

Docosahexaenoic acid and other fatty acids induce a decrease in pH_i in Jurkat T-cells

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1 Docosahexaenoic acid (DHA) induced rapid ($t_{1/2} = 33$ s) and dose-dependent decreases in pH_i in BCECF-loaded human (Jurkat) T-cells. Addition of 5-(*N,N*-dimethyl)-amiloride, an inhibitor of Na^+/H^+ exchanger, prolonged DHA-induced acidification as a function of time, indicating that the exchanger is implicated in pH_i recovery.

2 Other fatty acids like oleic acid, arachidonic acid, eicosapentaenoic acid, but not palmitic acid, also induced a fall in pH_i in these cells.

3 To assess the role of calcium in the DHA-induced acidification, we conducted experiments in Ca^{2+} -free (0% Ca^{2+}) and Ca^{2+} -containing (100% Ca^{2+}) buffer. We observed that there was no difference in the degree of DHA-induced transient acidification in both the experimental conditions, though pH_i recovery was faster in 0% Ca^{2+} medium than that in 100% Ca^{2+} medium.

4 In the presence of BAPTA, a calcium chelator, a rapid recovery of DHA-induced acidosis was observed. Furthermore, addition of $CaCl_2$ into 0% Ca^{2+} medium curtailed DHA-evoked rapid pH_i recovery. In 0% Ca^{2+} medium, containing BAPTA, DHA did not evoke increases in $[Ca^{2+}]_i$, though this fatty acid still induced a rapid acidification in these cells. These observations suggest that calcium is implicated in the long-lasting DHA-induced acidosis.

5 DHA-induced rapid acidification may be due to its deprotonation in the plasma membrane (flip-flop model), as suggested by the following observations: (1) DHA with a $-COOH$ group induced intracellular acidification, but this fatty acid with a $-COOCH_3$ group failed to do so, and (2) DHA, but not propionic acid, -induced acidification was completely reversed by addition of fatty acid-free bovine serum albumin in these cells.

6 These results suggest that DHA induces acidosis *via* deprotonation and Ca^{2+} mobilization in human T-cells.

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Keywords: Docosahexaenoic acid; Jurkat T-cells; intracellular pH_i ; calcium

Abbreviations: AA, arachidonic acid (20:4 n-6); BAPTA-AM, bis-(*o*-aminophenoxy)ethane-*N,N,N',N'*-tetra-acetic acid tetra-acetoxymethyl ester; BSA, bovine serum albumin; $[Ca^{2+}]_i$, free intracellular calcium concentration; DHA, docosahexaenoic acid (22:6 n-3); DMA, 5-(*N,N*-dimethyl)-amiloride; EPA, eicosapentaenoic acid (20:5 n-3); NHE, Na^+/H^+ exchanger; OA, oleic acid (18:1 n-9); PA, propionic acid; pH_i , intracellular pH_i ; PUFA, polyunsaturated fatty acid; TCR, T-cell receptor; VPATPase, vacuolar-type H^+ -ATPase

Introduction

Under physiological conditions, activation of T-cells is initiated by an interaction between an antigen-specific T-cell receptor (TCR) and a foreign antigen associated with a membrane-bound major histocompatibility complex molecule (Grinstein & Dixon, 1989). This event triggers a number of intracellular biochemical processes that assure T-cell cycle progression (Grinstein & Dixon, 1989; Germain, 1994). The early stages of T-cell activation, initiated within 1–100 s of T-cell engagement, involve mobilization of intracellular calcium, activation of protein kinase C (PKC), expression of immediate early genes, interleukin-2 synthesis and its secretion (Cantrell, 1996). The late events that assure transition of cells from S phase to the G_2/M phase of the cell cycle are sustained

responses, implicating the prolonged PKC activation and the expression of receptors for interleukin-2, IL-2R (Ward, 1996). Regulation of cytosolic pH_i is crucial to many cellular functions, particularly for T-cell proliferation (Grinstein & Dixon, 1989). A rapid increase in intracellular pH is associated with the activation of a cascade of second messengers during T-cell proliferation (Gerson *et al.*, 1982). Although changes in pH_i and Ca^{2+} have been observed in many cell systems, it is not yet clear which of these ionic events, if any, are necessary steps in leading to DNA synthesis (Ives & Daniel, 1987).

Increase in cytosolic pH is reported to be due to the functioning of one of the membrane-bound transporters, the Na^+/H^+ exchanger (NHE), which mediates an electroneutral exchange between extracellular Na^+ and intracellular H^+ , and plays a major role in the regulation of cell volume (Parker & Castranova, 1984) and pH homeostasis (Grinstein & Dixon, 1989) in several cell types. The exchanger is pH_i dependent and

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activated allosterically by intracellular protons. Indeed, acidic intracellular pH activates it, but the exchanger remains relatively inactive at neutral pH_i (Orlowski & Grinstein, 1997). NHE1 isoform has been identified in human T-cells and is one of the main mechanisms responsible for proton extrusion in these cells (Siczkowski *et al.*, 1994). However, studies on pH_i changes have yielded to conflicting results on whether Na^+/H^+ activation in stimulated T-cells was dependent on Ca^{2+} influx (Rosoff & Cantley, 1985).

During the recent past, there has been an upsurge of information on the role of polyunsaturated fatty acids (PUFAs) in the regulation of immune cell functions (Khan & Hichami, 2002). The PUFAs of n-3 series, eicosapentaenoic acid (EPA, 20:5 n-3) and docosahexaenoic acid (DHA, 22:6 n-3), have been considered as immunosuppressors, whereas PUFAs of n-6 series (viz arachidonic acid, AA, 20:4 n-6) have been reported to be immunostimulators (Calder, 1996). The n-3 PUFAs have been shown to exert protective effects against rheumatoid arthritis (Kremer *et al.*, 1990), psoriasis (Bittiner *et al.*, 1988) and multiple sclerosis (Bates *et al.*, 1989). Moreover, nutritional interventions, conducted in mice which were fed an EPA- and DHA-rich diet, have shown that n-3 PUFAs diminished IL-2 secretion associated with a lesser expression of the α chain of the IL-2R in T-cells (Jolly *et al.*, 1998). As far as pH_i homeostasis is concerned, AA has been shown to induce a decrease in pH_i in the cytoplasm and nucleoplasm of rat cerebellar granule cells (Chen *et al.*, 2001), rat cardiac myocytes (Wu *et al.*, 2000), platelets (Cavallini *et al.*, 1996) and rat thymocytes (Astashkin *et al.*, 1993). Nevertheless, no detailed information is available on the role of n-3 PUFAs in the modulation of intracellular pH in human T-cells. Therefore, the present study was designed in order to shed light on the role of DHA, an n-3 PUFA, in the modulation of pH_i in (Jurkat)T-cells.

