

Disruption of the Actin Cytoskeleton Induces Fluorescent Glucose Accumulation on the Rat Hepatocytes Clone 9

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Key Words

2-NBDG • Clone 9 hepatocytes • Confocal microscopy
• F-actin • Glucose metabolism

Abstract

Background: Glucose transport and metabolism are highly specialized in hepatocytes. Actin cytoskeleton is fundamental to the maintenance of their morphology as well as to ensure their functionality. Here we study the effect of the actin disrupting natural compounds cytochalasin B and latrunculin A on the glucose metabolism of the Clone 9 rat hepatocytes once the glucose molecule is inside them and the effects of two hormones which main function is regulating the glucose metabolism on the actin cytoskeleton of Clone 9 cells. **Methods:** F-actin was labeled by using Oregon Green 514 ® phalloidin and glucose inside cells was monitored with the fluorescent D-glucose derivative; 2-NBDG. **Observations and measurements** were carried out by using a confocal microscope. **Results:** Nor insulin neither glucagon was able to induce any significant effect in the quantity of F-actin present on Clone 9 cells. But insulin triggers a strong reorganization on the pattern of distribution of F-actin. However, the actin cytoskeleton disruption

induced by CB and more efficiently by Lat A caused accumulation of 2-NBDG in cells. **Conclusions:** These results state that disruption of the actin cytoskeleton induces fluorescent glucose accumulation on the rat hepatocytes Clone 9 suggesting that actin disrupting agents cause a blockage in the glycolytic pathway of Clone 9 hepatocytes.

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Introduction

Actin is one of the most conserved proteins in eukaryotic cells [1, 2] and its importance can be divided in two groups of activities; structural and functional. Actin cytoskeleton is essential to cytokinesis, cell adhesion, cell movement and migration, cytosolic traffic and maintenance of the cellular shape [3]. Many natural compounds that interact with the actin cytoskeleton have been discovered in the last years. These molecules disturb the cell integrity and functions [4].

Latrunculin A (Lat A) is a macrolide isolated from the marine sponge *Negombata magnifica*. This compound disrupts the actin cytoskeleton by sequestering G-actin (monomeric) and avoiding its polymerization [5-

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7]. Cytochalasins constitutes another group of actin disrupting compounds. These fungal metabolites inhibit actin polymerization by capping the barbed-end of the actin filament [8]. Therefore, both macrolides disrupt actin cytoskeleton by using different action mechanisms. Many other macrolides affects actin cytoskeleton such as halichondramide, swinholide or pectenotoxin [4, 9-13].

On the other hand, D-glucose uptake and metabolism is one of the most studied routes in cells. Any failure in this metabolic pathway frequently entails severe disorders in the organism. Recent investigations discovered that 2-NBDG (2-{N-[7-nitrobenz-2-oxa-1, 3-diazol 4-yl] amino}-2-deoxyglucose), a fluorescent analogue of the D-glucose that gets into cells by using the same transporters [14-18], could be used to monitor the rate of glycogen synthesis in cells [19].

D-glucose uptake by cells and the actin cytoskeleton have been previously related. Actin disruption inhibits D-glucose uptake in most of cells by avoiding GLUT 1 and 4 to reach the plasmatic membrane because they are stored in vesicles inside the cell [20, 21]. However so far there is no information about how drugs that modify cytoskeleton could affect glucose metabolism in hepatocytes.

In this study, we evaluated the effect of compounds that disrupt the actin cytoskeleton on the metabolism of D-glucose. In order to do that, rat hepatocytes from a cell line (Clone 9) were exposed to Lat A or cytochalasin-B (CB) and the consequent changes in the actin cytoskeleton and D-glucose metabolism were evaluated.

Materials and Methods

Latrunculin A (Lat A) derived from *Negombata magnifica* and Cytochalasin-B (CB) from *Drechslera dematioidea* were purchased from Sigma-Aldrich (Madrid, Spain).

