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Editorial

## The PGE<sub>2</sub>-Stat3 connection in cardiac hypertrophy

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See article by Frias et al. [6] (pages 57–65) in this issue.

### 1. Stimulation of cardiomyocyte hypertrophy

In vivo cardiac hypertrophy is a slow process in which the myocytes increase in size in response to increased workload due to either an increase in hemodynamic load or to a loss of functional myocytes. Although the mechanical load has long been recognized as the most powerful hypertrophic stimulus, its signal transmission from the cell surface to the nuclear transcription activities has largely remained elusive. Apart from the mechanical stress, the hypertrophic reaction is monitored by a large number of neurotransmitters, hormones, growth factors, and cell mediators that under normal conditions stay in balance within a limited range of variation [1]. However, under chronic mechanical overload, ischemic conditions, or late-onset genetic diseases, all of which may be accompanied by inflammation, the ensemble of these factors becomes imbalanced, affecting the gene expression program and thus triggering the remodeling process that eventually leads to fatal cardiac dysfunction. The signaling cascades involved in the hypertrophic reaction include the mitogen-activated protein kinase (MAPK) pathway with its four branches (i) ERK1/2 (extracellular signal-regulated kinases), (ii) p38 MAPK, (iii) JNK (c-Jun N-terminal kinase) and (iv) BMK1 (big MAPK1=ERK5); furthermore, the cAMP-dependent PKA pathway, the PI3K–PKB/Akt pathway, the calcineurin — NFAT–MEF2 pathway, and the JAK/Stat (Janus kinase/signal transducer and activator of transcription) pathway are also involved [1]. The latter JAK/Stat signaling cascade is typically activated during inflammation by the interleukin-6-related cytokines

(IL6, cardiotrophin-1=CT1, and the leukemia inhibitory factor=LIF), which are potent inducers of cardiac hypertrophy [2]. In addition, prostaglandin-E<sub>2</sub> (PGE<sub>2</sub>) has been reported to be involved in cardiomyocyte hypertrophy, specifically by stimulation of its G-protein-coupled receptor subtype EP4 involving transactivation of the epidermal growth factor receptor (EGFR) and ERK1/2 [3,4] (Fig. 1). PGE<sub>2</sub> is synthesized by an enzyme cascade localized to the nuclear envelope comprising phospholipase-A<sub>2</sub> (PLA<sub>2</sub>), cyclooxygenase-2 (COX2), and prostaglandin-E synthases (PGES). PGE<sub>2</sub> may act extracellularly and intracellularly since its EP4 receptor is localized to the sarcolemma as well as to the nuclear envelope [5].

### 2. Stat3 here — who's knocking?

The group of Ursula Lang elegantly shows in this issue of the Journal (Frias et al., ref. [6]) that 1 μM PGE<sub>2</sub> induces an increase in cell size and stimulates protein synthesis in spontaneously beating ventricular neonatal rat cardiomyocytes (VNRC) in culture by signaling primarily via its EP4 receptor subtype involving successive phosphorylation of MEK1/2 (mitogen-activated ERK activating kinases) and ERK1/2 and of Stat3 at Tyr705 in the C-terminal transactivation domain. Upon stimulation with 100 nM PGE<sub>2</sub> the degree of phosphorylation of MEK1/2 and ERK1/2 reaches a transient maximum after 2–5 min while in Stat3 the maximum occurs only after 60 min. Inhibition of the ERK1/2 pathway by U0126 (inhibiting both MEK1 and MEK2) and application of the tyrosine kinase inhibitor genistein both abolished PGE<sub>2</sub>-induced tyrosine phosphorylation of Stat3 and its binding to the sis-inducible element in the promoter region of the *c-fos* gene as assessed by electrophoretic mobility shift assay. Contributions by p38 MAPK or PKA signaling could be excluded by using various

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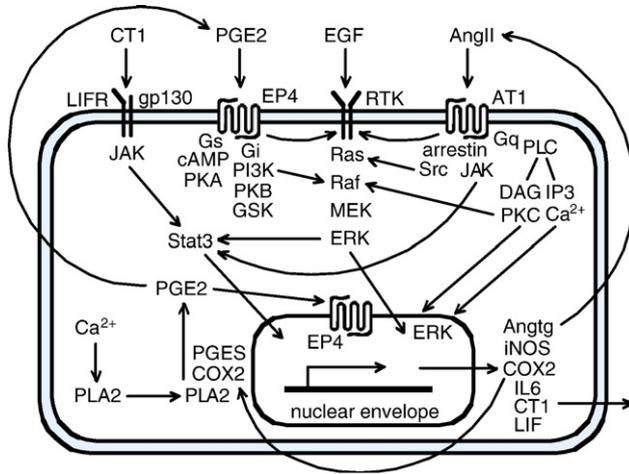


