# Regulatory effect of various steroid hormones on the incorporation and metabolism of [<sup>14</sup>C]stearate in rat hepatoma cells in culture

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## Abstract

We have examined the incorporation and metabolism of [<sup>14</sup>C] stearic acid within the total lipids of HTC rat-hepatoma cells in suspension culture in presence and in absence of steroidal hormone stimulation. Both, glucocorticoids (dexamethasone, cortisol and corticosterone) and mineralocorticoids (deoxycorticosterone and aldosterone) as well as the estrogen  $\beta$ -estradiol and the androgen testosterone enhanced the extent of  $\Delta 9$  desaturation to oleic acid of the saturated precursors, whereas only the two mineralocorticoids affected the incorporation rate of the exogenous acid into total cellular lipids, thus promoting a little stimulation. Furthermore, all the hormones tested increased the radiolabelling of the total cellular phospholipids except deoxycorticosterone and testosterone, the former having no effect and the latter exerting a moderate inhibition. On the other hand, the incorporation of <sup>14</sup>C into neutral lipids was stimulated by testosterone, in contrast to the inhibition of this parameter observed exclusively with either the mineralocorticoids or the estrogen. Within the phospholipid subclasses, the radiolabelling of phosphatidylcholine was augmented by means of all the steroids tested save deoxycorticosterone and testosterone, whereas phosphatidylethanolamine exhibited a decrease only in the presence of testosterone. In a similar fashion, within the neutral lipids, the predominating triglyceride fraction was preferentially labelled – at the expense of other subclasses of lesser abundance – upon treatment with the steroids except aldosterone, which exerted no effect. The results obtained were correlated with those changes observed in the mass distribution of the different lipid subclasses either with or without prior hormonal stimulation. (Mol Cell Biochem **145**: 1–9, 1995)

Key words: rat-hepatoma, steroidal hormones, lipid metabolism

Abbreviations: ALD – aldosterone; ANOVAR – analyses of variance; CE – cholesterol esters; Cl – cortisol; CLN – cardiolipin; Cst – corticosterone; CH – cholesterol; DG – diacylglycerides; DOC – 11-deoxycorticosterone; Dx – dexamethasone phosphate salt;  $\beta$ E – 17- $\beta$ -estradiol; FAME – fatty acid methyl esters; GLC – gas-lipid chromatography; HEPES – N-2-hydroxyethyl-piperazine-N-2-ethanosulfonic acid; HTC – hepatoma tissue culture; IMEM-Zo – improved minimal essential medium-Zinc option; LPC – lysophosphatidyl-choline; LPL – lysophospholipids; MG – monoacylglycerides; NEFA – free fatty acids; PC – phosphatidyl-choline; PE – phosphatidyl-ethanolamine; PG – phosphatidyl-glycerol; PI – phosphadyl-inositol; PS – phosphadidyl-serine; SM – sphingomyelin; TG – triacylglycerides; TLC – thin-layer chromatography; Tst – testosterone

## Introduction

It is well known that palmitic and stearic acids are desaturated

at the  $\Delta 9$  position (probably by two independent enzymes) to give palmitoleic and oleic acids, respectively [1]. Each one of these monounsaturated acids then becomes the initial pre-

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cursor for the biosynthesis of an independent family of polyunsaturates [2, 3]. The  $\Delta 9$  desaturase may also be considered as a lipogenic enzyme, whose function is to desaturase dietary or newly synthesized saturated fatty acids for deposition primarily in triacylglycerides [1, 4, 5]. Other studies have indicated that palmitoyl- and stearoyl-CoA desaturases play a relevant role in providing monoenoic acids for export as lipoprotein components [6, 7]. These families of desaturating enzymes also contribute to maintaining the appropriate fluidity in the phospholipids of biological membranes which structures in turn, regulate a wide variety of membrane-associated events [6, 8–10].

