Abstract
The effect of oleic, linoleic and γ-linolenic acids on ROS production by 3T3 Swiss and Rat 1 fibroblasts was investigated. Using lucigenin-amplified chemiluminescence, a dose-dependent increase in extracellular superoxide levels was observed during the treatment of fibroblasts with oleic, linoleic and γ-linolenic acids. ROS production was dependent on the addition of β-NADH or NADPH to the medium. Diphenyleneiodonium inhibited the effect of oleic, linoleic and γ-linolenic acids on fibroblast superoxide release by 79%, 92% and 82%, respectively. Increased levels of p47phox phosphorylation due to fatty acid treatment were detected by Western blotting analyses of fibroblast proteins. Increased p47phox mRNA expression was observed using real-time PCR. The rank order for the fatty acid stimulation of the fibroblast oxidative burst was as follows: γ-linolenic > linoleic > oleic. In conclusion, oleic, linoleic and γ-linolenic acids stimulated ROS production via activation of the NADPH oxidase enzyme complex in fibroblasts.

Introduction
Plasma fatty acid levels are elevated during conditions such as diabetes and inflammation, which are commonly associated with ROS production and the development of excess fibrous connective tissue in organs or tissues due to fibroblasts [1], [2], [3], [4].

ROS modulate the activity of signalling pathways involved in fibroblast migration, proliferation, connective tissue deposition, vascular tone and senescence [5], [6]. ROS play a role in the activation of NF-κB, phospholipase D, protein kinase C (PKC) and mitogen-activated protein kinase (MAPK) in response to various agonists [5], [7]. ROS are also required for the activation of Akt, p70S6K and G-proteins, such as Gi, Go, and for the expression of early growth response factor-1 (EGFR) and vascular endothelial growth factor (VEGF). The activation of signalling cascades by low levels of ROS results in increased cell cycle progression. For example, the proliferative state of fibroblasts is tightly associated with intracellular ROS levels. Low ROS levels lead to cell growth arrest, which is induced by contact inhibition. On the other hand, the overproduction or insufficient scavenging of ROS can result in enhanced fibrosis and oxidative stress, which have been implicated in several diseases [2], [8], [9].

The regulation of the intracellular redox state by fatty acid-induced changes in NAPDH oxidase activity is thought to have an important impact on redox-sensitive signalling cascades. NADPH oxidase was originally believed to be present in phagocytes only; however, its expression has been demonstrated in several non-phagocytic cell types such as vascular smooth muscle cells [10], pancreatic β cells [11] and fibroblasts [12], [13]. ROS are produced by NADPH oxidase homologues (known as NOX) in non-phagocytic cells. The NADPH oxidase subunits gp91phox and p22phox are integral membrane proteins [2]. These subunits form heterodimeric flavocytochrome b558, which forms the catalytic core of the enzyme. However, this enzyme exists in an inactive state in the absence of the other subunits. Additional subunits are required for regulation and are located in the cytosol during the resting state. These subunits include the proteins p67phox, p47phox and p40phox and the small GTPase Rac. p47phox is phosphorylated at multiple sites by a number of protein kinases, including members of the PKC family, and it is important in the regulation of NADPH oxidase activity. Cell stimulation leads to the phosphorylation and translocation of p47phox to the membrane. In the membrane, p47phox and p67phox directly interact with and activate NOX [2], gp91phox, p67phox and p47phox expression has been reported in fibroblasts from different species [12], [13].