Methods

Chemicals

The culture medium RPMI-1640, L-glutamine, HEPES buffer, streptomycin/penicillin and fetal calf serum (FCS) were purchased from Biowhitaker, Belgium. 2',7'-bis-(carboxyethyl)-5(6')-carboxyfluorescein (BCECF)/acetoxymethylester (AM) and Fura-2/AM were obtained from Molecular Probes, Eugene, OR, U.S.A. Unless otherwise stated, all other chemicals including DHA (22:6 n-3), DHA methyl ester, AA (20:4 n-6), EPA (20:5 n-3), oleic acid (OA, 18:1 n-9), palmitic acid (16:0), nigericin, bumetanide, oligomycin and bafilomycin were purchased from Sigma (St Louis, MO, U.S.A.).

Cell culture and lipid analysis

The human (Jurkat) T-cells were routinely cultured in RPMI-1640 medium supplemented with 10% FCS, 2 mM L-glutamine, 50 $\mu\text{g ml}^{-1}$ penicillin-streptomycin and 20 mM HEPES at 37°C, in a humidified chamber containing 95% air and 5% CO_2 . Cell viability was assessed by trypan blue exclusion test. Cell numbers were determined by a haemocytometer.

Fatty acid derivatives were analysed by using high-performance liquid chromatography (HPLC). Cells were

incubated for 1 h in the presence of (^{14}C)DHA (10 μM) and then washed three times in RPMI 1640 medium (0.2% bovine serum albumin (BSA)). Total lipids were extracted by chloroform/methanol/NaCl 2 M (1:1:0.9, v/v/v) and neutral lipids were separated by thin-layer chromatography in the following solvent system: hexane/diethyl ether/acetic acid (90:30:1). Radio-labelled spots corresponding to fatty acids were scraped off and extracted by hexane. Fatty acid classes were separated by reverse-phase HPLC using a 5 μM LiChrosorb RP 18 and the following solvents: acetonitrile/water (95:5, v/v). A volume of 1 ml min^{-1} fraction was collected, dried, and the radioactivity of each fraction was quantified. The retention times of fatty acids were determined using authentic standards.

Measurement of intracellular pH (pH_i)

For these experiments, cells ($2 \times 10^6 \text{ ml}^{-1}$) were loaded with the pH-sensitive fluorescent probe BCECF-AM (3 μM) for 30 min at 37°C under a 95% air–5% CO_2 atmosphere, in a nominally bicarbonate-free medium containing: 140 mM NaCl; 5 mM KCl; 1 mM MgSO_4 ; 1 mM CaCl_2 ; 20 mM HEPES; 1 mM NaH_2PO_4 and 5.5 mM glucose, pH 7.4. After loading, the cells were washed three times (720 g \times 10 min) to remove the extracellular dye, and made to remain suspended in the identical buffer. A sample (4×10^6 cells) was then transferred to a magnetically stirred cuvette in a PTI spectrofluorometer. BCECF fluorescence was detected in the ratio mode at the excitation wavelengths of 440 and 506 nm, and at an emission wavelength of 535 nm. Calibration of fluorescence vs pH was performed by equilibration of external and internal pH with nigericin (10 μM) in a high K^+ buffer and by adding aliquots of propionic acid (PA; 0.1 M). The relative fluorescence ratio values were plotted against corresponding pH_i values, which allowed determination of the unknown pH_i .

For the experiments conducted in the absence of external calcium (0% Ca^{2+}), CaCl_2 was replaced by 1 mM EGTA in the buffer. All test molecules were added in small volumes with no interruption in recordings. Fatty acids were dissolved in ethanol (w/v, 0.1%) and used immediately or kept at -20°C , tightly sealed under the stream of nitrogen.

Measurement of free intracellular Ca^{2+} concentrations, $[\text{Ca}^{2+}]_i$

The cells ($2 \times 10^6 \text{ ml}^{-1}$) were washed with phosphate-buffered saline, pH 7.4, and then incubated with Fura-2/AM (1 μM) for 60 min at 37°C in a loading buffer containing: 110 mM NaCl; 5.5 mM KCl; 25 mM NaHCO_3 ; 0.8 mM MgCl_2 ; 0.4 mM KH_2PO_4 ; 0.33 mM Na_2HPO_4 ; 20 mM HEPES; 1.2 mM CaCl_2 , and the pH was adjusted to 7.4.

After loading, the cells were washed three times (720 g \times 10 min) and made to remain suspended in the identical buffer. The fluorescence intensities were measured in the ratio mode in a PTI spectrofluorometer at 340 and 380 nm (excitation filters) and 510 nm (emission filters). The cells were continuously stirred throughout the experiment. The intracellular concentration of free Ca^{2+} , $[\text{Ca}^{2+}]_i$, was calculated using the equation: $[\text{Ca}^{2+}]_i = K_d \times (R - R_{\min}) / (F_{\max} - F) (\text{Sf2/Sb2})$. A value of 224 nM for K_d was added into the calculations. R_{\max} and R_{\min} values were obtained by addition of ionomycin (5 μM) and MnCl_2 (2 mM), respectively.

Experiments were also conducted in Ca^{2+} -free medium (0% $CaCl_2$ in the phosphate-buffered saline was replaced by EGTA (1.2 mM)). The test molecule DHA was added into the cuvettes in small volumes, with no interruptions in the recordings.

Confocal microscopy

The cells ($2 \times 10^6 ml^{-1}$) were loaded with the pH-sensitive fluorescent probe BCECF-AM ($3 \mu M$), as described for pH_i measurements. The cells were transferred to the microscopic slide that was examined by an oil-immersion objective ($\times 63$) of Leica TCS 4D confocal microscope, equipped with a numerical aperture of 1.4. The cells were excited by the laser beam at 488 nm and the excited fluorescence was detected at 515 nm. The X – Y plane images (512 pixels \times 512 pixels) were generated from fluorescence emission images collected with a band pass filter.

Statistical analysis

Results are shown as mean \pm s.e.m. for a given number of experiments (n). The data were analysed by using Statistica (4.1 version, Statsoft, Paris, France). The significance between mean values was determined by analysis of variance one way, followed by a least-significant-difference (LSD) test, with a value of $P < 0.05$ being considered statistically significant.

Results

DHA induces a decrease in intracellular pH in a dose-dependent manner

The resting level of pH_i in (Jurkat) T-cells in the nominally bicarbonate-free medium was 7.22 ± 0.03 ($n = 30$). DHA induced a rapid ($t_{1/2} = 33$ s) and dose-dependent decrease in pH_i that plateaued at the concentration of $10 \mu M$. During the time course of the experiments (20 min), complete recovery of the resting pH_i could not be observed. In the subsequent experiments, we used DHA at $10 \mu M$ (Figure 1a,b). Figure 1c shows that addition of DHA induces acidosis only in the cytoplasm, without significantly influencing the BCECF fluorescence in the nucleoplasm, of Jurkat T-cells.

