

**Univerzita Karlova v Praze**  
**1. lékařská fakulta**

Studijní program: Biomedicína  
Studijní obor: Fyziologie a patofyziologie člověka



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Regulace receptorů spřažených s G proteiny

*Studie muskarinových a  $\beta$ -adrenergických receptorů u  $M_2$ KO myši*

Regulation of G protein-coupled receptors

*Study of muscarinic and  $\beta$ -adrenoceptors in  $M_2$ KO mice*

Disertační práce

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Praha, 2014

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JAN BENEŠ

## **Poděkování**

Rád bych poděloval především svému školiteli prof. MUDr. Jaromíru Myslivečkovi, Ph.D za odborné vedení a pomoc, kterou mi věnoval jak při pregraduálním tak postgraduálním studiu a při přípravě této práce.

Dále bych chtěl poděkovat svému školiteli konzultantovi doc. MUDr. Vojtěchu Melenovskému CSc. za veškerou pomoc, zejména pak za cenné rady a zkušenosti týkající se tvorby odborného textu.

### **Identifikační záznam**

BENEŠ, Jan. *Regulace receptorů spřažených s G proteiny: Studie muskarinových a  $\beta$ -adrenergických receptorů u  $M_2KO$  myší. [Regulation of G protein-coupled receptors: Study of muscarinic and  $\beta$ -adrenoceptors in  $M_2KO$  mice.]* Praha, 2014, 93 s, 0 příl. Disertační práce (Ph.D.). Univerzita Karlova v Praze, 1. lékařská fakulta, Fyziologický ústav. Školitel Mysliveček, Jaromír, školitel konzultant Melenovský, Vojtěch.

## **Použité zkratky**

AC .....	adenylyl cyclase
AR.....	adrenergic receptor, adrenoceptor
$\beta$ -AR .....	$\beta$ -adrenergic receptor, $\beta$ -adrenoceptor
CaMKII.....	calmodulin activated protein kinase II
Df.....	degree of freedom
EGF .....	epidermal growth factor
EGFR .....	epidermal growth factor receptor
Epac .....	exchange factor directly activated by cAMP
GPCR .....	G protein-coupled receptor
GRK .....	G protein receptor kinase
HR .....	heart rate
IVSd.....	interventricular septum thickness in diastole
JAK .....	Janus kinase
LVIDd .....	left ventricle end-diastolic diameter
MAPK .....	mitogen activated protein kinase
MMP.....	matrix metalloproteinase
MR .....	muscarinic receptor
NOS .....	NO synthase
NTS .....	nucleus tractus solitarii
PKC.....	proteinkinase C
PLAX .....	parasternal long axis
PLB, PLC .....	phospholipase B, C
PSAX .....	parasternal short axis
PTCH .....	patched
PYK2.....	proline-rich tyrosine kinase
Rap .....	Ras-related protein
RTK .....	receptor tyrosine kinase
SMO .....	smoothed
Smad.....	intracellular proteins that transduce signals in TGF $\beta$ pathway
Src .....	non-receptor protein tyrosine kinase (derived from the word „sarcoma“)
TGF $\beta$ .....	transforming growth factor $\beta$

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## **Souhrn (česky):**

Cílem práce bylo podrobně analyzovat myši s vyřazeným  $M_2$ -receptorem ( $M_2$ KO) jednak za klidových podmínek a dále po působení stresového podnětu (chladový stres). Zaměřili jsme se zejména na roli antagonistických receptorů (adrenoceptorů) v srdci a jejich roli v adaptaci na chybějící  $M_2$ -receptor. Provedeny byly vazebné studie, studie genové exprese, echokardiografie, telemetrický monitoring srdeční činnosti, tělesné teploty a aktivity, analýza variability tepové frekvence, analýza biorytmů, analýza srdeční činnosti po podání agonistů a antagonistů muskarinových receptorů a adrenoceptorů, stanovení aktivity adenyllyl cyklázy a NO syntázy, stanovení sérové koncentrace katecholaminů a genové exprese enzymů syntetizujících katecholaminy. Vyřazení  $M_2$ -receptorů vedlo ke kompenzatornímu snížení  $\beta_1$  a  $\beta_2$ -adrenoceptorů doprovázené snížením jejich genové exprese, down-regulovány byly rovněž  $M_3$ -receptory. Chybějící  $M_2$ -receptory byly funkčně nahrazeny  $\beta_3$ -receptory, které byly upregulovány (nikoli kardioinhibičními  $M_4$ -receptory). Signalizace přes  $\beta_3$ -receptory vedla dále cestou adenylátcyklázy a nikoli NO syntázy. Všechny tyto změny byly zjištěny pouze v levé komoře, odpovědným mechanismem se tedy zdá být heterologní regulace. Mírně vyšší bazální frekvence a nižší tělesná teplota  $M_2$ KO myši podtrhuje důležitost telemetrické monitorace. I přes všechny pozorované rozdíly, celková srdeční funkce u  $M_2$ KO myši byla zachována.

**Klíčová slova:**  $M_2$ KO myši, adrenoceptory, muskarinové receptory, srdce, stres, adaptace

**Abstract (in English):**

The aim of the work was to perform in-depth analysis of M<sub>2</sub>KO mice both at baseline and upon a challenge with a cold stress and to explore the role of opposing receptors (i.e. adrenoceptors) in adaptation to lacking M<sub>2</sub>-receptors in the heart. We have performed receptor binding studies, study of receptor gene expression, echocardiography, telemetric monitoring of heart rate, body temperature and activity, heart rate variability and biorhythm analysis, analysis of heart rate response to the application of drugs (carbachol, atropine, isoprenaline, propranolol), assessment of adenylyl cyclase and NO synthase activity, measurement of catecholamine blood concentration and gene expression of catecholamine-synthesizing enzymes. We have found that the disruption of M<sub>2</sub>-receptor gene caused a compensatory decrease of cardiostimulatory  $\beta_1$ -adrenoceptors and  $\beta_2$ -adrenoceptors with corresponding down-regulation of the gene expression, M<sub>3</sub>-receptors were down-regulated as well. Missing M<sub>2</sub>-receptors were functionally replaced by the main cardioinhibitory  $\beta_3$ -adrenoceptors that were up-regulated, not by cardioinhibitory M<sub>4</sub>-receptors.  $\beta_3$ -adrenoceptors were found to signal through adenylyl cyclase instead of NO synthase. All these changes were found in the left ventricle only, so heterologous regulation is likely to be the responsible mechanism. Slightly higher basal heart rate and lower basal body temperature of M<sub>2</sub>KO animals underscore the importance of telemetric measurement to avoid biased recordings. Despite all the changes found the overall cardiac function was unaltered.

**Key words:** M<sub>2</sub>KO mice, adrenoceptors, muscarinic receptors, heart, stress, adaptation

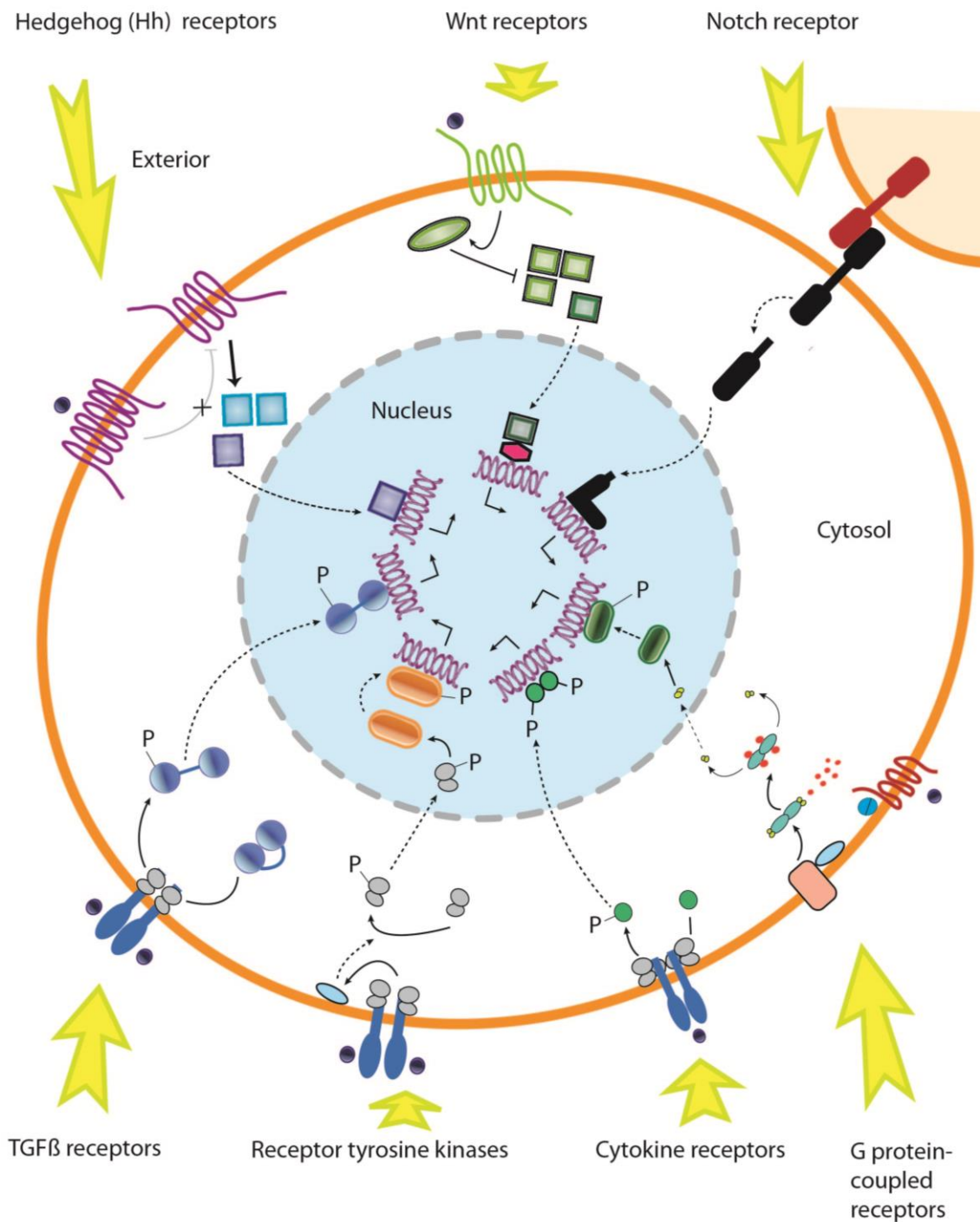
## **1. Introduction**

### **1.1. Cellular signalization**

In a multicellular organism, all cells are involved in cooperation, which requires mechanisms for cell communication. There is either a direct cell-to-cell signaling or cells communicate at the longer distances through „signaling molecules“ that control metabolic processes within cells, the synthesis and processing of proteins, the growth and differentiation of tissues and the composition of both intracellular and extracellular environment.

Signaling molecules are synthesized and released by signaling cells and either influence just neighboring cells (paracrine signaling, synaptic transmission) or travel throughout the body (hormones) but induce a specific response only in target cells that have appropriate receptors. Substances used for intercellular signaling encompass a huge variety of items including small molecules (derivatives of amino acids, lipids, acetylcholine, steroids), oligopeptides (vasopressin) or large proteins (growth hormone, TSH etc.).

Principally, there are two different receptor systems. Receptors can be localized either intracellularly (in this case, they usually bind hydrophobic molecules such as steroids that diffuse spontaneously through the plasma membrane) or they are located in the plasma membrane. Ligands for these receptors bind to a structurally complementary site on the extracellular or membrane-spanning domains of the receptor, which causes conformational changes in the cytosolic domain of the receptor, which subsequently induces specific changes within the cell - a process called signal transduction. Signals from outside the cell can induce either changes in the activity or function of preexisting proteins or they can induce a de-novo production of proteins (proteosynthesis). Although there are many signaling pathways that enable the cell to accomplish a myriad of cellular functions there are principally only a few major classes cell surface receptors. G protein-coupled receptors comprise the largest family (with a several variants represented by hedgehog and Wnt receptors), catalytic receptors (cytokine receptors, TGF $\beta$  receptors, receptor tyrosine kinases) and Notch receptors (Alexander et al., 2011). These are depicted in Fig. 1.1.



**Fig. 1.1. The overview of major types of surface receptors**

Major classes of cell-surface signaling receptors are depicted.

**Hedgehog pathway** is shown in the upper left corner. Binding a ligand (e.g. sonic hedgehog) on a receptor (Patched-1, PTCH1) leads to the inhibition of Smoothened (SMO), which releases the inhibition of SMO upon GLI transcription factors. GLI then migrates into the

nucleus where it regulates the expression of target genes. Hedgehog pathway is particularly important in development.

**Wnt pathway** is activated by binding of Wnt-protein ligand to a Frizzled family receptor, which passes the signal to the Dishevelled protein inside the cell. There are three Wnt pathways: the canonical Wnt pathway (using  $\beta$ -catenin signaling protein) and two noncanonical pathways (planar cell polarity pathway, Wnt/calcium pathway) that work independently of  $\beta$ -catenin. Wnt pathways are crucial in embryonic development, axis patterning, cell fate specification, cell proliferation, cell migration and insulin sensitivity.

**Notch signaling** usually occurs between neighboring cells. One expresses the Notch ligand (e.g. Delta), the adjacent one then Notch receptor. Binding of the ligand promotes two steps involving proteolysis; as a result, intracellular domain is liberated and enters the nucleus where it regulates the gene expression. Notch signaling is important in multiple processes including neuronal and cardiovascular development.

**TGF $\beta$  signaling pathway** begins with binding a ligand to TGF $\beta$  type II receptor (sometimes type III receptor that presents the ligand to the type II receptor). Type II receptor then recruits and phosphorylates type I receptor, which releases the inhibition of type I receptor kinase activity. Activated type I receptor phosphorylates Smad protein. Two phosphorylated Smad proteins associate with Co-smad protein and enter the nucleus, where they regulate the expression of target genes.

**Receptor tyrosine kinases (RTKs)** are receptors that have intrinsic protein tyrosine kinase activity in their cytosolic domain. Upon binding a ligand they dimerize, which causes the phosphorylation of cytosolic receptor tyrosine residues. Phosphorylated tyrosine residues then act as substrates for other proteins (e.g. Son-of-sevenless, Sos) that transmit the signal further in the cell. Growth factors (epidermal, fibroblast, vascular endothelial) and insulin typically signal through RTKs.

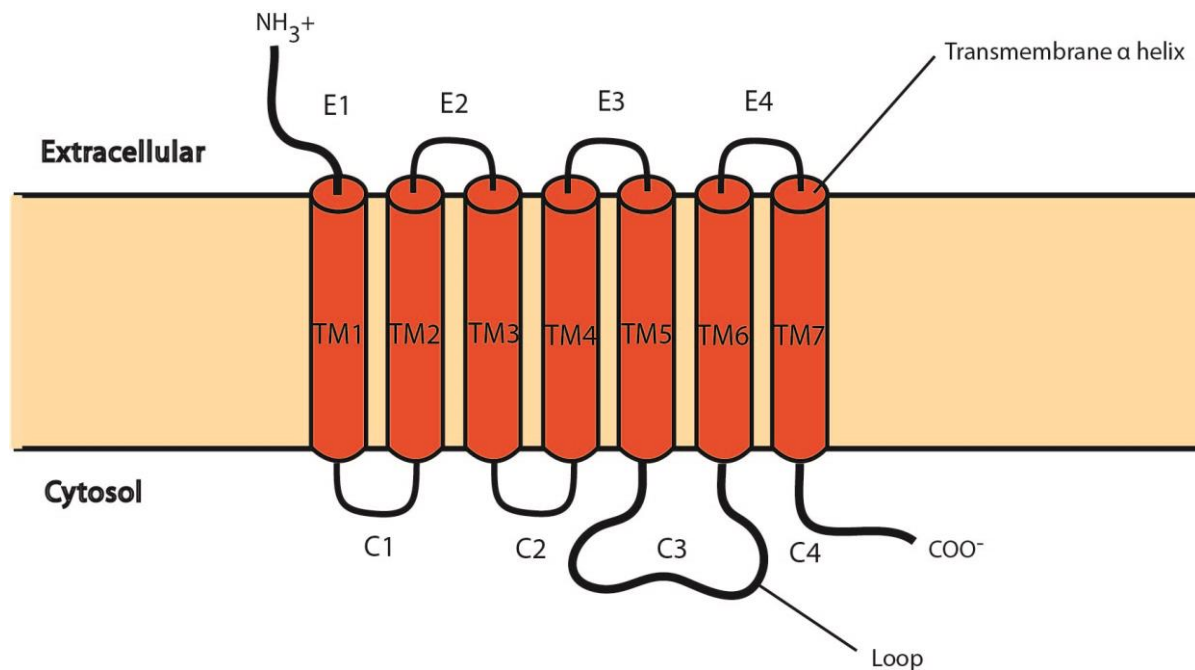
**Cytokine receptors** share many similarities with RTKs. Upon binding a ligand, cytokine receptor dimerizes, which brings Janus kinases (JAKs) located in the intracellular portion of the receptor to the close proximity. They phosphorylate the inner portion of the receptor leading to the activation of STAT proteins that dimerize and enter the nucleus, where they control a transcription of target genes.

Signaling through **G protein-coupled receptors** will be thoroughly reviewed in the following sections. Adapted after (Brivanlou and Darnell, 2002).

## 1.2. G protein-coupled receptors

G protein-coupled receptors (GPCR) are one of the major receptor systems in organisms. They transfer various messages as different as photons, organic odorants, nucleosides, nucleotides, peptides, lipids and proteins. There are at least six families of G protein-coupled receptors showing no similarity in their sequence. Similarly, considerable differences exist concerning the receptor binding sites. Adrenoceptors belong to the group Ib (Bockaert and Pin, 1999).

According to their name, these receptors signal through G proteins. From the structural point of view all G protein-coupled receptors contain seven membrane-spanning regions having their N-terminal segment on the extracellular side and their C-terminal segment on the cytosolic side of the plasma membrane. The structure of the GPCR is in Fig. 1.2.



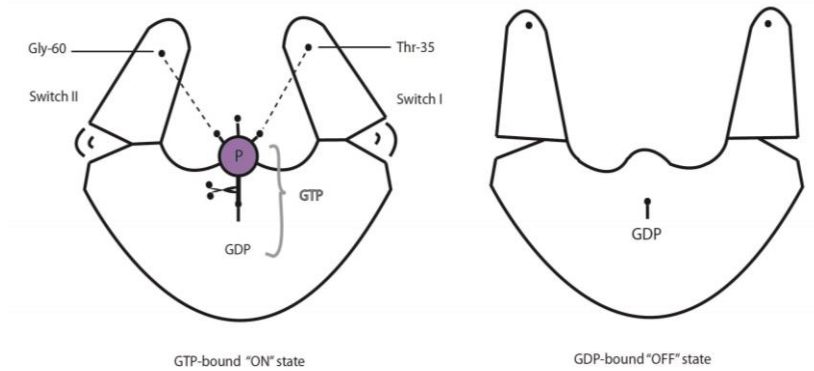
**Fig 1.2.: The structure of the typical G protein coupled receptor**

G protein-coupled receptors have four extracellular segments (E1-E4) seven transmembrane domains (TM1-TM7) and four cytosolic segments (C1-C4). The carboxy-terminal segments C4 and C3 loop (in some receptors also C2 loop) are involved with the interaction with G protein. Adapted after (Lodish H, 2000).

## 1.3. G proteins

G protein-coupled receptors belong to a family of proteins that transmit the signal through binding of GTP and its subsequent hydrolysis to GDP and phosphate (GTPase superfamily).

GTP hydrolyzing proteins are either monomeric (e.g. Ras protein) or trimeric, consisting of  $\alpha$ ,  $\beta$ , and  $\gamma$  subunit (G proteins). Both monomeric and trimeric GTPase superfamily proteins exist in two distinct forms – an inactive form with bound GDP and active form with bound GTP.

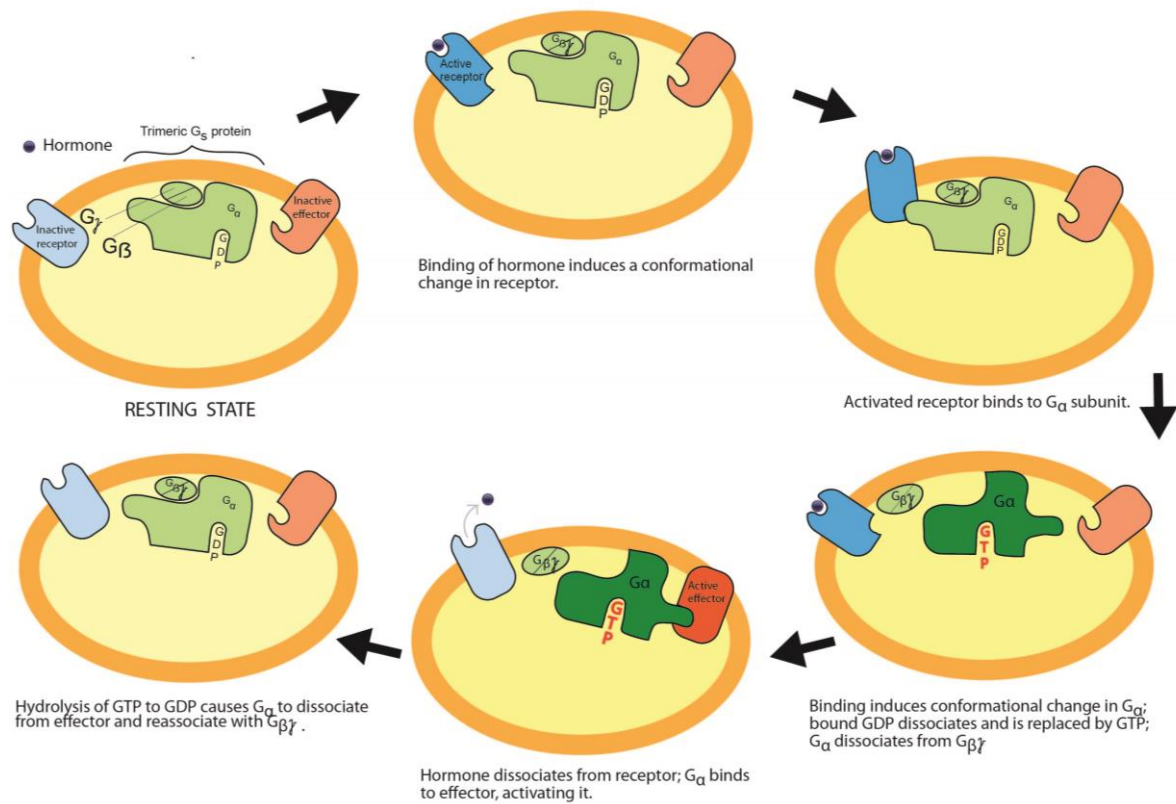


**Fig. 1.3.1: Switching mechanism for GTPase superfamily proteins**

In order to be able to transduce the signal, G protein needs to be able to switch between two states (GTP-bound “on” state, GDP-bound “off” state). In the active (“on”) state, two domains (named switch I and II, respectively) interact with the terminal  $\gamma$ -phosphate of the bound GTP through the interactions with conserved threonine and glycine residues. Upon GTP hydrolysis,  $\gamma$ -phosphate disappears leading to the different conformation of both switch I and II- inactive (“off”) state. Adapted after (Vetter and Wittinghofer, 2001).

