

Cyclooxygenase pathways

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This review compiles the current knowledge on the effects of prostanoids — arachidonic acid metabolites — on their own synthesis, activity and degradation. Interaction mechanisms between the receptors for the relevant compounds are presented, in particular with regard to the cooperation between a thromboxane A₂ and prostaglandin I₂ receptors. The questions of desensitization and internalization of receptors are discussed. The stages of the inflammatory response and tumor progression are analyzed against the background of the disruption of the synthesis of prostanoids. Special attention is given to the significance of 15-deoxy-Δ^{12,14}-prostaglandin J₂ in the regulation of the synthesis of prostanoids and its role as an anti-inflammatory agent. Ultimately, therapeutic approaches as used in various treatments are discussed in the light of the available knowledge.

Key words: cyclooxygenase, prostaglandin, 15-deoxy-Δ^{12,14}-prostaglandin J₂, receptor desensitization

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BIOSYNTHESIS AND BIOLOGICAL IMPORTANCE OF PROSTANOIDS

Prostanoids are tissue hormones synthesized from long-chain polyunsaturated fatty acids, mainly arachidonic acid (AA) (Sales *et al.*, 2008). Phospholipase A₂ (PLA₂), and to be more precise cytoplasmic phospholipase A₂ (cPLA₂), is the first enzyme engaged in synthesizing these compounds, as it specifically releases AA from lipids in the cell membrane, which then enters cyclooxygenase pathways (Fig. 1) (Linkous & Yazlovitskaya; 2010).

The two main COX isoforms are cyclooxygenase-1 (COX-1) and cyclooxygenase-2 (COX-2). Both transform AA into prostaglandin H₂ (PGH₂), which then again transformed by proper synthases, mainly cytosolic prostaglandin E synthase (cPGES), microsomal prostaglandin E synthase-1 (mPGES-1), microsomal prostaglandin E synthase-2 (mPGES-2), prostaglandin I synthase (PGIS), and thromboxane synthase (TxS) into prostaglandins (PGs) and thromboxanes.

The *COX-1* gene is principally constitutive in function and possesses a typical, GC-rich housekeeping promoter. In contrast, the *COX-2* gene resembles an early response gene. It is strongly induced by mitogenic and proinflammatory stimuli, superinduced by inhibitors of protein synthesis, and acutely regulated at both transcriptional and posttranscriptional levels (Lasa *et al.*, 2000).

The prostanoids of noteworthy biological significance include: prostaglandin E₂ (PGE₂), prostaglandin F_{2α} (PGF_{2α}), prostaglandin D₂ (PGD₂), 15d-PGJ₂, PGI₂, and

TxA₂. PGE₂ is an important proinflammatory prostaglandin revealing antiapoptotic properties, which plays essential functions in renal physiology, the physiology of the immune system and is a developmental and growth factor associated with neoplasms (Harris, 2013). While PGF_{2α} is synthesized in the uterine wall and plays a crucial role not only in the physiology of this organ, it is also an autocrine growth factor in connection with endometrial adenocarcinoma (Sales *et al.*, 2008; Woodward *et al.*, 2011). A dehydrated derivative of PGD₂ (15d-PGJ₂) is a principal prostaglandin which inhibits prostanoid synthesis and the intensity of inflammatory reactions (Surh *et al.*, 2011). Other important products of arachidonic acid transformations include TxA₂ and PGI₂, synthesized by COX-1 and COX-2, respectively (Catella-Lawson *et al.*, 1999). Both these prostanoids play important roles in the physiology and pathology of blood vessels (Caughey *et al.*, 2001; Debey *et al.*, 2003; Meyer-Kirchrath *et al.*, 2004). TxA₂ causes the aggregation of blood platelets and the contraction of blood vessels, whereas PGI₂ has opposite properties.

