# Relationship between (Na + K)-ATPase activity, lipid peroxidation and fatty acid profile in erythrocytes of hypertensive and normotensive subjects

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**Abstract** Oxidative stress may play a role in the pathogenic mechanism of essential hypertension. Lipid peroxidation can alter the cellular structure of membranebound enzymes by changing the membrane phospholipids fatty acids composition. We investigated the relationship between (Na + K)-ATPase activity, lipid peroxidation, and erythrocyte fatty acid composition in essential hypertension. The study included 40 essential hypertensive and 49 healthy normotensive men (ages 35-60 years). Exclusion criteria were obesity, dyslipidemia, diabetes mellitus, smoking, and any current medication. Patients underwent 24-h ambulatory blood pressure monitoring and blood sampling. Lipid peroxidation was measured in the plasma and erythrocytes as 8-isoprostane or malondialdehyde (MDA), respectively. Antioxidant capacity was measured as ferric reducing ability of plasma (FRAP) in the plasma and as reduced/oxidized glutathione (GSH/GSSG ratio) in erythrocytes. (Na + K)-ATPase activity and fatty acids were determined in erythrocyte membranes. Hypertensives had higher levels of plasma 8-isoprostane, erythrocyte MDA, and relative percentage of saturated membrane fatty acids, but lower plasma FRAP levels, erythrocyte GSH/GSSG ratio, (Na + K)-ATPase activity and relative percentage of unsaturated membrane fatty acids, compared with normotensives. Day-time systolic and diastolic blood pressures correlated positively with lipid peroxidation parameters, but negatively with (Na + K)-ATPase activity. These findings suggest that the modulation of (Na + K)-ATPase activity may be associated with changes in the fatty acid composition induced by oxidative stress and provide evidence of a role for this enzyme in the pathophysiology of essential hypertension.

**Keywords** Lipid peroxidation  $\cdot$  Essential hypertension  $\cdot$  (Na + K)-ATPase  $\cdot$  Fatty acids

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# Introduction

Reactive oxygen species (ROS) have been shown to be involved in the pathogenesis of hypertension [1, 2]. Hypertension causes 7.1 million deaths annually [3] and is a major risk factor for cardiovascular disease mortality [4]. An imbalance between the ROS generation and antioxidant defense systems in the body resulting in oxidative stress has been reported in spontaneously hypertensive rats [5] and human essential hypertension [6]. ROS production could damage arterial walls including an impairment of the endothelium-dependent vasodilation or endothelial dysfunction [7].

The interaction of ROS with biological membranes produces a variety of functional modifications due to either direct interaction with the molecular cell machinery and/or oxidative modification of the environment of biological



macromolecules [8]. Lipid peroxidation contributes to the loss of cellular functions through the inactivation of membrane enzymes and cytoplasmic proteins. In erythrocytes, increased lipid peroxidation generates changes in lipid composition, thus, altering the phospholipid fatty acid profile and membrane fluidity [9]. The validity of erythrocyte fatty acid composition as a reliable biomarker of that in other organs has been supported by recent studies showing a significant correlation of erythrocyte polyunsaturated fatty acids (PUFAs) composition with that found in the liver [10] and muscles [11].

Lipid composition of the erythrocytes has been implicated in the modulation of membrane-bound enzymes, such as (Na + K)-ATPase, Ca<sup>2+</sup>-ATPase, Mg<sup>2+</sup>-ATPase, and acetylcholinesterase [12]. (Na + K)-ATPase activity depends on its close interaction with phospholipids rich in PUFAs. Since PUFAs are the major targets of ROS renders the membrane phospholipids particularly sensitive to metabolic conditions associated with oxidative stress. Consequently, an increase in PUFAs enhances the vulnerability of membranes to undergo modifications in fatty acid composition [13], thus, impairing the (Na + K)-ATPase micro-environment needed for optimal enzyme activity [14]. (Na + K)-ATPase inhibition triggers Ca<sup>2+</sup> entry and increases the myogenic tone and contractility in arterial smooth muscle cells [15]. This results in an increase in peripheral vascular resistance, the hemodynamic hallmark of hypertension [16]. The aim of the present study was to test the hypothesis that increased lipid peroxidation leads to changes in cell membrane fatty acid composition that are functionally related to blood pressure elevation through an impairment of (Na + K)-ATPase activity.

