

REVIEW

Involvement of Membrane Fluidity in Endogenous Protective Processes Running on Subcellular Membrane Systems of the Rat Heart

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Summary

Membrane fluidity is a widely recognized biophysical variable that provides information about structural organization of the subcellular membranes exhibiting physical characteristics of liquid crystals. The term "fluidity" reflects in this case the tightness in packing of acyl parts of the membrane phospholipid molecules, a feature that may influence considerably the molecular mobility and *via* that also the sensitivity and reactivity of membrane-bound transporters, receptors and enzyme systems. Data presented in this review are aimed to demonstrate the substantial role of changes in membrane fluidity occurring in the processes associated with endogenous protection observed in cardiac sarcolemma and mitochondria in diverse pathologies, particularly in diabetes and hypertension.

Key words

Rat heart sarcolemma • Mitochondria • Membrane fluidity • Diabetes • Hypertension • Endogenous protection

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Introduction

Responsible body of the membranes consist of an inversely oriented bi-layer of parallel arranged

phospholipid molecules with their non-flexible polar heads facing the aqueous environment on either side of the membrane. Because of their structural organization the subcellular membranes behave like a liquid crystal. Receptors, various transporters and systems involved in the transfer of signals are embedded in the membranes – some of them even span the lipid bilayer completely. However, proper function of these systems requires adequate mobility of their molecules. This mobility is secured by flexibility of acyl chains of the membrane phospholipids. The less tightly are packed the acyl chains of phospholipids the more fluid is the lipid bilayer of membranes and the more free may move the molecules of embedded systems. This makes membrane fluidity a potent modulator of membrane functions.

Role of membrane fluidity in function of subcellular membranes and metabolism

It has been accumulated much evidence that diverse impulses targeting the membranes may increase or decrease MF. Chemical compounds, including pharmaceuticals, may immerse into the membrane, depending on hydrophobicity of their molecules and, modulate the MF. Finally, spontaneous or evoked changes in the chemical composition of membranes may also alter the MF. High sensitivity of MF to various influences as well as the differences in specific properties of different subcellular membranes represent the reasons

why the physiological range of changes in MF was not yet demarcated sufficiently.

In the past 30 years a considerable amount of information concerning the involvement of different membrane systems such as sarcolemma, sarcoplasmic reticulum, and mitochondria in heart function and metabolism has appeared in the literature. It became absolutely clear that these membrane structures not only represent morphological barriers between various myocardial cell compartments but they are also intimately associated with a wide variety of molecular, biochemical and biophysical events occurring both in the cell compartments and in the membranes itself (Ziegelhoffer *et al.* 2009).

Control over the ionic fluxes as well as the neurohormonal and calcium-mediated cell signaling also belongs to unique properties of the cardiac sarcolemmal membranes (Dhalla *et al.* 1977, Pierce and Dhalla 1985).

Mitochondria are most frequently discussed because of their association with respiration and oxidative phosphorylation. Less frequently is mentioned that they also provide the control over the above processes (Sordahl and Schwartz 1967, Schwartz 1971/72). More recently it was shown that mitochondria may also exert control over formation of the ephemerical energy transition pores, also termed as mitochondrial contact sites, in their own membranes (Knoll and Brdiczka 1993, Bakker *et al.* 1994, Ziegelhoffer-Mihalovičová *et al.* 1997, 2002, Ziegelhoffer *et al.* 2009).

Role of membrane fluidity in physiological and pathophysiological processes

It was confirmed that regardless of the type of membrane, the MF may have effective control over many physiological processes (Shinitzky 1984, Gennis 1989). Since MF is reflecting the interplay between the membrane lipids and proteins it represents a variable with virtually unlimited range of influence on the milieu in which the membrane-bound systems are acting. Because of its basic dependence on a specific composition of lipids, changes in MF may also be induced by alterations in the diet or metabolism (Waczulíková *et al.* 2010). Perturbations that increase the area per lipid molecule, such as higher degree of phospholipid (PL) unsaturation, lower average length of the acyl chains, particular shifts in the lipid profile, decreased level of cholesterol etc., all result in an increase in MF (Gennis 1989, Watala 1993). Free radicals and diverse chemical compounds, mainly pharmaceuticals, may interact with membranes and