The fluorescent dye Oregon Green® 514 Phalloidin for F-actin labeling and 2-NBDG (2-{N-[7-nitrobenz-2-oxa-1, 3-diazol 4-yl] amino}-2-deoxyglucose) were from Molecular Probes (Leiden, The Netherlands).

Cell culture: Nutrient mixture F-12 Ham Kaighn's modification, streptomycin sulphate salt and penicillin G potassium salt were purchased from Sigma (Madrid, Spain) and foetal porcine serum was from Gibco (Barcelona, Spain).

Bovine Serum Albumin (BSA) was purchased from ICN Biomedicals, Inc. (Ohio, U.S.A.).

Drugs: Glucagon was from Novo Nordisk Pharma S.A. (Madrid, Spain) and insulin from Intervet S.A. (Salamanca, Spain).

All other chemicals were reagent grade and purchased from Sigma-Aldrich (Madrid, Spain) or Panreac (Barcelona, Spain).

Buffer solution for confocal assays (in mM): 138 NaCl, 5.5 D-glucose, 5 KCl, 1 MgCl₂, 1.5 CaCl₂ and 10 Hepes. pH 7.4.

Cell culture

Rat hepatocytes from the cell line Clone 9 (ECACC N° 88072203) were grown on 60 mm tissue culture plates in F-12 Ham Kaighn's modification supplemented with 2.5 g L⁻¹ NaHCO₃, 28 mg L⁻¹ streptomycin sulphate salt, 17 mg L⁻¹ penicillin G potassium salt and 10% fetal bovine serum pH 7.2. The hepatocytes were grown in a humidified atmosphere with 5% CO₂ at 37 °C [22]

Hepatocytes were seeded on coverslips, which were placed in 8-well sterile plates. Coverslips were used when the cells reached confluence. In this moment cells are ready to one of the following protocols.

F-actin cytoskeleton staining

Once hepatocytes reach confluence, cells were incubated with drugs for 3 h in the culture medium. After that, hepatocytes were washed with PBS and fixed with 4% paraformaldehyde solution (10 min). Then, cells were permeabilized for 5 min with PBS-0.1% Triton X-100 and incubated with 0.165 µM Oregon Green® 514 Phalloidin for F-actin labeling for 20 min. Coverslips were mounted on slides with 1:1 glycerol-PBS and sealed with nail varnish in order to preserve fluorescence and stored at 4 °C [9, 11]. Control cells were incubated in the same conditions with the toxin vehicle, dimethylsulfoxide (DMSO).

Confocal microscopy for visualizing morphology, actin cytoskeleton distribution and measuring

Confocal imaging was registered with a 60x oil immersion objective of a Nikon Eclipse TE2000-E inverted microscope attached to the C1 laser confocal system (EZC1 V.2.20 software; Nikon Instruments Europe B.V., The Netherlands). A 488nm Argon laser was used to excite Oregon Green® 514 phalloidin.

Fluorescent images shown as volume-render projections belong to Z-stacks acquired at 0.5-µm intervals at 512x512 pixel resolution.

Fluorescence measurements corresponding to F-actin labeling were registered and calculated with quantification software from Z-stack sections. Values from all independent experiments were averaged for a single data point.

Results are presented as the percentage of the mean value ± standard error of the mean (s.e.m.) of fluorescence emitted by cells treated with drugs, versus controls, with n = 3.