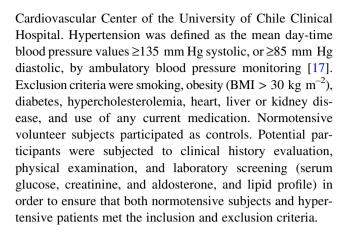
# Materials and methods

# Study design

A cross-sectional design applied to 40 hypertensive and 49 healthy normotensive subjects. The study protocol was approved by the local Ethics Committee of the University of Chile Clinical Hospital, and performed in accordance with the Helsinki Declaration II. All participants signed written consent and no complications were encountered during the study.

# **Patients**

We recruited essential hypertensive patients (stage 1, Seventh Joint National Committee) [3] who were not under pharmacological treatment. Inclusion criteria were male sex and age between 35 and 60 years. Potential participants were recruited from outpatients consulting at the



# Ambulatory blood pressure monitoring

Blood pressure levels were determined through ambulatory blood pressure monitoring on a regular workday (during 24 h from 8:30 AM) using an oscillometric monitor (SpaceLabs 90207, SpaceLabs Inc., Redmond, WA), previously checked for accuracy against simultaneous measurements by mercury sphygmomanometer. The mean day-time value of blood pressure was registered. This device fulfills the validation criteria's of the British Hypertension Society protocol [18] and the Association for the Advancement of Medical Instrumentation (AAMI) for studies in ambulatory condition [19]. The oscillometric accuracy, assessed by Spacelabs-intra-arterial average differences, was  $-0.6 \pm 5.9$  and  $0.9 \pm 6.4$  mm Hg (means  $\pm$ standard deviation), for systolic and diastolic pressures, respectively, which are within the AAMI accuracy standard. The estimated oscillometric reproducibility was  $-0.3 \pm 3.2$ and  $0.1 \pm 3.5$  mm Hg (means  $\pm$  standard deviation), for systolic and diastolic pressures, respectively [20, 21].

# Blood samples

Venous blood samples (<10 ml) were collected in chilled vacutainers containing disodium EDTA (1.5 mg ml<sup>-1</sup>). Plasma and red blood cell lysates, were stored at  $-70^{\circ}$ C. Samples for 8-isoprostane measurement were collected in plastic tubes previously treated with antioxidant butylated hydroxytoluene (final concentration 1 mmol l<sup>-1</sup>). Erythrocyte membranes were isolated by ultra-centrifugation (Sorvall Inc. Newton, Connecticut, USA) at 100,000~g for 45 min and stored in eppendorf tubes containing histidinesucrose buffer (pH 6.8) at  $-80^{\circ}$ C until used for (Na + K)-ATPase and Mg<sup>2+</sup>-ATPase activity measurements.

Ferric reducing ability of plasma (FRAP)

Plasma antioxidant status was assessed by measuring its ability to reduce ferric to ferrous iron (FRAP) with a detection limit  $10 \mu mol l^{-1}$  [22].



### Glutathione

Reduced glutathione (GSH) and glutathione disulfide (GSSG) were assayed by fluorometry [23]. The GSH/GSSG ratio was also determined, as a parameter of intracellular redox status.

# Malondialdehyde (MDA)

Lipid peroxides were assayed spectrophotometrically at 532 nm by the thiobarbituric acid reaction at pH 3.5, followed by solvent extraction with a mixture of n-butanol/pyridine (15/1, v/v) [24]. Tetramethoxy-propane was used as the external standard and the level of lipid peroxides was expressed as nmol malondialdehyde (MDA)/g Hb.

# 8-Isoprostane

8-Isoprostane concentration in plasma, recognized as a reliable biomarker of lipid peroxidation in vivo [25], was determined by using ELISA kit (Cayman, Ann Arbor, MI) and results expressed as pg/ml.