The desaturation rate at  $\Delta 9$  position would also seem to be highly dependent on the hormonal state of the animal. In microsomal preparations of rat or avian livers  $\Delta 9$  desaturase activity is significantly stimulated by insulin [1, 5, 11] or thyroxin [1, 11, 12] administration. Moreover, the effect of the natural estrogen 17-\beta-estradiol was investigated by Lippiello and co-workers in rooster liver [6] and by Kasturi and Joshi in the murine 3T3-L1 cell line [5]. These researchers have reported that the biosynthesis of monoenoic fatty acids was stimulated by estradiol supplementation. In addition, we have also reported that  $\Delta 9$  desaturase was strongly stimulated by a soluble protein induced by dexamethasone [13] or 11-deoxycorticosterone [14] treatment. By contrast the effect of other steroid hormones on this desaturase enzyme has not been investigated in mammals. The study of the hormonal regulation of  $\Delta 9$  desaturase activity in whole animal is often complicated by the occurrence of indirect changes in the concentration of hormones and other biological molecules in blood. The use of isolated cells maintained in a chemically defined culture medium, i.e. without serum, would thus prove of fundamental importance in the system under investigation in order to determine the primary physiological effect of different steroid hormones on the regulation of either lipid metabolism in general or stearoyl-CoA desaturase activity in particular. Of further advantage was the fact that the HTC-cell system used for these studies responds to physiological concentrations of various steroids of different chemical structures and diverse biological activities.

## Materials and methods

#### Chemicals

[1–<sup>14</sup>C]Stearic acid (58.9 mCi/mmol, 99% pure) was from New England Nuclear Corp., Boston, MA, USA. Unlabelled stearic and eicosa-11-enoic acids were provided by Nu-Chek Prep, Elysian, MN, USA. The acids were stored in benzene under an atmosphere of nitrogen at 20°C. Concentrations and purities were confirmed by both liquid-scintillation counting and the gas-liquid chromatography (GLC) of fatty acid methyl esters prepared in the presence of appropriate internal standards. Neutral lipids and phospholipids standards for thinlayer chromatography (TLC) were obtained from Serdary Research Laboratory Inc., London, Ontario, Canada. Fattyacid methyl-ester mixtures for GLC identifications, defatted bovine serum albumin and HEPES (N-2-hydroxyethyl-piperazine-N-2-ethanosulfonic acid) were from Sigma Chemical Company, St. Louis, MO, USA. Dexamethasone (sodium phosphate salt) was from Merck, Sharp and Dhome, Buenos Aires, Argentina. Other steroid hormones used were obtained from Sigma Chemical Company. All chromatographic solvents were RPE grade and obtained from Carlo Erba, Milan, Italy. Other chemicals used were reagent grade and provided by commercial sources.

#### Cell culture

The hepatoma cells used in these experiments were from the Morris minimal deviation hepatoma number 7288c, a solid hepatoma from which the ascites-tumor line (HTC cells) had been derived [15]. HTC cells were maintained and grown *in vitro* at 37°C in surface culture on Swim's 77 (S-77) medium supplemented with 10% (v/v) calf serum, under conventional sterile conditions as previously described [16].

#### Assay procedure

Monolayers of HTC cells in the logarithmic growth phase (approximately 72 h) were detached from the growing surface with 0.1% (w/v) trypsin [17]. They were then suspended in modified IMEM-Zo medium [18] in which the CaCl, concentration was reduced to 1.5 mM, linoleic acid was omitted and HEPES was added to 5-mM final concentration. Soybean trypsin inhibitor, type I (0.1%, w/v) was then added during the following washing steps. The cell suspension was sedimented twice by centrifuging in 20 ml of modified IMEM-Zo medium at 4–5°C for 10 min (500  $\times$  g). The final pellet was resuspended in the same medium and counted in a hemocytometer. Aliquots (5  $\times$  10<sup>6</sup> cells) were finally incubated at 37°C under an atmosphere of 95% oxygen and 5%, (v/v) carbon dioxide in a metabolic shaker (70 rpm). The incubation volume was 5 ml in 25 ml siliconized Erlenmeyer flasks. Cell viability (> 96%) was assessed by the criterion of trypan-blue dye exclusion [19]. Two types of experiments were designed in order to study the effect of various steroid hormones on the incorporation and desaturation of stearic acid as well as the distribution of the endogenously generates oleate among the different lipid subclasses. In these experiments, the cells were incubated for 12 h in the presence of dexamethasone (Dx), cortisol (Cl), corticosterone (Cst),

11-deoxycorticosterone (DOC), aldosterone (ALD),  $17-\beta$ estradiol (BE) or testosterone (Tst), at 1.0 µM concentration. Three hours before the end of the incubation period, the media were supplemented with a mixture of labelled and unlabelled stearic acid 80 µM (0.5 µCi/flask). The acid was added as the sodium salt bound to defatted bovine serum albumin, according to Spector et al. [20] in a ratio of 6 nmol fatty acid to 1 nmol albumin. One set of control cells was supplemented with delipidated albumin at the same concentration as the fatty-acid-supplemented group; while another group of controls was treated identically, but with the omission of hormones. These latter flasks were, however, supplemented with the vehicle in which the hormones were dissolved (ethanol at 20 mM). In all instances, at the end of the incubation period, the cell suspensions were transferred to icecold tubes and centrifuged at 500 × g for 5 min. Each pellet was washed three times with 5 ml of cold 0.85% (w/v) NaCl and resuspended in the same solution. An aliquot of the cell suspension was used to determine cell viability and the amount of cellular protein [21]. The rest was centrifuged at  $2,000 \times g$  for 10 min, the supernatant was discarded, and each cell pellet was processed separately for lipid analysis.