G proteins are localized at the intracellular side of cytoplasmic membrane. All G proteins contain three subunits designated as  $\alpha$ ,  $\beta$  and  $\gamma$ .  $\beta$  and  $\gamma$  subunits remain bound together and are referred to as  $G\beta\gamma$  subunit. The  $G\alpha$  subunit is a GTPase switch protein that cycles between an inactive (off) and active (on) state. Upon binding a ligand, a conformational changes in the receptor occurs, which leads to the binding of activated receptor to G protein ( $G\alpha$  subunit). This leads to a conformational changes in  $G\alpha$  and is followed by a dissociation of GDP and binding of GTP (that is more abundant than GDP in the cell).  $G\alpha$  and  $G\beta\gamma$  subunits then dissociate and activated  $G\alpha$  (in some cases also  $G\beta\gamma$ ) transmit the signal further on other messengers. It is due to 3D-conformation when the third phosphate of GTP causes a conformational change and two domains (termed switch I and switch II) interact with the terminal  $\gamma$ -phosphate of GTP. Subsequently, intrinsic GTPase activity causes the hydrolysis of GTP to GDP and  $P_i$ , which leads to the release of  $\gamma$ -phosphate and to a conformational change and a separation of both switch I and II with the loss of the ability to transmit the signal. Both

subunits ( $G\alpha$  and  $G\beta\gamma$ ) then reassemble and are ready for the next round of signal transmission (Fig. 1.3.2).



**Fig. 1.3.2. Signal transduction through G protein**

Adapted according to the (Lodish H, 2000).

Signal transduction occurs at multiple levels (receptor ligand, receptor, G protein, adenylyl cyclase, cAMP, protein kinase A, activated enzymes) and at all levels the signal is usually greatly amplified. For example, a single molecule of a ligand activates one hundred molecules of G protein, leading to several hundred molecules of cAMP, each activating protein kinase A that subsequently activates several hundred molecules of the target protein. Thus, the plasmatic adrenaline concentration as low as  $10^{-10}$  mmol.l<sup>-1</sup> can induce substantial biological effects. This cascade leads not only to a mere amplification of the signal, but also allows an entire group of proteins involved in the same biological process to be regulated in a coordinated manner.

There are multiple G proteins encoded in genomes of organisms. Among the 3 subunits ( $\alpha$ ,  $\beta$  and  $\gamma$ ) only  $\alpha$  subunits directly interact with G protein-coupled receptors. The most important for the purpose of this thesis is  $G_s$  that activates adenylyl cyclase,  $G_i$  that inhibits it and  $G_q$



that activates phospholipase C. Compared to the enormous number of GPCR the number of genes encoding  $\alpha$  subunit of G protein is very small (Hurowitz et al., 2000). Based on their sequence morphology, it has been long thought that there were only 4 major classes of  $G\alpha$  proteins (Gs, Gi, Gq and  $G_{12}$ ) (Downes and Gautam, 1999). Each class can be subdivided into 2-4 families: the Gs class contain  $G\alpha_s$  and  $G\alpha_{olf}$ ; Gi comprises  $G\alpha_i$ ,  $G\alpha_o$ ,  $G\alpha_t$  and  $G\alpha_z$ ; Gq encompasses  $G\alpha_q$ ,  $G\alpha_{11}$ ,  $G\alpha_{14}$  and  $G\alpha_{15/16}$ ; and  $G_{12}$  contains  $G\alpha_{12}$  and  $G\alpha_{13}$  (Downes and Gautam, 1999). Only recently the fifth class of  $G\alpha$  proteins (Gv) was discovered (Oka et al., 2009). This fifth type was lost in many lineages including mammals.

#### **1.4. Types of G protein-coupled receptors (relevant for the topic of the thesis)**

##### **1.4.1. Adrenoceptors**

Adrenoceptors are membrane receptors located in both neuronal and non-neuronal tissue. They are responsible for mediating response to endogenous catecholamines noradrenaline (NA) and adrenaline (A). Noradrenaline is released from the noradrenergic postganglionic nerve terminals and both NA and A are secreted from the adrenal medulla. They are responsible for controlling many important physiological functions (cardiovascular, respiratory, neuronal, pupil contraction and dilatation, digestive, energetic metabolism, endocrine) (Calzada and de Artinano, 2001).

An observation that there must be more than one type of adrenoceptors was published in 1948, when Ahlquist observed differences in responses evoked by synthetic amines. He concluded that these differences must be due to the presence of different receptors, which he termed  $\alpha$  and  $\beta$  (Ahlquist, 1948).  $\alpha$ -adrenoceptors have mostly excitatory effects (vasoconstriction, uterine musculature contraction, urethra contraction, pupil dilation etc.) with the exception of intestinal relaxation. On the contrary,  $\beta$ -adrenoceptors have mostly inhibitory effects (vasodilatation, uterine musculature relaxation, bronchodilatation etc.) with the exception of cardiac stimulation (Calzada and de Artinano, 2001).

##### **1.4.1.1. $\alpha$ -adrenoceptors**

$\alpha$ -adrenoceptors divide into  $\alpha_1$  and  $\alpha_2$  subtypes. Both subtypes have other three subtypes ( $\alpha_{1A}$ ,  $\alpha_{1B}$ ,  $\alpha_{1D}$  for  $\alpha_1$  and  $\alpha_{2A}$ ,  $\alpha_{2B}$  and  $\alpha_{2C}$  for  $\alpha_2$ -subtype). Formerly,  $\alpha_{1C}$  was also described but it was shown later that this receptor is encoded by the same gene as  $\alpha_{1A}$ -AR and now it is considered as low affinity subtype of  $\alpha_{1A}$ -AR (Hiraizumi-Hiraoka et al., 2004, Muramatsu et al., 2005).

All  $\alpha_1$  subtypes are activated by both adrenaline and noradrenaline and blocked by the  $\alpha_1$ -adrenoceptor inhibitor prazosine. The subtypes vary in their amino acid sequence, tissue distribution and signaling pathways they activate. Recent findings on adrenoceptor signaling revealed many features that have challenged a classical linear paradigm of G protein-coupled receptor signaling. Adrenoceptors were proved to couple to multiple G proteins and even signal through G protein independent pathways, switch their signaling pathways over time and interact within or between their receptor subfamilies (Xiao et al., 2006). Thus paradigm of linear G protein-coupled signaling has gradually yielded to the model of „signalome“ where an individual adrenoceptor associates with multiple signaling pathways in time and spatially specific manner (Park and Palczewski, 2005). Moreover, adrenergic receptors were found to possess constitutive activity (Kjelsberg et al., 1992) and to form both homo- and heterodimers (Hague et al., 2004, Uberti et al., 2005). Having this in mind, it can be stated that according to the general paradigm,  $\alpha_1$ -AR couple to Gq protein and activate phospholipase C $\beta_1$ , which leads to diacylglycerol (DAG) and inositoltriphosphate (IP $_3$ ) production and subsequent activation of protein kinase C (PKC). The role of calcium release due to increased IP $_3$  concentration in cardiac myocytes is controversial. The  $\alpha_{1A}$ -subtype was shown to mediate cardioprotection through the activation of ERK. Under certain circumstances,  $\alpha_{1B}$ -AR can couple to Gi protein (Olshansky et al., 2008).

$\alpha_2$ -AR couple to Gi protein and inhibit adenylyl cyclase (Gibson and Gilman, 2006). The expression and function of both  $\alpha_1$  and  $\alpha_2$  adrenoceptors in the heart will be discussed in a separate chapter. Both  $\alpha_1$  and  $\alpha_2$  subtypes are present postsynaptically in blood vessels, where they mediate vasoconstriction (Langer and Hicks, 1984).  $\alpha_1$ -adrenoceptors are expressed in the prostate gland and are used as a therapeutic target in treating symptoms with benign prostate hyperplasia (Roehrborn and Schwinn, 2004).  $\alpha_2$ -adrenoceptors are widely expressed throughout the central nervous system and the periphery. In the central nervous system, the area with a major density of  $\alpha_2$ -adrenoceptors is the locus coeruleus, a region that contains the cell bodies for the ascending and descending noradrenergic neurons (MacDonald and Scheinin, 1995).

$\alpha_{2A}$ -subtypes are present in the prefrontal cortex and  $\alpha_{2A}$  agonists are therapeutically used in the treatment of attention deficit hyperactivity disorder (Childress and Sallee, 2012).  $\alpha_{2A}$ -subtype plays a major role in regulating vegetative nervous system; they are present on presynaptic noradrenergic nerve terminals where they inhibit the release of noradrenaline and adrenaline and thus inhibiting sympathetic nerve output (Hein et al., 1999).  $\alpha_{2A}$ -adrenoceptors were also found in platelets (Shekim et al., 1994), kidney and gastric mucosa (Gyires et al.,

2007).

#### **1.4.1.2. $\beta$ -adrenoceptors**

As already mentioned, the presence of distinct adrenergic receptors (termed  $\alpha$  and  $\beta$ ) was first described in 1948 (Ahlquist, 1948). In 1967, the evidence about distinct types of  $\beta$ -adrenoceptors was given. The first subtype mediated heart rate response to catecholamines; the second one then mediated bronchodilatation and vasodepression (Lands et al., 1967). Later on, the third subtype of  $\beta$ -adrenoceptor was found (Emorine et al., 1989). Moreover, at the end of 20<sup>th</sup> century, a fourth  $\beta$ -adrenoceptor was described ( $\beta_4$ -AR) (Kaumann and Molenaar, 1997) but similarly as in the case of  $\alpha_{1C}$  it was found later that this  $\beta_4$ -AR is only a low affinity state of  $\beta_1$ -AR (Granneman, 2001, Kaumann et al., 2001). However, the role of this receptor both in physiological processes as well as in pathology is estimated (Myslivecek et al., 2003, Mallem et al., 2004).

$\beta$ -adrenoceptors play multiple roles in the organism. Both  $\beta_1$ - and  $\beta_2$ -adrenoceptors are expressed in blood vessels; the proportion of both receptor subtypes varies in different vessel types (Chruscinski et al., 2001).  $\beta_2$ - and  $\beta_3$ -AR are expressed in bone marrow stroma cells where they have specific functions and collaborate during progenitor mobilization (Mendez-Ferrer et al., 2010). The presence of  $\beta$ -adrenoceptors has been suggested on nerves,  $\beta$ -adrenoceptors have been revealed in perivascular sympathetic nerves near the adventitial-medial border (Briones et al., 2005), which is consistent with a nurturing function of catecholamines in the repair and maintenance of nerves.  $\beta_2$ -AR are widely expressed in bronchial tissues, but the exact localization of them has been challenged recently (Davies, 2009). It has been long assumed that  $\beta_2$ -AR are on smooth muscle cells and they are responsible for removing asthma-induced contraction and mediate bronchodilatation. However, other cell types including the pulmonary airway epithelium have been proposed as central factors in airway reactivity and remodeling (Daly and McGrath, 2011).  $\beta$ -adrenoceptors also play an important role in mediating various metabolic processes; they play a role in insulin secretion from the pancreatic  $\beta$ -cells as adrenergic tone is one of the mechanisms regulating insulin secretion:  $\beta_2$ -AR activation stimulates insulin secretion and  $\alpha$ -AR activation inhibits it. This effect, is however species specific. Human  $\beta$ -cells express  $\beta$ -adrenoceptors but  $\beta$ -cells of a rat do not (Lacey et al., 1990).  $\beta$ -adrenoceptors are also expressed in the liver where they play a critical role in the regulation of blood sugar and glycogen metabolism. Adrenaline and  $\beta$ -AR are the most important mediators of glycogen regulation (Nonogaki, 2000).  $\beta_2$ -adrenoceptor stimulation induces glycogen breakdown and

accelerates the rate of glycolysis. Moreover, it also stimulates a glucagon release (Shulman, 1999). Effects of  $\beta$ -agonist on liver beyond the glycogen regulation have been described.  $\beta_2$ -stimulation was shown to be protective against hepatocyte apoptosis; an effect that was mitigated by  $\beta_2$ -AR inhibition (Andre et al., 1999).  $\beta_2$ -agonists were also shown to improve muscle glucose transport in rat model of obesity and diabetes (Pan et al., 2001). The  $\beta_2$ -AR response has been showed to be critical in mediating hypoglycaemia awareness. Defective tissue response to glucagon and adrenaline is responsible for this syndrome leading to  $\beta_2$ -AR insensitivity. After a careful avoidance of hypoglycaemia, the restored sympathetic response occurs primarily by improved  $\beta_2$ -AR sensitivity (Fritsche et al., 2000).  $\beta_2$ -adrenoceptors have been also proved to play a role in mediating a parathyroid hormone action on bone formation. Intermittent parathormone treatment increases bone mass formation, but for a proper functioning of this axis the presence of  $\beta_2$ -receptors is critically important (Hanyu et al., 2012). Recently, it has been found that  $\beta_2$ -adrenoceptors (through both Gs-PKA and  $\beta$ -arrestin pathways) trigger DNA damage and suppress p53 levels and thus mediate the effect of chronic stress (Hara et al., 2011).

Adipocytes are another cell population largely influenced by  $\beta$ -adrenoceptors.  $\beta_3$ -adrenoceptors have been identified as critical receptor system involved in mediating adrenergic action upon adipocytes (Lipworth, 1996). Among other,  $\beta_3$ -adrenoceptors are playing a dominant role in regulating the release of leptin from the adipocytes. Noradrenaline and  $\beta_3$ -agonists inhibit a leptin release out of the adipocytes (Nonogaki, 2000).

#### **1.4.2. Muscarinic receptors**

Similarly to  $\beta$ -adrenoceptors, muscarinic receptors (MR) are metabotropic receptors that are coupled to G proteins. There are five subtypes of MR, which are distinct gene products. Analogously to  $\beta$ -adrenoceptors, recent research on MR signaling revealed the amazing complexity of signaling pathways originating on MR (Nathanson, 2000). However, according to their prevailing signaling pattern, muscarinic receptors can be divided into two major classes: odd numbered subtypes ( $M_{1-}$ ,  $M_{3-}$ , and  $M_{5-}$ -receptors) and even numbered subtypes ( $M_{2-}$  and  $M_{4-}$ -receptors). Odd numbered subtypes generally activate phospholipase C (PLC), employing pertussis toxin-insensitive G proteins of the Gq superfamily, and do not inhibit adenylyl cyclase. Even numbered subtypes inhibit adenylyl cyclase through pertussis toxin-sensitive G proteins of the Gi family. However, both  $M_{2-}$  and  $M_{4-}$ -MR can potentially activate PLC when expressed at high levels in certain cell types due to the release of  $\beta\gamma$  subunits (Katz et al., 1992). Under certain circumstances, both Gi and Gq-specific MR can

be potentially coupled to Gs and stimulate adenylyl cyclase activity (Migeon and Nathanson, 1994) but it is not the only way how MR can activate adenylyl cyclase. By increasing intracellular calcium levels, Gq-coupled MR can activate calmodulin-sensitive adenylyl cyclase and thus increase the level of cAMP (Choi et al., 1992). Moreover, various isoforms of adenylyl cyclases can be phosphorylated (and thus regulated) by protein kinase A (PKA), protein kinase C (PKC) and calmodulin-dependent protein kinases (CaMKs) (Hurley, 1999). Last but not least, MR can regulate tyrosine kinase pathway and mitogen-activated protein kinase (MAPK) pathway employing multiple mechanisms. Gq-coupled receptors can activate proline-rich tyrosine kinase 2 (PYK2) through calmodulin-dependent protein kinases and PKC-mediated mechanism. PYK2 can activate c-src to cause the activation of the ERK pathway (Gutkind, 1998, Gudermann et al., 2000).

Muscarinic receptors mediate a plenty of physiological and pathophysiological processes throughout the body. In the central nervous system, all subtypes of muscarinic receptors are involved. M<sub>1</sub>-MR (Yamasaki et al., 2010), M<sub>2</sub>-MR and M<sub>4</sub>-MR (Ohno et al., 1994, Tzavara et al., 2003) have been implicated in attention, arousal and cognitive processes. M<sub>2</sub>-MR mediate tremor, hypothermia and analgesia (Gomez et al., 2001). M<sub>3</sub>-MR are involved in certain types of learning (Poulin et al., 2010) and are required for the proper functioning of hypothalamic GHRH neurons (Gautam et al., 2009). M<sub>4</sub>-MR are involved in the inhibitory control of striatal projecting neurons, where they control D<sub>1</sub>-dopamine receptor-mediated locomotor stimulation (Gomez et al., 2001). M<sub>5</sub>-MR are likely involved in the acetylcholine-mediated vasodilatation of cerebral arteries and arterioles. Moreover, M<sub>5</sub>-MR are expressed in the ventral tegmental area where they are involved in mesolimbic dopaminergic pathway for the regulation of morphine reward and withdrawal symptoms (Yamada et al., 2003). The role of muscarinic receptors in cardiovascular processes will be thoroughly discussed later. M<sub>3</sub>-MR have been found to mediate a contraction of urinary bladder and selective M<sub>3</sub> antagonists are employed therapeutically to treat an overactive bladder syndrome (Chapple, 2004). The role of muscarinic receptors has been shown in mediating many pathophysiological processes including an involvement in cancer growth. M<sub>2</sub>-MR have been shown to inhibit the growth of glioblastoma cell lines (Ferretti et al., 2012), M<sub>1</sub>- and M<sub>3</sub>-MR antagonist has been shown to inhibit a growth of non-small cell lung cancer cells (Hua et al., 2012). M<sub>3</sub>-MR were also shown to play a role in proliferation and perineural invasion of cholangiocarcinoma cells (Feng et al., 2012).

## **1.5. G protein coupled receptors in the heart**

### **1.5.1. $\alpha$ -adrenoceptors**

The major subtype playing the role in the heart is  $\alpha_1$ -subtype. In healthy human myocardium,  $\alpha_1$ -adrenoceptors constitute about 10% of total adrenoceptor binding with  $\alpha_{1A}$  and  $\alpha_{1B}$  being the predominant subtypes (Jensen et al., 2009a) whereas  $\alpha_{1D}$  is the predominant subtype in epicardial coronary arteries and smooth muscle cells (Jensen et al., 2009b).

The important feature concerning both  $\alpha_{1A}$  and  $\alpha_{1B}$  subtypes is that the amount of their binding sites is not diminished in heart failure both in left and right ventricles and their function in this conditions is maintained. This is in contrast with  $\beta$ -adrenoceptors that are down-regulated and also functionally uncoupled in heart failure (Jensen et al., 2009a) leading to diminished functional activation of potentially beneficial pathways including those mediating a positive inotropy.

### **1.5.2. $\beta_1$ -adrenoceptors**

$\beta_1$ -AR constitute about 70-80% of all  $\beta$ -AR in the heart (Brodde, 1991). They are localized widely through cardiomyocytes without particular local inhomogenities. When stimulated, they elicit full contractile response. Upon stimulation, Gs protein is activated and further activates adenylyl cyclase (AC). AC converts adenosinotriphosphate (ATP) into cyclic AMP (cAMP). cAMP further activates protein kinase A (PKA) resulting in phosphorylation of multiple proteins including those involved in coupling between excitation and contraction. This classical paradigm is true for only intermittent or short-term  $\beta_1$ -AR activation. Prolonged  $\beta_1$ -AR stimulation leads to G protein receptor kinase (GRK)-mediated receptor phosphorylation, which leads to the recruitment of  $\beta$ -arrestin to phosphorylated  $\beta_1$ -AR.  $\beta$ -arrestin plays two different roles in  $\beta_1$ -AR signaling: it works as a sterical inhibitor that blocks further interaction between the receptor and G protein and targets the receptor for internalization (Lefkowitz, 1998). Second, it serves as a scaffold protein that enables the interaction of other members of signaling cascade. Prolonged  $\beta_1$ -AR activation leads to the activation of calmodulin activated protein kinase II (CaMKII) that is responsible for clinically unfavorable effects of  $\beta_1$ -AR activation (i.e. increased apoptosis) that is observed in untreated heart failure (Yoo et al., 2009). The exact mechanism how  $\beta_1$ -AR is linked to CaMKII is still not completely resolved; however, the recent work has showed that  $\beta$ -arrestin ( $\beta$ -ARR) is required as a scaffold protein for an aggregation of CaMKII and Epac (exchange protein directly activated by cAMP) bringing them to the close proximity to cAMP generation site. Upon stimulation by cAMP, Epac exhibits guanine nucleotide exchange factor activity

toward Rap1. The exact mechanism how Epac activates CaMKII remains unknown but it seems to require phospholipase- $\epsilon$  (PLC- $\epsilon$ ) and protein kinase- $\epsilon$  (PKC- $\epsilon$ ). As a result, CaMKII is activated and transmits the signal further in the cell (Mangmool et al., 2010). However, signaling through  $\beta_1$ -AR can have a protective role as well as it has been showed that  $\beta_1$ -AR can also signal through  $\beta_1$ -arrestin that further leads to matrix-metalloproteinase (MMP)-mediated cleavage and extracellular shedding of heparin-binding epidermal growth factor (HB-EGF) that ultimately leads to EGFR transactivation (Noma et al., 2007). This pathway requires specific G protein-coupled receptor kinases 5 and 6. It has been showed that this pathway confers cardioprotection.

### **1.5.3. $\beta_2$ -adrenoceptors**

Compared with  $\beta_1$ -AR,  $\beta_2$ -AR constitute only about 20-30% of total  $\beta$ -adrenoceptors in the heart (Brodde, 1991). There are multiple aspects of how  $\beta_2$ -signaling differs from  $\beta_1$ -signaling. First,  $\beta_2$ -ARs are not uniformly distributed throughout the cardiomyocyte, but they are enriched in caveolae with particular spatial distribution. Second, although  $\beta_2$ -AR couple to Gs protein (which is similar to  $\beta_1$ -AR signaling), they are also able to couple to Gi, which elicits the contrary effect.  $\beta_2$ -AR coupling to Gi is preceded by PKA-mediated phosphorylation of  $\beta_2$ -AR. The effect of Gi can possibly occur since prolonged  $\beta_2$ -AR stimulation induces the translocation of  $\beta_2$ -AR out of caveolae. Moreover, prolonged agonist stimulation activates PI3-kinase and functionally further inhibits Gs. It has been proposed that  $\beta_2$ -adrenoceptor signaling hinders apoptosis and functionally oppose  $\beta_1$ -signaling. Furthermore, only  $\beta_1$ -AR and not  $\beta_2$ -AR can activate CaMKII. The reason for this lies in the different amino-acid sequence of the C-terminal tail of the receptor, which has been showed in chimera studies (Mangmool et al., 2010). C-tail of  $\beta_1$ -AR mediates a unique conformation of  $\beta$ -arrestin that allows for scaffolding of Epac and CaMKII to form a stable complex, leading to subsequent cAMP/Epac-mediate activation of CaMKII. Both  $\beta_1$ -AR and  $\beta_2$ -AR work usually in a complementary manner; however, under certain circumstances their function might be opposing. It has been showed that the stimulation of  $\beta_1$ -adrenoceptors promotes apoptosis (Communal et al., 1999, Zaugg et al., 2000, Zhu et al., 2003) whereas the stimulation of  $\beta_2$ -adrenoceptors has anti-apoptotic effects both in vitro (Communal et al., 1999) and in vivo (Chesley et al., 2000). These findings might have a therapeutic potential in treatment of cardiac disorders (e.g. heart failure).