Oleic (C18:1), linoleic (C18:2) and γ-linolenic (C18:3) acids are abundant fatty acids in human and rat plasma [14]. These fatty acids have the same carbon atom chain length but different degrees of unsaturation and positions of the double bonds in their molecules. Although fatty acids have been demonstrated to...
activate NADPH oxidase in leukocytes [15] and pancreatic β cells
[16], the effect of oleic, linoleic and γ-linolenic acids on NADPH
oxidase in fibroblasts has not yet been investigated. Given that
fibroblasts exert profound effects on the progression of inflamma-
tory chronic diseases, the goal of the present study was to
investigate the effect of fatty acids on intracellular and extracellular
ROS levels in cultured fibroblasts (the 3T3 Swiss and Rat 1 cell
lines) using three techniques: lucigenin-amplified chemilumines-
cence, reduction of hydroethidine and phenol red reduction. The
levels of p47phox phosphorylation and p47phox mRNA expression in
response to the addition of oleic, linoleic and γ-linolenic acids were
detected using Western blot analysis and real-time PCR,
respectively.

Materials and Methods

Dulbecco’s modified Eagle’s medium, HEPES, ampicillin,
streptomycin, Trizol reagent, random pd(N)6 primers, DNase I,
Superscript II RT and Taq DNA polymerase were purchased from
Invitrogen Life Technologies (Carlsbad, CA, USA). Fatty
acids, lucigenin, hydroethidine, hydrogen peroxide, β-NADH,
NADPH, and phenol red were supplied by Sigma Chemical Co.
(St. Louis, MO, USA). The p47phox rabbit antibody was obtained from Santa Cruz Biotechnology (Santa Cruz, CA, USA). The fatty
acids were dissolved in ethanol. The final concentration of ethanol
in the assay medium did not exceed 0.05%. At this concentration,
a preliminary experiment demonstrated that ethanol is not toxic to
fibroblasts and that it does not interfere with the obtained results.
Reagents, water and plastic wares used in the experiments were all
endotoxin free.

2.1. Preparation and treatment of the fibroblasts

Swiss 3T3 and Rat 1 fibroblasts, which were obtained from the
American Type Culture Collection, were maintained at 37°C in
Dulbecco’s modified Eagle’s medium supplemented with 25 mM
HEPES (pH 7.4), 10% (v:v) foetal bovine serum, 25 µg per mL
ampicillin, and 100 µg per mL streptomycin (culture medium).
For all experiments, cells were used between passages 5 and 7.
When the cells reached confluency, adherent fibroblasts were
treated with trypsin, gently washed, and resuspended in PBS
buffer. The cells were maintained on ice until they were assayed.
For the chemiluminescence assay, 2.5×10⁶ cells per mL were
added to 0.3 mL of pre-warmed (37°C) PBS buffer containing
lucigenin (1 mM), phenol red (0.28 mM) or hydroethidine (1 µM).
The cells were treated with various concentrations (0, 5, 25, 50,
100, or 200 µM) of oleic, linoleic or γ-linolenic acids. ROS release
was monitored for 30 min after the addition of the fatty acids. The assays were performed in PBS buffer supplemented with CaCl₂
(1 mM), MgCl₂ (1.5 mM), and glucose (10 mM) at 37°C in a final
volume of 0.3 mL.

2.2. Cell viability

The treated cells were centrifuged at 1,000× g for 15 min (4°C),
and the resultant pellet was resuspended in 500 µL phosphate-
buffered saline (PBS). Afterwards, 50 µL of propidium iodide
solution (50 mg/mL in PBS) was added, and the cell viability was
analysed using a FACSCalibur flow cytometer (Becton Dickinson,
San Juan, CA, USA). Propidium iodide is a highly water-soluble
fluorescent compound that cannot pass through intact mem-
branes, is generally excluded from viable cells and binds to DNA
by intercalating between the bases. Fluorescence was measured
using the FL2 channel (detection from 564–606 nm). Ten
thousand events were analysed per experiment. Cells with
propidium iodide fluorescence were then evaluated using the Cell
Quest software (Becton Dickinson, San Juan, CA, USA). Viability
was always greater than 98.5%, as indicated by flow cytometry.