#### Cellular-lipid extraction and separation

Lipids were extracted from the cell pellets by means of the method of Folch et al. [22]. The organic phases, containing more than 97% of the total radioactivity, were further separated by two chromatographic procedures. In the first system, the neutral lipids were separated by TLC on 0.25 mm thick silica gel G-60 plates (heat-activated) in a solvent system containing hexane:diethylether:acetic acid (80:20:1, v:v:v) [23]. Under these conditions, the phospholipids remained at the origin. The individual neutral lipids separated in this run were identified by comparison with a standard mixture containing (Rf values in parenthesis) monoacyl-glycerides (MG) (0.10), diacylglycerides (DG) (0.20), cholesterol (CH) (0.37), free fatty acids (NEFA) (0.40), triacylglycerides (TG) (0.72), fatty-acid methyl esters (FAME) (0.81), and cholesterol esters (CE) (0.90), which pure species were applied to each plate. Phospholipids were separated in a second TLC system [24] by two successive developments in the same direction. The first mobile phase was chloroform:methanol:ammonium hydroxide:water (70:25:3.5:1.5, v:v:v:v) while the second one was chloroform:methanol:acetic acid:water (80:10:2:0.75, v:v:v:v). The different subclasses of phospholipids were identified by reference to the following standard mixture developed in parallel with the samples: lysophosphatidyl choline (LPC) (<0.13), phosphatidylserine (PS) (0.19), phosphatidylinositol (PI) (0.25), sphingomyelin (SM) (0.30), phosphatidylcholine (PC) (0.39), phosphatidylethanolamine (PE) (0.55), cardiolipin (CLN) (0.69), and phosphatidyl-glycerol (PG) (0.71).

#### Lipid analysis

The plates were firstly developed and then dried under nitrogen. Lanes containing samples were covered with a glass plate, while the lipid markers were briefly sprayed with a tincture of iodine 1.5% (w:v) in methanol. Areas of interest were scraped off the plates and isolated lipids recovered by elution with chloroform/methanol, (1:1, v:v). Once dried, the samples were dissolved in 500 µl of cold chloroform/methanol, and aliquots were taken for liquid-scintillation counting in a Wallac 1214 Rackbeta liquid scintillation counter (Pharmacia, Turku, Finland) (97% efficiency for <sup>14</sup>C), interfaced to an Olivetti M-240 computer system. The plates were also scanned for radioactivity detection (radio-TLC) by means of a TLC-Proportional Radioactivity Scanner (Berthold LB-2832, Wildbad, Germany) equipped with a Hewlett-Packard 3396-A Data Station. No significant differences were found between radioactivity quantification by TLC-scanning or by liquid-scintillation counting. Some plates were charred [25] and quantified densitometrically (Karl Zeiss, Germany) according to the method of Katz et al. [26]. Aliquots of the original lipid extracts were dried and subjected to trans-esterification by heating in 2.5 ml 4 N HCl/ methanol at 68°C for 3 h under nitrogen. The resulting FAME were extracted three times with 2 ml of hexane. Organic phases from each tube were pooled and evaporated under nitrogen. The residue were prepared for GLC analysis and dissolved in 20-30 µl of hexane containing 1 µg/tube of eicosa-11-monoenoic acid as an internal standard. The GLC of FAME was performed in an Acromat GC-100 apparatus (Redhill, Surrey, England) with a 6 ft glass column packed with 10% SP-2330 on 100-200 mesh Chromosorb WAW-DMCS (Supelco Inc, Bellefonte PA), USA. After 4 min initial hold, the oven was programmed from 140-220°C at 1.5°C/min. The distribution of radioactivity among the FAME was determined by means of a Packard 894 Gas-Flow Proportional Counter (Downers Grove, Ill, USA) interfaced to the chromatograph through a nominal 10:1 splitter. A linear dual-pen recorder (LKB-2210 Two-Channel Recorder, Bromma, Sweden) was used to obtain simultaneously radioactivity and mass tracings for each chromatographic separation. The individual FAME were identified by comparison of their relative retention times with authentic standards, and the mass distribution was calculated electronically by quantification of the peak areas.