#### 1.5.4. $\beta_3$ - adrenoceptors

Challenge about the existence of another  $\beta$ -AR came at the end of the last century, when the 3<sup>rd</sup> subtype of  $\beta$ -adrenoceptor ( $\beta_3$ -AR) was cloned (Emorine et al., 1989). Messenger RNA (mRNA) of this receptor was identified in various tissues (adipose tissue, small and large intestine, gall and urinary bladder and myometrium) (Rouget et al., 2005) including the heart (Gauthier et al., 2000, Myslivecek et al., 2006). The  $\beta_3$ -AR protein has been also identified using immunohistochemical detection (De Matteis et al., 2002).

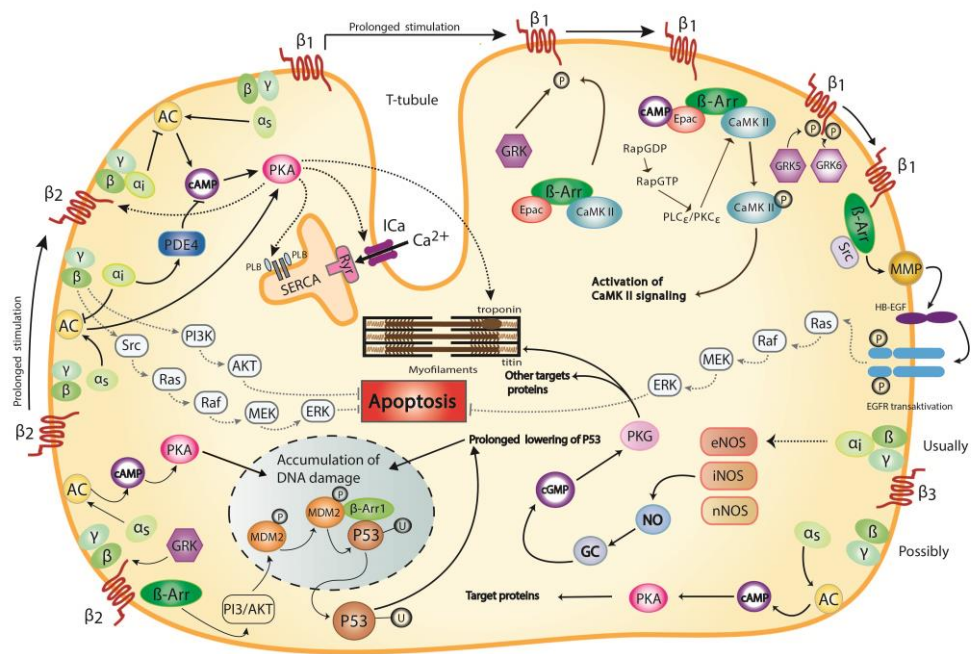
The activation of  $\beta_3$ -AR can produce different effects depending upon the heart chamber-atria vs. ventricles (Brixius et al., 2004, Spasojevic et al., 2011) and upon species (Gauthier et al., 2000, Skeberdis et al., 2008). In human ventricles, stimulation of the  $\beta_3$ -AR produces a marked decrease in cardiac contractility (Gauthier et al., 1996), which is in contrast to the other  $\beta$ -AR subtypes. It has been supposed that  $\beta_3$ -AR serves as a „brake“ during excessively increased sympathetic overstimulation since it is activated at high catecholamine concentrations and counteracts the effects of  $\beta_1$  and  $\beta_2$  activation by producing a negative inotropic effect (Moens et al., 2010). On the other hand, it has been shown that in human atria  $\beta_3$ -AR stimulation increases contractility (Skeberdis et al., 2008). It has been observed as well that the expression of  $\beta_3$ -AR is increased in the left ventricles of patients with heart failure (Moniotte et al., 2001) suggesting the role of  $\beta_3$ -AR up-regulation in pathological conditions.  $\beta_3$ -AR is unique since it can couple both to Gs and Gi without the requirement for receptor phosphorylation (Liggett et al., 1993, Soeder et al., 1999). Compared to  $\beta_1$ - and  $\beta_2$ -AR, the  $\beta_3$ -AR presents a relative in vitro and in vivo lack of desensitization following activation with agonists, which is supposed to be due to the lack of the phosphorylation sites for protein kinase A and  $\beta$ -adrenoceptor kinase ( $\beta$ -ARK) that are critical players for mediating receptor desensitization (Nantel et al., 1993). However, most recent work suggests that in human embryonic kidney (HEK) cells,  $\beta_3$ -adrenoceptor desensitization occurs, but it does not involve reduced receptor numbers or significant reduction of Gs expression. Instead, its mechanism seems to be changes at the level of adenylyl cyclase function (Michel-Reher and Michel, 2013). In human ventricles, the activation of  $\beta_3$ -AR causes an activation of the nitric oxide (NO) pathway. It has been thought long that the NOS isoform involved is endothelial NO synthase (eNOS) (Brixius et al., 2004) that has a crucial role in regulating L-type calcium channel activity in cardiac tissue. Protein-kinase G (PKG) is a downstream effector of the eNOS/NO signaling cascade; it phosphorylates L-type  $\text{Ca}^{2+}$  channels, which deactivates them (van der Heyden et al., 2005). However, recent studies have showed that eNOS is not the only



NO isoform that mediates signaling through  $\beta_3$ -AR as NO production via the neuronal NOS (nNOS) has been demonstrated (Amour et al., 2007). The role of inducible NOS (iNOS) has not been fully clarified yet but it is likely that NO production through this isoform may also contribute to NO production followed by  $\beta_3$ -AR activation (Maffei et al., 2007).

#### **1.5.5. $\beta_4$ -adrenoceptors**

At the end of 20<sup>th</sup> century, a fourth  $\beta$ -adrenoceptor was described ( $\beta_4$ -AR) (Kaumann and Molenaar, 1997) but later it was found out that this  $\beta_4$ -AR is only a low affinity state of  $\beta_1$ -AR (Granneman, 2001, Kaumann et al., 2001). However, some studies have shown a role of this receptor both in cardiac (Myslivecek et al., 2003) and vascular physiology (Malleme et al., 2004)- this receptor was found to mediate vasodilatation in normotensive but not hypertensive rats, which might have therapeutic consequences. The current concept of signaling through  $\beta$ -receptors is in Fig. 1.5.1



In non-cardiomyocytes

### **Fig. 1.5.1. The current concept of signaling through $\beta$ -receptors**

The predominant signaling through  $\beta_1$ -AR occurs through activation of Gs protein and subsequent activation of adenylyl cyclase (AC) that activates protein kinase A. Protein kinase A then phosphorylates its target proteins including calcium channel, phospholamban (PLB) and troponins. Upon persistent stimulation, G protein receptor kinase (GRK) is recruited leading to  $\beta_1$ -AR receptor phosphorylation, which results in  $\beta$ -arrestin ( $\beta$ -ARR) recruitment. This allows a conformation change to occur in  $\beta$ -arrestin molecule enabling stable interaction with calmodulin-activated protein kinase II (CaMKII) and exchange protein directly activated by cAMP (Epac).  $\beta$ -arrestin-mediated translocation of CaMKII and Epac to the  $\beta_1$ -AR complex bring these molecules in close proximity to the location of cAMP generation by AC. cAMP directly binds to and stimulates Epac, which leads to CaMKII activation via mechanism that requires Rap (Ras-related GTP binding protein), PLC $\epsilon$  and PKC $\epsilon$ , ultimately leading to phosphorylation of downstream effectors. Activation of CaMKII plays a role in various cardiac pathologic states and is considered to be maladaptive. Under different circumstances, persistent stimulation of  $\beta_1$ -AR leads to the receptor phosphorylation by specific GRK: GRK5 and GRK6, which leads to  $\beta_1$ -AR phosphorylation and recruitment of  $\beta$ -arrestin.  $\beta$ -arrestin further recruits Src (non receptor protein tyrosin kinase) that leads to matrix metalloproteinase activation (MMP) activation and cleavage of heparin bound-epidermal growth factor (HB-EGF). This leads to its binding to the respective receptor (EGF-R) and its transactivation. Downstream signaling cascade is activated eliciting cardioprotective action (e.g. apoptosis inhibition).

Signaling through  $\beta_2$ -AR predominantly occurs via Gs protein as well, leading to cAMP formation and PKA activation. However, upon persistent stimulation, PKA phosphorylates  $\beta_2$ -receptor inducing its coupling with Gi protein. Signaling through Gi protein directly inhibits the activation of AC; moreover, signaling through  $\beta\gamma$  subunits leads to the activation of PI3-kinase pathway and Src pathway mediating cardioprotective action (e.g. apoptosis inhibition).

In non-cardiomyocyte cells (germline cells), chronic activation of  $\beta_2$ -AR leads to the recruitment of  $\beta$ -arrestin 1, which facilitates Akt-mediated activation of MDM<sub>2</sub> (mouse double minute 2 homolog, also known as E3 ubiquitin-protein ligase). MDM<sub>2</sub> targets p53, leading to its degradation. In this process,  $\beta$ -arrestin-1 acts as a molecular scaffold. Prolonged lowering of p53 then leads to DNA damage accumulation.

$\beta_3$ -AR signaling was believed to occur through Gi protein activation subsequently leading to the NO synthase (NOS) activation. All isoforms (iNOS, eNOS, nNOS) are likely to be

involved. NOS activation leads to NO production with the subsequent activation of protein kinase G (PKG) that phosphorylates its target proteins (incl. titin). The data from this thesis support the hypothesis that  $\beta_3$ -AR signaling occurs also through Gs protein leading to PKA activation.

### **1.5.6. Muscarinic receptors**

In the heart, the dominant muscarinic receptor (MR) subtype is  $M_2$ . When stimulated,  $M_2$ -receptors activate Gi protein that inhibits adenylyl cyclase and thus counteracts cAMP-PKA-dependent signaling pathways. Gi also directly increases the activity of potassium channel, which results in hyperpolarization of the sinoatrial and atrioventricular node (Szabo and Otero, 1989). Moreover,  $\beta\gamma$  subunits directly activate the potassium channel operated by muscarinic receptor (Logothetis et al., 1987). Additionally, the stimulation of  $M_2$ -receptors can inhibit L-type calcium channel and thus directly reduce contractility of ventricular cardiomyocytes (Valenzuela et al., 1997). As a consequence, the stimulation of  $M_2$ -receptors markedly slows the heart, shortens atrial action potentials, increases smooth muscle contraction, inhibits the funny current ( $I_f$ , mixed sodium-potassium current responsible for starting the diastolic depolarization phase) and directly decreases contractility (Olshansky et al., 2008). The disruption of  $M_2$ -mediated signaling may have adverse consequences like heart failure (Zhang et al., 2002) or arrhythmias (Baba et al., 2004). The nature of non- $M_2$ -subtypes in the myocardium is still a matter of debate and the results published so far are contradictory. Binding studies performed on rats suggested that the minor subtype comprise of  $M_1$ - and  $M_5$ -subtypes (Myslivecek et al., 2008). However, other studies suggested that the major non- $M_2$ -subtypes were  $M_1$ ,  $M_3$  and  $M_4$  (Wang et al., 2004, Li et al., 2010).  $M_3$ -receptors were showed to mediate cardioprotective role of acetylcholine on myocardium; they were showed to ameliorate ischaemia-induced arrhythmias (Wang et al., 2012) and decrease the infarct size and mortality rate employing the  $M_3$ /NF- $\kappa$ B/miR-376b-5p/BDNF axis (Pan et al., 2012).

### **1.6. Regulation of G protein-coupled receptors**

The regulation of G protein-coupled receptors is very complex with many new findings emerging every year. Principally, receptor function can be regulated either on the level of their number present at the membrane or they can be regulated through post-translational modifications (e.g. phosphorylation). Signaling events induced by a certain receptor can lead to the regulation of the same receptor (i.e. homologous regulation) or a different receptor (i.e. heterologous regulation, cross-talk, cross-regulation).

### **1.6.1. Phosphorylation-dependent regulation of G protein-coupled receptors**

G proteins can be phosphorylated either by so-called G protein receptor-coupled kinases (GRK) or second messenger-dependent kinases (most often protein kinase A and protein kinase C, PKA and PKC, respectively). Both kinase systems phosphorylate serine and/or threonine residues of G protein-coupled receptors (GPCR), but these phosphorylation sites are different for each of them. Even among GRK, different GRK phosphorylate GPCR on different sites, which ultimately leads to the employment of different downstream signaling pathways.

The family of GRK consists of 7 homologs. GRK 1 and 7 phosphorylate exclusively visual opsins; on the contrary, GRK 2-6 phosphorylate various members of GPCR as well as other surface receptors (Lefkowitz and Shenoy, 2005). The important feature of GRK phosphorylation is that GRK phosphorylate only receptors with bound agonists. When stimulated by an agonist, GPCR undergo conformational changes and expose their binding sites for heterotrimeric G proteins.  $G\alpha$  subunits of G proteins exchange GTP for GDP leading to G protein activation and downstream signaling. Moreover, receptors with bound agonists immediately become a substrate for GRK-mediated phosphorylation, which leads to the recruitment of the cytosolic adaptor proteins called  $\beta$ -arrestins. The family of arrestins comprises 4 members (Lefkowitz, 1998, Gurevich and Gurevich, 2006). Traditionally, the main function of  $\beta$ -arrestin was to mediate the receptor desensitization but the knowledge about the myriad of arrestin roles has expanded enormously in recent years (Ibrahim and Kurose, 2012, Hu et al., 2013, Kang et al., 2013). Generally, besides desensitization,  $\beta$ -arrestins mediate endocytosis and signaling. The endocytosis requires the involvement of clathrin and  $\beta$ -arrestin-clathrin interaction. Moreover, studies conducted in recent years revealed that additional binding partners are necessary for efficient receptor internalization; these include the clathrin adaptor protein 2 (AP2), N-ethylmaleimide-sensitive fusion protein (NSF), the small G protein ARF6 and PI4P kinase (Nelson et al., 2008).  $\beta$ -arrestins also mediate scaffolding of the signaling kinases including MAP kinase, MAP kinase kinase and MAP kinase kinase kinase, but also Src and CaMKII (Engelhardt, 2007, Noma et al., 2007, Mangmool et al., 2010).

Prolonged desensitization of GPCR leads to receptor internalization, which is a process when phosphorylated arrestin-bound receptors are being removed from the plasma membrane to a membrane-associated intracellular compartment. The total amount of binding sites on the whole-cell level is unchanged, just the membrane fraction decreases (Tomankova and Myslivecek, 2011). Following internalization, GPCR can undergo different pathways. They

can be either lysosomally degraded (which leads to a diminished amount of binding sites; termed receptor down-regulation), can be recycled back to the plasma membrane (resensitization).  $\beta$ -arrestins can undergo a large variety of modifications that influence their subsequent signaling role. These modifications comprise phosphorylation, ubiquitination and S-nitrosylation. In terms of GPCR regulation, ubiquitination is of particular interest since an ubiquitinated  $\beta$ -arrestin can form a stable complex with GPCR that co-internalize into endocytotic vesicles. These complexes can be associated with activated protein kinase (e.g. MAPK) and form a so called „signalosomes“ spreading the signal further into the cell (Lefkowitz and Shenoy, 2005).

When a cell is facing a situation that a receptor is not adequately stimulated, the receptor can undergo a process of sensitization and up-regulation. Moreover, further class of proteins regulating the function of G-proteins was discovered- regulator of G protein signaling (RGS) proteins (Willars, 2006). These proteins work both as GTPase activating proteins (GAPs) (De Vries et al., 2000), but also regulate GPCR signaling independently of the GAP activity by modulation protein-protein interactions, protein translation or intracellular localization of signaling molecules (Abramow-Newerly et al., 2006). Some proteins of this family (RGS4, RGS6 etc.) play an important role in regulating cardiovascular function as well and dysfunction of these proteins may contribute to various pathological states including chronotropic incompetence or AV conduction block (Fu et al., 2007).

### **1.6.2. Phosphorylation-independent regulation of G protein-coupled receptors**

It has been also proved that besides the phosphorylation-dependent regulation of G protein coupled receptors, certain GRK promote a signaling that does not require a GPCR phosphorylation. This process has been studied the most for the interaction of mGlu1 receptor and GRK2. GRK2 directly binds to the second intracellular loop and the C-terminal tail of mGlu1a receptor but only to the second intracellular loop in the case of mGlu1b (Dhami et al., 2005).

### **1.7. The role of G protein-coupled receptors in autonomic nervous system**

The autonomic nervous system is the part of the peripheral nervous system that mediates an involuntary control over various mainly visceral organ and thus contributes significantly to the control of homeostasis. Among others, it exerts a control over the heart rate, blood pressure, the response of gastrointestinal tract to food, thermoregulation, contraction of the urinary bladder, focusing of the eyes and sexual arousal. Similarly to other parts of the

nervous system, autonomic nervous system has an afferent, central (integrative) and efferent part. It consists of two main branches: sympathetic and parasympathetic (McCorry, 2007). The enteric nervous system is also sometimes considered the part of the autonomic nervous system but sometimes it is considered to be independent. Sympathetic nervous system mediates a body response to a possibly dangerous event; it can be characterized as quick response mobilizing system. On the contrary, parasympathetic nervous system is activated more slowly and is active when the organism is relaxed. Usually, both systems work in opposition (e.g. in the cardiovascular system) but they may also work synergically (e.g. during sexual arousal and orgasm, wherein both play a role). Noradrenaline is the main mediator of sympathetic nervous system and acetylcholine is the main mediator of parasympathetic nervous system. The effect of both systems on the cardiovascular function occurs via G protein coupled receptors ( $\beta$ - adrenoceptors for sympathetic nervous system and muscarinic receptors for parasympathetic nervous system). Both systems exert a tonic influence and the resulting function (heart rate, blood pressure etc.) is the result of the net balance between the systems (McCorry, 2007). It should be mentioned that the proper function and balance between both systems is important in human physiology and disturbance in both sympathetic and parasympathetic function play a role in many disorders including high blood pressure and heart failure (Port and Bristow, 2001, Olshansky et al., 2008), which has important clinical implications.

### **1.8. The role of stress**

Stress is a response of the body to inner or outer strain that is able to elicit a response characterized by stress response elements (i.e. by the pathways and their activation that is evoked by a stressor). The responses to stressors are considered as specific. Therefore, every response that consists of a part of the specific pathway could be understood as stress response element. The main stress response elements are: specific activation of prefrontal/orbitofrontal cortex and limbic system, activity of neurons in nucleus tractus solitarii (NTS), activation of catecholaminergic groups in brainstem/medulla) and can have both positive (eustress) or negative impact (distress) on the body. Since stress provokes a distinct response of the organism, it is frequently used in research. Stressors can be divided according to their nature into physical, chemical, psychological, social, stressors disturbing cardiovascular/metabolic homeostatis and those affecting multiple systems (Kvetnansky et al., 2009).

We have used cold stress in our experiments since it is natural event (compared with immobilization or myocardial infarction), can be very well dosed and measured and can be used over a longer period of time thus enabling to investigate the effect of prolonged stress.

### **1.9. The importance of gene-knockout organisms for studying physiologic functions**

In the few past decades, the techniques of molecular genetics enabled to prepare mice where a particular gene from the genome was selectively suppressed - so called gene „knock-outs“. The particular gene can be switched off either on the whole-body level or selectively in a particular organ (i.e. the heart or the central nervous system). Knock-out studies thus provide a valuable insight into the function of many genes. It has been found that organisms may be viable although they have a knock-out of a gene that has been thought to be indispensable for survival. For example, animals with whole-body (Hrabovska et al., 2010) as well as CNS specific (Farar et al., 2012) knock-out of acetylcholinesterase (key enzyme degrading acetylcholine in the synapse) are viable and even (as in the case of CNS specific knock-out) have only mild functional impairment. This observation is in the striking contrast with the clinical experience of organophosphate poisoning. If acetylcholinesterase is inhibited postnatally (through an application of an organophosphate), such treatment is not tolerated and is always lethal. Moreover, even genes encoding proteins with elementary signaling function have been proved not to be necessary for survival as mice with whole-body c-fos knock out have been prepared (Johnson et al., 1992). However, the absence of such an important gene leads to severe functional impairment (Paylor et al., 1994). The experiments with gene knock-outs are facing many challenges too, though. First, due to functional impairment, it is sometimes challenging to breed the genetic line and maintain the population (Fritz and Robertson, 1996). Second, a severe genetic hit can trigger multiple adaptation changes that do not have to be easily interpreted (Benes et al., 2012a, Benes et al., 2013). One have to be cautious when using gene knock-out as a disease model as the possible adaptation changes induced by the knock-out can distort the role of the simulated disease itself (Sharma et al., 2004). Despite these limitations, knock-out models can provide a valuable tool to study the gene function. Therefore, main knock-out models related to adrenoceptors and muscarinic receptors will be briefly discussed below.

#### **1.9.1. $\beta_1$ -knockout mice**

Mice without functional  $\beta_1$ -adrenoceptors display a severe prenatal lethality (lethality rate about 90%) (Rohrer et al., 1996). Those surviving to adulthood have a grossly normal cardiac



morphology with similar heart weight/body weight ratios and similar myocardial catecholamine content.  $\beta_1$ KO mice have also slightly but significantly diminished binding of myocardial  $\beta_2$ -receptors. The activity of adenylyl cyclase is similar both under basal and forskolin-stimulated conditions. When measured for a short time, both heart rate and blood pressure were normal (Rohrer et al., 1996). However, the long-term heart rate measurement revealed that  $\beta_1$ KO mice have actually a decreased heart rate. A spectral analysis of heart rate variability measured as RR intervals directly from ECG tracing revealed that an absence of  $\beta_1$ -adrenoceptors increases a total autonomic variability through both time and frequency indexes (Ecker et al., 2006). When these mice were challenged by non-specific  $\beta$ -agonist isoprenaline, the increase in heart rate that is normally seen in WT mice was absent in KO. On the contrary,  $\beta_2$ -mediated hypotensive response was preserved both in WT and KO (Rohrer et al., 1996).  $\beta_1$ -receptors are also responsible for mediating a sympathetic nervous system mediated adaptive thermogenesis as  $\beta_1$ KO mice develop a hypothermia when challenged with a cold-stress. These mice are unable to perform a brown adipose tissue response upon the stimulation with dobutamine (selective  $\beta_1$ -agonist) or noradrenaline (Ueta et al., 2012).

### **1.9.2. $\beta_2$ -knockout mice**

Contrary to  $\beta_1$ KO mice, lacking of  $\beta_2$ -AR is not associated with prenatal lethality (Chruscinski et al., 1999). Adult  $\beta_2$ KO mice are normal and fully fertile. They have a normal resting heart rate and blood pressure and normal chronotropic response to the  $\beta$ -AR agonist isoprenaline (both  $\beta_1$  and  $\beta_2$  agonist). However, the hypotensive effect of isoprenaline is significantly attenuated in  $\beta_2$ KO mice. On the contrary, they have a more pronounced hypertensive response to adrenaline (Chruscinski et al., 1999). The spectral analysis of heart rate variability (HRV) revealed that  $\beta_2$ -receptors play (together with  $\beta_1$ -receptors) a role in mediating sympathetic regulation of HRV. The absence of  $\beta_2$ -receptors increases total autonomic variability through the frequency-domain indexes, total power and ultra-low frequency power (Ecker et al., 2006). When challenged with exercise test,  $\beta_2$ KO mice have higher total exercise capacity but their blood pressure during exercise is markedly higher. Finally, at comparable workload,  $\beta_2$ KO mice have a lower respiratory exchange ratio suggesting a difference in energy metabolism (Chruscinski et al., 1999).