DHA exerts additive effects in the presence of a weak acid

We employed PA, known to induce the cytosolic acid load by directly entering the cells and without accumulating in the phospholipid bilayer (Chen *et al.*, 2001). Addition of PA evoked an intracellular acidification of 0.32 ± 0.03 pH units ($n = 8$) (Figure 2b). When DHA was added after PA, the former induced a slight acidification (0.13 ± 0.02 pH units) that was rapidly recovered ($0.0035 \pm 0.02 \Delta pH \min^{-1}$) (Figure 2b). When PA was applied after DHA, we observed an additive acidification of 0.16 ± 0.02 pH units (Figure 2a).

DHA induces an increase in $[Ca^{2+}]_i$: possible role of calcium in the DHA-induced intracellular acidification

In order to determine the contribution of Ca^{2+} in the DHA-induced intracellular acidification, we first measured

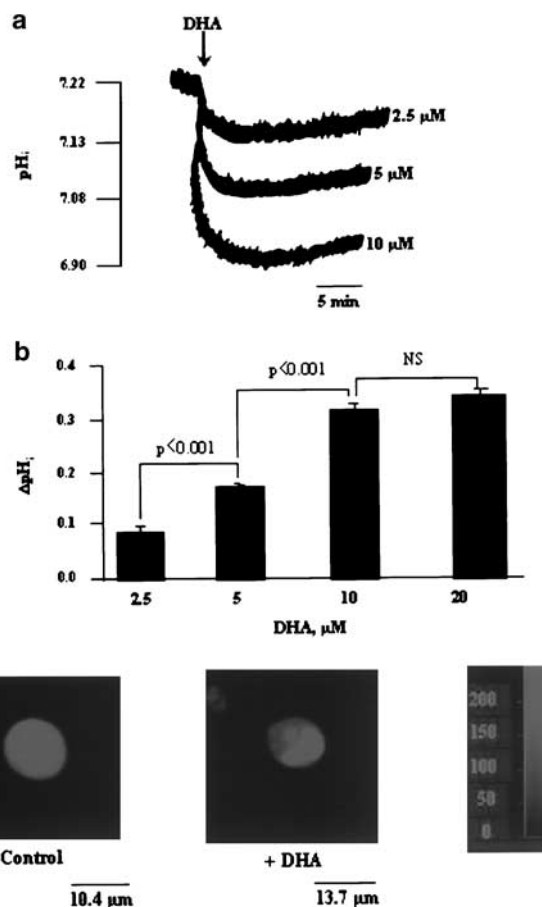


Figure 1 Effects of DHA on pH_i . Cells (4×10^6 assay $^{-1}$) were loaded with the fluorescent dye BCECF/AM, as described in Methods. (a) Intracellular acidosis induced by different concentrations of DHA, that is, 2.5, 5, 10 and 20 μM . (b) ΔpH_i represents the amplitude of the intracellular acidification produced. Values are expressed as mean \pm s.e.m. of independent experiments ($n = 30$). (c) Acidosis induced by DHA ($10 \mu M$), as visualized by confocal microscopy ($n = 6$). NS = insignificant differences.

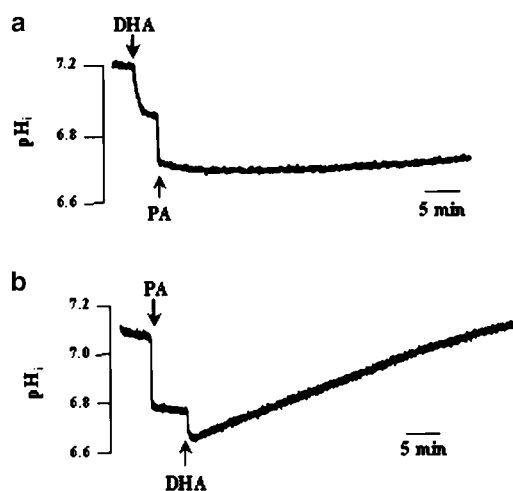


Figure 2 Effects of DHA and PA on pH_i . Cells (4×10^6 assay $^{-1}$) were loaded with the fluorescent dye BCECF/AM, as described in Methods. The arrowheads indicate the time when DHA ($10 \mu M$) and PA (2.5 mM) were added into the cuvette, without interruption in the recordings. The figure shows single traces of observations, which were reproduced independently ($n = 8$). All experiments were performed in the bicarbonate-free medium, pH 7.4.

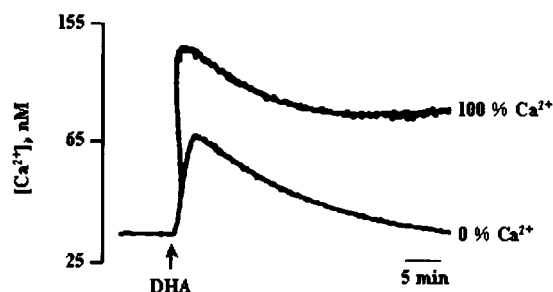


Figure 3 Effects of DHA on $[Ca^{2+}]_i$. Cells (4×10^6 assay $^{-1}$) were loaded with the fluorescent dye Fura-2/AM, as described in Methods, and then resuspended in the 100% or 0% Ca^{2+} buffers. DHA ($10 \mu M$) was added into the cuvette without interruptions in the recordings ($n = 4$).

intracellular free calcium after treatment with DHA. Addition of DHA induced an increase in $[Ca^{2+}]_i$ ($97.63 \text{ nM} \pm 0.03$) that was significantly curtailed when the experiments were conducted in 0% Ca^{2+} buffer ($28.5 \text{ nM} \pm 0.03$), as compared to that in 100% Ca^{2+} buffer (Figure 3). Indeed, the DHA-induced rise in $[Ca^{2+}]_i$ was inhibited by $70.8 \pm 1.2\%$ in 0% Ca^{2+} medium.

The role of calcium on the DHA-induced changes in pH_i was examined by conducting experiments in 0% Ca^{2+} buffer. In the 0% Ca^{2+} medium, DHA induced intracellular acidification to the same extent as in the 100% Ca^{2+} medium (0.27 ± 0.02 pH units); however, the recovery of the resting pH_i was faster in the former ($0.0093 \pm 0.01 \Delta pH \text{ min}^{-1}$) (Figure 4a). Then, we designed a Ca^{2+} -free/ Ca^{2+} reintroduction protocol that consisted of conducting the experiment in 0% Ca^{2+} buffer and adding, exogenously, first DHA and then $CaCl_2$. Hence, we observed that addition of $CaCl_2$ restored the DHA-induced prolonged decrease, as was observed in the 100% Ca^{2+} medium (Figure 4b). We, further, clamped intracellular free calcium, by using the calcium chelator bis-(*o*-aminophenoxy)ethane-*N,N,N',N'*-tetra-acetic acid tetra-acetoxymethyl ester (BAPTA-AM). T-cells were preincubated with BAPTA-AM ($50 \mu M$) for 15 min, then washed and resuspended in the 100% Ca^{2+} medium. Figure 5a shows that when intracellular free calcium is chelated, the recovery of pH_i is faster ($0.01 \pm 0.02 \Delta pH \text{ min}^{-1}$) than that in the medium without BAPTA ($0.003 \pm 0.01 \Delta pH \text{ min}^{-1}$). When BAPTA is added to 0% Ca^{2+} medium, DHA did not induce increases in $[Ca^{2+}]_i$, (Figure 5b), though this fatty acid was able to induce rapid acidification under such experimental conditions (Figure 5c).