2.3. Lucigenin-enhanced chemiluminescence assay

Lucigenin is extensively used to measure the production of
reactive oxygen species by chemiluminescence [15]. After being
excited by the superoxide anion, lucigenin releases energy in the
form of light. Lucigenin-amplified chemiluminescence is a specific
method for studying the kinetics of the superoxide production of
cells. Using this method, the response to xanthine/xanthine
oxidase results in a positive correlation with light measurements;
however, this method does not indicate augmented chemilumi-
nescence when peroxidase is added to the assay medium [17]. In
addition, chemiluminescence is inhibited in a dose-dependent
manner by scavengers of superoxide anions, e.g., SOD, but not by azide, catalase, mannitol or taurine. Thus, this is a specific method to measure superoxide anion production [18]. The chemiluminescence response was monitored for 30 min (37°C) using a microplate luminometer (model LB96V, EG&G Berthold; Bad Wildbad, Germany).

2.4. Flow cytometric measurements of reactive oxygen metabolites using hydroethidine

Hydroethidine (HE) has been widely used for the flow cytometric measurement of intracellular ROS levels. Hydroethidine, a reduced derivative of ethidium bromide, easily penetrates into cells and displays weak fluorescence when excited by light (480 nm). Hydroethidine is oxidised intracellularly by oxygen radicals and converted into ethidium bromide, which tightly binds to DNA and displays a strong red fluorescence [19]. One advantage of this method is the ability to evaluate the response of individual cells. Fibroblasts were treated as described above in the presence of hydroethidine (1 µM). Fluorescence was measured using the FL3 channel (detection above 670 nm) in a FACSCalibur flow cytometer (Becton Dickinson, San Juan, CA). Ten thousand events were analysed per experiment.

2.5. Hydrogen peroxide determination

For measuring hydrogen peroxide levels, we used a single, rapid and inexpensive method described by E. Pick et al. [20]. This assay is based on the horseradish peroxidase (HRP)-mediated oxidation of phenol red by H₂O₂. The reaction leads to the formation of a coloured compound whose maximum absorbance occurs at 610 nm [20]. The phenol red assay allows for the detection of reactive oxygen species both inside and outside of cells. Fibroblasts were treated as described above in the presence of phenol red (0.28 mM) and 1 U/mL peroxidase (horseradish Type II, [HRP]). The reaction was stopped by the addition of 10 µL of a 1 N sodium hydroxide aqueous solution.

2.6. Western blot analysis of p47phox phosphorylation

In the Western blot experiments, the total protein levels of each sample were quantified by the Bradford method. After this procedure, the same amount of protein was immunoprecipitated using the anti-phosphoserine antibody. After the immunoprecipitation, the samples were immunoblotted using anti-p47phox antibody. After incubating the cells at 37°C in the absence (0 min) or presence of fatty acids (100 µM – for 5, 10 or 15 min), they were homogenized in extraction buffer and prepared for Western blot analysis.