### **1.9.3. $\beta_3$ -knockout mice**

$\beta_3$ KO mice have a similar resting heart rate as WT mice but smaller left ventricle size and impaired LV relaxation (Varghese et al., 2000). When challenged with isoprenaline,  $\beta_3$ KO

mice are able to increase their heart rate to the similar extend as WT but they have much higher contractile response (Varghese et al., 2000). On the contrary, the cardioprotective properties are revealed during stress condition. In WT animals, nebivolol (clinically used  $\beta$ -blocker with  $\beta_3$ -agonistic effects) and CL316243 (specific  $\beta_3$ -agonist) were able to induce a cardioprotection during ischaemia/reperfusion, which was abolished in mice lacking  $\beta_3$ -adrenoceptor strongly suggesting a cardioprotective role of  $\beta_3$ -adrenoceptors (Aragon et al., 2011).

#### **1.9.4. $\beta_1/\beta_2$ -knockout mice**

Similarly to  $\beta_1$ -knockouts,  $\beta_1/\beta_2$ -knockout mice suffer with severe prenatal lethality (90-95%). Surviving  $\beta_1/\beta_2$ -knockout mice are not different both in baseline heart rate or mean arterial blood pressure (Rohrer et al., 1999). However, similarly to  $\beta_1$ KO, when monitored over a longer time period,  $\beta_1/\beta_2$ KO mice show a lower baseline heart rate and heart rate variability changes both in time and frequency domain correlating with effect of both  $\beta_1$  and  $\beta_2$ -receptors (Ecker et al., 2006). Upon a challenge with isoprenaline,  $\beta_1/\beta_2$ KO mice in comparison to WT have markedly attenuated both chronotropic and hypotensive effect of the drug. Moreover,  $\beta_1/\beta_2$ KO mice have significantly lower maximal heart rate during exercise and altered oxygen consumption and carbon dioxide production.  $\beta_1/\beta_2$ KO mice showed a mild but significant reduction in muscarinic receptors density (Rohrer et al., 1999).

#### **1.9.5. $\beta_1/\beta_2/\beta_3$ -knockout mice**

$\beta_1/\beta_2/\beta_3$ -triple knockout (TKO) mice are fully viable without apparent cardiovascular disturbances. However, they have lower heart weight/body weight ratio with lower cardiomyocyte size indicating cardiac hypotrophy. Furthermore, a lower resting heart rate and lower ejection fraction was found. These changes are accompanied by diminished activity of sarcoplasmic  $\text{Ca}^{2+}$ -ATPase (SERCA 2a) despite the protein expressions of SERCA 2a, ryanodine receptor and  $\text{Na}^+/\text{Ca}^{2+}$ -exchanger were similar. The protein expression of phospholamban was increased but its PKA-dependent phosphorylation at serine 16 was absent and CaMKII-dependent phosphorylation at threonine 17 was decreased. Despite these profound changes, there was no significant functional impairment of  $\beta_1/\beta_2/\beta_3$ -triple knockout mice in the running wheel tests. On the contrary, the mean velocity of running and the total trip distance were actually higher in TKO mice (Lee et al., 2010). Further experiments on these mice revealed normophagic obesity and cold-intolerance. Their brown fat has an impaired morphology and diminished expression of an uncoupling protein-1 (UCP-1)

following a challenge with cold. On the contrary, these mice have higher circulating levels of free fatty acids and glycerol both at basal and fasted states suggesting enhanced lipolysis (Jimenez et al., 2002).

#### **1.9.6. M<sub>2</sub>-knockout mice**

Mice lacking the gene for M<sub>2</sub>-receptor were generated in Wess laboratory (Gomez et al., 1999) and further characterized in subsequent studies. The knock-out of M<sub>2</sub>-receptor was found to have impact both on central nervous system as well as cardiovascular system. M<sub>2</sub>-receptor subtype was proved to play a key role in mediating whole-body tremor, hypothermia and analgesia- three of the most striking central muscarinic effects (Gomez et al., 2001). M<sub>2</sub>-receptors were showed to be involved in learning and memory; M<sub>2</sub>KO mice showed significant deficits in behavioral flexibility and working memory in the Barnes circular maze and the T-maze tests. These behavioral deficits were associated with profound changes in neuronal plasticity (measured at Schaffer collateral-CA1 synapse) and both short-term and long-term potentiation (STP and LTP, respectively) were severely impaired in hippocampi (Seeger et al., 2004). A study (LaCroix et al., 2008) examined the cardiovascular function of M<sub>2</sub>KO mice and showed that in spite of fully comparable baseline characteristics, after a week of chronic isoprenaline infusion followed by the bolus injection of phenylephrine, M<sub>2</sub>KO mice exhibited impaired ventricular function indicated by attenuated dp/dt and increased end-diastolic pressure. Furthermore, chronic isoprenaline infusion significantly increased matrix metalloproteinase (MMP) activity (LaCroix et al., 2008). Authors conclude that the deficiency of M<sub>2</sub>-receptors increases susceptibility to chronic adrenergic stress. Despite M<sub>2</sub>KO mice were so thoroughly investigated, virtually no attention has been paid to the role of other cardiac receptors.

## 2. Goals and Hypotheses

Despite M<sub>2</sub>KO mice were already a matter of investigation (discussed in the previous section (LaCroix et al., 2008), (Seeger et al., 2004)), very little is known about the adaptation mechanisms that take place after the knock-out of M<sub>2</sub>-gene. Similarly, the way they cope with stress is not well understood either.

The goal of the work was to perform in-depth analysis of M<sub>2</sub>KO mice both at baseline and upon a challenge with a cold stress and to explore the role of opposing receptors i.e. adrenoceptors in adaptation to missing M<sub>2</sub>-receptors. We have focused on the receptor level, subsequent changes in intracellular signaling as well as on the consequences in physiological functions on the whole-organism level.

We have hypothesized that:

1. There would be a down-regulation of cardiostimulatory  $\beta_1$ - and  $\beta_2$ -adrenoceptors based on the down-regulation of the  $\beta_1$ - and  $\beta_2$ -adrenoceptor gene expression. Since  $\beta_3$ -adrenoceptors have been proved to mediate cardioinhibitory function we have assumed that these receptors would be one of the receptors employed to overtake the missing function of M<sub>2</sub>.
2. M<sub>4</sub>-receptors would possibly take over the function of missing M<sub>2</sub>. Other non-M<sub>2</sub>-subtypes would not be significantly altered.
3. There would be no difference in the systolic and diastolic function of M<sub>2</sub>KO hearts.
4. There would be no difference in the heart rate, activity and basal temperature.
5. There would be no difference in the level of circulating catecholamines and the activity of catecholamine synthesizing enzymes.
6. The condition of prolonged (one week) stress would lead to different adaptation responses in M<sub>2</sub>KO mice with distinct adjustment in cardiac signaling.

## Rationale

Ad 1) A number of previous studies showed that there was a need for the homeostatic balance between sympathetic and parasympathetic nerve tonus, which means there is also a need for the balance between the number of muscarinic receptors and  $\beta$ -adrenoceptors as well as their functional output (Myslivecek et al., 1996, Garofolo et al., 2002, Brodde and Leineweber, 2004, Stavrakis et al., 2011). Based upon these studies, we have hypothesized that cardiostimulatory  $\beta_1$ - and  $\beta_2$ -adrenoceptors would be down-regulated. Conversely, since  $\beta_3$ -adrenoceptors are supposed to mediate cardioinhibitory function we have assumed that these receptors would be either without change or up-regulated to compensate for the missing function of  $M_2$ -receptors.

Ad 2) Although muscarinic signaling is complex, even numbered subtypes ( $M_2$ - and  $M_4$ -receptors) inhibit adenylyl cyclase through pertussis toxin-sensitive G proteins of the  $G_i$  family. Therefore we have assumed that  $M_4$ -receptors would possibly take over the function of missing  $M_2$ -receptors.

Ad 3) Systolic and diastolic function was invasively assessed in the study of (LaCroix et al., 2008) and no difference between wild-type and  $M_2$ KO mice was found.

Ad 4) At any given time point, heart rate reflects the dynamic balance between sympathetic and parasympathetic branch of the autonomic nervous system. Previously published studies did not report a difference in heart rate between WT and  $M_2$ KO mice (LaCroix et al., 2008), so we have not expected to find a difference either. On the central level, muscarinic receptors mediate hypothermia.  $M_2$ -receptor deficient mice have attenuated hypothermic response to oxotremorine (Bymaster et al., 2001) but their basal temperature was reported not to be altered. Similarly,  $M_2$ KO mice were reported to have similar locomotor activity as WT littermates (Gomez et al., 2001), so we have assumed that these parameters would be similar in our study as well.

Ad 5) The already published studies showed that  $M_2$ KO mice had similar basal heart rate and locomotor activity, which means they do not seem to be under chronic stress at baseline. Therefore we have supposed that the deletion of  $M_2$ -receptors would not result in the

difference in the overall sympathetic tonus and the level of circulating catecholamines with the respective activities of the enzymes would be similar.

Ad 6) Upon 1-week infusion of isoprenaline and subsequent increased afterload (increase in the arterial pressure by the infusion of phenylephrine), M<sub>2</sub>KO mice did not show an increase in contractility, which means they are more susceptible to stress (LaCroix et al., 2008). These differences are likely to be accompanied by altered signaling.

At the beginning, we have performed binding study and gene expression study. Then we have moved into studying physiologic parameters on the whole organism level and performed echocardiographic analysis, long-term telemetric measurement of heart rate, activity and body temperature, analysis of heart rate variability and pharmacological study (with muscarinic and adrenergic agonist and antagonist: carbachol, atropine, isoprenaline, and propranolol). Finally, we have focused on signaling changes- we have made an analysis of NO-synthase and adenylyl cyclase activity, measurement of circulating catecholamines and analysis of gene expression of catecholamine-synthesizing enzymes.

In the following sections, the methods and results will be discussed in this order.

### 3. Material and Methods

#### 3.1. Experimental animals

Mice lacking the gene for M<sub>2</sub>-muscarinic receptor clone were generated in Wess laboratory (Gomez et al. 1999) and then bred in our animal facility (Prague, Czech Republic).

M<sub>2</sub><sup>-/-</sup> (M<sub>2</sub>KO) mice (i.e., mice with complete deletion of M<sub>2</sub>-muscarinic receptor gene; genetic background: 129 J1× CF1) were generated as described previously (Gomez et al. 1999). In all experiments, age-matched wild-type (WT) mice of the corresponding genetic background were used as controls (M<sub>2</sub><sup>+/+</sup>, 129 J1×CF1). Animals were treated in accordance with the legislature of the Czech Republic and the EU, and the experimental protocol was approved by the Committee for the Protection of Experimental Animals of the 1<sup>st</sup> Faculty of Medicine, Charles University, Prague. The animals were maintained under controlled environmental conditions (12/12 light/dark cycle, lights on at 6 AM). Food and water were available *ad libitum*. The average ambient light phase temperature±SEM was 24.45±0.4 °C; the average ambient dark phase temperature±SEM was 24.22±0.4 °C. The temperature in the experimental room was checked every 10 min using a thermometric chip (Maxim Integrated Products, Inc., Sunnyvale, CA, USA). Male M<sub>2</sub>KO animals and their WT counterparts (weighing 24–38 g, 11–16 weeks old) were used in the study. The animals were housed at three individuals per cage. At the end of the experiments, the animals were sacrificed by decapitation; hearts were collected, connective tissue was carefully discarded, and the left and right ventricles were isolated, flash frozen, and kept at -80 °C until the experiments were performed.

#### 3.2. Cold stress

The animals (WT and KO) were exposed to cold for 1 day (24 h) or 7 days in conditions simulating natural situation. In the biological night (from 8 p.m. to 8 a.m.) the animals were exposed to the temperature 5.91± 0.38°C, while in the biological day the temperature rose to 9.73± 0.59°C. The light/dark cycle was the same as in control animals. The temperature in the experimental space was checked every 10 min using thermometric chip (Maxim Integrated Products, Inc., Sunnyvale, CA, USA). Animals were housed three per cage. At the end of experiment, animals were sacrificed by decapitation; hearts were collected, connective tissue was carefully discarded, the left and right ventricles were isolated, flash frozen and kept in -80°C until the experiments were performed.

### 3.3. Receptor binding

We performed receptor binding on membranes, which were prepared as follows: the tissue was weighed and homogenized in ice-cold Tris–EDTA buffer (Tris–HCl 50 mmol/l, EDTA 2 mmol/l, pH adjusted to 7.4) for determination of total  $\beta$ -adrenoceptors and muscarinic receptors (Myslivecek et al., 2007). For determination of  $\beta_3$ -adrenoceptors the following buffer was used: Tris-HCl 50 mmol/l,  $MgCl_2$ -100mmol/l, GTP-300 $\mu$ mol/l, Tiron-1 $\mu$ mol/l, ascorbic acid-1mmol/l as described previously (Muzzin et al., 1994). In all cases, the homogenate was centrifuged at 600g for 10 min at 4 °C, the supernatant was collected and the sediment was resuspended in the buffer and centrifuged again using the same conditions. The supernatants were mixed and centrifuged for 25 min at 32,000g at 4 °C. The supernatant was discarded; the sediment was re-suspended in the buffer and centrifuged again at the same velocity and time (25 min, 32,000g, 4°C). The amount of receptor binding sites was determined in duplicate using the following conditions: (a) total  $\beta_1$ -adrenoceptor +  $\beta_2$ -adrenoceptor binding sites ( $B_{max}$ ): concentration range 93.75–3,000 pmol/l [ $^3$ H]-CGP 12177, non-specific binding was determined with 50 $\mu$ mol/l propranolol; (b)  $\beta_3$ -adrenoceptor binding sites ( $B_{max}$ ): concentration range 4.68 - 150 nmol/l [ $^3$ H]-SB 206606, non-specific binding was determined with 1 mmol/l SR59230A; (c) muscarinic receptors binding sites: 32.5–2,000 pmol/l [ $^3$ H]-QNB, non-specific binding was determined with 5 $\mu$ mol/l atropine. The incubation was performed at 38 °C and lasted for 60 min with [ $^3$ H]-CGP 12177, 30 minutes for [ $^3$ H]-SB 206606 and 120 min with [ $^3$ H]-QNB. All incubation times were adjusted previously (Myslivecek et al., 2007). The amount of binding sites ( $B_{max}$ ) per milligram of protein (determined using BCA, a method kit; Sigma, Czech Republic) and the affinity constant ( $K_D$ ) were computed by non-linear regression using GraphPad Prism 5.01 program (GraphPad Software, San Diego, CA, USA). Affinity constants ( $K_D$ ) were used for the “single-point” measurement in order to determine the number of receptors, saving the amount of tissue needed, using the following saturating concentrations of radioligands (2,000 pmol/l [ $^3$ H]-CGP 12177, 120 nmol/l [ $^3$ H]-SB 206606 and 1,500 pmol/l [ $^3$ H]-QNB).  $\beta_1$ - and  $\beta_2$ -adrenoceptors were determined using co-incubation of membranes with 400 pmol/l [ $^3$ H]-CGP 12177 and with subtype-selective antagonists in triplicates ( $10^{-7}$  mol/l CGP 20712A for  $\beta_1$ -adrenoceptors, or  $10^{-8}$  mol/l ICI 118.552 for  $\beta_2$ -adrenoceptors).

### 3.4. Receptor gene expression

Total mRNA was isolated using the chloroform–isopropanol (RNA Bee, TelTest, TX, USA) method according to the manufacturer’s instructions. RNA yield and integrity was evaluated



spectrophotometrically (Tecan M200 Nanoquant; detection of 2µl of sample in 16 wells at the moment) at  $A_1=260$  and  $A_1/A_2=260/280$  nm, respectively. Samples with  $A_1/A_2$  between 1.6 and 1.9 were used for downstream procedures. Total RNA was purified to eliminate potentially contaminating genomic DNA using recombinant DNase (Ambion). Purified mRNA was subsequently transcribed into cDNA using Ready-To-Go You-Prime First-Strand Beads (GE Healthcare). qPCR was performed using TaqMan probes (Applied Biosystems, probe numbers: M<sub>1</sub>: Mm01231010\_m1, M<sub>2</sub>: Mm01167087\_m1, M<sub>3</sub>: Mm01338409\_m1, M<sub>4</sub>: Mm01331561\_s1, M<sub>5</sub>: Mm01701883\_s1,  $\beta_1$ -AR: Mm00431701\_s1,  $\beta_2$ -AR: m02524224\_s1) with Roche qPCRMasternmix (Roche) using the following protocol: 2 min at 50 °C, 10 min at 95 °C followed by 50 cycles for 15 s at 95 °C and 1 min at 60 °C in a final volume of 10µl. In order to determine the relative gene expression, the  $\delta$ - $\delta$ -C<sub>t</sub> method was used. The levels of the analyzed transcripts were normalized to the level of the reference gene transcript (18SrRNA, the level of which was practically the same in WT and KO). Then, for each sample, the following formula was used:  $\text{normalized ratio} = 2^{CT_1 - CT_2}$ , where the CT<sub>1</sub> is the CT of the reference transcript and the CT<sub>2</sub> is the CT of the target transcript; CT stands for the cycle threshold. No template controls and non-reverse transcription reactions were performed as negative controls.

### 3.5. Echocardiography

Animals were anesthetized with i.m. injection of Zoletil® 100, Rometar® 2 % 5:1, diluted ten times, 3.2 ml/kg. Chest hair was shaved and ventricular thicknesses and dimensions were measured by an experienced echocardiographer using a 14-MHz probe (Vivid 7 ultrasound system, GE Healthcare, Fairfield, CT, USA). Interventricular septum thickness, posterior wall thickness, left ventricle end-systolic, end-diastolic diameters and cycle lengths were measured from parasternal short axis projection. M-mode tracings were obtained close to the papillary muscle level. The probe position was adjusted in order to obtain a perfectly round shape of the ventricle. Subsequently, probe position was changed to obtain an apical four-chamber view (A4C), from which mitral valve E and A wave velocities, E/A ratio, mitral valve deceleration time and slope were measured using pulsed wave Doppler tracing mode. Probe position was slightly rotated then to obtain LV outflow tract measurements. Maximal velocity in the left ventricle outflow tract and left ventricle ejection time were measured. During the whole examination, care was taken to avoid excessive pressure, which could cause bradycardia. Finally, measured parameters were used to calculate the heart rate, left ventricle fractional

shortening, left ventricle mass, and left ventricular peak velocity of circumferential fiber shortening according to previously published equations (Tanaka et al., 1996).

### **3.6. Telemetric assessment of heart rate**

The telemetry system used was commercially available from Mini Mitter (Respironics, Andover, MA, USA). The transponders (E-Mitter, G<sub>2</sub>-HR) were implanted in the peritoneal cavity under the same anesthesia as used in echocardiography (Zoletil® 100, Rometar® 2 % 5:1, diluted ten times, 3.2 ml/kg); mice were left for 1 week to recover from the surgery and then used in the experiment. The sensor leads were used for heart rate data acquisition; other parameters (temperature and activity) were acquired directly from the transponders. Receivers were connected in series and connected directly to the PC into a single computer port, allowing for the determination of all parameters. The heart rate (HR) data were collected every 10 s. VitalView, Chronos-fit, and GraphPad software were used for the evaluation of data.

### **3.7. Heart rate variability**

Spectral analysis of heart rate variability obtained from telemetric recordings (heart rate collected every 10s) in intact animals was performed. Heart rate spectra were obtained using Fourier transformation. Subsequently, the area under each curve (AUC) was calculated and differences in AUC between WT and KO were compared.

### **3.8. Telemetric assessment of body temperature and animal activity**

For picture completion of biorhythm changes and possible correlation of increased animal activity with heart rate, body temperature and animal activity were also assessed using the same telemetric system. The data were collected every 10 s in case of temperature and every 60 s in case of body activity.

### **3.9. Biorhythm analysis**

The data collected by telemetry were grouped into half-hour sequences, the mean values were calculated and used for further analysis. The analysis was performed using Chronos-fit program (Arraj and Lemmer, 2006), employing Fourier analysis and stepwise regression technique. Then, the data were transferred into GraphPad Prism 5.04 program (San Diego, CA, USA) for further statistical analysis.

### **3.10. Drug application**

Animals were held in separate cages; every single animal had its own cage with free access to food and water. We maintained the same regular light/dark regimen as stated before. The sensor had been implanted in general anesthesia (see “Telemetric assessment of heart rate” for details), and after 1 week of recovery period the experiment was started. The application route was intraperitoneal (i.p.) and approximately 100 $\mu$ l (according to the weight) was applied to each animal. All of the drugs were administered at the same daytime (9:30 AM to 9:45 AM) in order to minimize various influences of circadian heart rate variation. The telemetric system (discussed earlier) was used for heart rate monitoring. Carbachol (carbamoylcholine chloride, 0.5 mg/kg), isoprenaline (isoprenaline hydrochloride, 50 mg/kg), atropine (atropine monohydrate sulfate, 30 mg/kg), and propranolol (propranolol hydrochloride, 20 mg/kg) were dissolved in saline, and the same volume of saline was applied to control animals. The doses were derived from dose–response curves in WT animals from preliminary experiments.

### **3.11. Adenylyl cyclase activity (basal and forskolin-stimulated)**

We have measured the basal level and forskolin stimulated activity of adenylyl cyclase using modified method of (Hoffert et al., 2005). Membranes were incubated in HBSS buffer (Hank’s Stock Solutions, i.e. measurement of basal cAMP level) with or without presence of 40  $\mu$ mol/l forskolin (stimulated adenylyl cyclase activity) for 10 min at 37°C. The reaction was stopped by adding 0.2 N HCl. Then, the mixture was incubated for 20 min at room temperature, which was followed by centrifugation at 10,000g for 10 min. Supernatants were saved for measurement of cAMP and pellets were used to measure protein content (BCA assay, Sigma). cAMP content was measured using an enzyme immunoassay kit (Cayman Chemical) at  $\lambda$ = 413 nm on a plate reader (Sunrise, Tecan). The final amount of cAMP was expressed as pmol/mg protein.

### **3.12. $\beta_3$ -adrenoceptor effect on adenylyl cyclase activity**

For the investigation of  $\beta_3$ -adrenoceptors ( $\beta_3$ -AR) mediated changes in adenylyl cyclase activity the tissue homogenates were pre-incubated with preferential  $\beta_3$ -AR agonist CL 316243 (100 nM) 10 min at room temperature. The activity of adenylyl cyclase was determined immediately using enzyme immunoassay kit as described above.

### **3.13. Determination of nitric oxide synthase activity**

The determination of nitric oxide synthase activity (NOS) in tissue homogenates was performed by the Ultrasensitive Colorimetric NOS Assay Kit (Oxford Biomedical Research, Inc., USA). The first phase of the experiment (in microcentrifuge tube) was proceeded with tissue homogenates (30 $\mu$ l) mixed with 50 mM HEPES with 0.5 mM EDTA (200 $\mu$ l), NADPH Part A containing NADP<sup>+</sup>, Glucose 6-Phosphate and L-Arginine (10 $\mu$ l), and NADPH Part B containing Glucose 6-Phosphate dehydrogenase (10 $\mu$ l). After mixing and incubating for 1 h at 37°C (water bath) the reaction mixtures were chilled on ice for 5 min and subsequently nitrate reductase (10 $\mu$ l) was added to each sample. After vortexing and incubating for 20 min at room temperature the reaction mixtures were centrifuged at 12,500 rpm for 5 min at 4°C. The amount of protein (determined using the Pierce BCA protein Assay Kit) in the reaction mixtures (total volume of 260 $\mu$ l) was within the range of 138–339 $\mu$ g. The second phase of the experiment (in flat-bottom microtiter plate) was performed with nitrite standard solutions (100 $\mu$ l) diluted from 100 to 0.5 $\mu$ M in duplicates and samples (100 $\mu$ l) also in duplicates followed by addition of the Griess reagents, SA in 3 N HCl (50 $\mu$ l), and NED (50 $\mu$ l). After shaking for 5 min at room temperature the absorbance was measured at 540 nm in multi-detection microplate reader (Synergy HT).