Effect of 5-(*N,N*-dimethyl)-amiloride (DMA), an NHE inhibitor, on DHA-induced intracellular acidification

NHEs are responsible for the regulation of intracellular pH and cell volume by extruding protons and taking up sodium ions into cells. To date, seven isoforms (NHE1–NHE7) have been identified and cloned. NHE6 and NHE7 are localized on recycling endosomes and the *trans*-Golgi network, respectively, whereas the other isoforms (NHE1–NHE5) are expressed in the cell plasma membrane (Grinstein *et al.*, 1989). Though immunoblotting experiments in human T-cells have shown that NHE1 is mainly responsible for pH_i regulation in these cells (Siczkowski *et al.*, 1994), no information is available, to

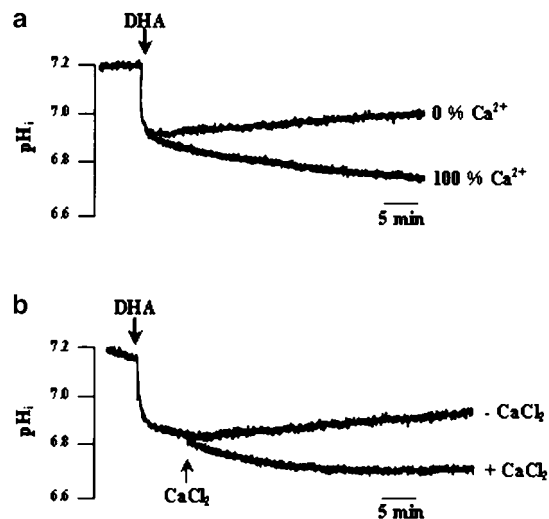


Figure 4 Role of external calcium in the DHA-induced decrease in pH_i . Cells (4×10^6 assay $^{-1}$) were loaded with the fluorescent dye BCECF/AM, as described in Methods. (a) Cells were resuspended in either 0% or 100% Ca^{2+} buffers. (b) Ca^{2+} -free/ Ca^{2+} -reintroduction experiment in 0% Ca^{2+} medium: $CaCl_2$ (2 mM) was applied after DHA ($10 \mu M$), as indicated in the figure. The figures show single traces of observations, which were reproduced independently ($n = 10$).

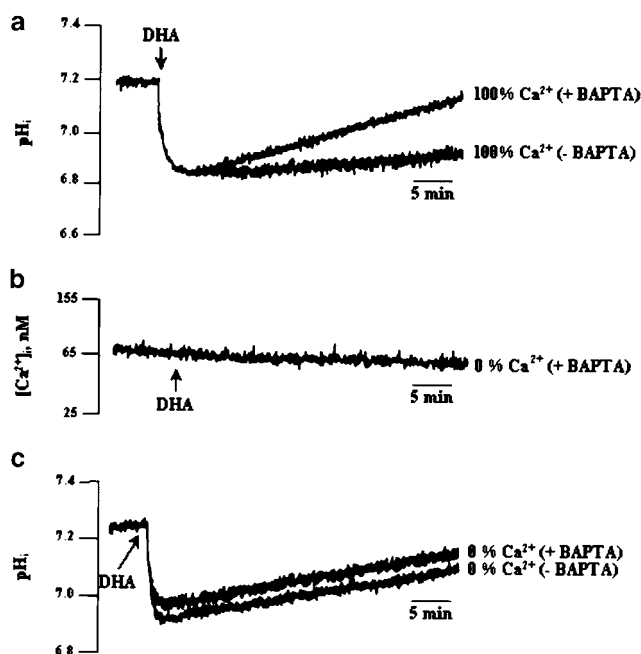


Figure 5 Effect of intracellular free calcium chelation on the DHA-induced decrease in pH_i . Cells (4×10^6 assay $^{-1}$) were loaded with the fluorescent dye BCECF/AM and preincubated with BAPTA/AM ($50 \mu M$, 15 min), as described in Methods. DHA ($10 \mu M$) was added at the end of the incubation. The figure shows single traces of observations, which were reproduced independently ($n = 6$).

our knowledge, whether this NHE subtype is implicated in the pH_i homeostasis in Jurkat T-cells. We, therefore, have termed it NHE.

It was of interest to assess whether the NHE was activated to counter the acidification evoked by DHA. We, therefore,

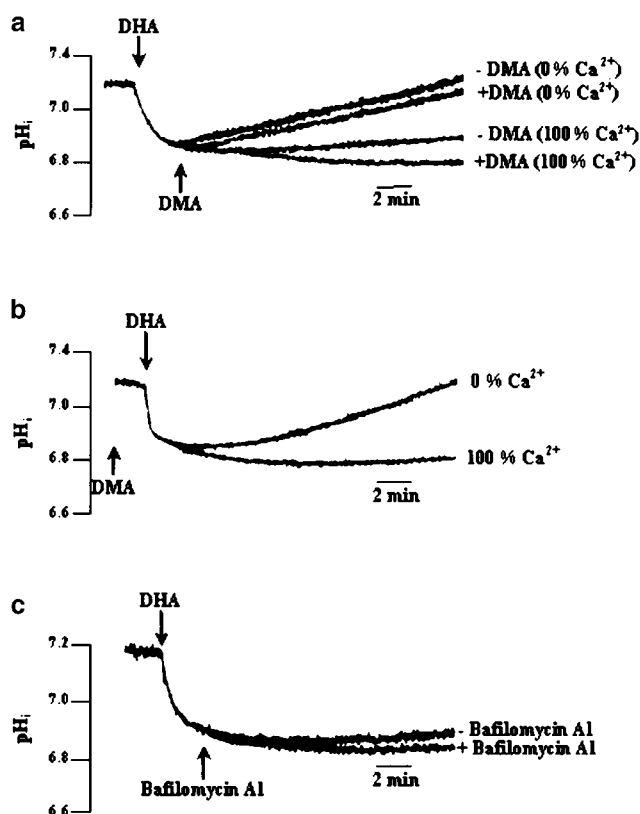


Figure 6 Effect of DMA and bafilomycin A1 on DHA-evoked decrease in pH_i . Cells (4×10^6 assay $^{-1}$) were loaded with the fluorescent dye BCECF/AM, as described in Methods. The arrow-heads indicate the time when the test molecules DHA ($10 \mu M$), DMA ($10 \mu M$) and bafilomycin A1 (50 nM) were added into the cuvette, without interruptions in the recordings. The figures show single traces of observations that were reproduced independently ($n = 11$). Experiments were performed both in the Ca^{2+} -containing (100% Ca^{2+}) and Ca^{2+} -free (0% Ca^{2+}) buffers.