The cells were homogenised in extraction buffer (100 mM Tris (pH 7.5), 10 mM EDTA, 100 mM NaF, 10 mM sodium pyrophosphate, 10 mM sodium orthovanadate, 2 mM phenylmethanesulfonyl fluoride, 0.01 mg/mL aprotinin) at 4°C for 30 s. After homogenisation, Triton X 100 was added to 1%, and the samples were incubated for 30 min (4°C). Samples were centrifuged at 13,000 g for 20 min (4°C). Each sample (250 µg) was immunoprecipitated using the anti-phosphoserine antibody (1:300 dilution). Immunoprecipitated samples were mixed with protein A-Sepharose for 4 h (4°C), subjected to electrophoresis and then immunoblotted using an anti-phosphoserine antibody. Briefly, the gel was transferred to a nitrocellulose membrane at 120 V for 1 h. The membrane was blocked with 5% defatted milk in a basal solution (10 mM Tris (pH 7.5), 150 mM NaCl, 0.05% Tween 20) at room temperature for 2 h. The membranes were washed 3 times (10 min for each wash) in basal solution and then incubated at room temperature for 3 h with an anti-p47phox antibody (1:3,000 dilution) in a basal solution containing 3% defatted milk. The membranes were washed again (3 washes, 10 min each) and incubated with an anti-IgG antibody (1:10,000 dilution) linked to horseradish peroxidase in a basal solution containing 1% defatted milk at room temperature for 1 h. Following the washings, the membranes were incubated with the substrate for peroxidase and a chemiluminescence enhancer (ECL).
Figure 3. Dose response treatment with the fatty acids. Superoxide anion levels in the incubation media for the Rat 1 (A, B and C) and 3T3 Swiss fibroblasts (D, E and F) (2.5 \times 10^6 cells per mL) containing \textit{b}-NADH (10 \muM), as measured by the lucigenin assay in the absence and presence of different concentrations of oleic (A and D), linoleic (B and E) or \textit{g}-linolenic (C and F) acid (0, 5, 50, 100 or 200 \muM). The results are presented as the mean \pm SEM of at least three experiments performed in triplicate. Presented as the effect of the fatty acids compared with the control, *p<0.05, ** p<0.01 and *** p<0.001.

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Western Blotting System Kit, GE Health Care, Little Chalfont, Buckinghamshire, England) for 1 min and immediately exposed to X-ray film. The films were then developed. The band intensities were quantified using optical densitometry.

2.7. Real-Time Reverse Transcriptase Chain Reaction-PCR

After being incubated at 37°C for 1 hour in the absence or presence of fatty acids (100 μM), RNA was obtained from Rat 1 fibroblasts using the Trizol reagent. The expression of NADPH oxidase component p47phox was evaluated using real-time PCR in a Rotor Gene 3000 PCR machine (Corbett Research, Mortlake, Australia) using the method described by Higuchi et al. (1992) [21]. cDNA probes were synthesised using 4 μg of total RNA and a mixture of the following reagents: 146 ng of “random primers”, 200 U of Superscript reverse transcriptase, 5× reaction buffer (50 mM Tris – HCl (pH 8.0), 75 mM KCl, 3 mM MgCl2), 5 mM DTT, and 500 μM dNTP in a final volume of 20 μL. The reaction was incubated for 2 min (25°C) to assemble the oligonucleotides and hybridise the RNA, which was followed by heating at 42°C for 50 min. The cDNA was stored at −20°C prior to the real-time PCR assay. To perform the real-time PCR reaction, 1 μL of cDNA was used in a final volume of 25 μL, which contained 100 μM of dNTPs, 10× reaction buffer (10 mM Tris–HCl, 50 mM KCl, 2 mM MgCl2, 1 U of Taq DNA polymerase, 0.1 μM of each primer (sense and antisense), and SYBR GREEN (1000× diluted) as the fluorescent dye. The primer sequences were designed using information from GenBank (National Center for Biotechnology Information (NCBI)). The quantification of gene expression was performed using the method described by Liu and Saint (2002) and the β2 myosin gene as an internal control [22].

2.8. Statistical analyses

Comparisons were performed using a one-way ANOVA and Dunnett’s Test. The significance was set at p<0.05. The results were obtained from 3 to 5 independent experiments and are expressed as the mean ± SEM.

Results

To assess any possible interference of the fatty acids with the ROS reaction involving the reagents, the appropriate controls were performed by using 10, 25, 50, 100, and 200 μM oleic, linoleic or γ-linolenic acids in the assays (lucigenin and phenol red) without cells. The three fatty acids did not directly affect the lucigenin and phenol red assays, as reported in our previous study [15]. Controls using the vehicle (ethanol) for the fatty acids were also provided.

In order to verify the influence of fatty acids on fibroblast viability, flow cytometric analysis was used to determine cell membrane integrity and DNA fragmentation of fibroblasts treated with oleic, linoleic and γ-linolenic acids (25, 50, 100 and 200 μM) for 30 minutes. No significant effect of treatments were found at experimental condition of this study (Figure 1).

