
DIRECTOR'S ADDRESS

Yet Another Decennium of the Institute of Physiology: A Dynamic Interplay of Innovative Approaches and 60 Years of Tradition

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I am pleased to introduce this special issue of *Physiological Research* published on the occasion of the 60th anniversary of the Institute of Physiology. It is only a second issue of this kind, the previous one being *Physiological Research* 53 (Suppl. 1) 2004. Since then, the Institute contributed its expertise to modern fields of physiology such as cardiovascular physiology, neurophysiology, energy metabolism, membrane transport, chronobiology, as well as relevant methodology. Diverse local and international collaboration has augmented such effort, as summarized in the attached *Synopsis* outlining the most significant achievements of Institute's departments during the past ten years. I very much hope that achievements of this kind will become Institute's tradition justifying at least equally optimistic forthcoming special issues in the decades to come.

Synopsis

The last decade has witnessed a considerable progress in experimental **cardiovascular research** focused on multifactorial polygenic diseases such as hypertension, ischemic heart disease, atherosclerosis, arrhythmias, heart failure or vascular disease. New approaches were also developed for the replacement of damaged blood vessels.

A great progress in the genetic analyses of complex pathophysiological traits in the BXH/HXB recombinant inbred (RI) strains developed from spontaneously hypertensive rat (SHR) and Brown Norway (BN) progenitors in the *Department of Genetics of Model Diseases*, has been achieved owing to an international collaborative effort supported by two EU project grants (EURATools and EURATRANS). Recent technical advances, including the next-generation sequencing technology, transcriptome analysis, genomic variation analysis, methylation-sequencing and ChIP-seq, quantitative proteomics, metabonomics, advanced bioinformatics as well as new tools for manipulating the rat genome, made it possible to identify the molecular basis of the first genetic determinants predisposing the SHR to hypertension (Pravenec *et al.* 2008), left

ventricular hypertrophy (McDermott-Roe *et al.* 2011) or cardiac fibrosis (Liška *et al.* 2014).

An increasing attention has been paid to the expression of genetic abnormalities into pathophysiological alterations responsible for abnormal vascular tone and high blood pressure. The estimation of a quantitative contribution of certain vasoactive systems to blood pressure control in the *Department of Experimental Hypertension* yielded a study evaluating the contribution of calcium influx and calcium sensitization to vascular tone control, the interplay of these two 'contractile' pathways and their modulation by distinct vasoconstrictor and vasodilator systems (Pintérová *et al.* 2010, Behuliak *et al.* 2013).

Considering high blood pressure as a major risk factor of the ischemic heart disease, the alterations of myocardial tolerance to ischemia-reperfusion injury in distinct forms of systemic hypertension have been studied in the *Department of Developmental Cardiology* (Neckář *et al.* 2012). In addition, a significant progress has been made in delineating the molecular mechanisms underlying high ischemic tolerance of immature and chronically hypoxic hearts, with a particular attention paid to the role of various mitochondrial proteins and redox-dependent protective signaling pathways (Borchert *et al.* 2011).

The *Department of Functional Morphology* studied whether a chronic administration of red palm oil (RPO, rich in antioxidants) and n-3 polyunsaturated fatty acids (n-3 PUFA, EPA/DHA Omacor) can ameliorate pathophysiological changes including cardiac arrhythmias and lipid level alterations induced by thyroid hormones. Using hyperthyroid and hypothyroid rats as “models of a diseased organism”, it has been established that RPO and n-3 PUFA can partly ameliorate changes in cardiac tissue remodeling, cell-to-cell communication, expression of Cx43 and protein kinases or in myosin heavy chain composition induced by thyroid hormone level alterations (Radosinska *et al.* 2013, Rauchová *et al.* 2013).

Developmental arrhythmology is a dynamically expanding field, in part owing to the advent of new tools such as high-speed cameras for optical mapping, as well as new markers of the developing cardiac conduction system. The analysis of several mutant mouse strains in the *Department of Cardiovascular Morphogenesis* shed new light on the function of the cardiac conduction system during embryonic development. Connexin40, a dominant connexin expressed in the His bundle, bundle branches and Purkinje fibers, has been found to be crucial for the right bundle branch conductivity only during later embryonic and postnatal stages (Sankova *et al.* 2012). An analysis of mice with myocardial-specific deletion of Pitx2C has demonstrated that their hearts have dual sinoatrial nodes, both acting as functional cardiac pacemakers (Ammirabile *et al.* 2012). Finally, overexpression of a dominant negative mutant form of Kcnq1 potassium channel in a mouse model of human long QT syndrome results in ventricular conduction system dysmorphogenesis and dysfunction (de la Rosa *et al.* 2013).