employed the NHE1 inhibitor, DMA, that has a high affinity for the exchanger, a low cell permeability and only exhibits nonspecific effects at concentrations 10–100-fold higher than that used in the present study (Kleymann & Cragoe, 1988). When DMA was applied before or after DHA, in 100% Ca^{2+} medium, the DHA-induced acidification was enhanced (0.11 ± 0.04 pH units) and little recovery was observed (Figure 6a,b). However, incubation of cells with DMA in 0% Ca^{2+} medium did not affect the fast pH_i recovery observed in these calcium depletion experiments (Figure 6a). We were also tempted to assess whether the vacuolar-type H^+ -ATPase (VPATPase) was implicated in proton extrusion in T-cells. VPATPase has been described in a variety of cell types, being responsible for the acidification of the intracellular compartments, such as lysosomes and endosomes (Forgacs, 1989). In addition to their endomembrane distribution, they have also been detected in the plasma membranes of various cell types (i.e. macrophages), thereby maintaining the cytosolic pH by extruding protons out of the cells (Bidani & Brown, 1990). Figure 6c shows that addition of bafilomycin A1 (at 50 nM), a potent inhibitor of VPATPase, failed to prolong the DHA-induced acidification (with bafilomycin 0.35 ± 0.03 vs without bafilomycin 0.29 ± 0.02 pH units).

DHA-induced acidosis: lack of involvement of the $Na^+ - K^+ - 2Cl^-$ symport, free radical production, NADPH oxidase or mitochondrial H^+ leakage

We investigated whether the $Na^+ - K^+ - 2Cl^-$ symport, mainly responsible for chloride and sodium influx, was implicated in DHA-induced decrease in pH_i . We, therefore, used a specific inhibitor of this symport, bumetanide. Table 1 shows that prior addition of bumetanide had no effect on the DHA-evoked intracellular acidosis (with bumetanide 0.28 ± 0.05 pH units vs without bumetanide 0.31 ± 0.02 pH units). The involvement of NADPH oxidase and free radical production was also tested. Prior addition of $CdCl_2$ or $ZnCl_2$, the inhibitors of the NADPH oxidase, and of superoxide dismutase (SOD) failed to influence DHA-induced intracellular acidification. Carbonyl cyanide *m*-chlorophenyl-hydrazone (mCICCP), which dissipates H^+ gradient across the inner mitochondrial membrane and oligomycin, a F_0F_1 -ATPase/ H^+ pump inhibitor, were used in order to determine the contribution of mitochondrial H^+ in the DHA-induced response. When mCICCP or oligomycin were applied before DHA, no enhanced acidification was observed (Table 1). It is important to mention that any of the molecules tested modified neither the initial rate of acidification nor the resultant fall in pH_i , induced by DHA.

DHA-induced decrease in pH_i is reversed by BSA

DHA-induced response could be a result of simple diffusion (flip-flop) of the fatty acid, demonstrated to occur in phospholipid bilayers and responsible for intracellular acidification (Kamp & Hamilton, 1992). Albumin has been shown to compete with PUFAs bound to the plasma membrane and to extract them from the lipid bilayer, thus decreasing the rate of intracellular proton influx (Kamp & Hamilton, 1993; Kamp *et al.*, 1995; Hamilton, 1998). If flip-flop were to occur in our system, on addition of fatty acid-free BSA and in presence of DHA, a rise in pH_i should be observed. When 0.2% fatty acid-free BSA was added after DHA, a rapid recovery ($0.16 \pm 0.01 \Delta pH \text{ min}^{-1}$) of the pH_i was observed (Figure 7a). This effect of BSA on pH_i recovery was specific to the intracellular acidification evoked by DHA, since no effect was seen on the acidosis induced by PA (Figure 7b). Similarly, the addition of DHA methyl ester ($-COOH$ group replaced by $-COOCH_3$) had no effect on pH_i (Figure 7c).

Table 1 Intracellular acidosis induced by the addition of DHA under various conditions

Treatment	ΔpH_i	N
DHA alone	0.31 ± 0.02	18
$CdCl_2$ + DHA	0.29 ± 0.05	4
$ZnCl_2$ + DHA	0.27 ± 0.03	4
Bumetanide + DHA	0.28 ± 0.05	4
SOD + DHA	0.27 ± 0.03	4
Oligomycin + DHA	0.28 ± 0.02	4
mCICCP + DHA	0.28 ± 0.02	4

SOD, superoxide dismutase (0.25 U ml^{-1}); $CdCl_2$ and $ZnCl_2$ were used at 1 mM, bumetanide at $100 \mu M$, oligomycin and mCICCP at $10 \mu M$; n , number of experiments performed. The ΔpH_i under the above conditions did not differ significantly ($P > 0.05$) as compared to the control (DHA alone).

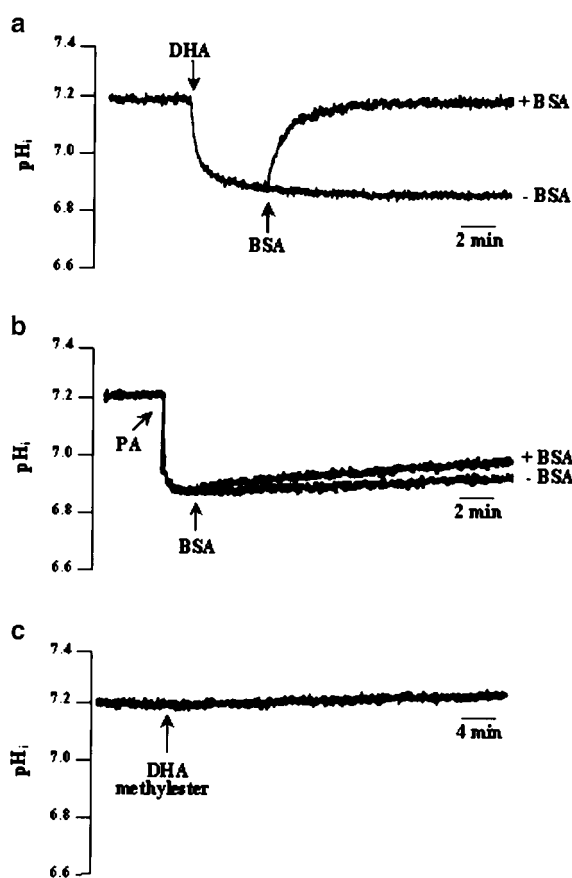


Figure 7 Effect of BSA and DHA methyl ester on pH_i. Cells (4×10^6 assay⁻¹) were loaded with the fluorescent dye BCECF/AM, as described in Methods. The arrowheads indicate the time when the test molecules DHA ($10 \mu\text{M}$), BSA ($0.2\% \text{ w v}^{-1}$), PA (2.5 mM) and DHA methyl ester ($10 \mu\text{M}$), were added into the cuvette, without interruptions in the recordings. The figures show single traces of observations that were reproduced independently ($n = 4$). All experiments were performed in the bicarbonate-free medium, pH 7.4.