In the presence of β-NADH (10 μM), a substrate for the oxidases [23], [24], the oleic, linoleic and γ-linolenic acids increased the extracellular superoxide anion (O2−) level, as indicated by the lucigenin-amplified chemiluminescence assay (Figure 2). Significant superoxide levels were immediately detected after the fatty acid treatments and remained elevated for at least 20 min in the treated fibroblasts. In the absence of β-NADH or NADPH, there was no chemiluminescence above the background level. The kinetic studies indicated that the induction of superoxide production in the fibroblasts is a fast event that occurs within minutes after treatments with the oleic, linoleic and γ-linolenic acids (Figure 2). Our results demonstrate that the three fatty acids increased fibroblast oxidative bursts as follows: γ-linolenic > linoleic > oleic. The period for maximal superoxide production was 8.2±2.6 min for oleic acid, 3.8±1.6 min for linoleic, and...
3.1 ± 1.2 min for γ-linolenic. An interesting aspect is that as the number of double bonds in the 18-carbon fatty acid molecule increased, so did its ability to stimulate oxidative bursts in the 3T3 Swiss and Rat 1 fibroblasts.

There was a positive correlation between fatty acid concentration (zero, 5, 25, 50, 100, or 200 μM) and superoxide production (Figure 3). The Pearson correlation coefficient determined for the Rat 1 fibroblasts was: r = 0.97 and p = 0.001 for oleic acid, r = 0.97 and p = 0.0008 for linoleic acid and r = 0.89 and p = 0.015 for γ-linolenic acid. The Pearson correlation coefficient for the Swiss 3T3 fibroblasts was: r = 0.63 and p = 0.1 for oleic acid, r = 0.75 and p = 0.08 for linoleic acid and r = 0.49 and p = 0.3 for γ-linolenic acid. No detectable superoxide production occurred when the fibroblasts were treated for 10 min with oleic, linoleic or γ-linolenic acid without the addition of β-NADH.

The addition of SOD (150 U/mL), a scavenger of the superoxide anion, to the assay system partially inhibited the chemiluminescence induced by the fatty acids. The responses of the 3T3 Swiss and Rat 1 fibroblasts were similar. The SOD control was performed by using inactive SOD (a heat-killed preparation), and the chemiluminescence induced by the fatty acids was not modified (data not shown).

The effect of the oleic, linoleic and γ-linolenic acids on fibroblast superoxide production was severely attenuated, i.e., reductions of 79%, 92% and 82%, respectively, when the cells were pre-treated for 5 min with diphenyleneiodonium (DPI), a specific inhibitor of NADPH oxidase (Figure 4).

The effect of oleic, linoleic and γ-linolenic acids on the phosphorylation of the NADPH oxidase component p47phox in Rat 1 fibroblasts was investigated using Western blot analysis. The phosphorylation of p47phox is a rapid event, and the activation of p47phox was analysed after 5, 10 and 15 min in Rat 1 fibroblasts incubated in the presence of oleic, linoleic or γ-linolenic acid (Figure 5). An increase in p47phox mRNA expression following fatty acid treatment was observed by real-time PCR (Table 1).

### Table 1. Effect of oleic, linoleic and γ-linolenic acids on the mRNA expression of NADPH oxidase component p47phox in Rat 1 fibroblasts.

<table>
<thead>
<tr>
<th>p47phox mRNA (relative units)</th>
<th>(mean ± SD)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Control</td>
<td>1 ± 0.04</td>
</tr>
<tr>
<td>Oleic acid</td>
<td>1766 ± 176***</td>
</tr>
<tr>
<td>Linoleic acid</td>
<td>318 ± 31***</td>
</tr>
<tr>
<td>γ-Linolenic acid</td>
<td>408 ± 314***</td>
</tr>
</tbody>
</table>

Presented as the effect of the fatty acids compared with the control, *p < 0.05, ** p < 0.01 and *** p < 0.001.