PROSTANOID SYNTHESIS IN THE COURSE OF INFLAMMATORY RESPONSE

Prostanoids play a chief role in inflammatory reactions. The synthesis of these compounds differs depending on the inflammatory reaction phase. During the first hours after the activation by a the proinflammatory sub-

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Abbreviations: 15-PGDH, 15-hydroxyprostaglandin dehydrogenase; 15d-PGJ₂, 15-deoxy-Δ^{12,14}-prostaglandin J₂; AA, arachidonic acid; AC, adenylate cyclase; AP-1, activator protein 1; CaMK-II, Ca²⁺/calmodulin-dependent protein kinase II; CBP, CREB-binding protein; COX, cyclooxygenase; COX-1, cyclooxygenase-1; COX-2, cyclooxygenase-2; cPLA₂, cytoplasmic phospholipase A₂; CRE, cAMP response element; CREB, cAMP response element-binding protein; CRTH2, chemoattractant receptor-homologous molecule expressed on Th2 cells; DP, prostaglandin D₂ receptor; EGFR, epidermal growth factor receptor; eNOS, endothelial nitric oxide synthase; EP, prostaglandin E₂ receptor; ERK1/2, extracellular-signal-regulated kinases 1 and 2; FP, prostaglandin F_{2α} receptor; GRK, G protein-coupled receptor kinases; IKK, IκB kinase; IKKβ, IκB kinase subunit β; IP, prostaglandin I₂ receptor; IP₃, inositol trisphosphate; JNK, c-Jun N-terminal kinase; KO, knockout; LPS, lipopolysaccharides; MAPK, mitogen-activated protein kinase; mPGES, microsomal prostaglandin E synthase; MRP4, multidrug resistance-associated protein 4; NF-κB, nuclear factor κB; NSAID, non-steroidal anti-inflammatory drug; PG, prostaglandin; PGIS, prostaglandin I synthase; PGT, prostaglandin transporter; PI3K, phosphatidylinositol-4,5-bisphosphate 3-kinase; PKA, protein kinase A; PKB, protein kinase B; PKC, protein kinase C; PLA₂, phospholipase A₂; PLCβ, phospholipase Cβ; PPARγ, peroxisome proliferator-activated receptor γ; PTEN, phosphatidylinositol-3,4,5-trisphosphate 3-phosphatase; ROS, reactive oxygen species; sPLA₂, secreted phospholipase A₂; TP, thromboxane A₂ receptor; TPα, thromboxane A₂ receptor α isoform; TPβ, thromboxane A₂ receptor β isoform; TxA₂, thromboxane A₂; TxS, thromboxane synthase.

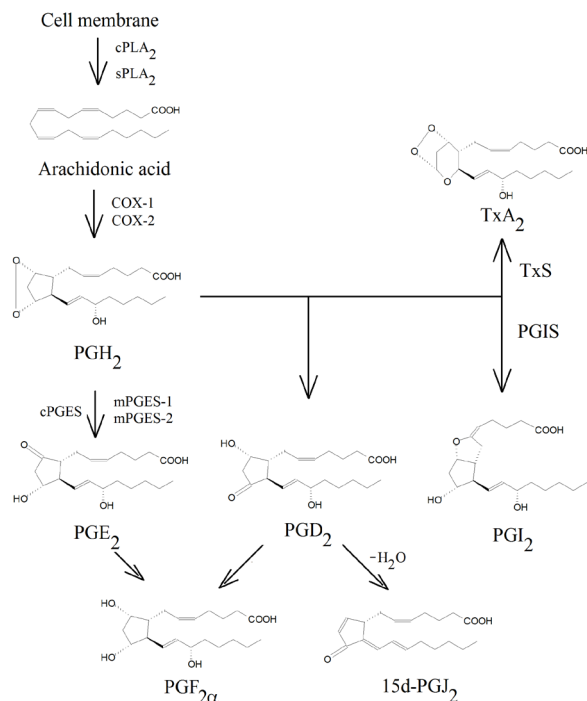


Figure 1. COX pathways.