All results were expressed as the means and standard error calculated from 3-5 independent analyses. Paired or unpaired student's *t* test and 'ANOVAR' were used for statistical evaluations. p< 0.01 was considered significantly different when compared to the corresponding control group.

## **Results and discussion**

We have firstly examined the effect of physiological concentrations of various steroidal hormones on the uptake and  $\Delta 9$ desaturation of [14C]stearic acid in suspensions of HTC cells (Fig. 1). Although the effect of Tst was not statistically significant, this androgen exhibited a marked tendency to inhibit the uptake of the exogenous acid. Furthermore, from all the hormones tested in this system, only DOC and ALD increased the uptake of labelled stearic acid in a significant fashion (Fig. 1, solid bars). Such increment in the incorporation of this exogenous metabolic precursor could be explained by an increase in its subsequent utilization. Mineralocorticoids have been reported to produce a general activation of lipid metabolism, and ALD, in particular, was shown to increase the degradation of specific phospholipid fatty acids [27, 28] and to stimulate oleic-acid metabolism [29] as well as phospholipase A, activity [30] in urinary bladder. These effects, occurring within 30 min of the beginning of the hormone treatment and persisting for up to 6 h, lead to a depletion in the intracellular level of stearic acid [29]. By contrast Tst caused a decrease in the level of sterified fatty acids in human aortic smooth-muscle cells along with a reciprocal increment in their free-fatty acid content [31]. If the androgen acts in a similar fashion in our system, the resulting major availability of free fatty acids might well explain the decreased uptake of exogenous stearic acid that we observed (Fig. 1).

Despite several papers about the stimulation of lipid metabolism by glucocorticoids, specifically an increase in the biosynthesis of phosphatidylcholine [32, 33] and triacylclycerides [34], the uptake of labelled stearate into cellular lipids was not significantly modified under Dx, Cl or Cst treatments in our studies (Fig. 1). In view of these findings, we decided to investigate the effect of glucocorticoids on the incorporation of [<sup>14</sup>C]18:0 into neutral or polar lipid subclasses; the results obtained will be discussed below.

All the hormones tested evoked a significant increase in cellular  $\Delta 9$  desaturation (Fig. 1, hollow bars) in contrast to the results obtained for the fatty acid uptake. The increase in oleic acid formation upon exposure of the HTC cells to hormones was 50% with glucocorticoids, 65% with Tst or mineralocorticoids, and ca. 100% with BE with respect to control values. A similar stimulation of  $\Delta 9$  desaturase activity was previously found in our laboratory with liver microsomal preparations from rats injected with physiological doses of Tst [35] or Dx phosphate [13]. Moreover, González Rodríguez and co-workers demonstrated that the activity of the liver microsomal  $\Delta 9$  desaturase was markedly enhanced after the injection of  $\beta E$  to ovariectomized rats [36]. Lipiello et al. [6] reported that the same estrogen, when administered to roosters, significantly increased the activity of the enzyme with stearic acid as substrate. The stimulatory effects caused



Fig. 1. Effect of the treatment of HTC cells with various steroid hormones on the uptake ( $\blacksquare$ ) and  $\Delta 9$  desaturation ( $\bullet$ ) of stearic acid. Aliquots of HTC cells suspended in IMEM-Zo medium (1. 106 cells/ml) were incubated for 12 h at 37°C in the presence or absence of different steroid hormones at a final concentration of 1 µM. During the last 3 h of incubation a mixture of labled and unlabelled sodium stearate (80  $\mu$ M, 0.50  $\mu$ Ci/ flask) was added to all flasks. At the end of the experiment, the cells were centrifuged (10 min at 500  $\times$  g) and washed by a second sedimentation. The pellets were extracted by the method of Folch et al., and the lipid extracts were processed to determine the uptake of  $^{14}C$  and  $\Delta 9$  desaturation of the substrate. For details see Materials and methods section. The results were expressed as the percent change with respect to the mean value for the two controls:  $40.0 \pm 0.7$  nmol of stearic acid incorporated per mg of cellular protein and  $33.2 \pm 0.7$  nmol of oleic acid produced per mg of cellular protein. Each value was the mean of 4-5 independent determinations. Vertical lines represent 1 SEM. Asterisks indicate values significantly different from controls at p < 0.01.

by the other steroidal hormones studied in the present experiments (Fig. 1) have not been previously reported.