### **3.14. $\beta_3$ -adrenoceptor effect on NOS activity**

For the investigation of  $\beta_3$ -adrenoceptor ( $\beta_3$ -AR) mediated changes in NO-synthase (NOS) activity the tissue homogenates were pre-incubated with preferential  $\beta_3$ -AR agonist BRL 37344 (1 mM) 10 min at room temperature. The time dependence (0, 5, 10, 15, 30 min) of NOS activation by BRL 37344 was determined in preliminary experiments showing 10 min as the start of peak level of activation. The activity of NOS was determined using the Ultrasensitive Colorimetric NOS Assay Kit (Oxford Biomedical Research, Inc.).

### **3.15. Catecholamine blood concentration**

Blood samples were collected in general anesthesia (see “Telemetric assessment of heart rate” for details), which was administered 25 min prior to sampling. Blood was drawn from the tail vein and the amount was more than 100 $\mu$ l (the plasma amount used for subsequent analysis was 50 $\mu$ l). Catecholamines were determined by enzyme immunoassay for the quantitative determination of Adrenaline (A) and Noradrenaline (NA) using 2-CAT Research ELISA kit (DEE5400, Demeditec Diagnostics GmbH, Germany). Briefly, A and NA were extracted using a cis-diol specific-affinity gel, acylated and then converted enzymatically. The solid

phase-bound antigen and A or NA in the samples/standards/controls competed for a fixed number of antiserum (rabbit) binding sites. After the system reached equilibrium, free antigen and free antigen–antiserum were washed out. Antirabbit IgG and peroxidase conjugate detects solid-phase bound antiserum. After converting TMB (substrate), the reaction was stopped and absorbance was read at 450 nm.

### **3.16. Gene expression of catecholamine-synthesizing enzymes**

In order to assess if there is no chronic adrenergic stress in KO animals, the gene expression of catecholamine-synthesizing enzymes was measured. Total RNA was prepared from frozen mice adrenal medullae using the RNazo1™ reagent (Tel-Test, Friendswood, TX, USA) according to the manufacturer's instructions. Reverse transcription was performed from 1.5µg of total RNA using Ready-To-Go You-Prime First-Strand Beads (Amersham Bioscience, Buckinghamshire, UK) and pd(N)6 primer according to the manufacturer's protocol. To determine mRNA levels in the adrenal medullae of mice, the following primers were used:

tyrosinhydroxylase (TH):

For: 5'-GAAGGGCCTCTATGCTACCCA-3', Rev: 5'-TGGGCGCTGGATACGAGA-3';

dopamine-β-hydroxylase (DBH): For: 5'-GACTCAACTACTGCCGGCACGT-3',

Rev: 5'-CTGGGTGCACTTGTCTGTGCAGT-3',

phenylethanolamine N-methyltransferase (PNMT):

For: 5'-TACCTCCGCAACAACACTACGC-3', Rev: 5'-AAGGCTCCTGGTTCCTCTCG-3',

glyceraldehyde-3-phosphate dehydrogenase (GAPDH):

For: 5'-AGATCCACAACGGATACATT-3', Rev: 5'-TCCCTCAAGATTGTCAGCAA-3'.

After the initial denaturation at 94 °C for 5 min, each cycle of amplification consisted of 60 s at 94 °C, 30 or 60 s at the respective annealing temperature and with a specific number of cycles (Benes et al., 2012b), and 60s at 72°C. Final polymerization was performed for 7 min at 72 °C. All reactions were performed in the linear phase of amplification. PCR products were separated on 2 % agarose/ethidium bromide gels. The density of individual bands was evaluated by STS 6220I system (Ultra-Lum, Inc., BLD Science, Garner, NC, USA), and relative expression values were calculated by the analysis of band intensities using PCBAS 2.08e software (Raytest, Inc., Dusseldorf, Germany). Semiquantitative values were expressed relative to the housekeeping gene (GAPDH).

### **3.17. Statistical analysis**

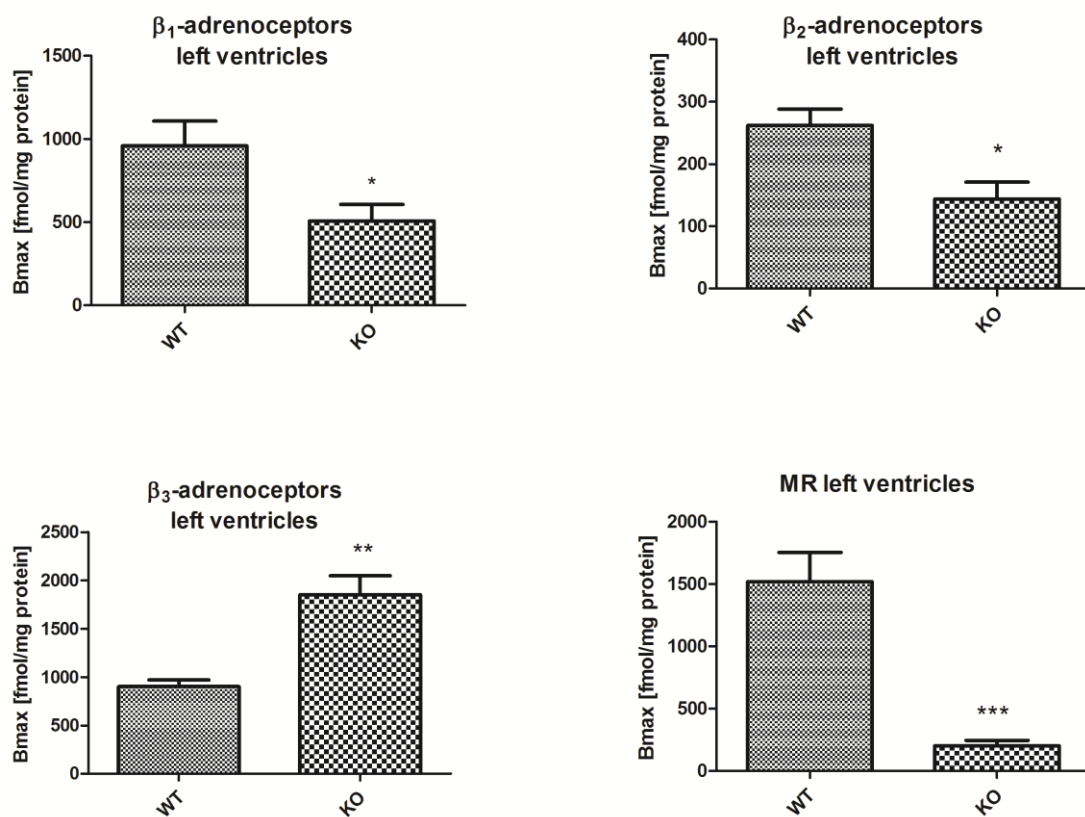
Results are presented as mean $\pm$  S.E.M. and each group represents an average of six to eight animals. In some cases we had to pool the tissue; in these cases, the number of experimental objects in the group is lower. Statistical differences among groups were determined by 1-way analysis of variance (ANOVA). For multiple comparisons, modified Newman–Keuls correction was used. The data from drug application experiments were evaluated using 2-way ANOVA (repeated measures; factors: treatment, phenotype). Heart rate variability data were analyzed using SPSS 19 software (IBM, Inc.) and GraphPad Prism 5.01 (GraphPad software, Inc.). The comparison of two groups was performed using Student t-test or Mann–Whitney test (if data did not reveal Gaussian distribution as assessed by F test). Values of  $p < 0.05$  were considered to be significant.

## 4. Results

We have initially performed binding experiments and subsequently gene expression experiments both in left and right ventricles. In right ventricles, no difference in binding and gene expression in all analyzed muscarinic and  $\beta$ -receptor subtypes were found between WT and KO animals except for binding of muscarinic receptors that was diminished to the similar extent as in left-ventricle (6% of WT). Therefore, other signaling parameters (adenylyl cyclase, NO-synthase) were analyzed in left ventricles only. All data presented below thus pertain to the situation in left ventricles.

### 4.1. Binding characteristics, basal state

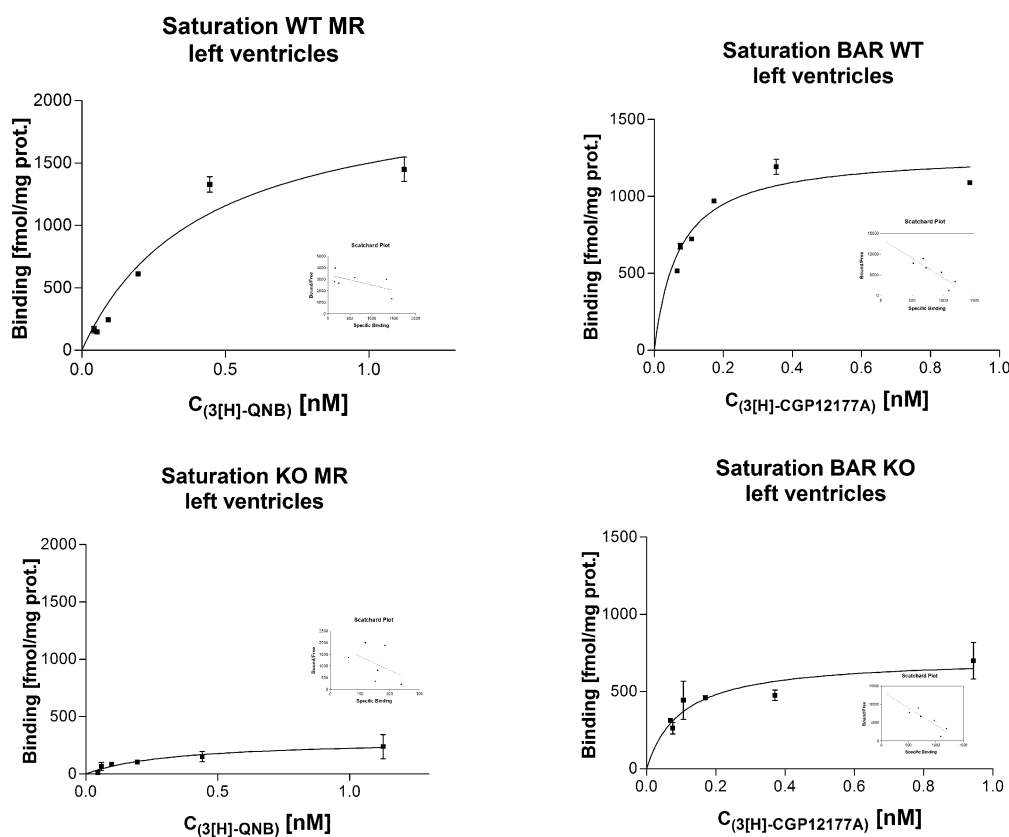
Gene disruption of  $M_2$ -receptor gene changed the amount of all  $\beta_1$ -,  $\beta_2$ -, and  $\beta_3$ - adrenoceptor binding sites (Fig. 4.1.1). The cardiostimulatory  $\beta_1$ - and  $\beta_2$ -AR were decreased in comparison to WT. On the contrary, the amount of  $\beta_3$ -AR binding sites was increased in  $M_2$ KO mice. The binding of muscarinic receptor specific radioligand in  $M_2$ KO mice was not completely abolished with 13 % of remaining binding sites.



**Fig. 4.1.1. The changes in the number of  $\beta_1$ -,  $\beta_2$ -,  $\beta_3$ -adrenoceptor binding sites and muscarinic receptor binding sites in  $M_2$ KO mice.**

$\beta_1$ - adrenoceptor binding sites (top left),  $\beta_2$ - adrenoceptor binding sites (top right),  $\beta_3$ -adrenoceptor binding sites (bottom left) and muscarinic receptor binding sites (bottom right). \* $p < 0.05$  different from WT, \*\* $p < 0.01$  different from WT, \*\*\* $p < 0.001$  different from WT,  $n = 3-4$  ( $\beta_1$ - and  $\beta_2$ -AR) and  $n = 6$  (MR and  $\beta_3$ -AR).

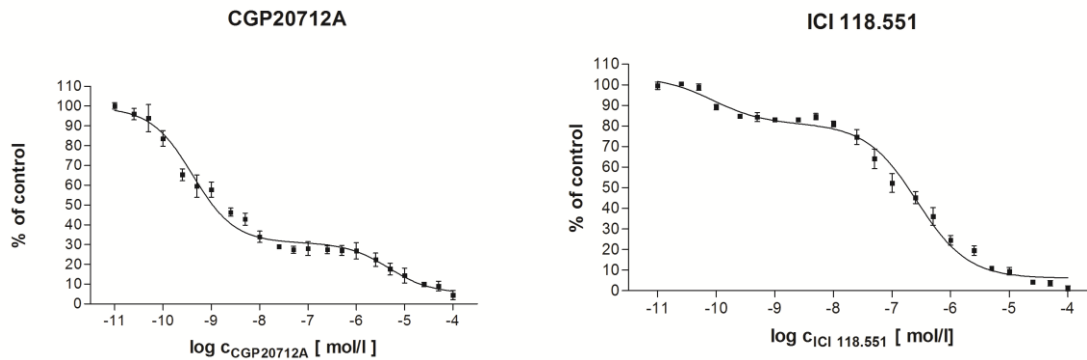
The experiments with saturation binding and competition with CGP20712A and ICI118.151 proved that the affinity of receptors ( $K_D$ ) was not affected by  $M_2$ -receptor knockout ( $155 \pm 25$  pmol/l vs.  $208 \pm 48$  pmol/l for  $\beta$ -AR WT and KO, respectively;  $50.1 \pm 2.5$  pmol/l vs.  $52.2 \pm 4.5$  pmol/l for MR in WT and KO, respectively, Fig. 4.1.2). Similarly, saturation experiments showed that  $K_D$  was similar for  $\beta_3$ -AR WT and KO mice ( $1030 \pm 169.6$  pmol/l vs.  $760.3 \pm 134.9$  pmol/l,  $p = n.s.$ ). These data suggest that observed changes occur in the amount of binding sites rather than in the affinities of receptors to ligands.





**Fig. 4.1.2. Saturation binding to muscarinic (left) and  $\beta$ -adrenoceptors (right) in WT (top) and KO (bottom) animals.**

Abscissa: concentration of free radioligand [nmol.l<sup>-1</sup>]. Ordinate: B<sub>max</sub> [fmol.mg.prot<sup>-1</sup>]. Insert: Scatchard plot of binding. MR, Muscarinic receptors; BAR,  $\beta$ -adrenoceptors.



**Fig. 4.1.3. Competition binding curves for CGP 20712A ( $\beta_1$ - AR antagonist) and ICI 118.551 ( $\beta_2$ -AR antagonist) with <sup>3</sup>H-CGP12177.**

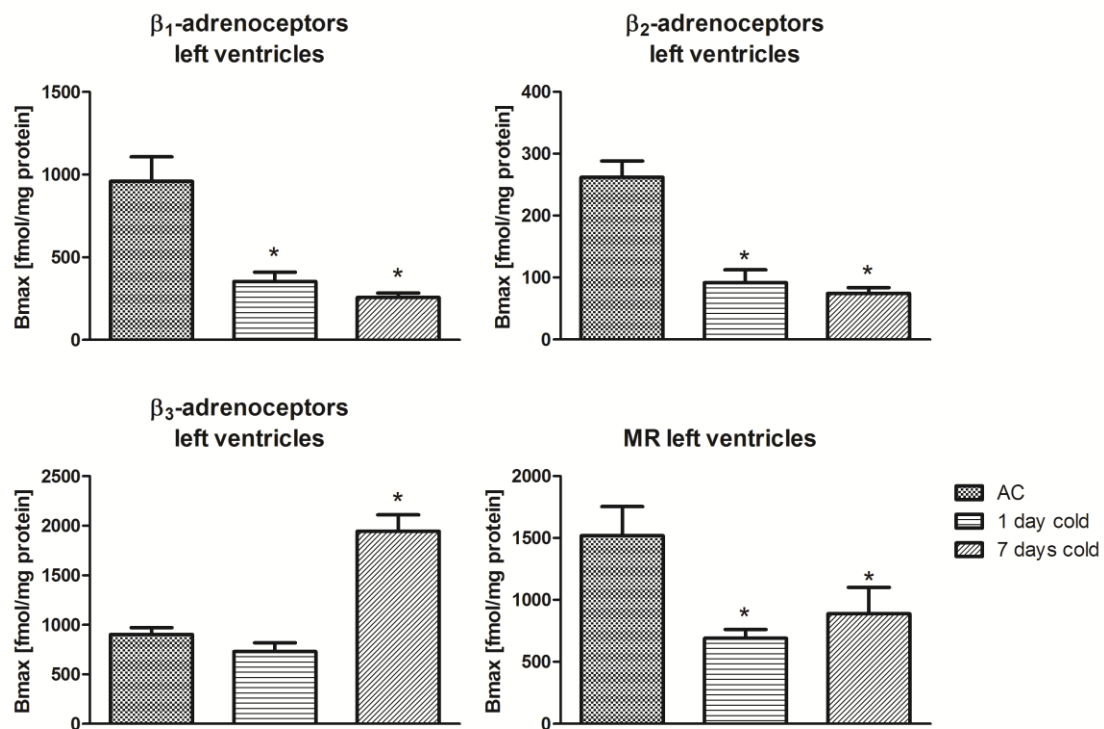
Average data from 4 independent experiments. Abscissa: log of antagonist concentration. Ordinate: Bound [% of binding in the absence of antagonist].

#### 4.2. Receptor binding upon stress

In WT animals, the cold stress resulted in a decrease in  $\beta_1$ - and  $\beta_2$ -AR (both after 1 day stay in the cold and after 7 days in cold) and also in a decrease in MR (see Fig. 4.2.1a, b).  $\beta_1$ -AR decreased to 37% of control values after 1 day of cold and further decreased to 27% of control values after 7 days of cold (1-way ANOVA, P= 0.0007, F= 20.94, df= 10).  $\beta_2$ -AR were diminished to 35% after 1 day of cold and to 28% after 7 days in cold, respectively (1-way ANOVA, P= 0.0003, F= 27.33, df= 10). On the contrary,  $\beta_3$ -AR increased (to 216% of control values) after 7 days of cold (1-way ANOVA, P=0.0001, F= 31.80, df= 17), Fig. 4.2.1. Muscarinic receptors decreased to 46% of control values after 1 day of cold and the changes prevailed till the day 7 (when 58% of control values were measured; 1-way ANOVA, p= 0.0182, F= 5.298, df= 17). The magnitude of changes expressed as percent of WT unstressed animals is shown in Tab. 4.2.3.

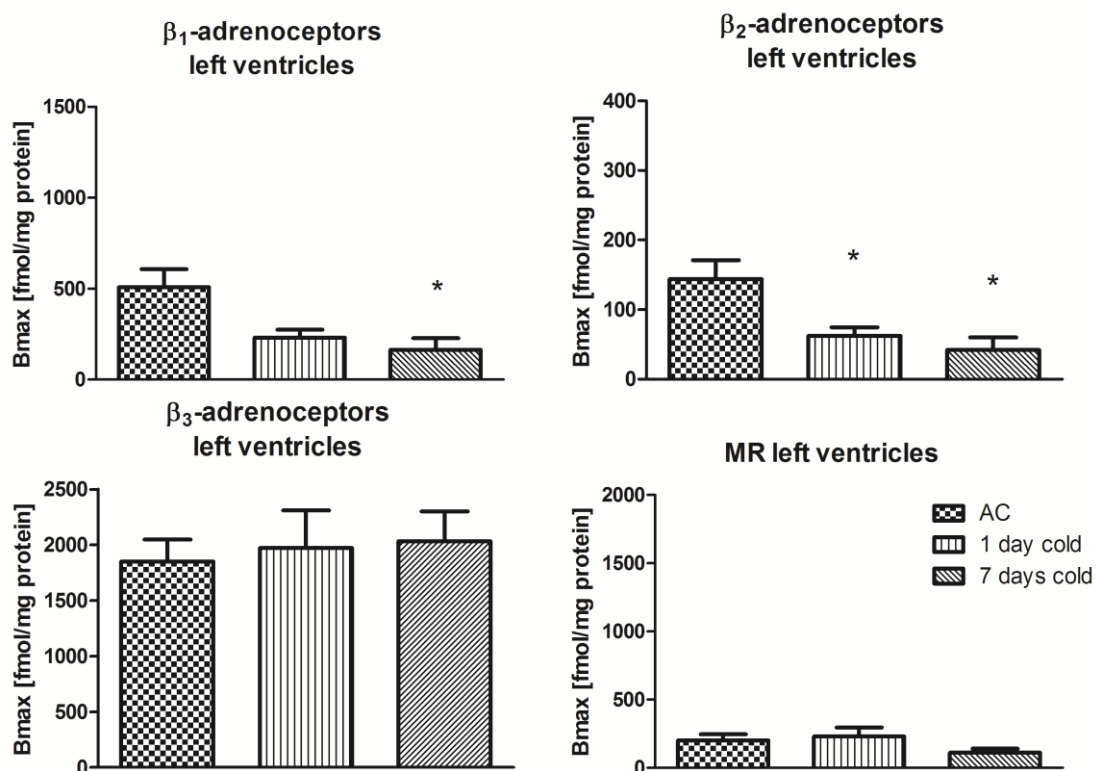
The reaction of cardiostimulatory  $\beta_1$ - and  $\beta_2$ -AR to cold was similar in KO animals. In KO animals, there was a decrease in  $\beta_1$ -AR number after 1 day of cold (to 45% of KO control values) and further persisted- 32% of KO control values were measured after 7 days of cold. The changes in  $\beta_2$ -AR were almost the same as those in WT:  $\beta_2$ -AR decreased to 44% and to 29% in KO animals (after day 1 and day 7, respectively), Fig. 4.2.2.

In contrast to cardioinhibitive receptors in wild types,  $\beta_3$ -AR in KO animals (which were already increased at baseline compared to WT) did not further change in reaction to cold (Fig. 4.2.2.). The magnitude of changes expressed as percent of WT unstressed animals is shown in Table 4.2.3. Muscarinic receptors (that were greatly attenuated at baseline in KO) did not further change upon cold stress (Tab. 4.2.3.).



**Fig. 4.2.1. Changes of receptor binding during exposure to cold in wild type animals in left ventricles.**

Abscissa: number of days in cold. Ordinate: receptor binding expressed as B<sub>max</sub> [fmol/mg. prot.<sup>-1</sup>]. a, b, c, d: appropriate type of receptor is marked in the heading.



**Fig 4.2.2: Changes of receptor binding during exposure to cold in knockout animals.**

Abscissa: number of days in cold. Ordinate: receptor binding expressed as B<sub>max</sub> [fmol/mg prot.<sup>-1</sup>]. a, b, c, d: appropriate type of receptor is marked in the heading.