modulate MF (Taam *et al.* 1979). In addition, physiological and pathological impulses, including seasonal changes, which target the membranes, may also change MF. According to our findings, elevated MF of rat heart mitochondria observed in the winter/spring season was accompanied with a considerable increase in the mitochondrial Mg²⁺-ATPase activity (Mujkošová *et al.* 2008). On the other hand, lowered MF of the diabetic rat heart sarcolemma was found to be associated with a decrease in Ca²⁺ influx (Götzsche 1991, Ziegelhoffer *et al.* 1996, 2003b, Ziegelhoffer-Mihalovičová *et al.* 2003) as well as in the Na,K-ATPase activity (Ziegelhoffer *et al.* 1997, 2003a, Ziegelhoffer-Mihalovičová *et al.* 2003). Finally, estimation of MF parallelly with one or more purposefully chosen dominant membrane functions, enabled to reveal that some diabetes-induced alterations in the heart sarcolemma and mitochondria, previously thought to be pathologic, belong, in reality, to the mechanisms of endogenous protection (Ziegelhoffer-Mihalovičová *et al.* 2003, Ziegelhoffer 2005, Ferko *et al.* 2006).

Aim of the present paper is to document the regulatory role of MF for the function of membrane-bound systems in the heart sarcolemma and mitochondria in both health and disease. It provides a selection of findings obtained with biochemical and physiological techniques completed with spectroscopical estimation of MF. Presented data extend the previous knowledge and/or revise some older conceptions about the role of MF in various physiological and pathological situations.

Methods for studying subcellular membranes

Membrane dynamics covers virtually all modes of molecular motion in the membrane, among which CH₂-bond vibrations, lipid rotation (about the long axis), lateral “hopping” of lipids and proteins and rotation of protein are discussed most frequently (Gennis 1989). In selection of adequate technique for measurement of the above properties the rate of their molecular motion should be taken into account. If the components move slowly with respect to the speed of measurement, the technique yields a “static”, so called frozen picture of the membrane event. When the molecules move rapidly, then a time-resolved fluorescence spectroscopy provides the appropriate tool to track fine features of the movement. Quick, cost-effective and therefore more frequently used is the steady-state fluorescence anisotropy method. It

offers a time-averaged picture of movements of sensed components, since the speed of membrane motions is considerably higher than the time used for their detection.

In studying cardiac cell and membrane functions in both health and pathological conditions we utilized the following methods:

i) Fluorescence spectroscopy and fluorescence or confocal laser scanning microscopy, with employing exogenous fluorescent probes (Fig. 1). These techniques have proved useful in the investigation of membrane fluidity, protein-lipid interactions, membrane permeability and transport across the membranes (Ziegelhöffer *et al.* 1997, 2000, Šikurová *et al.* 1997, 2000, Máleková *et al.* 2007, Waczulíková *et al.* 2008) as well as selected electrical properties of the membranes (Waczulíková *et al.* 2000, 2007, Zvarík and Šikurová 2008). For the accurate cell edge extraction in cell-volume determination from microscopic images, a parametric active contour algorithm by Kass *et al.* (1988) was used.

ii) Voltamperometry, to investigate some membrane-related functions of cardiac myocytes and their mitochondria such as respiratory chain function and oxidative phosphorylation rate (Ziegelhöffer 2005, Ferko *et al.* 2006, Uličná *et al.* 2008);

iii) Physiological methods such as isolated perfused “Langendorff” heart, either for testing the effect of changes in subcellular membranes on functional parameters of the heart (Ravingerová *et al.* 1997, 2000, Dobaczewski *et al.* 2006) or to isolate cardiomyocytes (Cagalinec *et al.* 2006, Waczulíková *et al.* 2007);

iv) Biochemical, particularly enzymological methods (Ziegelhöffer *et al.* 2003a, Breier *et al.* 1995, Ferko *et al.* 2008), techniques of cryoelectron microscopy (Waczulíková *et al.* 2007) and methods of biostatistics (Waczulíková *et al.* 2002, 2007, Waczulíková and Slezák 2009). All these methods are utilized in characterization of alterations in the activity of membrane-bound enzymes and ion transport systems associated with changes in MF.