# ATPases

Activities of (Na + K)-ATPase and Mg<sup>+2</sup>-ATPase were measured by the method of Katz and Epstein [26]. The assay mixture consisted in (mM) of 100 NaCl, 20 KCl, 6 ATP (vanadium free), 6 MgCl<sub>2</sub> and 10 imidazole buffer (pH 7.8), was pre-incubated at 37°C for 1 min, and the reaction was started by the addition of ATP. After 15 min reaction was stopped by the addition of 1 ml ice-cold 25% (w/v) trichloroacetic acid. Inorganic phosphate was measured in the supernatant by the method of Taussky and Shorr [27]. (Na + K)-ATPase activity was calculated from the difference between the amount of inorganic phosphate released in the presence and in the absence of K+ in the incubation medium. Mg<sup>+2</sup>-ATPase activity was calculated from the difference between the amounts of inorganic phosphate released in the absence of K+ at zero time. Protein concentration was measured by the method of Lowry et al. [28]. The specific activities were expressed as µmol inorganic phosphate released per milligram protein per hour.

Extraction and separation of phospholipids from erythrocyte membranes

Blood samples collected in syringes containing 5% (w/v) EDTA as anticoagulant were centrifuged at 2,500 rpm for 15 min at 4°C to separate erythrocytes. Erythrocyte membranes were isolated according to Huertas et al. [29]. Membrane lipids were extracted as described by Bligh and Dyer [30].

Preparation and analysis of fatty acid methyl esters (FAME)

Fatty acids from erythrocyte membranes were methylated. The phospholipids were eluted from silica gel with two 15 ml portions of chloroform/methanol/water (10:10:1, by vol.). Solvent was evaporated in a stream of nitrogen, and 10 mg tricosaenoic acid (23:0, internal standard) was added prior to the esterification with 0.2 N sodium-methanol for 30 min at 40°C, and then, with H<sub>2</sub>SO<sub>4</sub>– methanol as described for alkaline methylation [31]. Samples were cooled and the fatty acid methyl esters (FAME) were extracted with 0.5 ml hexane and analyzed by gas–liquid chromatography.

A Hewlett–Packard gas chromatograph (model 6890)(Palo Alto, CA, USA) equipped with a capillary column (50 m  $\times$  0.22 mm BPX70; 0.25U QC 0.08 SGE) was employed to separate FAME. The temperature was programmed from 180°C to 230°C at 2°C min<sup>-1</sup> with a final hold, separating 12:0 to 22:6n–3. Both detector and injector temperatures were set at 240°C. Hydrogen was used as carrier gas, at a flow rate of 1.5 ml min<sup>-1</sup> and split ratio of 1:80. FAME was identified by comparing their retention times with individual purified standards and quantified using a Hewlett–Packard integrator (HP 3396 Series III).

# Materials

All reagents were purchased from Sigma-Aldrich (St. Louis, MO, USA), Merck (Darmstadt, Germany) and Riedel-de Häen (Germany), and were of the highest commercial grade available.

# Statistical analysis

Descriptive statistics of variables used the means and standard error of the means (SEM). The source of variation between normotensive and hypertensive subjects was assessed by unpaired Students t-test, for normally distributed parameters, with a P value < 0.05 for statistical significance. The association of variables was studied by Pearson correlation test due to their Gaussian distribution.

# Results

# Clinical characteristics

Clinical characteristics of the 89 subjects in the hypertensive and normotensive groups are shown in Table 1. All parameters were within the normal range and were not different between the two groups except for significant



**Table 1** Clinical characteristics of essential hypertensive patients (n = 40) and healthy normotensive subjects (n = 49)

Characteristic	Normotensive $(n = 49)$	Hypertensive $(n = 40)$	P Value
Age (year)	43.4 ± 1.1	44.6 ± 1.4	0.49
Body mass index (kg/m²)	$24.8 \pm 1.2$	$25.9 \pm 1.3$	0.53
Blood glucose (mmol/l)	$4.93 \pm 0.07$	$5.09 \pm 0.07$	0.11
Creatinine (µmol/l)	$79.6 \pm 15.0$	$81.4 \pm 9.7$	0.93
Total cholesterol (mmol/l)	$4.59 \pm 0.16$	$4.81 \pm 0.14$	0.31
High-density lipoprotein (mmol/l)	$1.26 \pm 0.04$	$1.18 \pm 0.05$	0.18
Low-density lipoprotein (mmol/l)	$2.71 \pm 0.07$	$2.86 \pm 0.09$	0.21
Triglycerides (mmol/l)	$1.34 \pm 0.06$	$1.49 \pm 0.08$	0.12
Awake systolic BP (mm Hg)	$118.5 \pm 1.3$	$138.1 \pm 2.1$	<0.001*
Awake diastolic BP (mm Hg)	76.6 ± 1.7	92.1 ± 1.4	<0.001*
Heart rate (beats/min)	$72.8 \pm 1.2$	$74.1 \pm 1.3$	0.47

