAS 1000 bioimaging analyser (Fuji, Tokyo, Japan) to determine the relative amounts of GLUT4 and LPL mRNAs.

Lipolysis. The glycerol concentration of the culture medium was used as a measure of lipolysis. The glycerol content was determined by an established bioluminescent method using ATP-linked firefly luciferase [27]. Glycerol concentrations are expressed a μ mol glycerol \cdot mg protein⁻¹ · 24 h⁻¹, but are given as percentage of glycerol concentrations of control cultures.

Statistical analysis

Results are expressed as mean \pm SEM of at least three experiments in duplicate. Analysis of variance was used for statistical analysis. Comparisons between TNF α -treated and untreated cultures were performed using Student's *t*-test for paired data. *p*-values less than 0.05 were considered as statistically significant.

Results

All experiments on the metabolic effects of TNFa were performed in cultured human adipocyte precursor cells from human adipose tissue after having acquired the adipocyte phenotype under serum-free, hormone-supplemented culture conditions. After 16 days in culture between 50 and 80% of the cells were differentiated as assessed by microscopic lipid accumulation (Fig. 1) and expression of glycerophosphate dehydrogenase (GPDH) activity. During the following days the functional state of the cells remained stable as assessed by GPDH activity, the morphological appearance of the cells became more spherical with fewer but larger lipid droplets (data not shown).

Effect of $TNF\alpha$ -treatment on 2-deoxy-D-glucose transport. 3 days prior to the glucose transport experiments the newly developed fat cells were changed to a medium devoid of cortisol and T₃ but containing



Fig.2(A, B). Effect of TNF α on basal and insulin-stimulated 2-deoxy-D-glucose transport in cultured human adipocytes. Cells were prepared for the transport study assay as described under Methods. 2-Deoxy-D-glucose uptake was measured with or without 15-min stimulation of 10^{-7} mol/l insulin after a 24-h (A) or 72-h (B) preincubation with 5 nmol/l TNF α Data are presented as mean ± SEM of four experiments in duplicate. * p < 0.01

 2×10^{-11} mol/l insulin. During the final 24 h cells were cultured in a completely insulin-free medium in the absence or presence of 5 nmol/l TNFa A 24-h exposure of the cells to TNFa had no effect on the basal rate of hexose transport but completely abolished the stimulatory action of 10^{-7} mol/l insulin on glucose uptake (Fig.2). A 72-h incubation of human fat cells with TNFa resulted in a significantly elevated basal rate of glucose uptake by $71 \pm 13\%$ (p < 0.01). Again, preincubation with insulin failed to increase the rate of 2-deoxy-D-glucose transport (Fig.2).

Effect of TNF α on the protein expression of the glucose transporters GLUT4 and GLUT1. To determine whether the reduced uptake of 2-deoxy-D-glucose in response to insulin might result from a reduced number of GLUT4 molecules, the effect of TNF α on the



Fig. 3. Effect of TNF α on the cellular content of GLUT4 and GLUT1 protein. Human adipocytes were incubated for 24 and 72 h, respectively, with 5 nmol/l TNF α . Total membrane fractions were subjected to immunoblotting with specific antisera, as described in Methods

cellular GLUT4 protein content was studied. Total membrane fractions from TNFa-treated and untreated cells were prepared and subjected to immunoblotting using a specific antiserum to identify GLUT4. The experiments revealed that TNFa caused a marked reduction in the number of glucose transporters (Fig.3). Densitometric analysis confirmed a significant decrease in GLUT4 protein. A 24-h exposure of human fat cells to 5 nmol/l TNFa resulted in a reduction of the GLUT4 amount by $49 \pm 3\%$, a 3-day exposure to 5 nmol/l TNF α decreased the amount of GLUT4 by approximately 82 ± 18 %. In contrast, TNF α treatment for 24 h did not significantly alter the amount of GLUT1 protein (data not shown). A 72-h exposure of fat cells to TNFα resulted in an almost twofold increase in the total amount of GLUT1 (Fig. 3).

Effect of TNF α on GLUT4 mRNA. To examine whether the decrease in the number of GLUT4 transporters in the presence of TNF α was due to a reduced expression of the specific mRNA, Northern blotting was performed to determine the level of GLUT4 mRNA. These experiments demonstrated that exposure of cultured human adipocytes to 5 nmol/l TNF α dramatically decreased the cellular amount of GLUT4 mRNA. As shown in Figure 4A, a 24-h as well as a 72-h incubation with the cytokine was associated with a disappearance of specific mRNA for GLUT4.