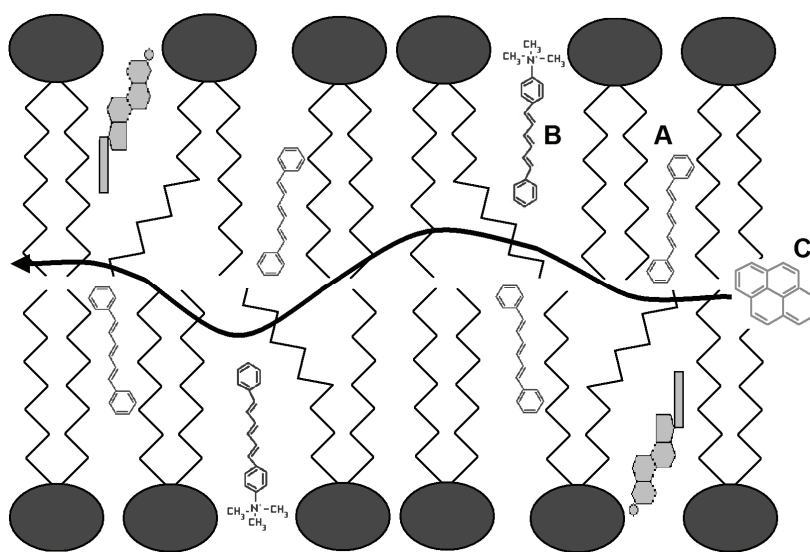


Fig. 1. Location of three of the most widely used fluorescent probes in membrane lipid bilayers, DPH (A), TMA-DPH (B) and pyrene (C). Adapted from Waczulíková *et al. Physiol Res* **59** (Suppl 1): S9-S17, 2010.

Measuring of membrane fluidity

MF is measured by recording a fluorescence signal which reflects the motion of membrane-permeable fluorescent probes, often with preferred orientations, in the respective parts of the membrane bilayer (Fig. 1). The probe should be chosen so that its emission maximum is out of range of the background signal, i.e. out of sample autofluorescence region. This indicates that interpretation of the obtained experimental data depends on both, the method and the probe applied.

Most measurements are made in steady-state

condition with continuous excitation and emission. Principles of the measurement may be briefly described as follows: Polarization P or anisotropy r of the light emitted by the fluorescent probe is dependent on the orientation of its molecular transition dipole moment with respect to the direction defined by the polarizers (parallel and perpendicular) used to record the fluorescence components. Both components of intensities emitted by the fluorescent probe, I_{\parallel} and I_{\perp} , are registered and their values are used to calculate the steady-state anisotropy value r_s :

$$r_s = \frac{I_{\parallel} - I_{\perp}}{I_{\parallel} + 2I_{\perp}} \quad (1)$$

Results represent an average value r_s , which is, however, related to both, the lifetime of the excited state and the rotational correlation time of the probe. This indicates that steady-state anisotropy measurements are sensitive to both the rate of motion and any constraints to that motion. Hence, the information about the dynamics and molecular order of the motion became intermixed. Nevertheless, the information on structural ordering of lipid molecules prevails (Shinitzky 1984, Gennis 1989). Details regarding the working protocols for measuring MF with the fluorescent probes are presented in our previous papers (Waczulíková *et al.* 2007, Waczulíková *et al.* 2008, Ziegelhöffner *et al.* 2009).

To evaluate MF in our experiments we used a parameter r_s (1) of a fluorescent probe 1,6-diphenyl-1,3,5-hexatriene (DPH) or its cationic derivative 1-[4(trimethylamino)phenyl]-6-phenyl-1,3,5-hexatriene (TMA-DPH). DPH partitions favourably into membranes, it has an intense fluorescence, it is assumed not to bind to proteins, and it is sensitive to the physical state of the membrane. Aqueous solution of DPH alone has a negligible fluorescence (Shinitzky 1984, Gennis 1989). Therefore, the fluorescence signal from labelled membranes could be solely ascribed to DPH molecules in the membrane. DPH incorporates into the hydrocarbon inside of membranes, whereas its cationic derivative TMA-DPH is located closer to the membrane surface (Fig. 1). Thus, the recorded values of fluorescence anisotropy report about MF in both parts the membrane lipid bilayer.