Data are expressed as means  $\pm$  SEM. BP, blood pressure

higher mean day-time systolic (SBP) and diastolic blood pressure (DBP) in the hypertensive group (P < 0.001),

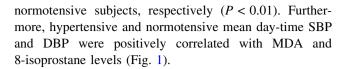
Antioxidant status and oxidative stress-related parameters

The antioxidant status measured in plasma (FRAP) and erythrocyte (GSH/GSSG ratio) as well as lipid peroxidation products (plasma 8-isoprostane and erythrocyte MDA) are shown in Table 2. FRAP levels and GSH/GSSG ratios were 27% and 31% lower (P < 0.001) in hypertensive compared to normotensives subjects, respectively. In addition, erythrocyte MDA and plasma 8-isoprostane levels were 23% and 43% higher in the hypertensives than

Table 2 Plasma and erythrocyte antioxidant status and lipid peroxidation-related parameters of the study participants

Parameter	Normotensive subjects $(n = 49)$	Hypertensive patients $(n = 40)$	P value
Plasma			_
FRAP (µmol/l)	$419.6 \pm 9.7$	$307.1 \pm 11.2$	<0.001*
8-isoprostane (pmol/l)	$78.7 \pm 2.8$	$112.7 \pm 3.2$	<0.001*
Erythrocytes			
GSH/GSSG ratio	$7.46 \pm 0.08$	$5.17 \pm 0.17$	<0.001*
Malondialdehyde (nmol/g Hb)	$301.9 \pm 1.1$	$372.2 \pm 1.4$	<0.02*

Data are expressed as means ± SEM. FRAP, ferric reducing ability of plasma; GSH, reduced glutathione; GSSG, oxidized glutathione. \* Significant difference by unpaired Student's *t*-test



Activity of ATPases, lipid peroxidation and systolic blood pressure

Activities of (Na + K)-ATPase and  ${\rm Mg}^{+2}$ -ATPase in the erythrocytes of both groups are shown in Fig. 2A. Erythrocyte (Na + K)-ATPase activity was 14% lower in the hypertensives compared to normotensives (P < 0.03). In contrast, no significant differences were found in the  ${\rm Mg}^{+2}$ -ATPase activity (P = 0.25). The correlation between erythrocyte MDA and plasma 8-isoprostane levels with (Na + K)-ATPase activity is shown in Fig. 2B and C, respectively. Erythrocyte (Na + K)-ATPase activity correlated negatively with both MDA and 8-isoprostane in hypertensive and normotensive subjects. In addition, mean day-time SBP and DBP showed a negative correlation with (Na + K)-ATPase activity (Fig. 3) but not with  ${\rm Mg}^{+2}$ -ATPase in both groups.

Erythrocytes fatty acid composition

The fatty acid composition of erythrocytes phospholipids is shown in Fig. 4. The 16:00, 18:0 and 24:0 fatty acid levels were higher and 14:1, 16:1, 20:1, 22:1, 20:4n–6, 22:2n–9, and 22:6n–3 levels were lower (P < 0.05) in hypertensive compared to normotensives. Whereas 14:00, 20:0, 22:0, 18:1, 24:1, 18:2n–6, 18:3n–6, 20:5n–3 showed no significant differences between the two groups. The relative percentage of saturated fatty acids (SAFAs) were 86% higher but that of monounsaturated fatty acids (MUFAs) and PUFAs were 29% and 21% lower in hypertensives than the normotensives, respectively (P < 0.001)(Fig. 4A).