These changes in desaturase activity observed in the presence of gluco- or mineralocorticoids would involve modifications in the fatty acid profiles of target tissues. Indeed, Lien et al. reported that ALD administration to toads simultaneously decreased the content of saturates and increased the levels of monoenoates in their urinary bladders [29]. We also observed similar alterations in the hepatic microsomal lipids obtained from rats treated with either ALD or DOC [37] as well as in the total lipids of HTC cells incubated with Dx phosphate [38]. In the present experiments, we also found that all the steroidal hormones studied were able to produce the same analytical changes in the saturated/monoenoic fatty acid ratio. All the hormones tested also modified the fatty acid profiles which were characterized by a ca. 40% increase in the amount of monounsaturates over control values (Data not shown).

The marked changes evoked by the steroidal hormones on the uptake and metabolism of the stearic acid and the important modifications in the fatty acid composition prompted us to examine the distribution of the labelled substrate into polar and neutral lipids (Fig. 2) and to characterize its distribution among the main lipid subclasses (Fig. 3). Accordingly, our study on the incorporation of [<sup>14</sup>C]stearate into polar lipids of HTC cells treated with various steroids demonstrated that



Fig. 2. Effect of the treatment of HTC cells with various steroid hormones on the incorporation of [<sup>14</sup>C]stearic acid into neutral lipids ( $\blacksquare$ ) or phospholipids ( $\square$ ). Suspensions of HTC cells were incubated under conditions similar to those described in Fig. 1. At the end of the experiment, the cells were centrifuged and washed. Cellular lipids were extracted by the method of Folch *et al.* Phospholipids and neutral lipids were separated by TLC. The plates were scanned for radioactivity detection; and then the lipids were scraped off the plates and reextracted in chloroform:methanol (1:1, v:v) mixture. Aliquots of these extracts were taken for liquid-scintillation counting. For details see *Materials and methods*. Each value was the mean of 4 independent analyses and is expressed as the per cent change with respect to the corresponding control:  $855.10^3$ and  $420.10^3$  dpm/mg of cellular protein for the uptake of total radioactivity into phospholipids and neutral lipids, respectively. Vertical lines represent 1 SEM.

all the hormones tested caused an increase in the incorporation of label ranging from 46-91% above control levels with the exception of DOC, which produced not significant changes, and Tst, which inhibited the uptake of [14C]stearate by about 30% (Fig. 2). With respect to neutral lipids, the glucocorticoid hormones (Dx, Cl or Cst) produced no significant modifications in the incorporation of labelled substrates (stearic acid and/or its direct metabolite oleic acid), while the mineralocorticoids, DOC or ALD, decreased the uptake of this radioactivity by ca. 40%. The effect caused by 17-bestradiol was opposite to that of testosterone, the androgen being the only steroidal species produces an enhancement of label in the HTC-cell phospholipids. Thus, while estrogen inhibited the uptake of <sup>14</sup>C by approximately 30%, Tst produced a 40% increase in the incorporation over the control data. Hence, the effect of testosterone was of particular interest because that hormone exhibited an action opposite to that seen with the others. In those cells exposed to testosterone, the ratio of <sup>14</sup>C incorporation into neutral vs. polar lipid fractions was 2 to 14-fold higher than the values obtained with any of the other steroid hormones studied.

The results shown in Fig. 2 could suggest that glucocorticoid hormones produce an increase in phospholipid biosynthesis with minor changes in the metabolism of neutral lipids. Since changes in the incorporation of <sup>14</sup>C into one particular lipid subclass upon hormonal treatment could be offset by variations in another one, it is important to consider the data



Fig. 3. Effect of steroid hormones on the incorporation of [14C]stearic acid into different cellular lipids. Aliquots containing 5.106 HTC cells suspended in IMEM-Zo medium, were incubated and harvested as described in the legend to Fig. 1. The lipids from each pellet were then extracted by the method of Folch et al., and the lipid subclasses were separated by TLC as stated in the text. The plates were next scanned for radioactivity detection and finally sprayed with iodine solution to reveal the position of each lipid species. After calculation of the respective Rfvalues, the spots were scraped off the plates and prepared for liquidscintillation counting. Results were expressed as the per cent change with respect to the corresponding control values (all expressed as dpm.103/mg of cellular protein, with the lipid subclasses in parenthesis): 314.6 (PE), 244.5 (PC), 42.7 (CLN+PG), 106.8 (NEFA), 243.6 (TG), and 52.5 (CE). Any lipid subclasses, not included in the figure, failed to show significant changes with respect to the controls. Each value is the mean of 3-4 independent determinations. Vertical lines represent 1 SEM.