Each newly designed experiment requires a series of measurements to investigate the homogeneity of the sample, because in colloidal solutions not only particle size, but also their concentration may induce scattering. This may be prevented by dilution of the sample to a level at which no significant light scattering can be observed. The time course of DPH (or TMA-DPH) incorporation is derived from consecutive measurements covering a time-range of at least 1200 s. High degree of fluorescence anisotropy indicates a high degree of structural order or vice versa, lower MF (Ziegelhöffner *et al.* 1997, Waczulíková *et al.* 2007, Slezák and Šikurová 2008).

Another fluorescence probe for estimating MF is pyrene (Fig. 1; Barenholz *et al.* 1996, Watala *et al.* 2002). Monitoring of pyrene mobility in the membrane or in

bilayers of lipid vesicles was preceded by pre-incubation of the membrane suspension with aqueous solution of the probe for 20 min at room temperature. Fluorescence of pyrene-labeled samples was excited at 339 nm and the emission was recorded in the range of 350-550 nm (2.5-nm slit). Pyrene mobility was reflected as the yield of the excimer formation (excimer fluorescence-to-monomer fluorescence ratio monitored at 395 nm and 470 nm, respectively), which directly depends on rate of the collision and on mutual distances of probe molecules in the lipid bilayer.

Methods used in investigation of endogenous protective mechanisms localized on subcellular membrane systems in the heart

Estimation of sarcolemmal Na,K-ATPase activity

Isolation of sarcolemmal membranes involved the treatments by hypotonic shock and with 0.6 mol.l⁻¹ sodium iodide. The membrane fraction obtained in this manner contained less than 3 % of sarcoplasmic reticulum and mitochondrial membranes. Na,K-ATPase activity is estimated on the base of inorganic phosphorus liberated by ATP splitting. For more details see Breier *et al.* (1995), Ziegelhöffner-Mihalovičová *et al.* (2003).

Perfusion of the isolated rat heart and induction of calcium paradox:

Rat hearts are quickly excised, cooled down, washed free of blood, placed in the perfusion device and “Langendorff”-perfused with Krebs-Henseleit (K-H) solution. After 15 min of stabilized perfusion at 37 °C the following variations in perfusion conditions are applied: i) Control hearts perfused for further 15 min with the basic K-H solution; ii) Hearts with calcium paradox perfused for 3 min with calcium free K-H solution (calcium depletion) and subsequently for 10 min with K-H solution containing 2.5 mmol.l⁻¹ Ca²⁺ (calcium repletion). The function of perfused hearts is evaluated by monitoring dp/dt_{max}, dp/dt_{min}. For more details see Ravingerová *et al.* (1997), Ziegelhöffner-Mihalovičová *et al.* (2003).

Estimation of mitochondrial Mg²⁺-dependent 2,4-dinitrophenol stimulated ATPase activity

Mg²⁺-dependent 2,4-dinitrophenol stimulated ATPase, also termed as the total ATPase was assessed in isolated mitochondria by estimation of inorganic phosphorus liberated from ATP splitting. Contamination

of the isolated mitochondrial fraction by other subcellular membranes was tested by estimation of marker ATPase activities of the sarcoplasmic reticulum and sarcolemma (Máleková *et al.* 2007) and it did not exceeded 2.2 %. For more details see Ferko *et al.* (2006), Mujkošová *et al.* (2008).

Estimation of conjugated dienes in cardiac mitochondrial membranes

Content of conjugated dienes in lipids of mitochondrial membranes was assessed by the method of Kogure *et al.* (1982) adapted to estimation of conjugated dienes in the membranes of heart mitochondria. Adaptation concerned the adjustment of optimal conditions for extraction of membrane lipids. For more details see Ziegelhöffer *et al.* (2009).