Fig. 1. Simplified scheme of the major routes connected with PGE<sub>2</sub> signaling. PGE<sub>2</sub> is synthesized by enzymes on the nuclear envelope surface. The PGE<sub>2</sub> receptor subtype EP4 locates to the cell membrane and to the nuclear envelope. The EP4 receptor is able to couple with either Gs or Gi proteins; when coupling to Gi, the heterodimeric Gβ/γ component is thought to activate the PI3K-PKB path [13]. The PI3K-dependent activation of the MEK-ERK pathway is not yet defined in detail. Angtg is intracellularly transformed to AngII before secretion. The AngII receptor AT1 may signal via Gq as well as by direct interaction with several adapter proteins including β-arrestin, Src, and JAK. Both EP4 and AT1 receptors can transactivate the EGF receptor (a receptor tyrosine kinase=RTK). ERK1/2 occurs in the cytosolic and nuclear fractions. The IL6-related cytokines may be secreted and act in a paracrine and autocrine manner while PGE<sub>2</sub> may also act in an intracrine manner on the nuclear EP4 receptors. For full transcriptional activity Stat3 requires phosphorylation at Tyr705 by JAK and at Ser727 by ERK. Abbreviations are in the text except for: DAG, diacylglycerol; GSK, glycogen synthase kinase; iNOS, inducible NO synthase; PI3K, phosphoinositide 3-kinase.

inhibitors and activators of these pathways. Furthermore, transfection of VNRC with small interfering RNA (siRNA) specifically targeting the rat Stat3 reduced expression of the latter by ~70% and inhibited the hypertrophic reaction. The requirement of 1 μM PGE<sub>2</sub> for eliciting the hypertrophic reaction observed in VNRC seems rather high in view of the ~10,000-fold lower physiological plasma levels (60–110 pM) in the rat [7]. The authors may argue, however, that the intracellular biosynthesis of PGE<sub>2</sub> could lead to significantly higher local concentrations than that found extracellularly in the plasma. Blockade of the delayed Tyr705 phosphorylation of Stat3 by the transcription inhibitor actinomycin-D as well as by the translation inhibitor cycloheximide [6] suggests that in VNRC stimulated by PGE<sub>2</sub> the phosphorylation of Stat3 at Tyr705 is not the direct target of ERK1/2 although Ser727 of Stat3 was shown to serve as substrate for ERK1/2 in other cell systems [2]. Bearing in mind that genistein abolishes Stat3 phosphorylation, it seems that an as yet undefined intermediate tyrosine kinase may originate from de novo protein synthesis. Obviously, the PGE<sub>2</sub> signaling in VNRC between EP4 and MEK1/2 as well as between ERK1/2 and Stat3 requires further exploration [6]. Here we tentatively

review possible candidates involved in knocking on Stats' door.

### 3. The JAK/Stat pathway

Most hypertrophic stimuli such as stretch, angiotensin-II (AngII), IL6, CT1, and LIF activate the latent cytoplasmic transcription factors Stat1 and Stat3. Downregulation of Stat3 is associated with end-stage heart failure, and its activation promotes cardiomyocyte survival and hypertrophy, while Stat1 correlates with pro-inflammatory responses and apoptosis [8,9]. In the canonical JAK/Stat signaling pathway, binding of the IL6-related cytokines induces dimerization of their receptors (gp130 and LIFR) and tyrosine phosphorylation in the cytoplasmic domain by the receptor-associated tyrosine JAK1/2 kinases for recruitment of specific Stats from the cytoplasm. After phosphorylation of Stat1 at Tyr701 and Stat3 at Tyr705, they dimerize (homo- or heterodimers) and translocate to the cell nucleus where they associate with the transcription machinery (Fig. 1). For full transcriptional activity Stat1 and Stat3 require additional phosphorylation of Ser727 in the transactivation domain, which represents a recognition site for ERK1/2, p38 MAPK, and other yet to be defined serine kinases [12]. In isolated mouse hearts rapid phosphorylation of Ser727 of Stat1/3 was observed resulting from the signaling cascade PKCε-Raf1-MEK1/2-ERK1/2 [10].

### 4. How may Stat3 be integrated into PGE<sub>2</sub> signaling for cardiac hypertrophy?

In the nucleus Stat3 stimulates the COX2 gene and COX2 protein expression, and the resulting increase in PGE<sub>2</sub> levels further stimulates expression of COX2 in a positive feedback loop (Fig. 1). This feedback loop may not only operate in paracrine or autocrine but also in the intracrine mode since both COX2 and the PGE<sub>2</sub> receptor EP4 locate primarily to the nuclear envelope [3,5]. Stat3 is also known to enhance expression of the angiotensinogen (Angtg) gene and subsequent generation of AngII [11,12]. AngII induces the release of the IL6-related cytokines (IL6, CT1, LIF), which, in turn, activate the canonical JAK/Stat signaling cascade leading to delayed tyrosine phosphorylation of Stat3, thus establishing an autocrine/paracrine loop (Fig. 1). AngII-induced hypertrophy is thought to primarily depend on the activation of its Gq-coupled receptor (AT1). However, the AT1 receptor is also able to activate several additional downstream signaling molecules including the Src family protein tyrosine kinases and ERK1/2 through G-protein-independent mechanisms [13]. In particular, AT1 may directly interact with JAK2, the SHP2 tyrosine phosphatase, phospholipase-C (PLC), and other proteins serving as signaling adapters and scaffold components. Furthermore, both AT1 and EP4 are reported to transactivate the EGFR, which may also signal via the ERK1/2 pathway (Fig. 1). Thus, multiple signaling tracks join the MEK-ERK pathway,

which emerges as the major communication route. Multifarious direct and indirect (involving de novo protein synthesis) signaling loops connecting the PGE<sub>2</sub>-EP4, JAK/Stat, and MEK-ERK systems may account for fast as well as delayed activation by phosphorylation cascades. Despite many advances in our understanding of the intracellular signaling network, cardiac hypertrophy and heart failure remain a formidable challenge since an effective causative treatment strategy is still lacking.

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