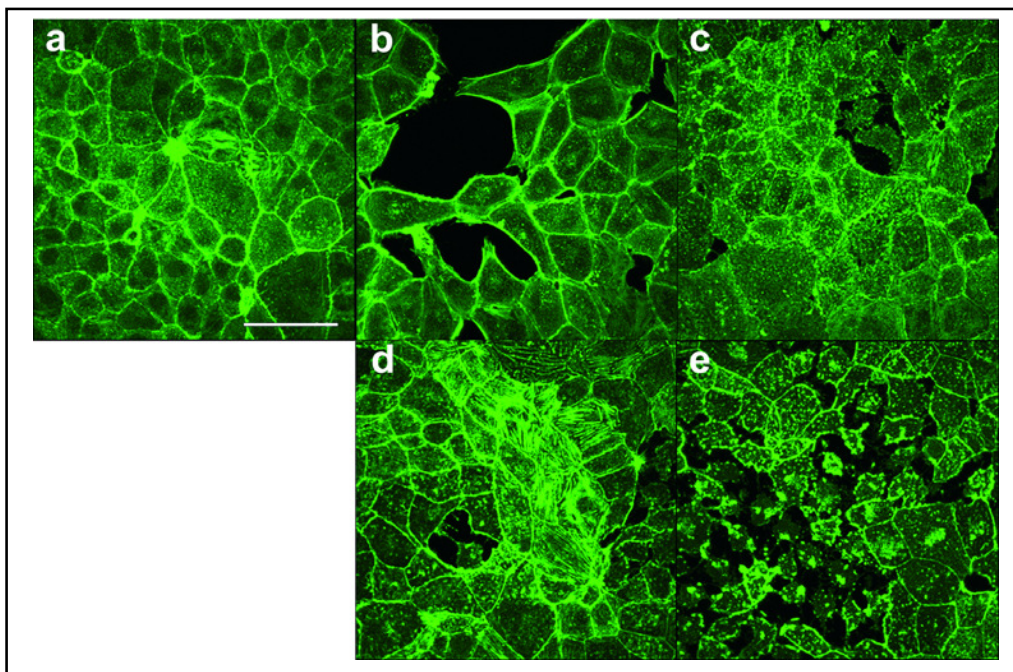
Fluorescent detection of D-glucose metabolism with 2-NBDG

Hepatocytes were washed 3 times with the buffer solution for confocal assays (cited above in Materials section) and then incubated in buffer solution with 500µM 2-NBDG at 37°C for 3 h. Then, fluorescent dye was removed by washing 3 times with the buffer solution for confocal assays. Then cells were incubated with drugs or vehicle for 3 h.

Confocal microscopy for visualizing and measuring

Fluorescence measurements and images were registered with a 60x oil immersion objective of a Nikon Eclipse TE2000-E

Fig. 1. Images of the effect of actin disrupting agents on the actin cytoskeleton of Clone 9 hepatocytes. Confocal imaging of F-actin labeled with Oregon Green® 514 phalloidin. Control (a) and Clone 9 cells incubated for 3h with 1 or 2 μ M CB (b and c respectively) 100 or 200 nM Lat A (d and e respectively). Images are representative from 3 independent experiments. Scale bar = 50 μ m.



inverted microscope attached to the C1 laser confocal system (EZC1 V.2.20 software; Nikon Instruments Europe B.V., The Netherlands). Fluorescent images shown as volume-render projections belong to Z-stacks acquired at 0.5- μ m intervals and 512x512 pixel resolution.

Fluorescent images are shown as volume-render projections that belong to Z-stacks. Z-stack is a collection of images superimposed taken by Z-scan of the sample at 512x512 pixel resolution. Z-scans are acquisitions taken each 0.5 μ m along the Z axis of the sample by confocal software with a motorized focus.

Fluorescence measurements corresponding to 2-NBDG labeling were registered and calculated with quantification software from Z-stack sections. Values from all independent experiments were averaged for a single data point.

Results are presented as the percentage of the mean value \pm standard error of the mean (s.e.m.) of fluorescence emitted by cells treated with drugs, versus controls, with $n = 3$.

Statistical analysis

Results were analyzed using the Students's t-test for paired data where appropriate. A probability level of $\alpha=0.05$ was set to indicate statistical significance.

Results

Clone 9 rat hepatocytes were incubated for 3 h with 1 or 2 μ M CB, 100 or 200 nM Lat A or vehicle (dimethylsulfoxide) and then stained for F-actin labeling with Oregon Green®-phalloidin as described in Materials and Methods section. 1 μ M Cytochalasin B induced slight changes in the pattern of distribution of F-actin when compared to control cells (Fig. 1B and 1A respectively).