Arachidonic acid is released from cell membrane by cPLA₂. It can then undergo transformations by cytochrome P450, lipoxygenases or COX. The latter enzymes transform arachidonic acid into PGH₂. The newly formed prostaglandin is transformed into other prostaglandins or TxA₂ by appropriate synthases.

stance COX-2 expression increases and cPLA₂ is activated, which leads to increased PGH₂ synthesis (Murakami *et al.*, 1997). However, a simultaneous lack of mPGES-1 expression, transforming PGH₂ into PGE₂, results in a coincident increase of PGD₂ synthesis (Xiao *et al.*, 2012). Nonetheless, the COX-2 expression in this phase does not depend on the autocrine activation by PGE₂ (Murakami *et al.*, 1997).

During the second phase, lasting for 12 to 48 hours after the stimulation, mPGES-1 and cPLA₂ expression takes place, which intensifies PGE₂ synthesis (Murakami *et al.*, 1997; Xiao *et al.*, 2012) with an unchanged synthesis of PGD₂ (Xiao *et al.*, 2012). Expression of COX-2 and cPLA₂ undergoes autocrine stimulation by PGE₂, but there is no simultaneous influence of this compound on the expression of mPGES-1 (Murakami *et al.*, 1997; Xiao *et al.*, 2012). The addition of inhibitors of COX-2 and/or cPLA₂ results in a simultaneously decreased expression of both enzymes (Murakami *et al.*, 1997; Xiao *et al.*, 2012), while adding PGE₂ suppresses the effect of these inhibitors on the COX-2 and cPLA₂ expression (Murakami *et al.*, 1997). Besides, in the course of this inflammatory phase, depending on its duration, the expression of COX-2 can be hampered by 15d-PGJ₂ (Inoue *et al.*, 2000).

DYSREGULATION OF THE PROSTANOID SYNTHESIS PATHWAY IN NEOPLASTIC DISEASES

PGE₂ plays an important role in cancer development. It shows antiapoptotic activity and supports angiogenesis, which is why it could be a promising target in an antineoplastic therapy (Greenhough *et al.*, 2009).

The pathways of prostanoid synthesis tend to become dysregulated with the progression of a neoplastic disease (Table 1). First of all, the expression of COX-2 and mPGES-1 is increased and consequently the intensity of PGE₂ production increased (Badawi & Badr, 2003; Nakanishi *et al.*, 2008). Apart from the amplifying inflammatory reactions, the anti-inflammatory mechanisms are dysregulated. Expression of peroxisome proliferator-activated receptor γ (PPAR γ), which participates in the inhibition of inflammatory reactions, is decreased due to an increased COX-2 expression (Inoue *et al.*, 2000). Also the synthesis of 15d-PGJ₂ decreases as a result of the rise in mPGES-1 expression (Badawi & Badr, 2003). The expression of mPGES-1 causes an increased PGE₂ synthesis, as mentioned above, at the cost of PGD₂, which leads not only to the deactivation of the function of 15d-PGJ₂, but also supports angiogenesis and the progress of neoplastic diseases (Murata *et al.*, 2011; Davoine *et al.*, 2013).

The changes in the prostanoid synthesis in neoplastic cells occur along with changes in the enzymes degrading those compounds. Neoplastic cells reveal an entire lack or a decreased expression of 15-hydroxyprostaglandin dehydrogenase (15-PGDH), a PGE₂ degrading enzyme (Wolf *et al.*, 2006; Jang *et al.*, 2008; Lee *et al.*, 2013). The mechanism underlying the expression disorders of 15-PGDH may partially result from the increased expression of COX-2 (Tong *et al.*, 2006). In colorectal neoplasia a lowered expression of the prostaglandin transporter (PGT) participating in uptaking PGs is often noted, as well as an increased expression of multidrug resistance-associated protein 4 (MRP4) taking part in the secretion of PGs from the cell (Holla *et al.*, 2008). Furthermore, the expression of prostaglandin receptors also increases. An increased expression of prostaglandin E₂ receptor-4 (EP₄) has been observed, among others, on colorectal neoplastic cells (Chell *et al.*, 2006). All these changes result in an increased concentration of PGE₂ in the surroundings of neoplastic cells. This causes an autocrine activation of neoplastic development and the progress of the disease.