Cytochemical detection and quantitative determination of energy transition pores (contact sites) in heart mitochondria

The method for cytochemical detection of energy transition pores (ETP) is based on determination of the octameric mitochondrial isoform of the creatine kinase. The reaction of this enzyme is coupled to a series of subsequent reactions, the last being the reduction of thiocarbonyl nitroblue tetrazolium salt. After this step the sample becomes dehydrated, embedded in Epon 12 and further investigated in electron microscope. Detected contact sites are quantified by means of a stereological method and expressed in units of surface density. For more details see Ziegelhöffer-Mihalovičová *et al.* (1997).

Experimental data and discussion

In last 15-20 years the role of MF became investigated more intensively also in the membrane systems of the myocardium. A fast accumulating number of studies confirmed the regulatory role of MF for function of cardiac subcellular membrane systems in physiological and also in pathological conditions. Nevertheless, these studies also revealed that the simplified standard conception: “increased MF leads to augmented function of the membrane-bound systems”, does not provide all information which may be obtained from MF investigation.

It was demonstrated that in streptozotocin-induced experimental diabetes the calcium uptake of the heart exhibits a dramatical decrease as soon as 96 h after the injection of streptozotocin (Fig. 2; Gøtzsche 1991).

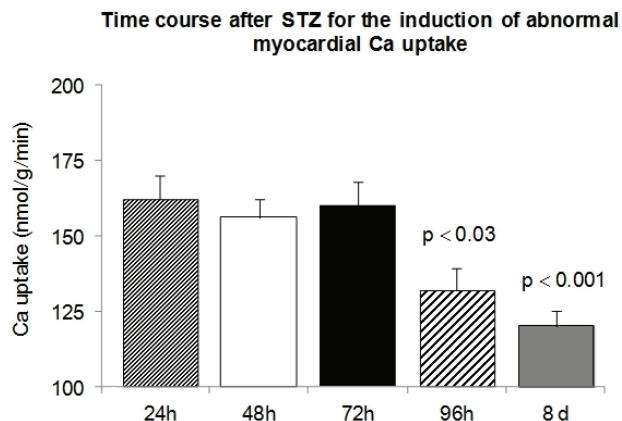


Fig. 2. Time course of myocardial calcium uptake after induction of experimental diabetes by application of streptozotocin. Data adapted, from the nomogram published by Gøtzsche (1991).

To explain the background of this phenomenon we investigated the fluidity of cardiac sarcolemma in similar conditions and revealed a marked 18.2 % ($p < 0.05$) increase in DPH anisotropy (Fig. 3) representing decreased MF that was also accompanied by a significant 33.7 % ($p < 0.05$) decrease in Na,K-ATPase activity (Ziegelhöffer *et al.* 1997, Ziegelhöffer-Mihalovičová *et al.* 2003). The last two findings were in accordance with the expectations. Nevertheless a parallel 25.2 % ($p < 0.05$) increase in sarcolemmal Mg/Ca-ATPase activity was also registered (Ziegelhöffer *et al.* 1996). The latter discordance might result from the fact that although MF is regulatory to function of sarcolemmal components it may be itself regulated by several other factors and the diverse influences may interfere.

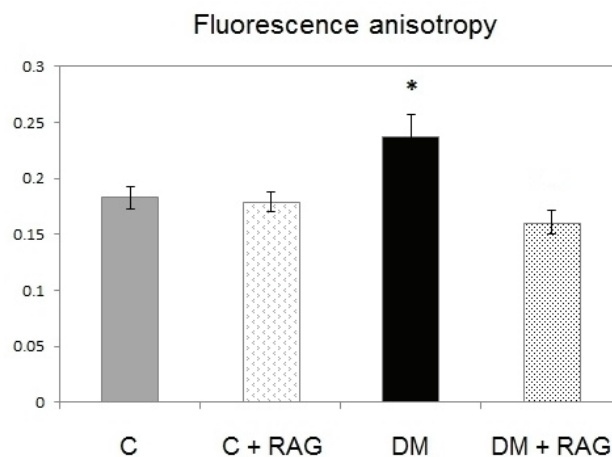


Fig. 3. Streptozotocin-diabetes and resorcylicidene amino-guanidine (RAG)-induced changes in DPH anisotropy of the rat heart sarcolemma. Groups: DM – diabetes; DM+RAG – diabetes + treatment with RAG; C – healthy controls. Groups consisted of 12-15 hearts.