# Discussion

The findings of the present study confirm previous data reporting association of blood pressure with both antioxidant status and oxidative stress-related parameters [2]. Furthermore, our data provides a new viewpoint related to fatty acid composition and (Na + K)-ATPase activity in patients with essential hypertension. This suggests an involvement of membrane lipid profile in the modulation of (Na + K)-ATPase activity and its role in the pathogenesis of blood pressure elevation.

The lower erythrocyte GSH/GSSG ratio and plasma FRAP levels (parameters indicative of the antioxidant defenses) in hypertensives, is consistent with the higher lipid peroxidation. Antioxidants normally contribute to the



<sup>\*</sup> Significant difference by unpaired Student's t-test

Fig. 1 Pearson correlation of plasma and erythrocyte lipid peroxidation parameters to mean day-time systolic ( $\mathbf{A}$ ,  $\mathbf{C}$ ) and diastolic ( $\mathbf{B}$ ,  $\mathbf{D}$ ) blood pressure of normotensive (n = 49) (open circles and dotted line) and hypertensive (n = 40) (solid circles and line) participants, respectively. Hemoglobin (Hb)

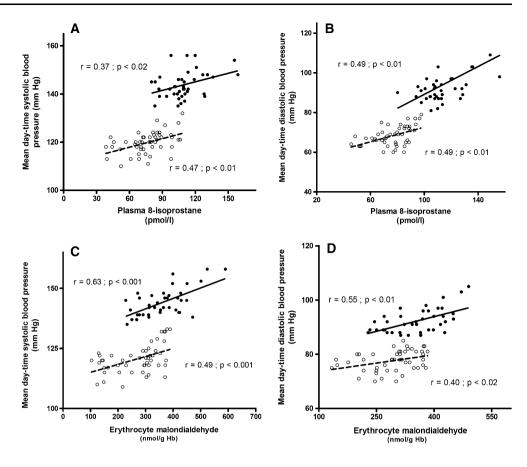
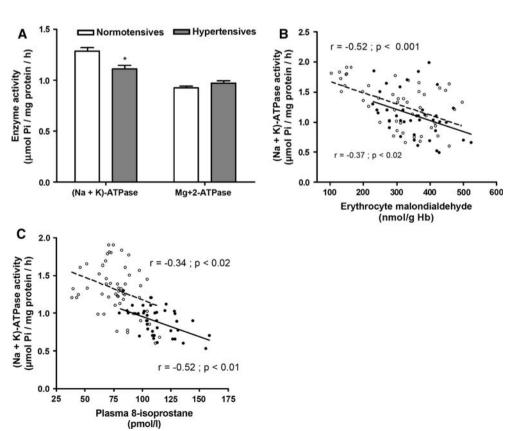
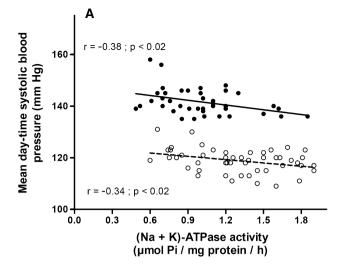
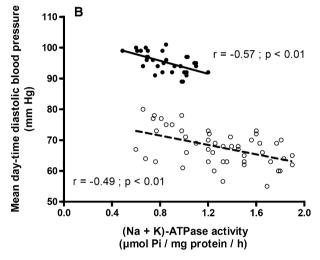


Fig. 2 Erythrocyte ATPase activity (mean  $\pm$  SEM) in both study groups. \* P < 0.03 vs. normotensives (A). Pearson correlation of (Na + K)-ATPase activity to malondialdehyde (B), 8-isoprostane (C) of normotensives (n = 49) (open circles and dotted line) and hypertensives (n = 40) (solid circles and line) participants









**Fig. 3** Pearson correlation of (Na + K)-ATPase activity to mean day-time systolic (**A**) and diastolic (**B**) blood pressure of normotensive (n = 49) (open circles and dotted line) and hypertensive (n = 40) (solid circles and line) participants, respectively