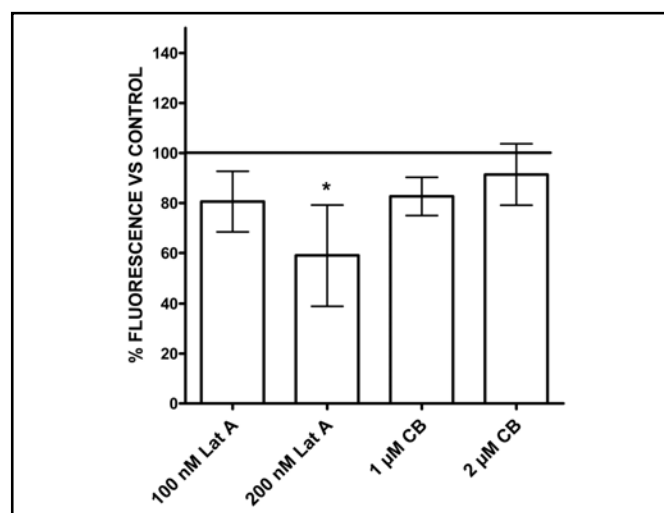
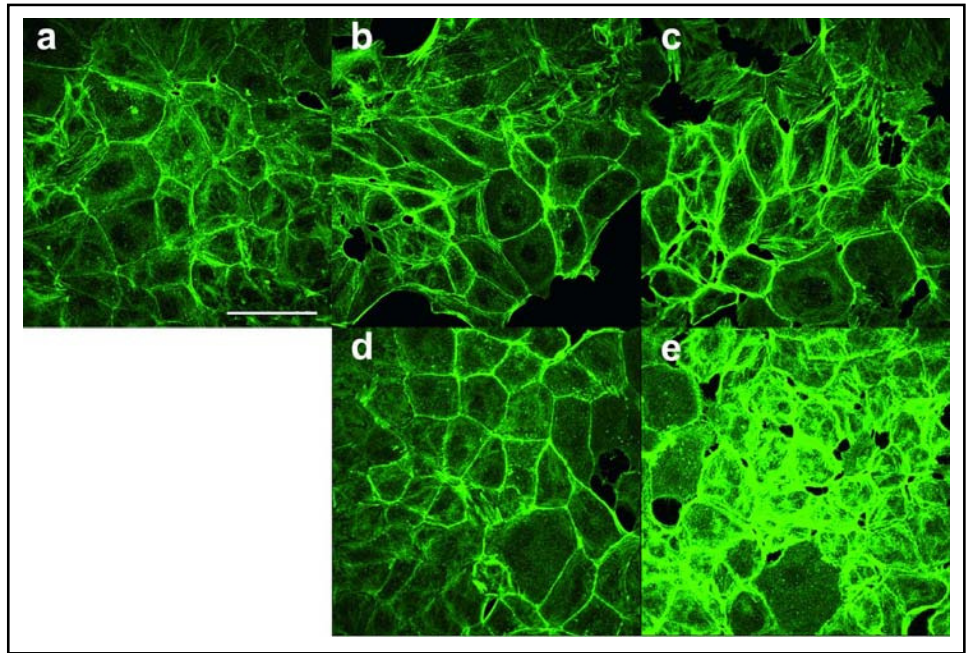


Fig. 2. F-actin fluorescence measurements of Clone 9 hepatocytes treated with actin disrupting agents. The present diagram represents the percentage (mean \pm s.e.m. for $n=3$ experiments) of fluorescently labeled F-actin in cells treated for 3h with 1 or 2 μ M CB either 100 or 200 nM Lat A relative to control cells (100%). Mean \pm s.e.m. of $n=3$ experiments. * $P<0.05$; * $P<0.01$, significantly different from control; Student's t-test.