reported in Fig. 3. In these experiments we investigated the distribution of <sup>14</sup>C within the principal lipid subfractions in HTC cells incubated for 3 h in the presence of [<sup>14</sup>C]stearic acid (80  $\mu$ M) after a 12 h treatment with steroid hormone at a concentration of 1.0  $\mu$ M. Glucocorticoids were found to produce an enhancement on <sup>14</sup>C incorporation into the PC fraction, with only minor changes occurring in the other polar lipid subclasses. A possible explanation for this fact are the following observations: (a) Glucocorticoids stimulated pulmonary PC biosynthesis [32, 33] as well as the activity of three enzymes related to phospholipid synthesis: glycero-





Fig. 4. Effect of steroidal hormones on the per cent weight distribution of phosphorylated lipid subclasses from HTC cells. Suspensions of the hepatoma cells were incubated under the conditions described in the legend to Fig. 3 except that the mixture of labelled and unlabelled stearic acid was replaced by unlabelled sodium stearate at a final concentration of 80  $\mu$ M. After separation of the phospholipids by TLC, the plates were charred and quantified densitometrically as described in *Materials and methods*. Each datum is the mean of 3–4 independent determinations and is expressed as the per cent change for the experimental group in question respect to the corresponding control densitometric values, with the latter being expressed as the per cent of the sum of the optical densities for all the lipid subclasses. The control data used for reference in this way are the following (lipid fraction in parenthesis): 4.0 (LPL), 8.1 (SM), 36.8 (PE), 10.5 (PI), 7.4 (PS), 28.6 (PC), and 4.6 (CLN+PG). Vertical lines represent 1 SEM.

phosphate acyltransferase [39], phosphatidate phosphotransferase [33, 39], and glycerol-phosphate dehydrogenase [40]. (b) Corticosteroids also induced fatty acid synthetase in fetal lung, the products from which pathway, either as free acid or incorporated into phospholipids, activates cholinephosphate cytidyltransferase [41]. By contrast, the synthetic Cl analogue, Dx, caused a small decrease (*ca.* 20%) in the labelling of the CLN + PG fraction (Fig. 3). Moreover, in a similar fashion to glucocorticoids, though in a lesser extent, ALD and  $\beta E$  also stimulated the incorporation of <sup>14</sup>C into the PC subclass. With  $\beta E$ , this increment is likely to be the consequence of the stimulation of two independent enzymic activities, both lead-



Fig. 5. Effect of the steroidal hormones on the per cent weight distribution of nonphosphorylated lipid subclasses from HTC cells. Suspensions of the hepatoma cells were incubated under the conditions described in the legend to Fig. 3 except that the mixture of labelled and unlabelled stearic acid was replaced by unlabelled sodium stearate at a final concentration of 80  $\mu$ M. After separation of the neutral lipids by TLC, the plates were charred and quantified densitometrically as described in *Materials and methods*. Each datum is the mean of 3–4 independent determinations and is expressed as the per cent change for the experimental group in question respect to the corresponding control densitometric values, with the latter being expressed as the per cent of the sum of the optical densities for all the lipid subclasses. The control data used for reference in this way are the following (lipid fraction in parenthesis): 0.1 (MG), 2.5 (DG), 22.2 (CH), 7.1 (NEFA), 57.6 (TG), and 10.5 (CE). Vertical lines represent 1 SEM.

ing to an increased overall capacity for PC synthesis: cholinephosphate cytidyltransferase [33] and glycerol kinase [43].

The comparison of the results shown in Figs 2 and 3 reveals a clear correlation between the effects of the gluco- and mineralocorticoids studied on the incorporation of <sup>14</sup>C into total phospholipids and into the PC subclass. Moreover the inhibition observed in the labelling of polar lipids in the presence of Tst (Fig. 2) could be adequately accounted for by the decreased capacity of the CLN + PG and PE subclasses to incorporate and/or metabolize stearic acid (Fig. 3), since PE is one of the principal fractions within the phospholipids of HTC cells [44, 45].