It was hypothesised, that the decreased permeability of sarcolemma to calcium ions observed in diabetes (Fig. 2) may be caused, at least in part, by non-enzymatic glycation of proteins (NEGP) and the related free radical production in the sarcolemma. It was assumed that prevention of NEGP would also prevent the decrease in MF and *via* it normalize the activities of sarcolemmal enzymes. Application of resorcyldene aminoguanidine (RAG), a non-toxic compound with anti-glycation and antioxidant effects (Čársky *et al.* 1978, Šikurová *et al.* 2000) indeed prevented completely the diabetes-induced decrease in MF and the sarcolemmal enzyme activities (Fig. 3; Ziegelhoffer-Mihalovičová *et al.* 2003). This testifies for modulation of MF by the processes of NEGP.

Preventing by RAG the diabetes-induced decrease in MF of the sarcolemma alleviated, to certain degree, the disturbances in cardiac function. But, on the other hand, it also weakened the capability of the heart to maintain adequate calcium handling.

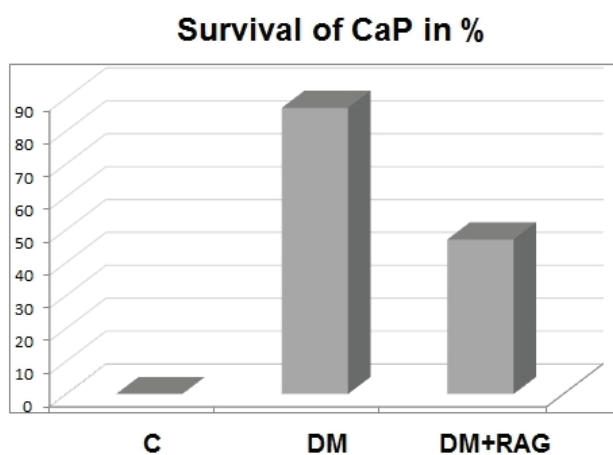


Fig. 4. Effect of calcium paradox and resorcyldene aminoguanidine (RAG) treatment on survival rate of isolated perfused diabetic hearts. Groups: D – diabetes; D+RAG – diabetes + treatment with RAG; C – healthy controls. Groups consisted of 12-15 hearts.

Isolated perfused hearts of rats with streptozotocin-diabetes showed massively depressed sensitivity to external calcium (Göttsche 1991). The latter feature was manifested in 83 % survival of an extremely severe form of calcium overload, the calcium paradox. On the other hand, the survival rate in the group of RAG-treated diabetic hearts, which exhibited better preserved MF and sarcolemmal enzyme activities, was diminished to 47 % only. None of the healthy control hearts survived the calcium paradox (Fig. 4). These

findings indicated that low MF of cardiac sarcolemma is favourable for survival of the diabetic heart. It proved to be a powerful means of endogenous protection of the myocardium against overload with external calcium (Ravingerová *et al.* 1996, Ziegelhoffer *et al.* 1996).

In contrast to sarcolemma with decreased MF, the mitochondria from streptozotocin-diabetic rat hearts exhibited a significant ($p < 0.05$), 83.3 % increase in fluidity of the bulk of their membrane lipids (Ferko *et al.* 2006). This state may result from different parallel running processes: i) remodeling of composition of membrane lipids (Wojtzak *et al.* 1969) that may be induced either by diabetes itself or by pseudo-hypoxia. The latter novel term characterizes a state when hypoxia is not caused by lack of oxygen but by insufficiency of the respiratory chain to utilize O_2 properly; ii) another process may be that the radicals produced in diabetic heart mitochondria are preferably extruded from mitochondria *via* the chloride channel (Máleková *et al.* 2007) instead of reacting with lipids inside the mitochondrial membrane; iii) third possible mechanism for keeping the MF of diabetic heart mitochondria elevated is associated with enhanced formation of energy transition pores (ETP) in mitochondrial membranes (Ziegelhoffer-Mihalovičová *et al.* 1997). These ephemeric structures, serving to high capacity energy transfer from the mitochondria to the cytoplasm, are bridging both mitochondrial membranes (Ziegelhoffer-Mihalovičová *et al.* 2002, Ziegelhoffer 2005, Ziegelhoffer *et al.* 2002, 2009). Their enhanced formation represents a compensatory response to increased energy demands of myocardial cells. This also indicates that ETP formation coupled with fluidization of the mitochondrial membrane belongs to mechanisms of endogenous protection of the myocardium. The process of MF elevation is initiated with the signal carried by increased calcium transients that leads to ETP formation in the mitochondrial membranes (Ziegelhoffer *et al.* 2002). ETPs are pulling both mitochondrial membranes close together and pushing aside the intramembrane fluid. The latter is then crowding in places between the ETPs and exerts elevated pressure on the lipid layers. Distances between the acyl chains of the membrane phospholipids increase and consequently the MF also increases (Ziegelhoffer 2005, Ziegelhoffer *et al.* 2009). This process can be demonstrated using the model of isolated perfused heart. Perfusion with elevated concentrations of calcium in perfusate increases considerably the number of mitochondrial ETP (Fig. 5) and subsequently also the MF (Fig. 6).