Effect of other fatty acids on intracellular pH

To get insight into the specificity of DHA on intracellular acidification, other fatty acids of the n-9, n-6 and n-3 families were tested. DHA and these fatty acids were used in the same batches of cells. OA ($18:1$ n-9), AA ($20:4$ n-6) and EPA ($20:5$ n-3) were all able to acidify the cytoplasm of Jurkat T-cells, when used at $10 \mu\text{M}$ (Figure 8a), the concentration that gave for each fatty acid an optimal response (Table 2). However, DHA-induced acidification (0.34 ± 0.02 pH units) was significantly higher than that triggered by AA (0.22 ± 0.03 pH

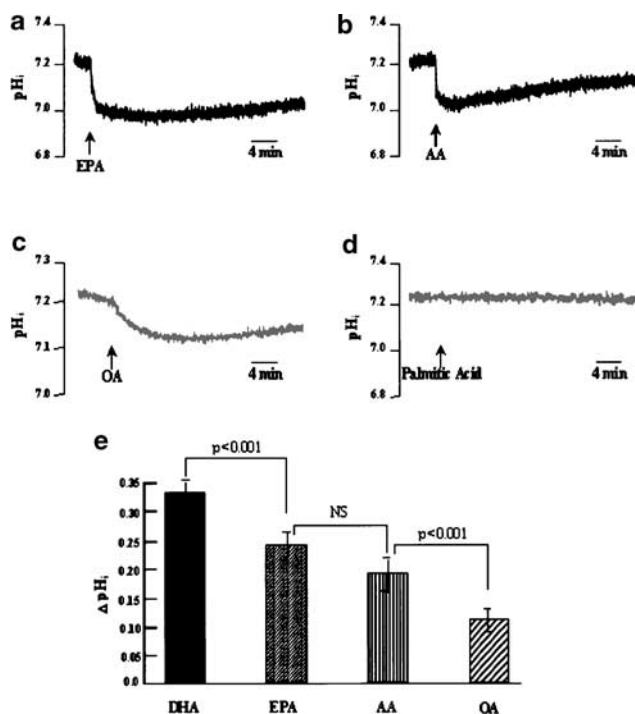


Figure 8 Effect of different fatty acids on pH_i. Cells (4×10^6 assay⁻¹) were loaded with the fluorescent dye BCECF/AM, as described in Methods. (a–d) Intracellular acidosis induced by different fatty acids: AA, arachidonic acid, EPA, eicosapentaenoic acid, OA, oleic acid and palmitic acid. All fatty acids were used at $10 \mu\text{M}$. (e) ΔpH_i represents the amplitude of the intracellular acidification produced by these agents. All fatty acids, including DHA, were used in the same batches of cells. Values are expressed as mean \pm s.e.m. of independent experiments ($n = 5$). NS = insignificant differences.

units) or OA (0.13 ± 0.04 pH units). EPA, the other important PUFA of the n-3 family, triggered a pH_i fall (0.25 ± 0.02 pH units) which was significantly lesser than the DHA-induced acidosis. EPA- and AA-induced responses were not significantly different (Figure 8a,b,e). In order to assess the implication of unsaturation of fatty acids, a saturated fatty acid was tested on pH_i. Addition of palmitic acid ($16:0$; at $10 \mu\text{M}$) to cells did not affect the intracellular pH. However, as palmitic acid has a low water solubility, we verified (not shown) using ¹⁴C-labelled palmitic acid that, under our experimental conditions, it did not precipitate, upon immediate addition, as the acid-soap in contact with the aqueous medium. This could have accounted for the palmitic acid inability to induce a pH_i drop. We observed that palmitic acid at $10 \mu\text{M}$ and even at $20 \mu\text{M}$ was almost completely soluble in the bicarbonate-free buffer (not shown).

Table 2 Intracellular acidosis (ΔpH_i) induced by AA, EPA and OA at different concentrations

Fatty acid	Concentrations (μM)		
	5	10	20
AA	0.12 ± 0.03	0.21 ± 0.03^a	0.24 ± 0.02^b
EPA	0.17 ± 0.03	0.27 ± 0.03^a	0.29 ± 0.02^b
OA	0.05 ± 0.03	0.14 ± 0.03^a	0.18 ± 0.02^b

^aRepresents significant values ($P < 0.001$), as compared to $5 \mu\text{M}$ concentration of the respective fatty acid.

^bRepresents insignificant values, as compared to $10 \mu\text{M}$ concentration of the respective fatty acid ($n = 7$).

Discussion

PUFAs of n-3 series, particularly DHA and EPA, have been shown to exert immunosuppressive effects (Calder, 1996); however, their mechanisms of action in T-cell activation have not been well elucidated. Intracellular pH being an important factor of T-cell proliferation, we undertook the present study to elucidate the role of an n-3 PUFA, DHA, on the modulation of pH_i . DHA, the end product of the metabolism of α -linolenic acid (Lands, 1991), evoked an intracellular acidification. Since AA has been shown to induce acidosis both in the cytoplasm and nucleoplasm in rat brain cells (Chen *et al.*, 2001), we used confocal microscopy to localize *in situ* acidification. DHA-induced acidification was only observed in the cytoplasmic area of Jurkat T-cells. In order to shed light on the mechanism(s) by which DHA could act, PA, a weak acid, was employed. When DHA was applied after PA or *vice versa*, additive effects were observed, indicating that these two agents act differently. Indeed, PA does not accumulate in the phospholipid bilayer and, hence, intracellular acid load depends on the amount of PA present in the extracellular medium and the amount able to enter and dissolve within the cell (Wu *et al.*, 2000).