Effect of $TNF\alpha$ on LPL mRNA. Since the amount of LPL plays a critical role in the uptake of triglycerides by fat cells, the effect of TNF α on the cellular content of LPL mRNA was investigated in additional experiments. After a 24-h incubation with 5 nmol/l TNF α the amount of LPL mRNA detected by Northern blotting was reduced by approximately 70%. A 72-h exposure of human adipocytes to the same concentration of TNF α resulted in an only moderately



Fig.4A,B. Effect of TNF α on GLUT4 (**A**) and LPL (**B**) mRNA content in cultured human fat cells. Adipocytes were exposed to 5 nmol/l for 24 or 72 h, respectively. Total RNA was electrophoresed, transferred to a nylon membrane and hybridized with specific ³²P-labelled probes. One representative experiment is shown

higher suppression of LPL mRNA levels by approximately 80 % compared to control cultures (Fig. 4B). These experiments clearly indicate that, similar to its effect on GLUT4, TNF α apparently regulates the amount of LPL at the level of gene expression.

Lipolysis. Another possibility by which TNFa could cause a loss of stored lipids in human adipocytes over time involves a stimulation of the lipolytic system. We therefore investigated the effect of $TNF\alpha$ on the glycerol concentrations in the culture medium. It became apparent that TNFa potently increased lipolysis. When cells were exposed for 24 h to varying concentrations of the cytokine in the presence of 20 pmol/l insulin an up to fivefold increase of the glycerol content was observed compared to control cultures (Fig. 5). Even at 0.1 nmol/l TNF α the glycerol content of the culture medium was increased by 3.5-fold. However, the lipolytic activity of TNFa became apparent only after 6 h suggesting that this effect does not occur via a rapid direct activation of the adenylate cyclase system and subsequent regulatory proteins (Fig. 6).

In all experiments with TNFo, treatment of cultures with the cytokine was not associated with de-



Fig. 5. Dose-response-relationship of the effect of TNF α on lipolysis in cultured human adipocytes. Cells were exposed to varying concentrations of TNF α for 24 h in the absence of insulin. The glycerol content of the culture medium was measured using a bioluminescence method. Results are expressed as percentage of control cultures (mean ± SD: 116 ± 17 µmol/mg protein/24 h = 100 %). Data represent mean ± SEM of five separate experiments in triplicate. * p < 0.05 vs cultures not exposed to TNF α



Fig.6. Time-course of the effect of TNF α on lipolysis. Human adipocytes were exposed to 5 nmol/l TNF α for the time periods indicated. The glycerol content of the medium was measured as described under Methods. Results are given as percentage of the respective control cultures. Data represent mean \pm SEM of four experiments in triplicate. * p < 0.05 vs cultures not exposed to TNF α

tectable changes of cell morphology. In particular, there was no morphological evidence for a cytotoxic effect of $TNF\alpha$ (data not shown).

Discussion

The results of this study clearly demonstrate that TNF α exerts diverse effects on the main metabolic functions in human adipocytes. TNF α abolished the stimulatory action of insulin on glucose uptake and decreased the cellular GLUT4 protein content. In addition, our experiments indicate that TNF α also reduces LPL mRNA levels and – in a dose-dependent manner – stimulates the mobilization of stored triglycerides. The combination of these effects may be sufficient to explain the marked delipidation of human adipocytes during long-term exposure to TNF α as reported recently [17].

To study the action of TNF α on human adipose tissue, we used a new cell culture model that is based upon the in vitro differentiation of stromal cells obtained from human adipose tissue samples. In material from young adults up to 80 % of these cells undergo differentiation and develop the biochemical and morphological characteristics of mature fat cells. They express high lipogenic enzyme activing, establish a glucose transport system consisting of GLUT1 and the insulin-responsive GLUT4 subtype and are able to release stored lipids upon stimulation by catecholamines (Hauner et al., unpublished observation). Although most cells exhibit a multilocular morphology, this culture system appears to be a suitable and reliable model to study human fat-cell metabolism for extended periods of time.