Control cells showed an F-actin meshwork distributed all along the cell body with intensely fluorescent bundles in the cell cortex corresponding to the stress fibers (Fig. 1a). F-actin labeling of 1 μ M CB treated cells seemed less intense and some punctuated fluorescence appeared (Fig.1b). This effect was more pronounced in cells treated

Fig. 3. Effect of agents that alter the glucose metabolism on the actin cytoskeleton of Clone 9 hepatocytes. Confocal imaging of F-actin labeled with Oregon Green® 514 phalloidin. Control Clone 9 cells (a) and Clone 9 cells incubated for 3h with 10 nM or 1 μ M glucagon (b and c respectively) 100 nM or 1 μ M insulin (d and e respectively). Images are representative from 3 independent experiments. Scale bar = 50 μ m.



with 2 μ M CB (Fig. 1c) but decrease in fluorescence value was no evident (Fig. 2). 100 nM Lat A was enough to change actin cytoskeleton on Clone 9 cells in a similar way. F-actin decreased and became punctuated (Fig. 1d). 200 nM Lat A disrupted F-actin acutely. Fluorescence disappeared in many parts of cells and F-actin meshwork and bundles was almost replaced by fluorescent dots (Fig. 1e). Fluorescence quantification revealed a significant decrease in the F-actin signal when cells were treated with Lat A, being more pronounced with the higher concentration of toxin (Fig. 2).

Then, Clone 9 cells F-actin distribution was studied after 3h incubation with 10 nM or 1 μ M glucagon either 100nM or 1 μ M insulin. We used these two hormones that modify in different ways the glucose metabolism to test their effect on the actin cytoskeleton of Clone 9 cells; insulin enhances the glycogen formation and the glucose uptake and glucagon decreases the glycogen formation and increases the glucose release in hepatocytes. Glucagon was not able to exert any significant effect on the F-actin pattern of distribution or in the quantity of F-actin in Clone 9 cells (Fig. 3b, 3c and 4). However, insulin seemed to induce a localization change on fluorescence since the cortical bundles that usually appear in the basal zone of the cells (the adherent zone) showed more intense (Fig. 3d and 3e). But this appreciation was neither reflected in the total volume of the cells nor on the fluorescence values (Fig. 4).

Finally, experiments were carried out to evaluate the effect of Lat A and CB on the D-glucose metabolism of Clone 9 cells. Cells were preincubated with 500 μ M

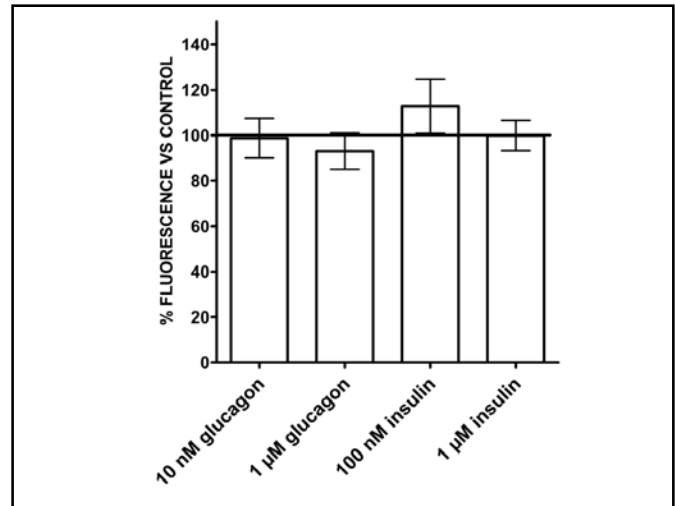
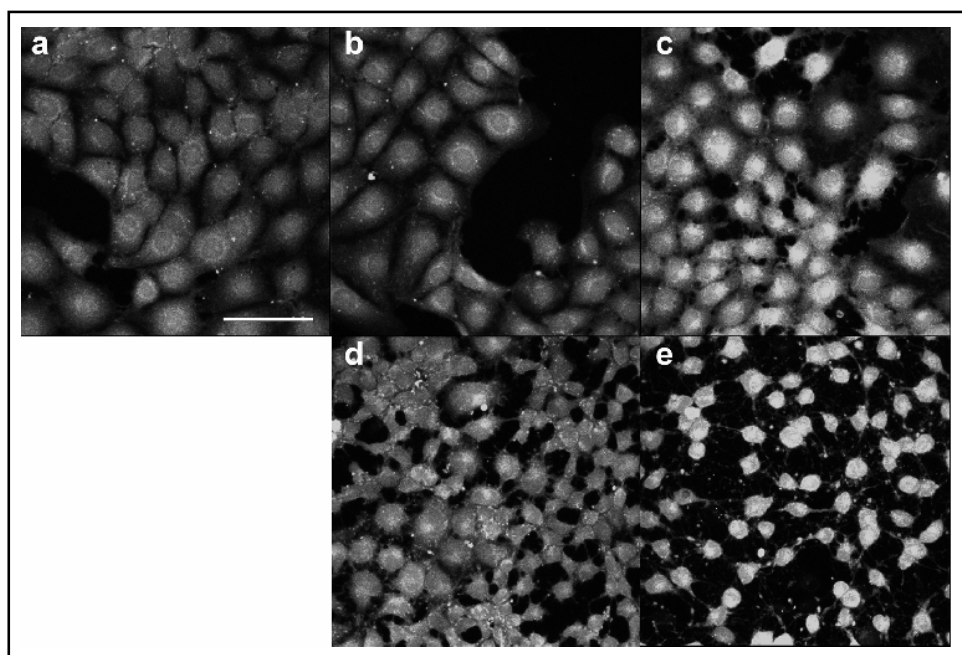