Regarding neutral lipids, glucocorticoids caused a significant decrease in the incorporation of labelled stearate into NEFA and CE (save in the latter subclass where inhibition in the presence of Dx failed to reach a significant level). These negative effects (Fig. 3) were, however, almost completely counterbalanced by a simultaneous stimulation of <sup>14</sup>C incorporation into TG (Fig. 3), with the end result of no significant change in the neutral lipid fraction (Fig. 2). These findings are consistent with the strong stimulation by glucocorticoids of TG biosynthesis observed in isolated hepatocytes [34]. Accordingly, the substantial increment in the labelling of the TG fraction (Fig. 3) would indicate an increased secretion of triacylglycerols in the form of particles having the properties of very low density lipoproteins [34]. The endogenous mechanism by which this particular class of steroids alters the rate of TG biosynthesis and/or secretion in ours and in other systems is, however, still unknown.

Mineralocorticoids also produced significant alterations in the neutral lipids of HTC cells (Fig. 3). DOC inhibited the incorporation of [<sup>14</sup>C]stearate into CE and stimulated the incorporation of labelled metabolites into TG and NEFA. Exposure to ALD also inhibited significantly the uptake of <sup>14</sup>C into CE without any change in the other neutral lipid subclasses. To the best of our present knowledge, these observations represent the first experimental evidence that mineralocorticoids markedly modify the neutral lipid metabolism.

Sexual steroids also caused changes in the neutral lipids of HTC cells. The uptake of 14C into TG was stimulated by both Tst of  $\beta E$  treatment. The androgen produced an increase in the incorporation of <sup>14</sup>C into NEFA, while the estrogen decreased the total radioactivity associated with CE. Previous reports had indicated that estrogens increased significantly the biosynthesis of TG in liver [46]. We found a similar behavior in HTC cells treated with  $\beta E$ . Exposure of a ortic smooth-muscle cells to estrogens produced a simultaneous inhibition of hydroximethylglutaryl-CoA reductase activity with a concomitant stimulation of the incorporation of exogenous cholesterol. These effects when combined produce an elevation in cellular CE levels [31], which enhancement is offset by Tst and thoroughly reversed by tamoxifen treatment [31, 46, 47]. Furthermore, earlier experiments reported by Nassen et al., had demonstrated that estrogen treatment also produced a marked decrease in the rate of cellular-cholesterol esterification [31]. These results were also obtained in our experimental system (Fig. 3).

The main parameter influencing the uptake and subsequent metabolic utilization of an exogenous fatty acid is the lipid subclass into which that precursor and its eventual metabolites are incorporated. Accordingly, the quantification of the different lipids in HTC cells, with and without hormonal treatment, is necessary for a meaningful interpretation of all experimental results presented above. For this reason, Fig. 4 shows the relative mass distribution of polar lipids in HTC cells in the presence and absence of treatment with different steroidal hormones. Similarly, Fig. 5 presents the corresponding results for neutral lipids in the cells.

As stated above, glucocorticoids stimulate the biosynthesis of PC in lung [32–34, 39–41]. We could speculate that this effect also occurs in HTC cells as shown in Fig. 3. Moreover, this supposition is supported by the increment in the relative amount of PC from those data in Fig. 4. The significant decrease observed in the percentage of PI under glucocorticoid treatment (Fig. 4) does not correlate with any other similar change in the uptake of <sup>14</sup>C into this lipid subclass (data not shown). A parallel behavior was exhibited by the PS fraction. The decreases observed in the relative amounts of CLN + PG fraction under Dx, Cl or Cst treatments were of similar magnitude, but only in the case of Dx the results here were correlated well with the decreased incorporation of <sup>14</sup>C seen in this lipid fraction (Fig. 3).

Mineralocorticoids produced a significant increase in the percentage of SM and a concomitant decrease in the relative content of PS (Fig. 4). Unfortunately, these lipid fractions showed a very poor capacity to incorporate labelled substrates so that no differences were found respect to control cells (data not shown). Thus, it is not possible to draw any comparison between the results from the experiments in Figs 3 and 4 for the SM or PS fractions. The hormonal effects of mineralocorticoids on these lipid fractions are of interest since they have not been previously described. The decreases produced by DOC over the relative amounts of the CLN + PG and PS subclasses would be evoked through the reported 'glucocorticoid-like' effect displayed by this hormone (but not by ALD) on hepatic tissue [47]. ALD, on the other hand, caused a significant increase in the percentage of PC, effect not seen with DOC (Fig. 4). Moreover, this increment correlates well with the uptake of radioactivity registered in this lipid subfraction after the 3 h labelling with [<sup>14</sup>C]stearate (Fig. 3). The biological effect of ALD over PC metabolism was entirely novel. Notwithstanding, previous studies carried out on amphibians, have indicated that this mineralocorticoid hormone increases the phospholipase A, activity and stimulates the acylation-deacylation cycle in the urinary bladder of the toad [27, 28]. Despite these results, we failed to detect any significant change in the level of the LPL fraction after ALD treatment (Fig. 4).