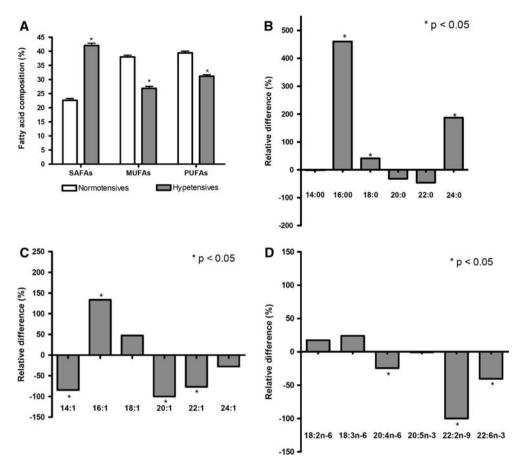
prevention of peroxidation of polyunsaturated fatty acids. Erythrocytes MDA and plasma 8-isoprostane levels were elevated in hypertensives (Table 2) and show a positive correlation with both mean day-time SBP and DBP (Fig. 1).

ROS are increasingly implicated in human hypertension, since they exert important effects in vascular biology [32]. This suggests that ROS are contributors in mediating vasoconstriction occurring in hypertensive patients [1, 32–34]. However, to the best of our knowledge, differences in fatty acid pattern and its functional relationship with oxidative stress-related parameters in essential hypertension has not been reported. The present findings of increased lipid peroxidation in hypertensives are consistent with specific changes in membrane fatty acid profile, showing that the relative proportion of SAFAs was increased and

those of MUFAs and PUFAs were decreased (Fig. 4A), this could cause structural changes in plasma membrane [35]. The physicochemical properties of plasma membranes are partly determined by the degree of fatty acid unsaturation [36]. The incorporation of PUFAs into membrane phospholipids has been suggested to damage cellular membranes by virtue of the susceptibility of highly unsaturated acyl fatty acid to peroxidation [37, 38]. Our findings would support the idea that the unsaturation grade of the membrane lipids is related to lipid peroxidation susceptibility, although the method used for measurement of fatty acids gives only a relative proportion of saturated, mono- and polyunsaturated fatty acids. All of the above mentioned membrane effects could contribute to modulate the functional activity of membrane-bound enzymes. Lipid peroxidation, contributes to the loss of some cellular function through the inactivation of membrane-bound enzymes and native ion channels [8]. Accordingly, previous studies have shown that oxidative stress causes an inhibition of Ca<sup>+2</sup>. Mg<sup>+2</sup>-ATP-ase in the erythrocytes [39] and sodium channel dysfunction in cardiomyocytes [40]. In agreement with this view, our data demonstrate that the impairment of erythrocyte (Na + K)-ATPase activity associated positively with lipid peroxidation (Figs. 2B and C). Impairment of (Na + K)-ATPase activity could be due to the loss of its optimal interaction with the membrane components, as a consequence of increased lipid peroxidation. Although a direct inhibition by peroxynitrite, as occurs in liver plasma membranes, should not be discarded [41]. In contrast, the lack of significant differences in the activity of erythrocyte Mg<sup>+2</sup>-ATP-ase between hypertensives and normotensives may indicate a lesser functional dependence of this enzyme activity, respect to (Na + K)-ATPase, upon the interaction with the membrane ROS targets. Therefore, it could be suggested that Mg<sup>+2</sup>-ATP-ase is causally unrelated to oxidative stress-mediated hypertension. This view is supported by the fact that Mg+2-ATP-ase in spontaneously hypertensive rats showed no relationship with blood pressure [42]. In contrast, (Na + K)-ATPase activity is expected to be influenced by the fatty acid unsaturation [13], in oxidative stress settings due to the vulnerability of PUFAs to ROS attack [43, 44]. Therefore, it is likely that the involvement of fatty acid composition in the modulation of membrane-bound (Na + K)-ATPase enzyme, thus, explaining the possible functional effect derived from alterations in fatty acid pattern. The above mentioned considerations could give an explanation to the finding of lower levels of (Na + K)-ATPase activity in hypertensive than normotensive participants (Fig 2A). Attempts to analyze the contribution of n-6/n-3 PUFA ratio in the modulation of both (Na + K)-ATPase activity and blood pressure were not carried out in the present study. Although it has been reported that n-6/n-3 PUFA ratio may be