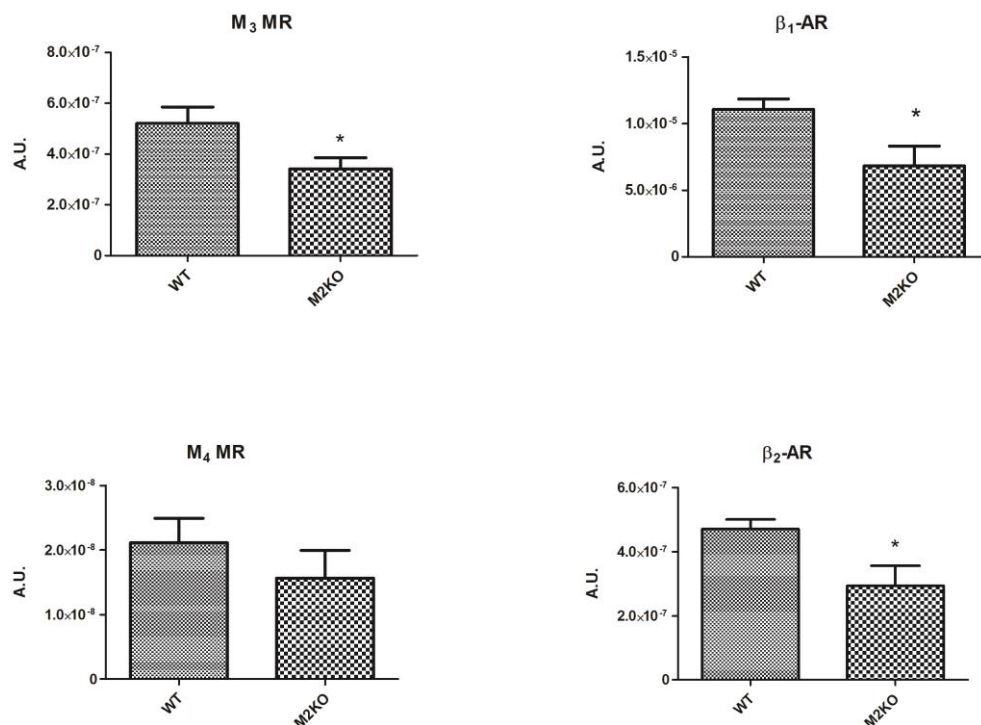
Genotype	Receptor subtype	Exposure to stress		
		Unstressed	1-day-stressed	7-day-stressed
WT	β <sub>1</sub> -AR	100 ± 15%	37 ± 6%*	27 ± 3%*
M <sub>2</sub> KO		53 ± 10%*	24 ± 5%*	17 ± 7%*
WT	β <sub>2</sub> -AR	100 ± 10%	35 ± 8%*	28 ± 4%*
M <sub>2</sub> KO		54 ± 10%	24 ± 5%*	16 ± 7%*
WT	β <sub>3</sub> -AR	100 ± 8%	82 ± 10%	219 ± 19%*
M <sub>2</sub> KO		208 ± 22%*	222 ± 38%	228 ± 30%
WT	MR	100 ± 15%	46 ± 5%*	58 ± 14%*
M <sub>2</sub> KO		13 ± 3%*	15 ± 4%	7 ± 2%

WT unstressed animals are considered as standard (100%), \**P* < 0.05 significant difference from unstressed WT

**Table 4.2.3. The proportional changes in binding sites in the left ventricles summarizing the magnitude of changes between groups**

### 4.3. Receptor gene expression, basal state

In the left ventricles of WT animals, M<sub>1</sub>- and M<sub>5</sub>-receptors were not expressed. The gene knockout of M<sub>2</sub>-receptors naturally abolished their gene expression. Moreover, M<sub>3</sub>-receptor gene expression was diminished in M<sub>2</sub>KO animals (Fig. 4.3.1.). M<sub>4</sub>-receptor gene expression remained unchanged in M<sub>2</sub>KO animals. The gene expression of both  $\beta_1$ - and  $\beta_2$ -adrenoceptors was decreased to approximately 60 % in KO animals (Fig. 4.3.1.).

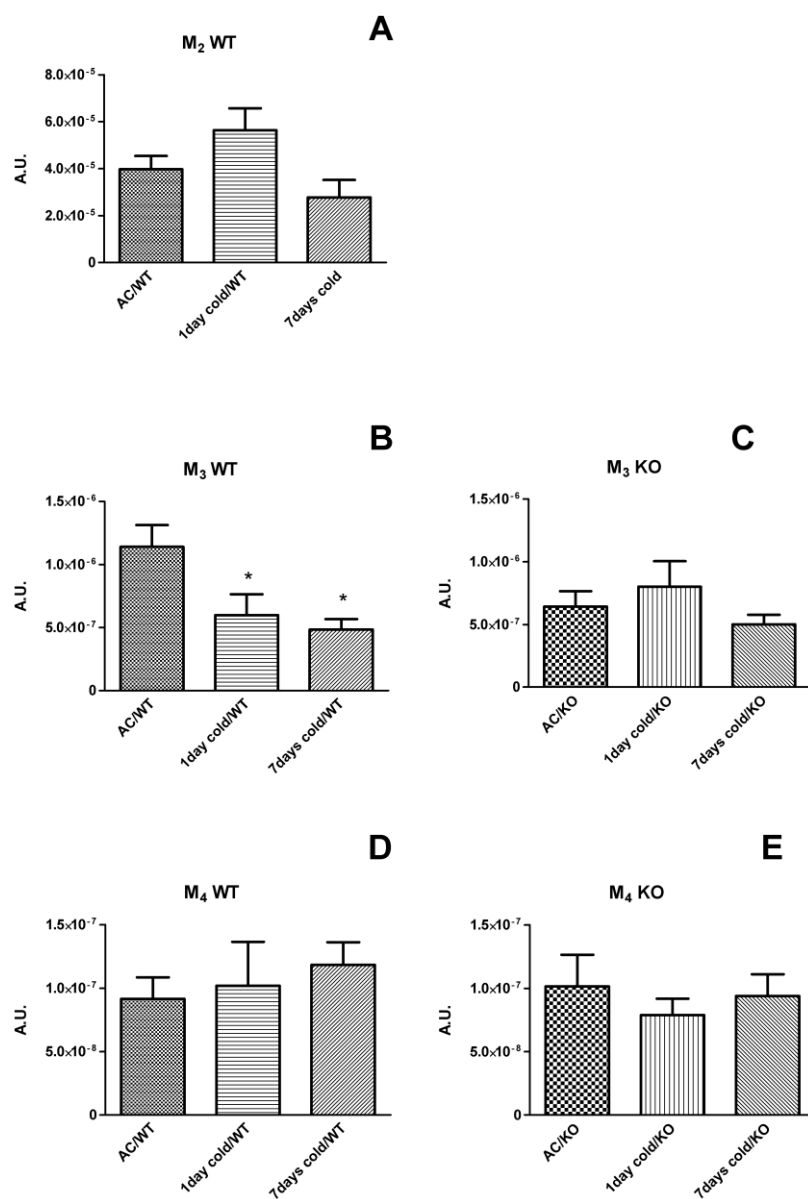


**Fig. 4.3.1. The changes in M<sub>3</sub>, M<sub>4</sub>, and  $\beta_1$ - and  $\beta_2$ -adrenoceptor gene expression in left ventricles in M<sub>2</sub>KO animals.**

\*p < 0.05 different from WT, n = 8 (WT and KO). M<sub>2</sub>-receptor gene expression was undetectable in KO animals, in addition to that, M<sub>1</sub>- and M<sub>5</sub>-receptor gene expressions were undetectable both in WT and KO mice.

#### 4.4. Receptor gene expression upon stress

Further, we have performed a gene expression assay determining the amount of mRNA for all MR subtypes ( $M_1$ - $M_5$ ) after the application of cold stress. As expected,  $M_2$ -receptors were not expressed in KO animals. In addition to that, the gene expression of  $M_1$ - and  $M_5$ -receptors was undetectable both in the unstressed and stressed wild types as well as in unstressed and stressed knockout animals. The expression of  $M_4$ -receptors were not changed in WT and KO upon stress (Fig. 4.4.1.d,e). On the contrary, the expression of  $M_3$ -receptors was decreased upon cold stress in WT but not in KO animals (Fig. 4.4.1.b,c). The magnitude of changes expressed as percent of WT unstressed animals is shown in Table 4.4.2.



**Fig 4.4.1: Changes of gene expression of muscarinic receptor subtypes (M<sub>1</sub>–M<sub>5</sub>) during exposure to cold in wild type and M<sub>2</sub> knockout animals.**

Abscissa: number of days in cold. Ordinate: the relative amount of mRNA expressed in A.U. [arbitrary units]. M<sub>2</sub> WT: M<sub>2</sub>-receptor gene expression, KO animals are not shown as M<sub>2</sub>-MR were deleted. M<sub>1</sub>- and M<sub>5</sub>-MR were not detectable. A: M<sub>2</sub> in WT, B: M<sub>3</sub> in WT, C: M<sub>3</sub> in KO, D: M<sub>4</sub> in WT, E: M<sub>4</sub> in KO.

Genotype	Receptor subtype	Exposure to stress		
		Unstressed	1-day-stressed	7-day-stressed
WT	M <sub>1</sub> MR	Non detectable		
M <sub>2</sub> KO				
WT	M <sub>2</sub> MR	100 ± 14%	142 ± 24%	70 ± 19%
M <sub>2</sub> KO		Non detectable		
WT	M <sub>3</sub> MR	100 ± 15%	52 ± 13%*	42 ± 7%*
M <sub>2</sub> KO		56 ± 11%*	70 ± 18%	44 ± 7%
WT	M <sub>4</sub> MR	100 ± 19%	111 ± 38%	129 ± 20%
M <sub>2</sub> KO		111 ± 27%	86 ± 14%	103 ± 19%
WT	M <sub>5</sub> MR	Non detectable		
M <sub>2</sub> KO				

WT unstressed animals are considered as standard (100%), \**P* < 0.05 significant difference from unstressed WT

**Table 4.4.2. The proportional changes in gene expression of M<sub>1</sub>-M<sub>5</sub> muscarinic receptor subtypes in the left ventricles summarizing the magnitude of changes between groups**

#### 4.5. Echocardiographic parameters

The echocardiographic parameters describing both LV systolic and diastolic function were similar both in WT and KO animals. The data are summarized in the Table 4.5.1 below.

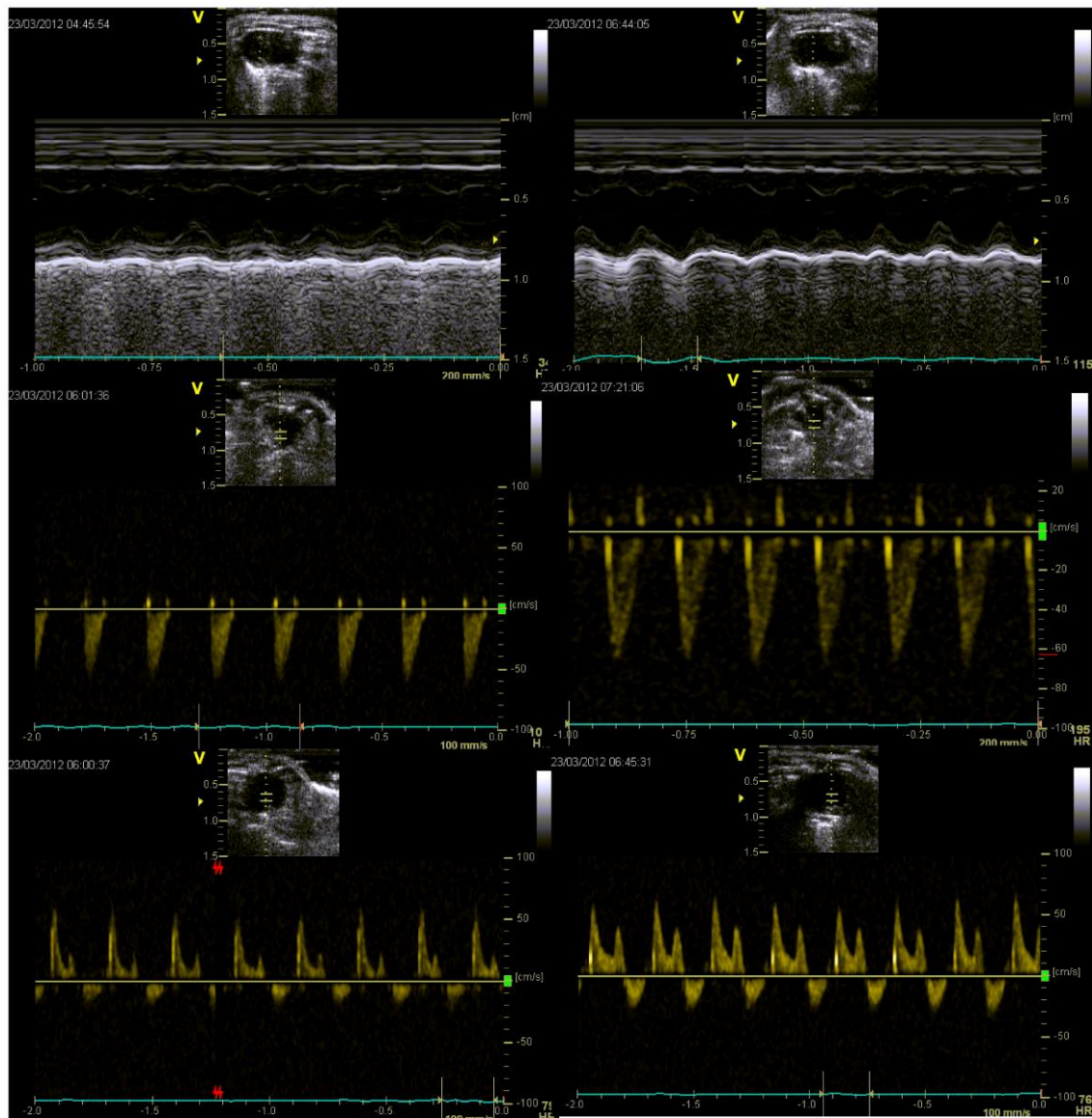
Representative echocardiograms of WT and KO animals are shown in Fig. 4.5.2.

<b>Parameter</b>	<b>WT (n=8)</b>	<b>KO (n=8)</b>	<b>Units</b>
<b>BW</b>	32.12±1.30	34.41±0.90	[g]
<b>M-Mode</b>			
IVSd-PSAX	0.98±0.08	0.87±0.05	[mm]
LVIDd-PSAX	3.24±0.17	3.62±0.16	[mm]
LVIDs-PSAX	1.95±0.19	2.09±0.18	[mm]
LVPWd-PSAX	1.12±0.14	0.83±0.05	[mm]
RR-PSAX	201.25±22.77	205.50±25.55	[ms]
<b>Doppler</b>			
MV E Vel	0.58±0.03	0.61±0.02	[m/s]
MV DecT	41.00±3.37	40.50±2.38	[ms]
MV Dec Slope	15.23±1.32	15.88±1.18	[m/s <sup>2</sup> ]
MV A Vel	0.26±0.02	0.30±0.04	[m/s]
MV E/A Ratio	2.29±0.15	2.40±0.35	
LVOT Vmax	0.66±0.03	0.60±0.04	[m/s]
LVOT maxPG	1.80±0.17	1.51±0.18	[mmHg]
LVET	86.13±5.26	86.63±5.73	[ms]
<b>Calculated parameters</b>			
HR	331.11±36.74	322.49±31.30	[min <sup>-1</sup> ]
LV FS	41.01±3.90	43.18±2.81	[%]
LV mass, echo	135.41±14.45	107.74±4.02	[mg]
Vcf	5.04±0.70	5.31±0.68	[circ/s]

**Table 4.5.1: Echocardiographic parameters in WT and KO animals**

The data are shown as mean±SEM. *IVSd-PSAX* interventricular septum thickness in diastole (measured in parasternal short axis, PSAX), *LVIDd-PSAX* left ventricle end-diastolic diameter measured in PSAX, *LVIDs-PSAX* left ventricle end-systolic diameter measured in PSAX, *LVPWd-PSAX* left ventricle posterior wall thickness measured in PSAX, *RR-PSAX* cycle

length measured in PSAX, *MV E Vel* mitral valve E wave velocity, *MV DecT* mitral valve deceleration time, *MV Dec slope* mitral valve deceleration time slope, *MVA Vel* mitral valve A wave velocity, *MV E/A Ratio* mitral valve E/A ratio, *LVOT Vmax* maximal velocity in the left ventricle outflow tract, *LVOT maxPG* maximal pressure gradient in the left ventricle outflow tract, *LVET* left ventricle ejection time, *HR* heart rate, *LV FS%* left ventricle fractional shortening, *LV mass*, echocardiographic calculation of LV mass, *Vcf* left ventricular peak velocity of circumferential fiber shortening



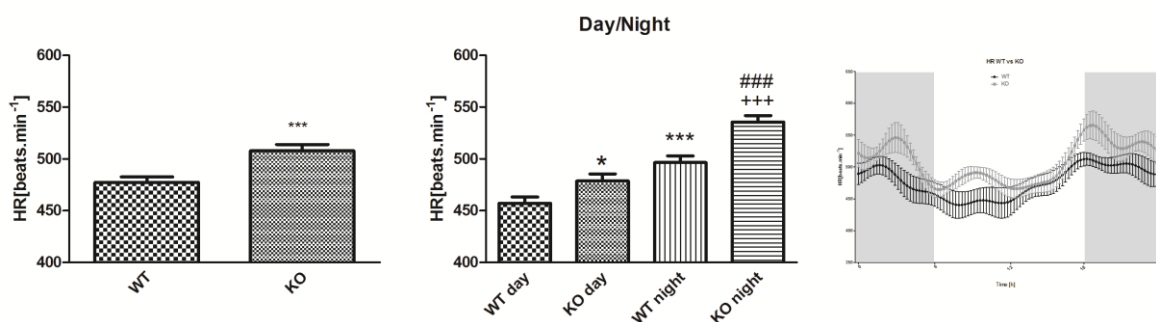
**Fig. 4.5.2. Representative echocardiograms of WT (left) and KO (right) animals.** Left: WT mice, right: KO mice. Top: Two-dimensional echocardiogram: end diastole, M-mode echocardiogram from the same mouse in the parasternal short-axis view. Middle: left



ventricle outflow tract (two-dimensional echocardiogram, pulsed wave Doppler tracing).  
 Bottom: mitral view (two-dimensional echocardiogram, pulsed wave Doppler tracing)

#### 4.6. Heart rate

When comparing means in WT and KO animals as averaged in 30-min intervals, we have found an increase in basal heart rate in M<sub>2</sub>KO animals (513.1±5.01 beats per minute, n=8 vs. 469.6±12.24, n=8, p=0.0054, KO vs. WT, i.e. an increase of 6 % in KO animals, Fig. 4.6.1. left). The difference between animals was also seen when comparing light phase and dark phase means; the increase was slightly higher in the nighttime (8% vs. 5% in the daytime; Fig. 4.6.1, middle). In order to better characterize the differences between wild-type and M<sub>2</sub>KO animals we have employed biorhythm analysis. Both WT and KO animals displayed a typical circadian pattern in HR with peak values in the dark span. However, WT and KO animals differed in biorhythm parameters (day mean 492.7±7.05 beats per minute, vs. 449.5±16.73, p=0.0320, KO vs. WT animals; night mean 530.6±10.61 vs. 488.5±9.03, p=0.0091; mesor 512.1±4.87, vs. 468.5±12.55, p=0.006; peak (i.e., maximal values) 585.1±19.28 vs. 531.6±7.51, p=0.0410; and minimal values 458.0±10.62 vs. 398.8±22.58, p=0.0240, n=6-8 for all parameters). In summary, the differences were between 8 and 15 %. The comparison of biorhythms in KO and WT animals is showed in Fig. 4.6.1. (right).

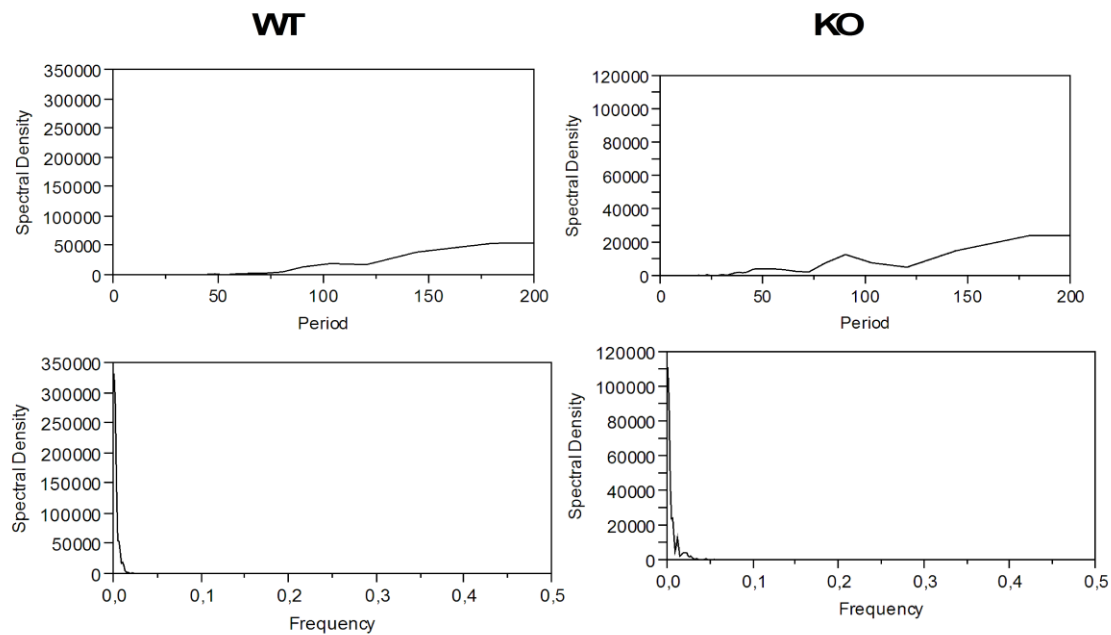


**Fig. 4.6.1. Heart rate in WT and M<sub>2</sub>KO mice.**

Average heart rate measured in 30-min intervals over a 24-hour period (left). Heart rate in the light and dark phase (middle). Comparison of biorhythm in WT and KO animals (right). \*p<0.05 different from WT day, \*\*\*p<0.001 different from WT day, +++p<0.001 different from WT night, ###p<0.001 different from KO day.

#### 4.7. Heart rate variability

Both animal groups (WT and KO) showed a similar distribution of power spectral densities (PSDs), Fig. 4.7.1. When areas under the curves (AUCs) were compared, Mann–Whitney test showed no difference between groups ( $6.03 \times 10^5 \pm 1.76 \times 10^5$  vs.  $5.84 \times 10^5 \pm 1.71 \times 10^5$ , WT vs. KO, respectively,  $p=0.916$ ).