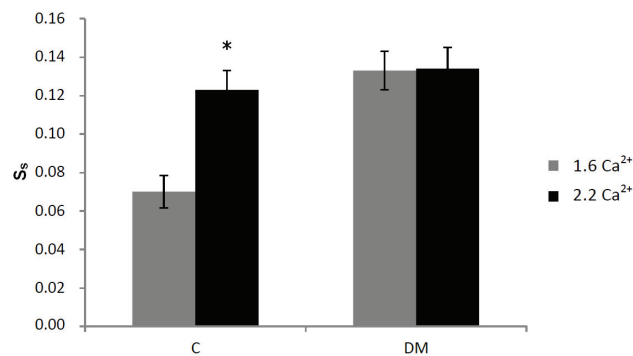


Fig. 5. Effect of calcium signaling on formation of energy transition pores in membranes of heart mitochondria. Groups: DM – diabetes; DM+RAG – diabetes + treatment with RAG; C – healthy controls. Results are mean \pm S.E.M. from 10 experiments. * significant at $p < 0.05$.

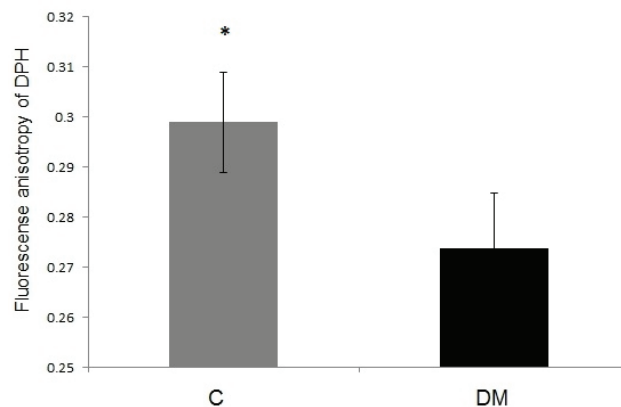


Fig. 6. Effect of calcium signaling on membrane fluidity of heart mitochondria. Groups: DM – diabetes; C – healthy controls. Groups consisted of 12-15 hearts. Results are means \pm S.E.M. from 10 experiments. * significant at $p < 0.05$.