The *Department of Biomaterials and Tissue Engineering* concentrates on the reconstruction of irreversibly damaged blood vessels. This is achieved by either innovating synthetic polymeric vascular prostheses currently used in clinical practice or by constructing new bioartificial vascular replacements using tissue engineering methods. The existing vascular prostheses are innovated mainly by coating their inner surface with specific protein layers, particularly fibrin-based ones, which is then followed by *in vitro* endothelialization (Filová *et al.* 2009). The construction of novel bioartificial blood vessels utilizes synthetic or natural polymers functionalized with nanoparticles or various ligands for cell adhesion receptors as well as nanofibrous or nanoporous scaffolds as carriers for the attachment, growth and phenotypic maturation of vascular endothelial

and smooth muscle cells (Bačáková *et al.* 2007, Novotná *et al.* 2013). A special attention has also been paid to the development of perivascular drug delivery system into blood vessels (Filová *et al.* 2011).

In the field of **energy metabolism**, mechanistic studies on specific aspects of mitochondrial energy conversion and its regulation have been extended to a research more strongly focused on either integrative cell biology or direct links to medicine, namely to mitochondrial diseases as well as obesity and diseases associated with it. Thus, recent studies of mitochondrial uncoupling proteins in the *Department of Membrane Transport Biophysics* revealed an antioxidant synergy of the mitochondrial uncoupling protein 2 (UCP2) and mitochondrial phospholipase iPLA_{2γ} preventing oxidative stress in tissues such as lung, and lipotoxicity in pancreatic β-cells. The role of mitochondria and redox regulations in glucose-stimulated insulin secretion upon cell hypoxic adaptation and in tumorigenesis is also subject to investigation (Jabůrek *et al.* 2013, Tauber *et al.* 2013).

Mitochondrial diseases, a previously underscored cause of fatal outcomes in childhood belong to the most severe inborn metabolic diseases. Identification of genes responsible for them, uniquely including also the mitochondrial genome, and elucidation of the relevant molecular pathogenic mechanisms had established the basis for the diagnostics and prevention of these yet untreatable diseases, and led to a significant progress in the understanding of mammalian mitochondrial biology. The *Department of Bioenergetics* has significantly contributed to the characterization of mechanisms of biogenesis of multi-subunit enzyme complexes of the mitochondrial oxidative phosphorylation system while studying inherited disorders of ATP synthase, the key component of mitochondrial energy provision. Identification of two new genes responsible for nuclear genetic defects of the ATP synthase, manifesting themselves as neonatal encephalocardiomyopathy, paved the way to the discovery of a novel biogenetic factor specific for higher eukaryotes (Čížková *et al.* 2008, Mayr *et al.* 2010).

Research conducted in the *Department of Adipose Tissue Biology* has focused on the systemic effects of the induction of mitochondrial oxidative capacity in white adipose tissue in response to omega-3 fatty acids and other factors. This line of investigation was prompted by the discovery that an enhanced energy expenditure in white adipose tissue induced by mitochondrial uncoupling in transgenic mice can

counteract obesity. Recent results documenting the role of mitochondrial oxidative phosphorylation in a healthy phenotype of white adipocytes revealed important details about the mechanisms of the beneficial health effects of omega-3 fatty acids, and suggested yet unexplored potential of omega-3 fatty acids in treating obesity and associated diseases (Jelenik *et al.* 2010, Flachs *et al.* 2013).