Fig. 4 Plasma membrane fatty acid composition of erythrocytes (means  $\pm$  SEM) in both study groups. \* P < 0.001vs. normotensives, (A). Relative differences (%) in the proportions of SAFAs (B). MUFAs (C) PUFAs (D) of hypertensive (n = 40) compared to normotensive (n = 49)participants. The positive or negative values indicate higher or lower proportions, respectively. SAFAs, saturated fatty acids; MUFAs monounsaturated fatty acids; PUFAs poly-unsaturated fatty acids



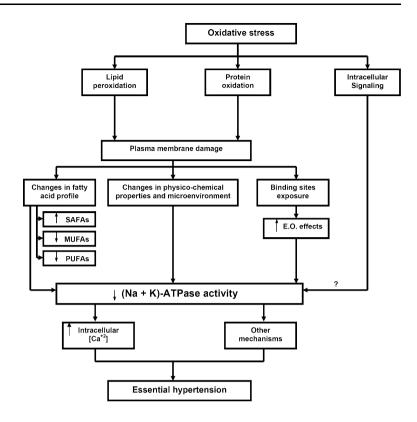
valuable in interpreting the biomarker data on fatty acid profile associated with prevention of coronary heart disease [45]. More recent studies based on large series of trials, concluded that focusing on the ratio distracts from the more important issue of the abundance of the *n*–3 PUFAs to be the single most important dietary change that can be made to improve cardiovascular health [46, 47].

The ubiquity of (Na + K)-ATPase in mammalian cells and its function in the vascular wall has suggested its role as a contributory factor in the pathogenesis of hypertension. Since the erythrocyte fatty acid profile changes reported in this study are expected to occur in other cell types also, it could be pointed out that the elevation of local Na + on the submembrane area due to the diminution of (Na + K)-ATPase activity, in arterial smooth muscle cells. could facilitate Ca<sup>2+</sup> entry through Na +/Ca<sup>2+</sup> exchanger type 1 (NCX1). Consequently, a rise in cytosolic Ca<sup>2+</sup> concentration would contribute in the elevation of blood pressure [48] Nevertheless, other effects likely to influence the modulation of (Na + K)-ATPase, due to increased lipid peroxidation and/or protein oxidation, should not be discarded. Thus, it could be speculated that the above mentioned structural membrane alterations may result in an increased exposure of the endogenous cardiac glycoside binding site of the (Na + K)-ATPase. The cardiac glycoside or endogenous ouabain (E.O.) is a ligand that behaves as a natural regulator of (Na + K)-ATPase in vivo [49, 50] and thus it may play a role in the mechanism of hypertension [51, 52]. The specific binding site of E.O. has a strong evolutionary conservation among all species [53]. This ligand has the only known receptor in (Na + K)-ATPase [54], whose  $\alpha_2$  isoform can mediate in the development of hypertension either through intracellular ion exchange or signaling cascades [55]. Functionally, E.O. can exert cardiotonic and vasotonic actions dealing to the development of human hypertension [56]. Consequently, it seems reasonable to assume that structural changes induced by lipid peroxidation in erythrocyte membrane, as well as in other cell types, could result in increased exposure of E.O. binding sites, thus, contributing to the elevation of blood pressure by the alternative above mentioned mechanism (Fig 3). The lack of these specific E.O. binding sites in the Mg<sup>+2</sup>-ATPase molecule prevents E.O.-enzyme interaction, thereby accounting for the differential response of erythrocyte (Na + K)-ATPase and Mg<sup>+2</sup>-ATPase activities in causing essential hypertension. A schema to explain this hypothesis is shown in Fig. 5.

In summary, these findings suggest that the modulation of (Na + K)-ATPase activity may be associated with changes in the fatty acid composition induced by oxidative



Fig. 5 Schema of a proposed hypothesis to explain the effects of oxidative stress on the cellular plasma membrane fatty acid composition and membrane-bound elements to account for the elevation of blood pressure through the modulation of (Na + K)-ATPase activity in essential hypertension. E.O., endogenous ouabain



stress or with oxidative stress directly, or both, and provide evidence of a role for this enzyme in the pathophysiology of essential hypertension.

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