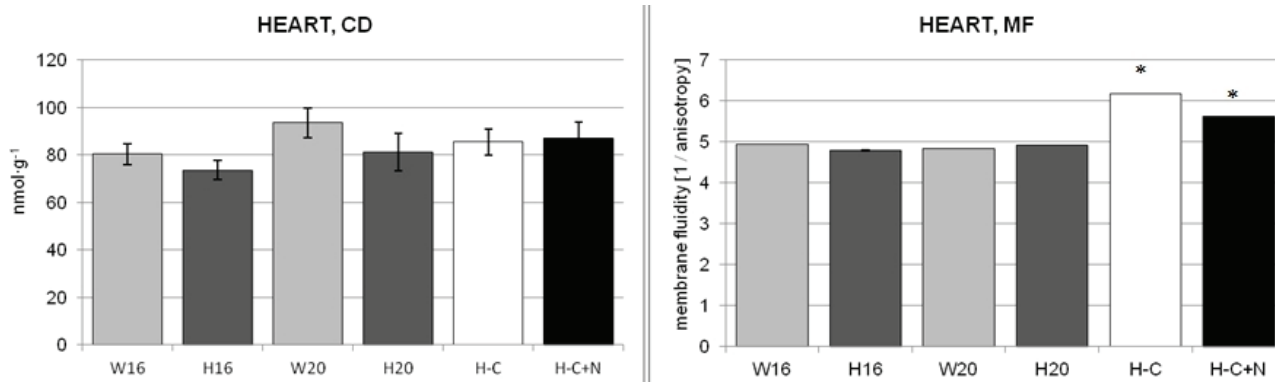


Fig. 7. Content of conjugated dienes (left hand panel) and membrane fluidity (right hand panel) in heart mitochondria of spontaneously hypertensive rats. CD – Conjugated dienes; MF – Membrane fluidity; W16 – 16 weeks old normotensive wistar controls; H16 – 16 weeks old hypertensive rats. Groups: W20 – 20 weeks old normotensive wistar controls; H20 – 20 weeks old hypertensive rats; H-C – 20 weeks old hypertensive rats treated with captopril for 4 weeks; H-C+N – 20 weeks old hypertensive rats treated with captopril + nifedipine for 4 weeks; * significant at $p < 0.05$ against H20.

Surprisingly little information was available about function of heart mitochondria in spontaneous hypertension. It could be assumed that mitochondria of hypertensive hearts are not limited in access to oxygen but they experience increased energy demands originating from elevated work-load of the heart. This might induce a compensatory elevation of mitochondrial functions. Our results confirmed this assumption. Hypertensive heart mitochondria exhibited significantly ($p < 0.05$ or more) increased Mg^{2+} -ATPase activity and elevated parameters of ATP production in oxidative phosphorylation (Mujkošová *et al.* 2010) but, unchanged membrane fluidity (Fig. 7, right hand panel). For the latter finding we have no special explanation at present. Nevertheless, a lack of increase in concentration of the conjugated dienes (Fig. 7, left hand panel) excludes any participation of free radicals-induced lipid oxidation in

this process.

On the other hand, treatment of hypertension with captopril or with captopril and nifedipine, both removed the hypertension efficiently and increased the MF of hypertensive heart mitochondria significantly (Fig. 7 right hand panel, $p < 0.05$ or more). Mechanism of this effect is still investigated.

Concluding remarks

In our research, we have been oriented to extension of knowledge about the regulatory role of subcellular membranes for function of the heart in health and disease. We paid particular attention to elucidation of mutual interplay between biophysical, biochemical and physiological processes securing the function of subcellular membranes in diverse physiological situations

and pathologies, particularly in diabetes.

In studies of abnormalities in calcium handling in the diabetic heart we have shown for the first time that decrease in fluidity of cardiac sarcolemma participates in protecting the diabetic heart against an overload with external calcium. This role makes modulation of MF to a means which may be utilized in endogenous protection of the myocardium.

We have reported for the first time the simultaneous measurements of membrane potential and dynamics as well as respiratory chain activities in the cardiac mitochondria under diabetic conditions. We have also shown that some of the processes regulating ATP production in the diabetic heart occur at the membrane level. Our results have pointed to the fact that not all changes triggered by some pathological impulse seem to be pathognomonic, but may represent compensatory and adaptive responses to the disease. It is necessary to emphasize that the latter assumption bears several important consequences: 1) the preconditioning-like protective and adaptive responses, discussed in detail by Ferdinandy *et al.* (1998), may be adequate to maintain the overall contractile performance of the diabetic heart at relatively normal levels, at least for a certain time; 2) the adaptive mechanisms start to develop in the cardiac myocytes; 3) it is highly probable that the observed fluidization of mitochondrial membranes with

concomitant increase in mitochondrial ATPase activity and changes in membrane potential serve adaptive functions in the mitochondria of diabetic heart. Nevertheless, the mechanism of partnership of these 3 factors in processes of endogenous protection still requires some more elucidation.

It may be concluded that focused coordination of biophysical, biochemical and physiological approaches may also in future open new insights in the regulatory role of subcellular membranes in heart function.

Conflict of Interest

There is no conflict of interest.

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