One of the best studied effects of $TNF\alpha$ in adipose tissue is suppression of LPL activity [3–8]. The reduced amount of LPL mRNA also found in our study may be due to a specific decrease of LPL gene transcription as originally shown by Zechner et al. [6] in 3T3 L1 adipocytes. However, previous studies on the effect of TNFa on LPL activity and gene expression in human adipose tissue have given conflicting results. In isolated fat cells kept in suspension culture, Kern [15] did not find a change in LPL mRNA induced by TNFo, whereas Fried and Zechner [16] reported a pronounced suppression of LPL synthesis and activity in cultured pieces of human adipose tissue. Our data obtained in fat cells newly developed from precursor cells clearly confirm that $TNF\alpha$ decreases LPL at the mRNA level. It is obvious that the suppression of LPL mRNA by TNFa administration may substantially contribute to the alterations in lipoprotein metabolism, particularly hypertriglyceridaemia, that are frequently seen in patients with chronic infectious diseases or cancer [1, 2].

An inhibitory effect of TNF α on GLUT4 expression was originally described in rat L6 myotubes [28]

and clonal preadipocytes [22, 23]. Our experiments clearly extend this observation to human fat cells kept in primary culture. However, the rapid loss of response to insulin seen in the 2-deoxy-D-glucose transport assay cannot be fully explained by the reduction of GLUT4 protein. Recent data indicate that TNFa may also interfere with the transmembrane signalling of insulin. In rat hepatoma cells, a 1-h incubation with TNFa was followed by a dramatic decrease in the insulin-induced tyrosine phosphorylation of both the insulin receptor β -subunit and IRS-1, its major cytosolic substrate without any reduction in insulin binding [29]. Similar results have been recently reported in 3T3 L1 and 3T3 F442A adipocytes [30]. We therefore assume that the loss of insulin responsiveness observed after a 24-h exposure to TNFa results from both suppression of GLUT4 gene expression and interference with the insulin signal transduction. However, whether the latter mechanism is active in human adipose tissue or whether other factors such as an increase of the lipolytic rate may be of importance for the loss of insulin action, remains to be demonstrated.

Interestingly, after a 3-day exposure to TNF α the basal rate of glucose uptake increased by almost twofold. In conjunction, we also measured a similar increase in the total amount of GLUT1 protein. Experiments in quiescent 3T3 fibroblasts indicate that TNF α can enhance glucose uptake by different mechanisms including stabilization of GLUT1 mRNA and alterations in the intrinsic activity of the glucose transporters [31, 32].

Previous studies have revealed that TNFa is also able to stimulate lipolysis in cultured 3T3 L1 adipocytes [4, 9–12]. We describe here for the first time that TNF α also promotes lipid release from human fat cells. Other evidence for a lipolytic action of TNF α in man was provided from an in vivo study, in which intravenous infusion of TNFa in patients with disseminated cancer resulted in an increase of nonesterified-fatty acid and glycerol levels [33]. The time-course and dose-response relationship for the TNFa effect on lipolysis in human fat cells was similar to those described in 3T3 L1 adipocytes [11]. The delayed stimulation of glycerol release by TNFa indicates that this effect may depend on new protein synthesis. A similar time-course for the effect of TNFa on lipolysis has been recently reported in cultured rat adipocytes with no effect on the expression of hormone-sensitive lipase, the rate-limiting enzyme of lipolysis [12]. A possible role of prostaglandins in mediating the lipolytic action of $TNF\alpha$ has been suggested [11], but this conclusion was withdrawn in a subsequent study [34].

The disturbances of carbohydrate and lipid metabolism observed in this in vitro study of human fat cells may be similar to those present in insulin-resistant states such as NIDDM and obesity. Recent studies in genetically obese rodents have suggested a link between an elevated expression of TNF α in adipose tissue and the development of insulin resistance and non-insulin-dependent diabetes. The importance of this association was substantiated by the finding that neutralization of TNF α by a soluble receptor significantly ameliorated insulin sensitivity on glucose utilization [23].

In conclusion, this study demonstrates profound effects of TNF α on metabolic functions of cultured human fat cells. This cytokine suppresses LPL mRNA, potently stimulates lipolysis and was found to abolish the insulin effect on glucose transport. The combination of these effects may be responsible for the delipidation of cultured human fat cells observed under chronic exposure to high concentrations of TNF α and may also contribute to the development of insulin resistance in states of increased TNF α -production.

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