Fig. 4. F-actin fluorescence measurements of Clone 9 hepatocytes treated with agents that alter the glucose metabolism. The present diagram represents the percentage (mean \pm s.e.m for n=3 experiments) of fluorescently labeled F-actin in cells treated for 3h with 10 nM or 1 μ M glucagon either 100 nM or 1 μ M insulin relative to control cells (100%). Mean \pm s.e.m of n=3 experiments.

2-NBDG for 3h. Then, cells were washed and incubation buffer was replaced by new solution without 2-NBDG. At this point, drugs were added and after 3h of incubation fluorescence values and images were registered. Cells incubated with 1 μ M CB maintained their shape and the fluorescence seemed to be unaltered (Fig. 5b). However, 2 μ M CB induced morphological changes in some cells

Fig. 5. Effect of actin disrupting agents on the glucose metabolism of Clone 9 hepatocytes. 2-NBDG was included in the incubation media of Clone 9 cells and then released. After, cells were exposed for 3h to 1 or 2 μ M CB (b and c respectively), 100 or 200 nM Lat A (d and e respectively) either drugs vehicle (a; control cells) for 3h. Images are representative from 3 independent experiments. Scale bar = 50 μ m.



appearing rounder than the control cells (Fig. 5c and 5a respectively). Moreover, the fluorescence intensity seemed increased as it could be seen in Fig. 6. However Lat A is the drug which has drastic significant effects. This compound triggered a sharp change in the morphology of the Clone 9 cells; cell bodies were retracted and rounded losing their leaf-like appearance (Fig. 5d and 5e). Also, fluorescence clearly increased when cells were treated with 100 nM Lat A, and even more acutely when treated with 200 nM (Fig. 6).

Discussion

A broad interaction between the actin cytoskeleton and glucose has been previously demonstrated. Uptake of D-glucose is mediated by the protein families GLUT and SGLT [23, 24]. GLUT 1 and GLUT 4 are stored in cytoplasmatic vesicles that are transported to the plasma membrane. Insulin is a hormone that enhances the glucose uptake by cells and the glycogen deposition mainly in hepatocytes. This compound was demonstrated to induce the translocation of the GLUT vesicles from the cytosol to the plasma membrane by a mechanism that implies a strong reorganization of the F-actin framework [25-27]. In this study, insulin did not produce significant changes in the quantity of F-actin. But, the appearance of very intensely labeled F-actin bundles (in the nearest plane to the substrate) shows that insulin induces evident changes in the pattern of distribution of F-actin on Clone 9 hepatocytes. However, glucagon, hormone that enhances