While studying the role of **membrane transporters** in cell cation and pH homeostases, a number of novel results have been obtained in the *Department of Membrane Transport*. A combination of molecular biology (cloning of genes and their heterologous expression), biophysics (measurements of relative membrane potential or intracellular pH and their changes) and chemistry (estimation of cation content and fluxes) made it possible to identify, clone and characterize new alkali-metal-cation transporters involved in the regulation of cation homeostasis in yeast cells (Petrezsélyová *et al.* 2013), to describe the structural properties of cation/proton antiporters (Kinclová-Zimmermannová *et al.* 2005, 2006) and to identify new regulatory proteins involved in cation homeostasis (Zahrádka *et al.* 2012).

Research of the *Department of Protein Structure* is focused on the molecular basis of the 14-3-3 protein-dependent regulation of three important 14-3-3 binding partners: the yeast neutral trehalase Nth1, the regulator of G-protein signaling 3 (RGS3) and phosducin (Veisova *et al.* 2012, Rezabkova *et al.* 2011, 2012, Macakova *et al.* 2013). The mechanisms under investigation feature a common aspect – a conformational change induced by binding to the 14-3-3 protein molecule takes place in regions that are remote from the segment containing the phosphorylated 14-3-3 binding motif(s). This confirms that the interactions between 14-3-3 and their ligands extend beyond the ligand-binding groove. This may explain the isoform-specific interactions between 14-3-3 proteins and their ligands. In addition, these studies have also revealed that the 14-3-3 proteins sterically block the binding surface of RGS3 and phosducin, thus inhibiting their interactions with other binding partners.

A significant progress has been made by several research groups at the Institute in the rapidly expanding field of **neurophysiology**. Recent evidence suggests that abnormal regulation of membrane receptors plays a fundamental role in the development of many neurological and psychiatric disorders, including Parkinson's, Alzheimer's, and Huntington's diseases, epilepsy, anxiety, depression, bipolar disorder,

schizophrenia, lupus erythematosus and ischemia. Proper understanding of the fundamental mechanisms regulating the membrane receptors in the mammalian nervous system is thus essential if novel approaches for treating these disorders are to be successfully developed.

Research into the mechanisms underlying the effects of so-called allosteric modulators affecting the activity of a receptor at a site different from the receptor's active site represent a topic of shared interest in the field of ion channel and G-protein coupled receptor research, in both peripheral and central nervous systems. Specifically, studies on the N-methyl-D-aspartic acid (NMDA) subgroup of ionotropic glutamate receptors have focused on the identification of clinically relevant antagonists capable of preferentially blocking the excitotoxic receptor activation, without interfering with its functions essential for a normal synaptic transmission and neural plasticity. Research at the *Department of Cellular Neurophysiology* has considerably extended the knowledge on function, structure, trafficking, molecular genetics, and pharmacology of ligand-gated ion channels including glutamate, acetylcholine, and pain-related transient receptor potential channels (Lindovsky *et al.* 2008, Boukalova *et al.* 2010, Borovska *et al.* 2012, Kaniakova *et al.* 2012, Marsakova *et al.* 2012).

The *Department of Neurochemistry* has contributed to the understanding of the allosteric pharmacology of muscarinic receptors (Jakubík *et al.* 2011, Janíčková *et al.* 2013) whereas the physiological role of purinergic receptors and identification of the molecular mechanisms underlying the action of ivermectin, a positive allosteric regulator of several ligand-gated ion channels including the P2X4 subtype, has been reported by the *Department of Cellular and Molecular Neuroendocrinology* (Jelínková *et al.* 2008, Bhattacharya *et al.* 2013). The progress in both fields may have implications for the development of new compounds for treating cognitive disorders and pain, and improving learning and memory.

The *Department of Functional Morphology* has demonstrated that transient receptor potential vanilloid receptors (TRPV1) located on presynaptic endings of primary afferents in the spinal cord dorsal horn play a significant role in the development of acute and chronic pain states. Their modulatory role is potentiated under pathological conditions when their activity may be affected by a number of cytokines, and their sensitivity to endogenous agonists is increased (Spicarova and Palecek 2009, Spicarova *et al.* 2011).

The *Department of Biochemistry of Membrane*

Receptors has reported that the plasma membrane-enriched fraction isolated from the cerebral cortex of rats exposed to increasing doses of morphine (10-50 mg/kg) for 10 days contained a high amount of adenylylase ACI and ACII. The other isoforms (ACIII-X) remained unaltered (Ujčíková *et al.* 2011). The increase of ACI and II hydrophobic membrane interior (Břejchová *et al.* 2012).