INFLAMMATORY REACTION POSITIVE FEEDBACK LOOP INVOLVING PROSTAGLANDIN E₂

COX-2 expression is in a positive feedback loop with its products a PGE₂ and PGF_{2 α} (Inoue *et al.*, 2000; Sales *et al.*, 2001; 2008). This process is of considerable significance as far as autocrine support of inflammatory reactions and activation of neoplastic cells' growth (Sales *et al.*, 2001; 2008). The mechanism associated with the way prostaglandins influence COX-2 expression consists in increasing the COX-2 mRNA stability and in activating the COX-2 gene promoter (Sales *et al.*, 2008). Both these processes depend on the type of cells (Inoue *et al.*, 2000; Sales *et al.*, 2001; 2008).

PGE₂ increases the expression of COX-2 acting through its receptors, prostaglandin E₂ receptor-2 (EP₂) and EP₄, which activate adenylate cyclase (AC), and this increases the concentration of cAMP in the cell (Sakuma *et al.*, 2004). In turn, cAMP increases the stability of COX-2 mRNA and activates COX-2 promoter by means of protein kinase A (PKA) and cAMP response element-binding protein (CREB) (Inoue *et al.*, 2000; Sales *et al.*, 2001; Fujino *et al.*, 2005; Díaz-Muñoz *et al.*, 2012).

Activated EP₄ also activates phosphatidylinositol-4,5-bisphosphate 3-kinase (PI3K), which causes a PKA-independent signal transduction to CREB (Fujino *et al.*, 2005).

Table 1. Prostanoid pathway compounds under three conditions.

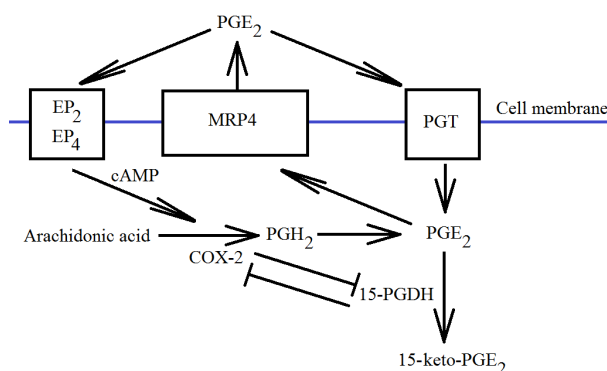
Comparison of three conditions: normal condition, inflammation caused by a pro-inflammatory factor, and cancer

Trait	Normal condition	Inflammation	Cancer cell
COX-2 protein expression	none	present	present
mPGES-1 protein expression	none	present	present
PGE ₂ level	low	high	high
15-PGDH protein expression	high	low	none
PGD ₂ level	normal	high	low
15d-PGJ ₂ level	normal	high	low

PGE₂ also increases COX-2 expression by activating mitogen-activated protein kinase (MAPK) cascades (Faour *et al.*, 2001; Rösch *et al.*, 2005). EP₄ uses PI3K to activate the extracellular-signal-regulated kinases 1 and 2 (ERK1/2) MAPK cascade, leading to increased COX-2 expression (Fujino *et al.*, 2003; 2005; Mendez & LaPointe, 2005; Rösch *et al.*, 2005; Sales *et al.*, 2008). Protein kinase C (PKC) and Ca²⁺ can also participate in the activity of PGE₂ (Rösch *et al.*, 2005). The additionally activated EP₄ receptor, through a pathway independent of PKA, activates the p38/MAPKAPK-2/hsp 27 cascade which is responsible for increasing the stability of COX-2 mRNA (Lasa *et al.*, 2000; Faour *et al.*, 2001; Rösch *et al.*, 2005). The stability of COX-2 mRNA is enhanced by preventing its degradation from the 3'-untranslated region of mRNA rich in AU nucleotides (Lasa *et al.*, 2000; Faour *et al.*, 2001).