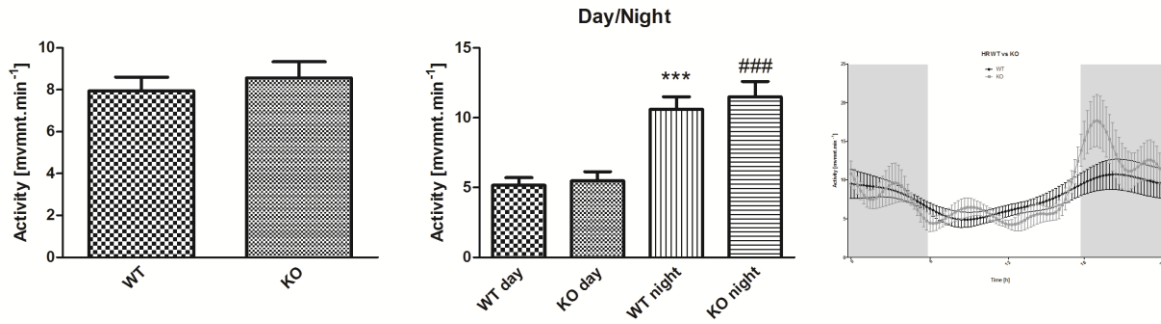


**Fig. 4.7.1. Representative figures of heart rate variability**

The representative figure of spectral density vs. period and spectral density vs. frequency in WT and KO animals. No difference was observed between KO and WT mice

#### 4.8. Activity

In order to determine whether the heart rate differences could be caused by an increase in activity (and therefore by higher demands for blood perfusion in KO animals), we have analyzed activity of WT and KO mice. Both WT and KO animals displayed a typical circadian pattern in their activity with peak values in the dark span. No difference in activity was observed both when the data were averaged in 30-min intervals over the whole 24-hours period as well as when the data from the light phase and the dark phase were treated separately. No difference in biological rhythm analysis was seen as well, Fig. 4.8.1.

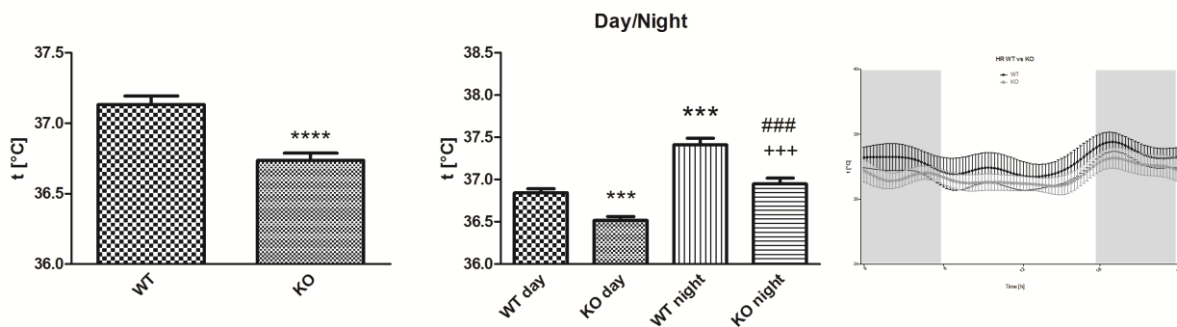


**Fig. 4.8.1. Activity**

Activity in WT and M<sub>2</sub>KO mice. Left: average activity. Middle: the activity in the light and dark phase (day or night). Right: comparison of biorhythm in WT and KO animals. \*\*\* $p < 0.001$  different from WT day. ### $p < 0.001$  different from KO day, 1-way ANOVA ( $p < 0.0001$ ,  $F = 15.83$ ,  $df = 93$ )

#### 4.9. Body temperature

The body temperature in M<sub>2</sub>KO animals was lower than in WT animals. When comparing the light and the dark phase, the same picture was obtained. The temperature of KO animals was lower both in the light phase and in the dark phase. In both groups (WT and KO) there was an increase in body temperature in the dark phase. Both WT and KO animals displayed a typical circadian pattern in temperature with peak values in the dark span. There was no difference in biological rhythm analysis (Fig. 4.9.1.).



**Fig. 4.9.1. Body temperature**

Body temperature in WT and M<sub>2</sub> KO mice. Left: average temperature. Middle: temperature in the light and dark phase (day or night). Right: comparison of biorhythm in WT and KO animals. \*\*\*\* $p < 0.0001$  different from WT, \*\*\* $p < 0.001$  different from WT day, ### $p < 0.001$



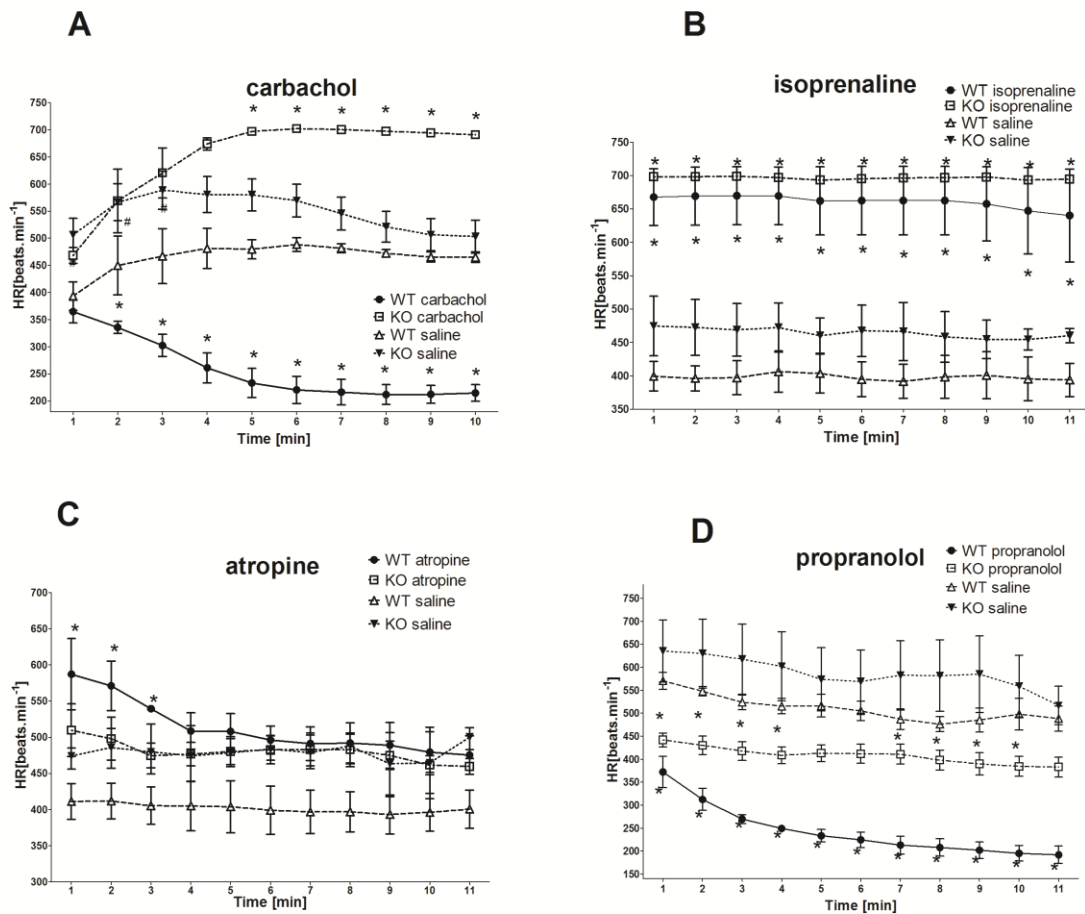
**Fig. 4.10.1.** The actograms of representative WT and KO mice showing increased values of respective parameters (black rectangles). **Abscissa:** time starting at 6:00 AM and showing two consecutive days in one row. **Ordinate:** The number of row representing two days. From top to bellow: WT activity, KO activity, WT HR, KO HR, WT temperature, KO temperature

#### 4.11. Drug application

Carbachol (see Fig. 4.11.1.a) caused expected effects of WT animals (bradycardia,  $257.4 \pm 18.03$  vs.  $464.5 \pm 8.645$  in WT animals treated with saline, i.e., decrease to 55 %). (The response of KO animals to saline treatment did not differ from the response in WT animals except for the first 3 min when the heart rate was slightly higher). In M<sub>2</sub>KO animals, carbachol induced tachycardia ( $651.4 \pm 24.56$  vs.  $546.9 \pm 10.91$ , i.e., increased by 19 %) when compared to saline-treated KO mice.

Isoprenaline (see Fig. 4.11.1.b) also caused expected effects in WT animals (tachycardia,  $687.1 \pm 6.593$  vs.  $390 \pm 3.29$ , i.e., 76 % increase compared to WT animals treated with saline). The same effects were observed in KO animals, where isoprenaline also evoked tachycardia although the effects were not as high as in WT animals ( $659.9 \pm 7.2$  vs.  $453.2 \pm 1.5$ , i.e., 46 % increase compared to saline-treated KO animals). There was also no difference in the response of KO animals to saline application when compared to WT counterparts.

Atropine (see Fig. 4.11.1.c) was able to increase the heart rate in WT animals ( $512.6 \pm 11.26$  vs.  $401.9 \pm 1.85$ ) but failed to affect the heart rate in KO mice ( $480.4 \pm 4.314$  vs.  $479.2 \pm 3.14$ ). Propranolol (see Fig. 4.11.1.d) was able to significantly decrease the heart rate in WT animals ( $243.1 \pm 16.88$  vs.  $510.4 \pm 8.662$ ). Similarly, in KO animals, the heart rate under propranolol treatment fell from  $586.6 \pm 10.27$  to  $408.2 \pm 5.56$ . Therefore, a decrease to 48 % of saline-treated animals was found in WT mice but only up to 70 % in KO counterparts.



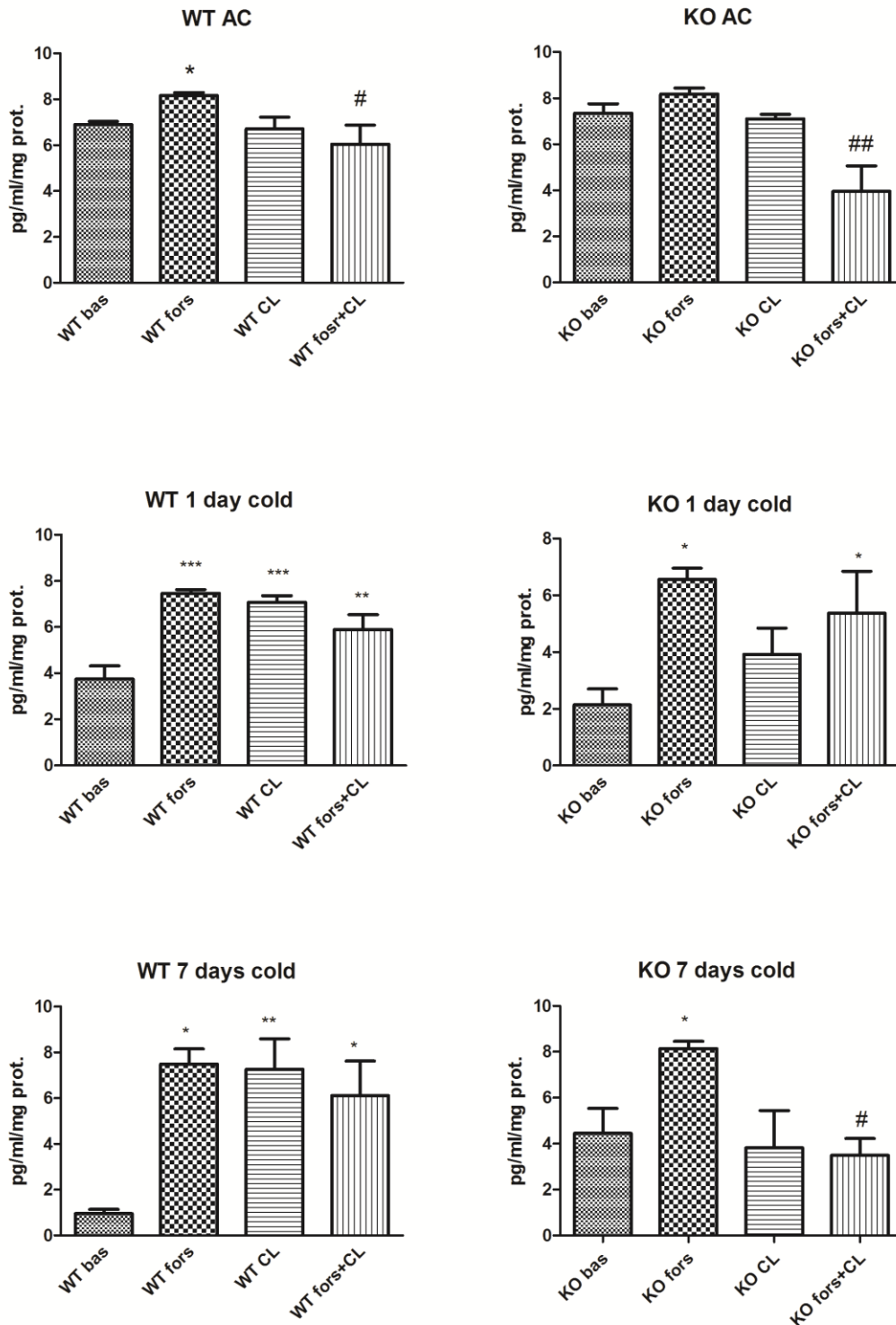
**Fig. 4.11.1. The effects of carbachol (0.5 mg/kg, a), isoprenaline (50 mg/kg, b), atropine (30 mg/kg, c), and propranolol (20 mg/kg, d) on heart rate as assessed by telemetric measurement.**

The drugs were applied i.p. and saline application served as control. Abscissa, time from the onset of drug effects; ordinate, heart rate [beats per min]; n=91–111. The onset of drug effects was: carbachol in the 1st minute, isoprenaline in the 21st minute, atropine in the 25th minute, propranolol in the 6th minute, respectively. The data were evaluated using 2-way ANOVA (repeated measures, factors: treatment, phenotype). The doses were derived from dose–response curves in WT animals from the preliminary experiments. \*p<0.05 from saline-treated animals, #p<0.05 from WT saline-treated animals. Higher significance (p<0.01, p<0.001) is not shown.

#### **4.12. Adenylyl cyclase activity**

Adenylyl cyclase activity increased in unstressed WT animals when it was stimulated by forskolin, which was abolished by CL 316243 (Fig. 4.12.1.a). CL 316243 alone had no effect on adenylyl cyclase activity in unstressed animals (Fig. 4.12.1.a, 1-way ANOVA,  $p= 0.0072$ ,  $F=5.347$ ,  $df= 23$ ). In contrast, CL 316243 was able to increase adenylyl cyclase activity after one (Fig. 4.12.1.c, 1-way ANOVA,  $p=0.0001$ ,  $F= 13.37$ ,  $df= 23$ ) and 7 days of cold (Fig. 4.12.1.e, 1-way ANOVA,  $P= 0.0087$ ,  $F= 5.63$ ,  $df= 18$ ). In stressed animals, forskolin also increased the amount of cAMP produced by adenylyl cyclase, but CL 316243 did not abolish these effects (Figs. 4.12.1.c, e).

In KO animals, forskolin was not able to increase the adenylyl cyclase activity in unstressed animals (Fig. 4.12.1.b), there was only difference between combined treatment with CL 316243 with forskolin and other groups (1-way ANOVA,  $P= 0.0005$ ,  $F= 9.26$ ,  $df= 23$ ). When stress was applied (after 1 or 7 days) there was marked increase after forskolin treatment (Fig. 4.12.1.d,f, 1 day: 1-way ANOVA,  $p= 0.0183$ ,  $F= 4.05$ ,  $df= 27$ ; 7 days: 1-way ANOVA,  $p= 0.0019$ ,  $F= 6.91$ ,  $df= 25$ ). In contrast to WT animals, CL 316243 had almost no effects on adenylyl cyclase activity in KO animals itself (Fig. 4.12.1.b, d, f); it only abolished the effects of forskolin after 7 days of cold.



**Fig 4.12.1 Changes of adenylyl cyclase activity during exposure to cold in wild type and knockout animals.**

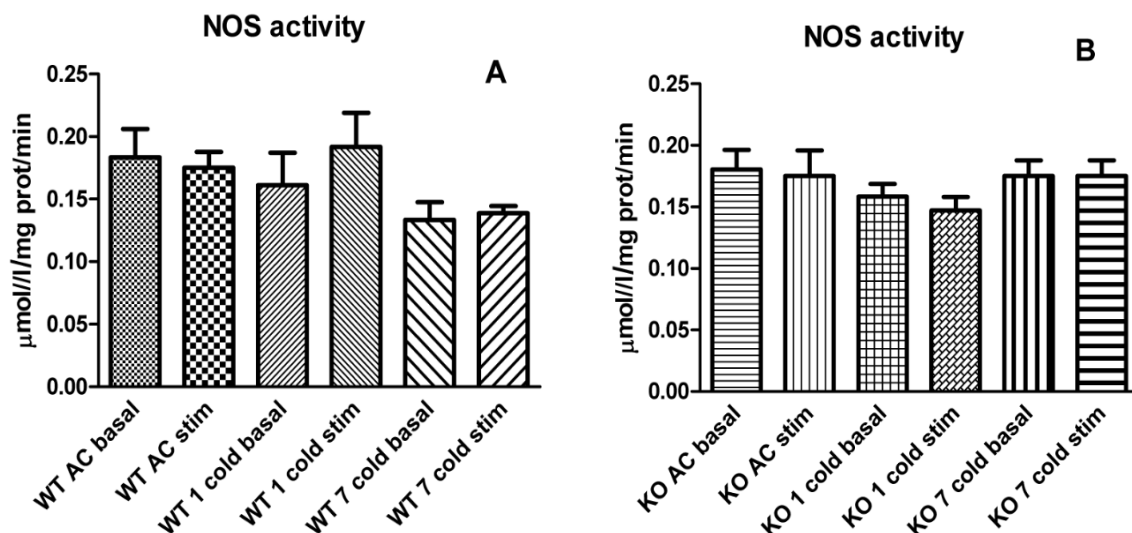
Abscissa: type of the drug added to the mixture (bas: basal level, i.e., no drug was added, fors: forskolin, CL: CL 316243, fors+CL: forskolin and CL 316243 were applied). Ordinate: the



amount of measured cAMP expressed as  $\mu\text{g}\cdot\text{ml}^{-1}\cdot\text{mg}$  of protein A: WT AC: unstressed WT animals, B: KO AC: unstressed KO animals, C: WT 1 day cold, D: KO 1 day cold, E: WT 7 days cold, F: KO 7 days cold.

#### 4.13. Nitric oxide synthase (NOS) activity

As it can be deduced from Fig. 4.13.1., BRL 37344 was not able to stimulate NOS activity both in WT animals (Fig. 4.13.1.a, 1-way ANOVA,  $p = 0.2352$ ,  $F = 1.451$ ,  $df = 35$ ) and KO animals (Fig. 4.13.1.b, 1-way ANOVA,  $p = 0.5565$ ,  $F = 0.8030$ ,  $df = 35$ ). Similarly, the stress had no effect on NOS activity in WT animals (after 1 or 7 days in cold) as well as in KO animals.



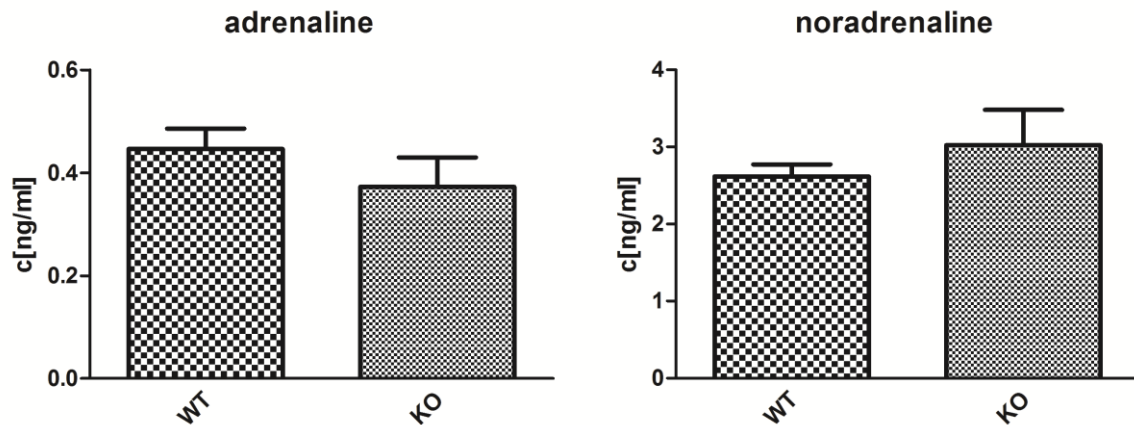
**Fig. 4.13.1. Changes of nitric oxide synthase activity during exposure to cold in wild type (a) and knockout (b) animals.**

Abscissa: type of stimulation used for NOS determination (basal basal level, i.e., no drug was added, stim: BRL 37344 treatment). Number of days in cold are shown as 1 or 7 cold (i.e., 1 day in the cold, 7 days in the cold). Ordinate: the amount of measured NO expressed as  $\mu\text{mol}\cdot\text{l}^{-1}\cdot\text{mg}\cdot\text{prot}^{-1}\cdot\text{min}^{-1}$

#### 4.14. Catecholamine blood concentration

Both adrenaline (A) and noradrenaline (NA) concentrations did not differ between KO and WT animals. Noradrenaline levels were  $2.26 \pm 0.15 \text{ ng}\cdot\text{ml}^{-1}$  in WT animals and  $3.03 \pm 0.46$

ng.ml<sup>-1</sup> in KO animals and adrenaline levels 0.45±0.04 ng.ml<sup>-1</sup> in WT animals and 0.37±0.08 ng.ml<sup>-1</sup> in KO animals (Fig. 4.14.1.).

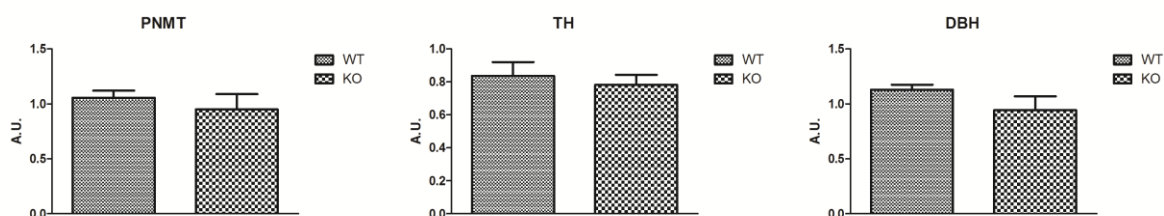


**Fig. 4.14.1. Concentrations of adrenaline and noradrenaline in the blood in WT and KO animals.**

Ordinate, catecholamine concentration [ng.ml<sup>-1</sup>], n= 3–10. No difference between WT and KO animals were observed.

#### 4.15. Gene expression of catecholamine-synthesizing enzymes

We have determined the gene expression of catecholamine-synthesizing enzymes (tyrosine-hydroxylase (TH), dopamine-β-hydroxylase (DBH) and phenylethanolamine-N-methyltransferase (PNMT)). No change in the gene expression of these enzymes between WT and KO animals was found (Fig. 4.15.1.).