In cognitive neuroscience, the *Department of Neurophysiology of Memory* has described an intriguing phenomenon of a continuous updating of a changing information, by employing a task involving avoidance of a small programmable robot, and the role of hippocampus in this behavior has been determined (Telensky *et al.* 2011). It has been also discovered that hippocampus is critical for recognition of objects projected onto a computer screen (such as "virtual reality in rats") (Levcik *et al.* 2013). In collaboration with other research groups, a novel neuroprotective steroidal derivative has been successfully patented (Rambousek *et al.* 2011).

The *Department of Computational Neuroscience* has mainly focused on neuronal coding and information processing. Both invertebrate and vertebrate olfactory sensory systems have been studied, and sensory-motor coupling in insect flight control analyzed. Various aspects of information representation in the temporal character of neuronal signals have been investigated in order to identify potential information coding modes (Kostal *et al.* 2007). Furthermore, methods enabling advanced statistical analysis of experimental data have been proposed (Kostal *et al.* 2013).

Using a complex approach, the *Department of Developmental Epileptology* has demonstrated that an intense epileptic activity leads to both acute and long-lasting morphological and functional alterations, often of progressive nature, in rats younger than two weeks (Kubová and Mareš 2013). Mechanisms responsible for the damage in immature brain include oxidative stress and mitochondrial dysfunction. Oxidative stress is apparently due to both increased free radical production and limited antioxidant defense (Folbergrová *et al.* 2012). Metabotropic glutamate receptors also play an important role in the generation of epileptic activity and neuronal damage. Agonists of groups II and III, as well as antagonists of group I of the metabotropic glutamate receptor exhibited an anticonvulsant and neuroprotective effect (Folbergrová *et al.* 2008, Lojčková-Janečková *et al.* 2009).

Recent progress in **chronobiology** is already having far-reaching consequences in other fields of

physiology. From a relatively specialized field focusing on the properties of the circadian clock in the brain and its output rhythms (mostly behavioral and humoral), the scope of chronobiology has broadened enormously owing to the discovery of the temporal regulation of cellular processes and its vital importance for cell survival. The cellular clocks are hierarchically organized at the systemic level into a circadian system. The disruption of communication among its individual components and/or the system's communication with the external environment result in the malfunction of the temporal regulation of physiological processes in general, with serious health consequences. Using the molecular biology tools, the *Department of Neurohumoral Regulations* discovered how the circadian system and its individual components develop during ontogenesis (Sumová *et al.* 2012). In collaboration with the *Department of Epithelial Physiology*, the studies on the circadian regulation in the gastrointestinal system revealed a specific function of the circadian clock in the colonic epithelial cells, and the mechanisms of entraining these clocks with the external environment (Sládek *et al.* 2007, Polidarová *et al.* 2011). Importantly, the clock malfunction has been associated with colorectal cancer development (Soták *et al.* 2013). Our recent studies in humans have revealed that neuropsychiatric disorders are associated with malfunctioning of the circadian system at the level of molecular clock regulation (Nováková *et al.* 2012). These findings may contribute to new chronotherapeutic approaches to various diseases.

A dramatic progress in the development of **new technologies and methodological approaches** has also been reflected by achievements of the Institute. Recent imaging and tissue preparation techniques provide high-resolution 3D image data. The *Department of Biomathematics* is engaged in the development of new microscopic visualization (Pelc *et al.* 2008) and image analysis methods, and their applications to microscopic architecture of various tissues such as blood capillaries in brain, muscles and placenta. Recent achievements are exemplified by a more efficient software for registration of neighboring physical slices (of microscopy specimens) suffering from discontinuities (Michálek and Čapek 2013), and a comparative study on length measurement of tubular structures in 3D (Kubínová *et al.* 2013).

A progress in analytical separation methods and their application to physiologically important compounds (e.g., steroids or pigments) at the *Department of Analysis of Biologically Important Compounds* is exemplified by the recent development of a gold nanoparticles-based

stationary phase approach (Mikšik *et al.* 2012, Pataridis *et al.* 2013). The proteomes of various biological material such as teeth, avian eggshells and human mummy, as well as non-enzymatic modifications of proteins have also been studied.

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