DEGRADATION PATHWAY OF PROSTAGLANDIN E₂

After synthesis PGE₂ is secreted outside the cell by MRP4. During the following phase, it can be uptaken by PGT and degraded in the cytoplasm (Fig. 2) (Veazza *et al.*, 2001). This particular transporter specifically transports not only PGE₂, but also PGD₂ and PGF_{2α} (Veazza *et al.*, 2001). This is followed by the degradation of PGE₂ in the cytoplasm by 15-PGDH into an inactive 15-keto-PGE₂ form (Holla *et al.*, 2008). Products of COX transformations take part in regulating the PGE₂ degradation pathway.

**Figure 2. PGE₂ degradation.**

PGE₂ is secreted from cells by MRP4. Outside the cells, PGE₂ activates its receptors EP₂ and EP₄, thus increasing COX-2 expression. It may also undergo uptake by means of PGT in the cytoplasm, PGE₂ is degraded by 15-PGDH to 15-keto-PGE₂. COX-2 and 15-PGDH, the key enzymes in the synthesis and degradation of PGE₂, mutually inhibit each other's expression, an increase in the expression of one enzyme leads to reduced expression of the other.

An activated prostaglandin F_{2α} receptor (FP) reduces the PGT activity, and hence the uptake of prostaglandin through this transporter is reduced (Veazza *et al.*, 2001). This regulation occurs through the G_s protein, but is independent of AC (Veazza *et al.*, 2001). Additionally, the levels of COX-2 and 15-PGDH expressions are directly related to each other. Increased expression of one enzyme inhibits the expression of the other one (Tong *et al.*, 2006; Liu *et al.*, 2008). Nevertheless, the dependence between the expression of both enzymes is independent of their activity. Adding an inhibitor of one enzyme does not affect the expression of the other (Tong *et al.*, 2006). The mechanism interconnecting the levels of COX-2 and 15-PGDH expression remains unclear. Most probably 15-PGDH is capable of binding 3'-untranslated region of mRNA COX-2 and, thereby destabilizing it (Tong *et al.*, 2006). The reciprocal influence of COX-2 on 15-PGDH expression, is even less well understood and is still being that examined in order to understand the regulation of expression of these enzymes in neoplastic cells, where COX-2 overexpression and reduced 15-PGDH expression is observed.

MECHANISMS OF PROSTAGLANDIN F_{2α}-INDUCED CYCLOOXYGENASE-2 EXPRESSION

PGF_{2α} is yet another prostanoid with a crucial role in the regulation of the synthesis of the prostanoids discussed in this review. It increases the expression of COX-2 through its receptor (FP) and hence increases its own synthesis (Sales *et al.*, 2008). The autocrine activation of this enzyme's expression by PGF_{2α} is of considerable importance for the growth and development of endometrial adenocarcinoma. When present in concentrations reaching 100 nM PGF_{2α} may even activate EP₂ (Fig. 3), leading to the same signal transduction as that caused by PGE₂ (Sales *et al.*, 2008). FP activates phospholipase Cβ (PLCβ) which releases inositol trisphosphate (IP₃) and diacylglycerol (Sales *et al.*, 2008; Woodward *et al.*, 2011) which activate PKC, however, this pathway will not induce the expression of COX-2 protein (Sales *et al.*, 2008). PGF_{2α} binds to FP, that caused signal transduction, which activates ERK1/2 MAPK and hence results in COX-2 expression (Jabbour *et al.*, 2005; Sales *et al.*, 2008).