The role of free intracellular calcium, $[\text{Ca}^{2+}]_i$, in the DHA-induced decrease in pH_i was also investigated. Hence, DHA induced increases in $[\text{Ca}^{2+}]_i$ which were significantly curtailed when experiments were conducted in 0% Ca^{2+} medium. It has been recently shown that DHA-induced increases in $[\text{Ca}^{2+}]_i$ are due to, in part, Ca^{2+} mobilization from reticulum endoplasmic stores, and the prolonged calcium response is contributed by calcium influx which takes place by opening of Ca^{2+} release-activated Ca^{2+} (CRAC) channels (Bonin & Khan, 2000). We observed that DHA-evoked intracellular acidification was modified when cells were resuspended in the 0% Ca^{2+} medium. Indeed, no significant difference in the amplitude of DHA-induced acidosis was seen between the two media of incubation (0 and 100% Ca^{2+} media); however, in the 0% Ca^{2+} buffer, the recovery of the resting pH_i was faster than that in 100% Ca^{2+} medium. Preincubation of BCECF-loaded cells with the calcium chelator BAPTA/AM induced a rapid recovery of the pH_i , as compared to that in 100% Ca^{2+} medium. Furthermore, addition of exogenous CaCl_2 to 0% Ca^{2+} medium diminished the DHA-induced rapid recovery. These observations suggest that calcium influx may also be implicated in the prolonged phase of DHA-induced acidification. Hence, we can hypothesize that DHA-induced increases in $[\text{Ca}^{2+}]_i$ could activate the $\text{Ca}^{2+}/\text{H}^+$ ATPase that drives Ca^{2+} out and H^+ in (Berthe *et al.*, 1991). Our hypothesis can be supported with the observations of Chien *et al.* (2001), who have demonstrated that in human peripheral T-cells, PMA-induced acidification occurs when intracellular concentrations of free calcium are increased, and this is due to Ca^{2+} efflux, followed by H^+ influx. Similarly, *N*-methyl-D-aspartate (NMDA)-induced Ca^{2+} -dependent intracellular acidosis is brought about by the activation of $\text{Ca}^{2+}/\text{H}^+$ ATPase in rat cerebellar granule cells (Wu *et al.*, 1999). To sum up, we can say that calcium is implicated in the maintenance of the DHA-evoked sustained decrease in pH_i . However, DHA-induced rapid acidification seems a calcium-independent phenomenon as in 0% medium, containing BAPTA/AM, DHA could not induce an increase in $[\text{Ca}^{2+}]_i$, though this fatty acid did rapidly acidify these cells under such conditions.

The role of DMA, a potent NHE1 inhibitor, in the DHA-induced acidosis was also investigated. Enhanced acidification was observed when DMA was applied before or after DHA and cells could not recover their diminished pH_i . Thus, NHE is implicated in proton extrusion and it is unlikely that DHA acidify by inhibiting the antiport. If such a process had occurred in our system, prior addition of DMA would have abolished the response of DHA. Surprisingly, preincubation of cells with DMA or subsequent addition of DMA after DHA, in 0% Ca^{2+} medium, could not delay the fast pH_i recovery observed after addition of DHA. In our study, the NHE1 isoform seems to be implicated, as Chambrey *et al.* (1997) have shown that NHE3 and NHE4 are amiloride-resistant isoforms, while NHE1 and NHE2 are the most sensitive isoforms to amiloride inhibition. The inhibition constants of DMA for these isoforms are as follows: NHE1 ($K_i = 0.023 \mu\text{M}$), NHE2 ($K_i = 0.25 \mu\text{M}$), NHE3 ($K_i = 14 \mu\text{M}$) (for a review, see Masereel *et al.*, 2003). Furthermore, NHE3 isoform is mainly expressed in high levels in the colon and small intestine, where it contributes to sodium absorption by the brush-border membrane in intestinal or renal epithelia (Masereel *et al.*, 2003). Furthermore, DMA was able to significantly inhibit NHE activity in the 100% Ca^{2+} medium. Thus, the fast pH_i recovery observed in the presence of DMA is not related to the existence of an amiloride-resistant NHE subtype in our cells.

The members of the NHE family display remarkable functionality. They are modulated by agents that primarily target tyrosine kinases, and also by agonists of Ser/Thr kinases like protein kinase A (PKA) and C (PKC). Bertrand *et al.* (1994) have demonstrated that calmodulin physically interacts with a particular subdomain of the NHE1 cytosolic region (Wakabayashi *et al.*, 1994), thus resulting in the activation of NHE1 by increases in intracellular calcium. This calmodulin-binding regulatory box is sufficient to account for the rapid and transient activation of NHE1 in response to growth factors and other Ca^{2+} -mobilizing agonists. An identical sequence is not found in NHE3 (Levine *et al.*, 1995). Indeed, Maly *et al.* (2002) have shown, by chelating the cytoplasmic calcium by BAPTA/EGTA, that cytoplasmic free calcium concentrations in epidermal growth factor receptor-6 (EGFR6) were critical for the regulation of the NHE. However, the results of the study of Chien *et al.* (2001) imply that in phytohemagglutinin (PHA)-stimulated human peripheral T-cells, the second messengers, for example, PKC and Ca^{2+} , may modulate pH_i in a dual-antagonistic mechanism, that is, PKC-mediated alkalization and Ca^{2+} influx-mediated acidification; hence, the latter observation corroborates our study. Bafilomycin A1, a potent and specific macrolide antibiotic that inhibits VPATPase (Crider *et al.*, 1994), was used in order to determine the contribution of VPATPase in proton extrusion. Bafilomycin A1 did not significantly enhance DHA-induced acidification, indicating that VPATPase does not participate to proton extrusion in T-cells. It has been shown that activation of the plasma membrane NADPH oxidase is associated with a rapid depolarization of the membrane potential that is followed by a slight fall in pH_i , and the generation of superoxide (O_2^-) (Henderson *et al.*, 1987). A study conducted on human resting neutrophils has shown that AA induced a pH_i fall as a result of the opening of the H^+ channel by this fatty acid. As described for H^+ conductance in snail neuron, the proton channel is inhibited by Cd^{2+} and Zn^{2+} ions (Henderson *et al.*, 1988). We,

therefore, used cadmium chloride (CdCl_2) and zinc chloride (ZnCl_2) to determine whether DHA activates the H^+ channel. Addition of these cations failed to diminish the DHA-induced intracellular acidification; thus, we can rule out the possibility that this fatty acid could activate the NADPH oxidase-associated H^+ channel. It also seemed unlikely that DHA could act by inhibiting the $\text{Na}^+ - \text{K}^+ - 2 \text{Cl}^-$ symport, mainly responsible for Na^+ and chloride cotransport, since prior addition of bumetanide, a specific inhibitor of this symport, did not affect DHA-induced acidification. Matsuyama *et al.* (2000) have recently shown that cytosol acidification is associated with mitochondrial-matrix alkalization, as a result of H^+ gradient dissipation across the inner mitochondrial membrane. We were tempted to elucidate if DHA-induced acidification was the result of mitochondrial- H^+ leakage. We, therefore, performed prior addition of oligomycin, a F_0F_1 -ATPase/ H^+ pump inhibitor, and carbonyl cyanide *m*-chlorophenyl-hydrazone (mCICCP), which dissipates H^+ gradient across the inner mitochondrial membrane. Again, we rule out the eventual contribution of mitochondrial H^+ in DHA-evoked response, since these two molecules failed to influence the decrease in pH_i induced by this fatty acid.