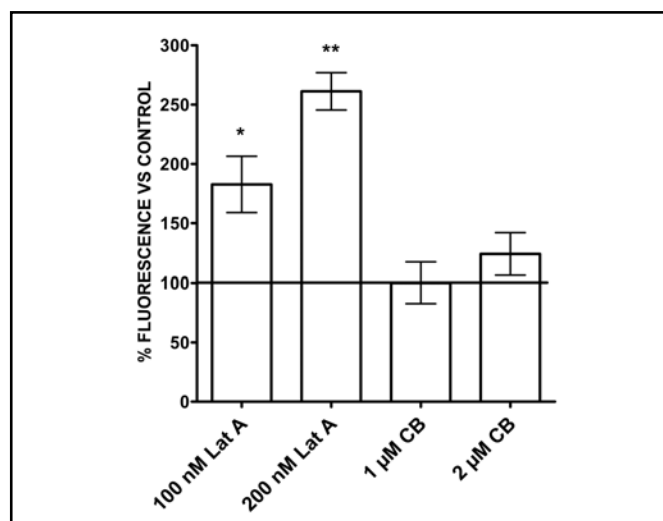


Fig. 6. 2-NBDG fluorescence measurements of Clone 9 hepatocytes treated with actin disrupting agents. The present diagram represents the percentage (mean \pm s.e.m for n=3 experiments) of 2-NBDG in cells treated for 3h with 1 or 2 μ M CB either 100 or 200 nM Lat A relative to control cells (100%). Clone 9 cells were preloaded with 2-NBDG by exposing them to the glucose indicator for 3h. Mean \pm s.e.m of n=3 experiments. *P<0.05; **P<0.01, significantly different from control; Student's t-test.

the glycolysis and inhibits the glycogen formation, did not change the pattern of distribution or quantity of the F-actin of Clone 9 cells.

Glycolysis is controlled by several mechanisms such as allosteric regulators as well as by reversible binding of glycolytic enzymes to the cytoskeleton. In fact, all the

glycolytic enzymes are described to bind to cytoskeleton with the exception of hexokinase that reversibly binds to mitochondria [28]. This mechanism has attracted much attention and has been identified as an important mechanism to generate ATP in the vicinity of the cytoskeleton modulating cellular functions and morphology. In our study, we reveal the effect of diverse actin disrupting agents in the glucose metabolism of Clone 9 hepatocytes by using 2-NBDG, a fluorescent D-glucose derivative that accumulates in living cells and not in dead cells. This dye was used in experiments of cell viability and glucose transport in different cellular models [14, 16, 29-34]. 2-NBDG is not cell permeable and enters in alive cells through the same transporters than D-glucose [14, 15, 35], then it is transformed into a still fluorescent 2-NBDG metabolite [16, 17]. Once this dye is inside the cells, it can follow the D-glucose degradation route or the glycogen formation pathway. In a previous work, we set up a method by using 2-NBDG that allows a noninvasive detection of glycogen being also a rapid testing of drugs that modify the incorporation or release of glucose from glycogen [19].

Cytochalasins are a family of fungal metabolites widely used to study the role of actin in cellular processes and to probe basic aspects of actin polymerization *in vitro*. In Clone 9 cells, CB enables 2-NBDG accumulation. However, CB and CD are potent inhibitors of glycogen synthesis in hepatocytes by an indirect mechanism that probably involves the actin disruption effect [36]. In fact, Baque et al. showed that disruption of actin cytoskeleton by CD enables a dramatic change in the cellular distribution of glycogenin, enzyme implicated in the synthesis of new glycogen particles *de novo* [37]. Therefore, CB may be avoiding normal degradation of 2-NBDG in hepatocytes by a direct or indirect pathway.