 $\beta E$  and Tst produced a marked inhibition in the proportion of PS (Fig. 4). The treatment with  $\beta E$  increased the percentage of PC and LPL, while the incubation in the presence of Tst caused an increment in the relative amount of PI. These results are in agreement with previous studies from other laboratories. For example,  $\beta E$  was reported to stimulate significantly the biosynthesis of PC in lung and the activity of cholinephosphate-cytidyltransferase [33]. Experiments carried out with muscle cells have also demonstrated that estrogen produces an anabolic effect on phospholipid metabolism [31]; whereas androgens either antagonize estrogenic action or produce no changes at all [31, 33].

Glucocorticoids injected to rats have been reported to stimulate the biosynthesis of TG and to produce fat liver [34]. In this work we have observed not only a stimulation in the uptake of labelled stearic acid into TG, (Fig. 3) but also a significant increase in the relative mass distribution of TG, DG and MG (Fig. 5). Hence, these stimulatory effects would be synergized by oleic acid derived from stearate by  $\Delta 9$ desaturation since previous reports have indicated that oleate can significantly stimulate the activity of diacylglyceroltransferase in isolated hepatocytes [39]. Figure 5 also shows that incubation in the presence of glucocorticoids resulted in a decrease in the amount of CH and CE with the effect of Cst being of lesser importance. On the other hand, all the glucocorticoids tested evoked a similar reduction in the content of NEFA (Fig. 5), effect that well correlated with the diminished capacity for <sup>14</sup>C uptake exhibited by this lipid fraction (Fig. 3).

Both mineralocorticoids, DOC and ALD, caused an increment in the level of the metabolic precursors of TG, (i.e.; MG and DG). The relative amount of TG was not, however, significantly modified. Furthermore, both steroid hormones increased the content of CH while they simultaneously decreased the level of CE (Fig. 5). A comparison of the results shown in Figs 3 and 5, reveals that the specific incorporation of <sup>14</sup>C in these latter lipid fractions was not markedly decreased under DOC or ALD treatment.

The relative contents of NEFA, DG, and CE were reduced after incubating the cells in the presence of  $\beta E$ . This steroid also produced an increment in the percentage abundance of TG, effect that is in agreement with those results previously obtained in aortic smooth-muscle cells [31]. In the latter cell type, however, the estrogen caused elevations in NEFA and CE, which changes were not observed with the HTC cells (Fig. 5). An adequate explanation for these discrepancies would no doubt be based on the elucidation of differences in other aspects of the total lipid metabolism of these two cell populations and must thus remain unresolved until such a time as those pertinent data became available for the aortic cells.

With Tst treatment, the relative abundance of both MG and DG decreased significantly, whereas the cellular TG content remained unchanged. This finding indicates that the principal pathway for the TG biosynthesis is depressed and that the metabolic precursors for the alternative biosynthetic process, MG and DG, are accordingly actively recruited to maintain the endogenous pool of TG at its normal physiologic levels. Figure 2 shows that Tst treatment produces the highest ratio of <sup>14</sup>C incorporation into neutral lipids *vs*. phospholipids, which result could be derived from the observed marked increase in the cellular NEFA content (*ca.* 140%), which effect would sequester <sup>14</sup>C labelling within the neutral-lipid fraction.

In conclusion, these experiments have demonstrated for the first time the effects produced by a wide variety of steroidal hormones on the incorporation and metabolism of stearic acid. It was also shown that  $\Delta 9$  microsomal desaturation is strongly stimulated and that the incorporation of labelled metabolites into different lipid fractions is modified by hormonal treatment. We would anticipate that the information derived from these studies will be useful in elucidating the mechanism underlying other metabolic changes arising from the effect of steroidal hormones, such as the regulation of water permeability by mineralocorticoids [27, 28], the action of glucocorticoids on the characteristics of the air-alveolar barrier [32, 33], the differences in lipid metabolism between males and females as attributed to sexual-hormone balance [48], and related actions involving alleviations of chronic diseases by means of the administration of natural or synthetic steroids.

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