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The Pearson correlation coefficient for the Swiss 3T3 fibroblasts was: r = 0.63 and p = 0.1 for oleic acid, r = 0.75 and p = 0.08 for linoleic acid and r = 0.49 and p = 0.3 for γ-linolenic acid. No detectable superoxide production occurred when the fibroblasts were treated for 10 min with oleic, linoleic or γ-linolenic acid without the addition of β-NADH.

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The effect of the oleic, linoleic and γ-linolenic acids on the phosphorylation of p47phox (an active catalytic subunit of NADPH oxidase) was investigated using Western blot analysis. Because the phosphorylation of p47phox is a rapid event, the activation of p47phox was analysed after 5, 10 and 15 min in Rat 1 fibroblasts incubated in the presence of oleic, linoleic or γ-linolenic acid (Figure 5). An increase in p47phox mRNA expression following fatty acid treatment was observed by real-time PCR (Table 1).
The treatment of fibroblasts with oleic, linoleic or γ-linolenic acid did not significantly increase intracellular levels of ROS, as indicated by hydroethidine reduction. In addition, these fatty acids did not significantly increase the basal levels of H$_{2}$O$_{2}$ in the 3T3 Swiss and Rat 1 fibroblasts, as indicated by the phenol red reduction assay under the conditions described herein (data not shown).

Discussion

In this work, evidence is presented that oleic, linoleic and γ-linolenic acids increase extracellular superoxide levels in fibroblast cell lines through the activation of p47phox and the stimulation of NADPH oxidase. We also observed that oleic, linoleic and γ-linolenic acids increase p47phox mRNA expression, suggesting that these fatty acids induce the synthesis of this regulatory subunit.

Our results demonstrate that the tested fatty acids increased the fibroblast oxidative burst as follows: γ-linolenic > linoleic > oleic acids. Studies involving fatty acid structure and function have demonstrated that, in general, as the number of double bonds in the fatty acid molecule increases, so does its ability to stimulate oxidative bursts in unstimulated neutrophils [15]. According to Hardy (1994), exogenous long chain and very long chain fatty acids use the same signal transduction pathways to stimulate ROS production by neutrophils [25].

Our results suggest that NOX and p47phox are implicated in ROS production by fatty acids treated fibroblasts; however in the cases of oleic acid and γ-linoleic acid, the inhibition is incomplete, suggesting that additional mechanisms may be at work. Superoxide can also be generated through the mitochondrial electron transport chain, xanthine – xanthine oxidase and cytochrome P450. Mitochondria generate superoxide mostly through the univalent reduction of oxygen in complexes I and III of the electron transport chain.

NADH cannot cross biological membranes under normal conditions. The cytosolic concentration of β-NADH is 270 μM, whereas the mitochondrial concentration is 638 μM [26]. If tissue damage occurs, for instance during periods of prolonged ischaemia and/or cellular death due to necrosis, a loss of plasma membrane integrity is observed [26]. Under these conditions, fibroblasts are able to produce ROS in response to fatty acids. These conditions are present during acute inflammation and in the skeletal muscle of athletes during high intensity exercises, such as the marathon and triathlon. We have demonstrated that pre-treating 3T3 Swiss and Rat 1 fibroblasts for 5–10 min with oleic, linoleic or γ-linolenic acid enhances fibroblast ROS production when compared with a β-NADH or NADPH treatment only.

The increased superoxide production observed when fibroblasts are treated with fatty acids plus β-NADH or NADPH may indicate an effect of metabolites as priming agents. Classical priming agents alone are not able to stimulate cell oxygen consumption and ROS production but lead to an incremental increase in the maximal rate of oxygen consumption when a second stimulus occurs [27], [28]. Hardy et al. (1994) demonstrated that pre-treating neutrophils with arachidonic, eicosapentaenoic or docosahexaenoic acids enhances their ability to respond to either fMLP or PMA, thereby producing more superoxide than when challenged with the stimulators only [25]. Recently, we demonstrated that the addition of PMA to an assay medium leads to an additive effect on the superoxide and hydrogen peroxide production induced by oleic, linoleic and γ-linolenic acids in neutrophils [15].