It has been reported that glycolytic enzymes are aggregated into multienzyme complexes and that these complexes may confer kinetic advantages, allowing for rapid reaction times and channeling of substrates. The binding of glycolytic enzymes to cytoskeletal structures may be important in this respect and may also facilitate the translocation of enzymes to different parts of the cell in response to changes in the metabolic demands. The glycolytic enzyme aldolase has recently been shown to be associated with the actin cytoskeleton in a reversible and dynamic process which is modulated by glycolytic inhibitors and also by CD [38]. However, CB has a number of effects not related to the F-actin inhibition of polymerization by capping. These include an F-actin severing effect, presumably leading to an increased

proportion of shorter actin filaments [39, 40], as well as an inhibition of cellular water permeability [41, 42], possibly via inhibition of the glucose transporter [43]. Furthermore, cytochalasin effects on F-actin in intact cells are highly variable, depending on the cell type in question, as well as on the cytochalasin concentration and congener used. Thus, a given cytochalasin treatment may reduce F-actin content in one cell type, but not in others [44] or differentially affect different F-actin domains within a given cell type [45, 46]. Those different CB effects make difficult to directly correlate the actin cytoskeleton disturbance found in hepatocytes with the glycolysis blockage. So, we used another kind of actin disrupting agent which specifically damage cells by disrupting their actin cytoskeleton; Lat A.

Lat A is a macrolide produced by the sponge from the Red Sea *Negomabata magnifica* [5] and a well known G-actin sequestering compound that binds to G-actin with a 1:1 stoichiometry [7]. This compound is a good tool to compare F-actin disrupting agents because its union to monomeric actin is very specific and it does not exert severing or capping activity [4, 6]. Due to its specific activity and significant results we found we can directly relate the effect of Lat A on the actin cytoskeleton of Clone 9 cells with the accumulation of 2-NBDG after its uptake and the glycolysis blockage.

According to research work of recent years, cytoskeleton plays an important physiology role mainly in the following three aspects: insulin secretion, insulin action and the intracellular distribution of glucose metabolism enzymes. In our study, when we use Lat A, we are showing the effect of the actin cytoskeleton disrupting in the glucose metabolism. Many previous and very recent studies have closely related the actin cytoskeleton with the glucose uptake by cells by using Lat A to avoid the actin cytoskeleton correct functioning [47-49]. However, there are not studies that directly correlate the glucose metabolism inside cells with the effect of a specific actin disrupting agent such Lat A. We load the cells with fluorescent glucose (2-NBDG) and observe its evolution with the treatment of cells with Lat A.

The 2-NBDG accumulation that we observed when Clone 9 cells were treated with Lat A could be the consequence of the glycogen formation or a glucose degradation blockage. Glycogen formation with 2-NBDG is characterized by the appearance of a dotted fluorescence pattern, while the disperse fluorescence disappears [19]. In cells treated with Lat A, fluorescence remains accumulated in cells but in a disperse pattern in the whole cell pointing to the glycolytic pathway blockage.

Moreover, it was previously demonstrated that the disassembly of the microfilament-microtubule system leads to a reduction in glycogen synthesis by inhibiting the translocation of glycogen synthase [36, 50].

In summary, we demonstrated that the disruption of the rat hepatocytes actin cytoskeleton, a cell model highly implicated in the glucose regulation in whole organisms, enables the glucose accumulation in cells by using, for the first time, a highly specific anti-actin drug, Lat A.

In conclusion, the impairment on glycolysis that the actin cytoskeleton disrupting agents tested induce in Clone 9 cells seems to be the consequence of a breaking-off in the connection between glycolytic enzymes and their substrates. This impairment of the glucose metabolism could lead to a serious failure in the normal activity of the cell, even to dead.

Abbreviations

2-NBDG (2-{N-[7-nitrobenz-2-oxa-1,3-diazol-4-yl]amino}-2-deoxyglucose); CB (cytochalasin B); CD

(cytochalasin D); F-actin (filamentous actin); Lat A (latrunculin A).

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