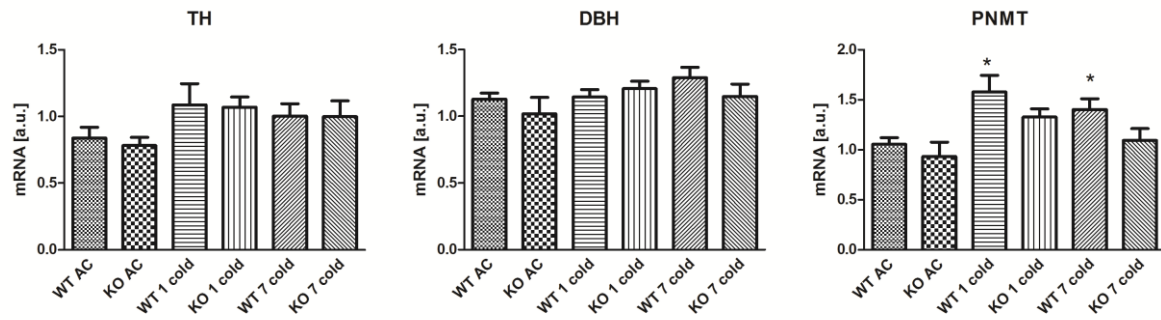


**Fig. 4.15.1: The gene expression of TH, DBH and PNMT in WT and KO animals.**

No change in the gene expression was observed. Ordinate: the relative amount of mRNA expressed in A.U. [arbitrary units].

#### 4.16. Changes in gene expression of catecholamine synthesizing enzymes

There was no difference in the gene expression of tyrosine hydroxylase and dopamine- $\beta$ -hydroxylase in both WT and KO and in basal state and upon cold stress. There was a higher level of gene expression of phenylethanolamine-N-methyl-transferase in WT animals after 1 and 7 days of cold (Fig. 4.16.1.).



**Fig. 4.16.1. The expression of catecholamine synthesizing enzymes**

Abscissa: animal type and number of days in cold. Ordinate: the relative amount of mRNA expressed in A.U. [arbitrary units]. \*  $p < 0.05$  from WT AC animals.

## 5. Discussion

In the presented work we show that the disruption of M<sub>2</sub>-receptor gene caused a decrease in muscarinic receptor binding as well as  $\beta_1$ - and  $\beta_2$ -adrenoceptor binding sites (to 52 and 55 %, respectively). In contrast to that,  $\beta_3$ -adrenoceptor binding was increased at the baseline. When challenged by cold-stress, both WT and KO animals showed a decrease in  $\beta_1$ - and  $\beta_2$ -adrenoceptor binding. WT animals showed a decrease in muscarinic binding as well; on the contrary, the remaining non-M<sub>2</sub>-binding preserved in M<sub>2</sub>KO mice was not further altered.  $\beta_3$ -AR binding (already increased at baseline in M<sub>2</sub>KO mice) continued to be increased to a similar extent upon stress stimulus (it was not further altered). WT mice showed eventually an increase in  $\beta_3$ -AR binding, but a prolonged stimulus was necessary to be applied (7-day cold stress), no changes were observed upon 1-day cold stress. M<sub>2</sub>KO mice had lower baseline gene expression of M<sub>3</sub>-receptors,  $\beta_1$ -adrenoceptors and  $\beta_2$ -adrenoceptors. Cold stress led to the decrease of gene expression of M<sub>3</sub>-receptors in WT but not KO mice. The expression of other receptor subtypes was not further altered. All these changes were chamber-specific (found in the left ventricle only). The overall cardiac function was not altered; no changes were found both in LV systolic and diastolic function as assessed by echocardiography (e.g. similar LV end-systolic and end-diastolic diameter, fractional shortening, mitral flow characteristics and maximal velocity in LV outflow tract). Only marginal changes have been found in physiological parameters; slightly higher basal heart rate during both dark and light phase in KO animals and slightly lower basal body temperature of M<sub>2</sub>KO animals have been found. Heart rate variability was similar in both groups either. The effects of isoprenaline and propranolol on HR were similar in WT and KO (in KO we have recorded the changes of lesser extent). Atropine was not able to increase HR in KO animals and carbachol decreased the HR in WT but increased the HR in KO. Our study suggests that the downstream signaling through  $\beta_3$ -AR occurs via adenylyl cyclase rather than NO synthase. There was a higher level of gene expression of phenylethanolamine-N-methyl-transferase in WT animals after 1 and 7 days of cold (and no difference in KO animals). The gene expression of tyrosine hydroxylase and dopamine- $\beta$ -hydroxylase was not altered both in WT and KO.

### 5.1. Gene expression and binding

Our results show that the disruption of M<sub>2</sub>-MR gene caused changes in the gene expression of M<sub>3</sub>-receptors,  $\beta_1$ -adrenoceptors and  $\beta_2$ -adrenoceptors together with the presumed decrease in muscarinic receptor binding sites as well as  $\beta_1$ - and  $\beta_2$ -adrenoceptor binding sites (to 52 and 55 %, respectively). The decrease of  $\beta_1$ -adrenoceptor and  $\beta_2$ -adrenoceptor binding sites was

caused by changes in the gene expression of the respective gene, as can be deduced from the gene expression study. In contrast to that,  $\beta_3$ -adrenoceptor binding was increased at the baseline. This phenomenon, i.e. that when one subtype receptor down-regulation occurs and it is followed by down-regulation of an antagonistic receptor as well was demonstrated multiple times (Paraschos and Karliner, 1994, Werry et al., 2003, Myslivecek et al., 2007) and it is fully in accordance with our hypothesis. Since the regulation is region specific (present in the left but not the right ventricle) heterologous regulation (i.e. local mechanism) is the most likely to be the responsible mechanism (Werry et al., 2003). This is an important fact suggesting that each ventricle has distinct mechanisms for receptor regulation.

We would like to point out that the binding of muscarinic-specific radioligand in  $M_2$ KO animals was not completely abolished, confirming the previous finding about another (minor) muscarinic population in the heart (Ito et al., 2009). On the level of gene expression, we were able to detect  $M_2$ -,  $M_3$ -, and  $M_4$ -receptor gene expression in WT animals;  $M_1$ - and  $M_5$ -receptors were not expressed. This is in a good agreement with some previously published data (Yang et al., 2005, Li et al., 2010, Pan et al., 2012). On the other hand, according to our previously published binding studies, the minor muscarinic population seemed to be  $M_1$  (Myslivecek et al., 2008). However, this previous study was performed on rats; on the contrary, this present study has been done with mice, so difference between species may play a role. Further, we have found a progressive decrease in  $M_3$ -gene expression upon cold stress in WT mice but not in  $M_2$ KO mice. The expression of  $M_4$ -receptors was not altered.  $M_3$ -receptors were showed to mediate cardioprotective role of acetylcholine on myocardium; they were showed to ameliorate ischaemia-induced arrhythmias (Wang et al., 2012) and decrease of infarct size and mortality rate employing the  $M_3$ /NF- $\kappa$ B/miR-376b-5p/BDNF axis (Pan et al., 2012). This supports the important role of non- $M_2$ -receptors for cardiac function. In our study, we have not found a compensatory up-regulation of  $M_4$ -receptors (that together with  $M_2$ -receptors inhibit adenylyl cyclase), so this hypothesis was not confirmed.

The receptor binding study on  $\beta_3$ -adrenoceptors showed a baseline increase in  $M_2$ KO that was sustained over the period of 7-day stress compared to WT mice, where an increase in  $\beta_3$ -AR appeared only after a rather prolonged stressful stimulus (7-day cold stress). This finding is in a good agreement with previously published studies and with the general paradigm on the function of  $\beta_3$ -adrenoceptors - they are perceived to be a „brake“ that is activated upon

excessively increased sympathetic overstimulation causing a negative inotropic effect counteracting the effects of  $\beta_1$  and  $\beta_2$  activation (Moens et al., 2010).

An application of 7-day cold stress was necessary to provoke an adrenergic response great enough to induce an up-regulation of  $\beta_3$ -AR. Our results show that CL 316243 (10,000 times more selective to  $\beta_3$ -adrenoceptors than to  $\beta_1$ - and  $\beta_2$ -adrenoceptors) was able to abolish the forskolin effects in WT unstressed animals. In contrast to that, CL316243 increases the amount of produced cAMP in cold, which suggests that  $\beta_3$ -adrenoceptors can switch from Gi coupling to Gs coupling. This is the property of  $\beta_3$ -adrenoceptors in the adipose tissue (Hamilton and Doods, 2008). Another possibility is that it can reveal the secondary state of  $\beta_3$ -adrenoceptors as it has been demonstrated earlier (Baker, 2005). Nevertheless, the increase of cAMP production by the  $\beta_3$ -AR activation is unobvious and can be seen in cold stressed WT animals only. Conversely, CL 316243 was not able to increase cAMP levels in KO animals, where  $\beta_3$ -AR are constantly increased. This fact gives evidence about the compensatory role of  $\beta_3$ -AR in the heart when  $M_2$ -MR are missing.

The possible interaction between increased level of cAMP and heart homeostasis should be evaluated further. As it has been demonstrated recently, there can be a connection of  $\beta_3$ -AR to  $Ca^{2+}$  signaling (Li et al., 2010) which could be also the possibility how to explain an increase in cAMP levels in WT only. In these experiments (Li et al., 2010),  $\beta_3$ -AR inhibits the  $Ca^{2+}$  currents induced by activation of  $\beta_1$ - and  $\beta_2$ -AR, which could explain the differences in CL 316243 effects in WT and KO animals (different amounts of  $\beta_3$ -AR binding sites). Concerning the downstream signaling pathway activated by  $\beta_3$ -AR, we have showed that the activity of adenylyl cyclase is importantly affected by  $\beta_3$ -adrenoceptor agonist, which suggests that adenylyl cyclase can be regulated during cold stress reaction by  $\beta_3$ -AR. This finding does not correlate well with the previous work suggesting that  $\beta_3$ -AR signaling occurs via the activation of NO synthase and subsequent production of NO. It has been thought long that the NOS isoform involved is endothelial NO synthase (eNOS) (Brixius et al., 2004) that has a crucial role in regulating L-type calcium channel activity in cardiac tissue. Protein kinase G (PKG) is a downstream effector of the eNOS/NO signaling cascade; it phosphorylates L-type  $Ca^{2+}$  channels which deactivates them (van der Heyden et al., 2005). However, recent studies show that eNOS is not the only NO isoform that mediates signaling through  $\beta_3$ -AR as NO production via the neuronal NOS (nNOS) has been demonstrated (Amour et al., 2007). The role of inducible NOS (iNOS) has not been fully clarified yet either, but it is likely that NO production through this isoform may also contribute to the NO production followed by  $\beta_3$ -AR activation (Maffei et al., 2007). Our work suggests that the

signaling through  $\beta_3$ -adrenoceptors can occur through the activation of cAMP as well. This illustrates the complexity of  $\beta$ -AR signaling and well supports the concept of „signalome“ where an individual adrenoceptor associates with multiple signaling pathways in time and spatially specific manner (Park and Palczewski, 2005).

Overall, since the main difference found was on the level of receptors, we propose that they serve as a tool to compensate  $M_2$ -knockout to preserve cardiac function.

## **5.2. Biorhythms (heart rate, temperature, activity)**

We should emphasize that although analyzed and discussed separately, there is a mutual connection between heart rate, activity and temperature (both the temperature of animals and ambient temperature). We have found that  $M_2$ KO mice have slightly but significantly increased heart rate ( $507.8 \pm 6.15$  vs.  $477.3 \pm 5.19$  beats.min<sup>-1</sup>, i.e., 6 %) both during light phase and dark phase with similar spectra of heart rate variability. Heart rate reflects the dynamic balance between sympathetic and parasympathetic branch of the autonomic nervous system, but the ratio between both of them might vary between species and other factors especially experimental conditions (ambient temperature) needed to be taken into account. In humans, the sinoatrial node is under the dominant influence of parasympathetic nervous system as the resting heart rate is much slower than the intrinsic rate of the sinoatrial node (Brubaker and Kitzman, 2011). On the other hand, it was long believed that in mice the intrinsic rate of the sinoatrial node is significantly lower than the actual heart rate (Gehrmann et al., 2000), which was interpreted that mice were under a dominant tonus on the sympathetic nervous system. It has been found recently that the ratio between sympathetic and parasympathetic tone depends greatly upon the ambient temperature. At 30°C, wild-type mice have the heart rate about 350 bpm whereas  $M_2$ KO mice 416 bpm (during the light phase) demonstrating that under these conditions the heart rate in mice is (similarly to humans) under dominant tone of parasympathetic nervous system (Swoap et al., 2008). However, when the ambient temperature drops to 21-23°C (normally used in animal facilities) the heart rate rises to cca 600 bpm (light phase) both in WT and  $M_2$ KO mice and the difference between groups disappears. Therefore, we have measured ambient temperature very carefully and have showed that in such well controlled environment (24.45 °C light phase/24.21 °C dark phase) it was possible to reveal a slight but significant increase in the heart rate in  $M_2$ KO animals, which is fully in accordance with previously published studies (Swoap et al., 2004, Swoap et al., 2008). On the other hand, some previously published work (Bymaster et al., 2001, LaCroix et al., 2008) reported no difference in the heart rate between WT and  $M_2$ KO mice.

However, certain factors have biased these observations. Both of these studies were performed on anesthetized animals, the experiments were performed during the day and the ambient temperature was not provided. Our analysis revealed that the difference in heart rate was greater during the night when mice are naturally more active. During the day the differences were more subtle and there were even intervals where there was virtually no difference at all. We believe we were able to record a difference between WT and M<sub>2</sub>KO mice because we used telemetric monitoring on wake animals and monitored the ambient temperature very carefully.

The long term telemetric analysis of activity (averaged values and biorhythm analysis) of the animals did not find any difference between WT and M<sub>2</sub>KO mice. This finding is in accordance with previously published data (Gomez et al., 2001). In this case, data obtained from long term telemetric monitoring are in accordance with the data obtained during the experiment. No difference in activity (KO vs. WT) together with lower body temperature in KO animals in relation to unchanged left ventricular function as assessed by echocardiography makes the possibility that the HR increase is caused by higher heart performance rather improbable. This data suggest that M<sub>2</sub>-receptors do not affect activity behavior in mice.

On the other hand, the long term measurement of the basal temperature of M<sub>2</sub>KO mice revealed that it was decreased (without change in biorhythm). This can explain the differences in oxotremorine-induced tremor and hypothermia in M<sub>2</sub>KO mice as published previously (Gomez et al., 1999).

### **5.3. Cardiac function**

No changes were found in any echocardiographic parameters (related to left ventricular systolic and diastolic function): there were no differences in left ventricular end-diastolic and end-systolic diameters, as well as in left ventricle hemodynamic indicators (maximal velocity in the left ventricle outflow tract, mitral flow characteristics—E velocity, A velocity, E/A ratio, left ventricle ejection time). Similarly, left ventricular peak velocity of circumferential fiber shortening showed no changes in KO animals. This observation is in agreement with previously published work (LaCroix et al., 2008) but in contradiction with study when M<sub>2</sub>-MR were knocked-out postnatally (a study using M<sub>2</sub>-MR coding plasmid DNA-immunized mice), where the impairment of heart function and changes in left ventricular echocardiographic parameters were found (Gimenez et al., 2005). These results indicate the



important differences between pre-natal and postnatal adaptation processes. Preserved systolic and diastolic function is fully in accordance with our hypotheses.

#### **5.4. Drug application**

The effects of carbachol in WT animals were expectable. On the other hand, the tachycardia in M<sub>2</sub>KO animals was not regularly observed. (Gomez et al., 1999) referred that incubation of atria derived from M<sub>2</sub>KO mice with the carbachol (1μM) had no significant effect on atrial beating frequencies. Similarly, *in vivo* study with carbachol in M<sub>2</sub>KO mice (Bymaster et al., 2001) showed no influence on the heart rate. On the other hand, (Kitazawa et al., 2009) described an increase in inotropy in the atria of M<sub>2</sub>KO animals. There are at least two explanations of this phenomenon that originate from our results: (a) the first possibility is that the decrease of β-adrenoceptors (to approximately one half of control values) when heart muscarinic receptors are almost absent in the heart is able to cope with basal conditions of the heart while it is not sufficient to cope with the sustained activation of muscarinic receptors; (b) the second possibility is that the effects of minor muscarinic receptor subtype (as described by (Willmy-Matthes et al., 2003, Wang et al., 2004, Wang et al., 2007, Wang et al., 2012) and considered as cardiostimulatory) can be revealed whilst it is hidden (in WT animals) by action mediated via major M<sub>2</sub>-receptors. According to our knowledge, this is the first observation of cardiostimulatory effects of carbachol *in vivo* in mice lacking the main cardioinhibitory muscarinic receptor subtype. This phenomenon further confirms the role of minor muscarinic receptor subtype in the heart. The third possibility is increased noradrenaline release from sympathetic nerve endings as described by (Trendelenburg et al., 2005) in M<sub>2</sub>/M<sub>3</sub>KO, but this rather would not be the case as we have not found a difference in blood catecholamines between WT and KO animals.

Upon the application of isoprenaline, we have observed an increase in HR both in WT and KO mice, but the effect on M<sub>2</sub>KO animals was attenuated. This observation is in the contradiction with the previously published data (LaCroix et al., 2008) that reported a significantly increased heart rate upon isoprenaline application in M<sub>2</sub>KO mice. The reason why we have observed opposite results is not clear; however, in our study all the measurements were performed by telemetry without using sedation, which we believe helped to obtain more reliable and unbiased results. The lesser extent of tachycardia in KO animals treated with isoprenaline can have the nature in having a decreased number of β-adrenoceptors (52% for β<sub>1</sub>-adrenoceptors and 55 % for β<sub>2</sub>-adrenoceptors). On the other

hand, the effects of  $\beta$ -adrenoceptor agonist persisted in KO animals, suggesting the compensatory role of receptor changes in heart physiology.

Atropine application increased the heart rate in WT animals but did not have this effect in  $M_2$ KO, confirming the absence of main cardioinhibitory receptors ( $M_2$ ) in the heart. It was reported previously that atropine had no effect in WT animals (LaCroix et al., 2008), which is in contradiction with our results. Again, we believe that the measurements performed by telemetry without using sedation helped us to obtain more reliable and unbiased results.

Propranolol decreased the heart rate both in WT and  $M_2$ KO animals; however, the drop in  $M_2$ KO animals was attenuated (a decrease to 48% in WT but only to 70% in  $M_2$ KO). This supports the hypothesis mentioned earlier that the decrease in  $\beta$ -adrenoceptors could be responsible for the diminished effects of  $\beta$ -adrenoceptor drugs in  $M_2$ KO animals. Once again, these data also support the hypothesis about the compensatory role of receptors in maintaining stable and physiological cardiac output.

### **5.5. Catecholamine blood concentration and gene expression of catecholamine-synthesizing enzymes**

In order to test if down-regulation of cardiac  $\beta$ -adrenoceptors (observed in our study) could be caused by the increased catecholamine release (carbachol decreases noradrenaline release from sympathetic nerves and this decrease is diminished in  $M_2/M_3$ KO animals (Trendelenburg et al., 2005)), we have measured the blood catecholamines. We have showed that the levels of circulating catecholamines were similar in WT and KO animals, which supports the hypothesis that heterologous regulation of  $\beta$ -adrenoceptors via intracellular signaling pathways is likely the responsible mechanism. The levels of circulating catecholamines were similar as referred by (Grouzmann et al., 2003). Similarly, the expression of the catecholamine-synthesizing enzymes (TH, DBH, and PNMT) showed that there was no increase in  $M_2$ KO animals indicating that  $M_2$ KO mice are not under chronic adrenergic stress. This is fully in accordance with our hypotheses.

### **5.6. Specific changes in cardiac autonomic receptor expression and function in $M_2$ KO mice**

The ability to manipulate the mice genes (knock-out and knock-in mice) provides the possibility for researchers to investigate the specific functions of the particular protein and to analyze its physiological roles more specifically than in the past. It was believed that this method was superior in ability to see the effect of a specific gene, but the initial hopes put into

the method were gradually corrected and now it is believed that it is rather an illustration of organism adaptation.

On the other hand, the knockout of M<sub>2</sub>-receptors is quite specific as the proportion of minor muscarinic heart receptors is minute, and therefore the general deletion of M<sub>2</sub>-receptor gene means that specifically in the heart main cardioinhibitory receptors are absent (minor subpopulation of cardioinhibitory M<sub>4</sub>-receptors are present). The presence of M<sub>2</sub>-receptors in peripheral tissue like in the intestine, lung, salivary glands and smooth muscles could not rather play a role in the regulation of heart function. On the other hand, the presence of M<sub>2</sub>-receptors on cardiac nerves (as autoinhibitory receptors), on autonomic ganglia (in a similar function, i.e., as autoreceptors), or in the hypothalamus (as receptors that can affect basal autonomic tonus) could change the picture of general knockout. Although there are some data about the presence of M<sub>2</sub>-MR in parasympathetic ganglia (or postsynaptic nerves) (Racke et al., 2006) and about the presence of MR on sympathetic ganglia (Fasano and Niel, 2009), which are rather M<sub>1</sub>-MR, their role is not strong enough to justify their importance in heart regulation. Moreover, our data suggest that they are not present in the heart (left and right ventricles) at all (missing M<sub>1</sub>-receptor gene expression in both WT and KO mice). The release-inhibiting heart receptors are more likely heterogenous (Zhou et al., 2002) with a defined role of M<sub>4</sub>-MR rather than M<sub>2</sub>-MR. On the other hand, MR are present in the medial preoptic area (Imeri et al., 1996), affecting the sleep–wake cycle and thermoregulation, and M<sub>2</sub>-MR appear to regulate adrenocorticotropin hormone responses to cholinergic stimulation (in males, not in females) (Rhodes et al., 2005). Moreover, little attention was paid to the counter-regulatory changes. Usually, the investigation concentrates on specific knockout changes - i.e., M<sub>2</sub>-receptors (Stengel et al., 2000),  $\beta_1$ -adrenoceptors (Rohrer et al., 1996, Ecker et al., 2006),  $\beta_2$ -adrenoceptors (Chruscinski et al., 1999, Ecker et al., 2006),  $\beta_3$ -adrenoceptors (Varghese et al., 2000),  $\beta_1/\beta_2$ -adrenoceptors (Kiriakis et al., 2008) or alternatively on total  $\beta$ -adrenoceptor knockout, i.e.  $\beta_1/\beta_2/\beta_3$ -adrenoceptor knockout (Lee et al., 2010). Only a few papers have described the alteration of counter-acting receptor responses in knockout animals. For example, (LaCroix et al., 2008) have noticed a changed response to isoprenaline in M<sub>2</sub>KO animals. Some authors have investigated the effects of downstream signaling molecules knockout (adenylyl cyclase, G proteins) on heart functions (Boknik et al., 2009, Gottle et al., 2009). According to our knowledge, our investigation is the first to emphasize the role of counter-regulatory receptors in the maintenance of heart functions.

## 6. Conclusion

In cardiac left ventricles, the disruption of M<sub>2</sub>-receptor gene caused a compensatory decrease of cardiostimulatory  $\beta_1$ -adrenoceptors and  $\beta_2$ -adrenoceptors with corresponding down-regulation of the gene expression, M<sub>3</sub>-receptors were down-regulated as well. Missing M<sub>2</sub>-receptors were functionally replaced by the main cardioinhibitory  $\beta_3$ -adrenoceptors that were up-regulated, not by cardioinhibitory M<sub>4</sub>-receptors.  $\beta_3$ -adrenoceptors were found to signal through adenylyl cyclase instead of NO synthase. All these changes were found in the left ventricle only, so heterologous regulation is likely to be the responsible mechanism. Slightly higher basal heart rate and lower basal body temperature of M<sub>2</sub>KO animals underscore the importance of telemetric measurement to avoid biased recordings. Despite all the changes found the overall cardiac function was unaltered.

## 7. References

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