In infectious/inflammatory processes, the source of fatty acids may be blood, extracellular fluids, bacterial cell membranes or infiltrating leukocytes. Under these conditions, levels of fatty acids and metabolites are increased, as are the number of dying cells, and hence, the β-NADH levels. In these situations, fatty acids released into the microenvironment can be an important mediator for the process of the resolution/progression of the damaged tissue through increased ROS production.

The high concentrations of fatty acids associated with the increased microvascular permeability observed in some diseases [29] may define specific loci, such as the interstitial space and body cavities, for fatty acid action. At these sites, fatty acids may activate neutrophils and macrophages and promote fibroblast ROS release. Fibroblast proliferation and fibrogenesis are important factors that lead to the complications of various diseases, such as atherosclerosis, rheumatic arthritis, diabetic nephropathy and retinopathy [30]. The data from this study indicate that oleic, linoleic and γ-linolenic acids are important inducers of ROS production by fibroblasts. This effect may be critical during certain physiological processes, such as wound healing. However, this effect could also be deleterious in proliferative diseases that involve fibroblast dysfunction, such as fibrosis, and by promoting uncontrolled inflammation [31]. These outcomes are more relevant in pathological conditions involving persistent increases in fatty acid serum levels, such as diabetes [32].

O’Donnell et al. (1996) demonstrated that fibroblasts treated with arachidonic, linoleic or (5S)-hydroxyeicosatetraenoic acid ([5S]-HETE) present enhanced superoxide generation. Their study suggests the involvement of 15-lipoxygenase on ROS production [26]. Maziere et al. (1999) reported that treating cultured human fibroblasts with oleic, linoleic or γ-linolenic acid increases their intracellular levels of ROS and lipid peroxidation products. This group also demonstrated the activation of the oxidative stress-responsive transcription factors AP1 and NF-κB [33].

Several studies have demonstrated the presence of NADPH oxidases in the homogenates and particulate fractions of endothelial and smooth muscle cells and that these NADPH oxidases are able to generate superoxide or hydrogen peroxide. Here, evidence has been presented that oleic, linoleic and γ-linolenic acids are potent inducers of ROS production in fibroblasts. These fatty acids stimulate ROS production via the dose-dependent activation of the NADPH oxidase complex. Excessive ROS production can damage cellular lipids, proteins and DNA, which impairs cell function. In fact, oxidative stress has been implicated in a number of human diseases and in the ageing process.

An increasing body of evidence indicates that in diseases such as metabolic syndrome [34], sepsis [35] and diabetes [36], the plasmatic concentrations of free fatty acids and ROS are higher. Considering the findings of the present study, we hypothesize that fatty acids, through their effects on fibroblasts, may contribute to the pro-oxidant state observed in these pathological conditions. Additionally, taking into account the fact that fibroblasts are essential cells in wound healing, a process that is impaired in diabetic individuals, fatty acids-induced production of ROS by fibroblasts may, together with changes in other cell types, be important in this context. In fact, ROS can activate neutrophils/macrophages to produce pro-inflammatory cytokines via NF-κB activation, and the pro-inflammatory status may contribute to chronic inflammation in non-healing wounds or in insulin resistance.

Author Contributions

Conceived and designed the experiments: EH AD HAA RC. Performed the experiments: EH AD AEH MARV. Analyzed the data: EH AD PN HAA RC. Contributed reagents/materials/analysis tools: ARC HAA RC. Wrote the paper: EH RC. Revised the article critically: EH AD HAA RC.
References