As we have seen in the previous paragraph that Ca^{2+} does play a role in the DHA-induced prolonged acidification, the mechanisms of action of DHA in transient ($t_{1/2} = 33 \text{ s}$) and immediate acidosis are not well understood. Hence, we would like to refer to the hypothesis of Kamp and Hamilton (Kamp & Hamilton, 1992; Kamp *et al.*, 1995), who have demonstrated that, in phospholipid bilayers, fatty acids can produce free intracellular H^+ by a simple diffusion. Indeed, the charged surface of the plasma membrane affects the ionization of fatty acids, thus, increasing their apparent pK (7.6) and the formation of nonionized fatty acids. As fatty acids have a high lipid solubility, all of them will bind to the lipid bilayer. At an extracellular pH of 7.4, fatty acids bind to the outer leaflet that consists of equal amounts of unionized fatty acids and ionized fatty acids. In all, 50% of the unionized fatty acid diffuses rapidly (within seconds) from the outer leaflet to the inner leaflet (flip) of the phospholipid bilayer, where half dissociates into an ionized form (FA^-), generating cytosolic H^+ (Kamp & Hamilton, 1992). The use of BSA has shown that such a mechanism is likely to occur in our system, as in the presence of albumin, the DHA-, but not the propionic-, induced acidification was totally reversed. The same has been observed for AA (Cavallini *et al.*, 1996; Wu *et al.*, 2000; Chen *et al.*, 2001). Hence, BSA extracts fatty acids from the inner leaflet and allows their transport in the reverse direction (flop). The notion that DHA could acidify the intracellular medium by deprotonation as a result of simple diffusion mechanism is further supported by the use of a DHA derivative in which the carboxylic end of the molecule was replaced by a methyl ester group. DHA methyl ester failed to induce intracellular acidification, and this in agreement with the work of Chen *et al.* (2001), who also observed that AA methyl ester had no effect on pH_i .

Long-chain fatty acids are important metabolic substrates for both energy production and lipid synthesis, as well as they participate in a variety of crucial cell-signalling cascades (Amri *et al.*, 1994; Tebbey *et al.*, 1994; Warnotte *et al.*, 1994). Civelek *et al.* (1996) have shown in adipocytes that exposure to lipolytic agents or external free fatty acids (FFA) results in a

rapid intracellular acidification that is reversed by metabolism of the FFA. As far as the metabolism/catabolism of DHA is concerned, we performed the analysis on the cells/assays during the time course of our study and we observed that this fatty acid was neither converted into docosapentaenoic acid or EPA nor acylated/esterified (results not shown). In fact, DHA is the end product of the metabolism of α -linolenic acid and it does not give rise to eicosanoids (Lands, 1991). Moreover, Jurkat T-cells do not possess the lipoxygenase and cyclooxygenase enzymes that account for lipid mediators (Kurland & Bockman, 1978; Goldyne *et al.*, 1984). Moreover, Kamp *et al.* (2003) have recently shown both in vesicles prepared from the plasma membrane of adipocytes and in the intact adipocytes that OA metabolism was very little during the time course of the experiment and it was not essential for the decreases in pH_i , observed after immediate addition of the fatty acid.

To determine whether the observed pH_i changes were unique to DHA, we tested other fatty acids of the n-9, n-6 and n-3 families, respectively, OA (18:1 n-9), AA (20:4 n-6) and EPA (20:5 n-3). All of these fatty acids induced a drop in pH_i , in the following order: $\text{DHA} > \text{EPA} > \text{AA} > \text{OA}$. The acidification induced by AA was lesser than that induced by EPA, though the difference was not significantly different. The dose-dependent experiments on AA, EPA and OA were performed for each fatty acid, in order to determine the concentration leading to an optimal response. Under our experimental conditions, all fatty acids exerted maximal decrease in pH_i at $10 \mu\text{M}$, thus underlying the specificity of the DHA-induced acidosis. Moreover, addition of palmitic acid (16:0) was without effect on pH_i , suggesting that fatty acid-mediated intracellular acidosis is not a so common feature. It seems to depend not only on the degree of fatty acid unsaturation, but also on the carbon chain length. However, the above observations could be limited by the differences in the water solubility that may exist between fatty acids, particularly between palmitic acid or OA and the others. Indeed, it is generally assumed that, at physiological pH ($\text{pH} = 7.0 - 7.4$), the fatty acids do not form soluble micelles and their equilibrium state is an insoluble aggregate, much like the phospholipid bilayer. In addition, water solubility increases with the degree of *cis*-unsaturation and carbon chain length of the fatty acids (Cistola *et al.*, 1986; Hicks & Gebicki, 1997). The solubility limit of fatty acids is difficult to measure (Richieri *et al.*, 1992; Vorum *et al.*, 1992). However, we noticed that palmitic acid, even at $20 \mu\text{M}$, was homogeneously distributed in the bicarbonate-free medium ($\text{pH} 7.4$); so, the observed effects of palmitic acid are not due to its aggregation. As also mentioned by Hamilton (1998), ethanolic solutions of fatty acids are generally used to achieve higher fatty acids concentrations for presentation to membranes. All fatty acids tested in the present study were dissolved in ethanol. We have not measured the amount of membrane-bound and unbound forms of each fatty acid, although one would expect, as indicated by the affinity constant, that almost all exogenously added free fatty acids adsorb quantitatively and rapidly to the plasma membrane phospholipid bilayer (Anel *et al.*, 1993). Gamberucci *et al.* (1997) have shown in Erlich tumour cells, treated with OA, that this fatty acid was taken up by cells at 1 min of incubation. In our system, at $10 \mu\text{M}$, fatty acids are likely to follow the same kinetics; thus, differences observed in the magnitude of fatty acid-induced pH_i fall might not be related to differences in the amount of fatty acid adsorption.

Though all of the unsaturated fatty acids induced a fall in pH_i, DHA was the most potent inducer of acidification and this difference, in part, may contribute to its mechanisms of action. Our study is certainly of physiological relevance as, under pathophysiological conditions, large amounts of free fatty acids can be released, as it is the case, during cardiac ischaemia, of AA whose concentrations are increased up to 50 µM (Nakamura *et al.*, 1989).

The present study has shown that treatment of human T cells with DHA, an n-3 PUFA, triggered a decrease in pH_i, which was due, in part, to simple diffusion of the fatty acid across the phospholipid bilayer and that the long-lasting acidification was

dependent on increases in [Ca²⁺]_i. As DHA has been shown to diminish T-cell proliferation (Bonin & Khan, 2000) and to induce apoptosis (Diep *et al.*, 2000), it would be of interest, in future, to determine whether DHA-induced long-term increase in intracellular H⁺ has a direct effect on immediate-early gene expression and apoptotic signalling pathways in human T cells, in normal or pathological conditions.

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