Whilst PLCβ activates ERK1/2 MAPK by means of epidermal growth factor receptor (EGFR), which leads to expression of COX-2 (Sales *et al.*, 2004; 2008), it seems that also PKA also participates in transducing signals from FP to ERK1/2 MAPK (Sales *et al.*, 2008). As for EGFR activation, c-Src and metalloproteinases may also participate in this process (Sales *et al.*, 2005). It has also been shown that the upregulation of COX-2 expression results from activating CREB and ac-

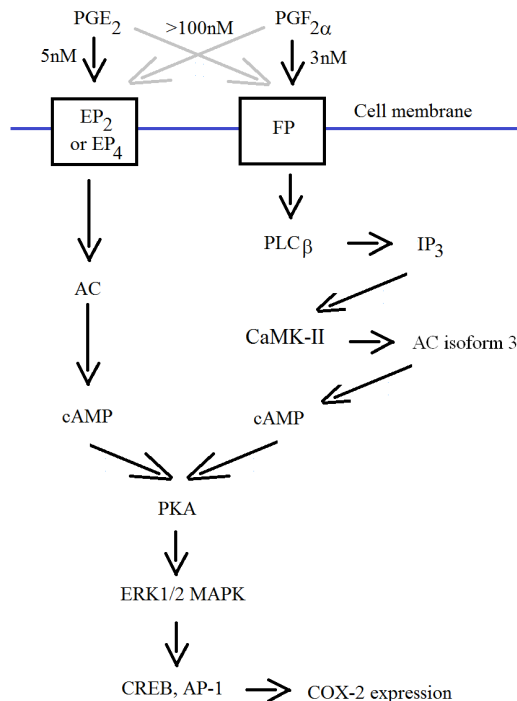


Figure 3. Mechanism of COX-2 expression stimulation by PGE₂ and PGF_{2α}.

Both prostaglandins activate their respective receptors: EP₂, EP₄ and FP. At larger levels of the order of 100 nM, PGE₂ is also capable of activating the FP receptor. EP₂ and EP₄ cause activation of AC and transduction of the signal to PKA. At the same time, FP activates PLCβ, products of which are involved in activation of PKC causing influx of calcium ions. This may have a synergistic effect in the EP₂ and EP₄-mediated activation of AC. Eventually, the signal is transduced to ERK1/2 MAPK, leading to activation of COX-2 promoter by CREB and AP-1.

tivator protein 1 (AP-1) by means of ERK1/2 MAPK (Sales *et al.*, 2008).

The affinity of PGE₂ for FP is lower (119 nM) than its affinity for EP₂ (4.9 nM) or EP₄ (0.79 nM) (Abramovitz *et al.*, 2000). In concentrations reaching 100nM, PGE₂ increases the expression of COX-2 not only by its EP₂ and EP₄ receptors but also through FP (Sales *et al.*, 2008). When increasing the expression of COX-2, activated EP and FP receptors act synergistically, as they activate ERK1/2 MAPK through PKA (Sales *et al.*, 2008). The calcium influx, dependent on IP₃, activates Ca²⁺/calmodulin-dependent protein kinase II (CaMK-II) and ultimately the transmission of signal to Ca²⁺ dependent AC isoform 3 (Abera *et al.*, 2010). The cross-activation of EP and FP receptor by PGE₂ and PGF_{2α} is crucial in neoplastic cells expressing FP as in endometrial adenocarcinoma (Sales *et al.*, 2008; Woodward *et al.*, 2011). The both prostaglandins increase COX-2 expression, which results in an autocrine upregulation of their level. This, again, leads to the development and growth of the neoplasm (Sales *et al.*, 2005; 2008).

15-DEOXY-Δ^{12,14}-PROSTAGLANDIN J₂ AS AN ANTI-INFLAMMATORY REGULATOR OF PROSTANOID SYNTHESIS

Apart from the above-mentioned PGE₂ and PGF_{2α}, other prostaglandins also exert an autocrine influence on prostanoid synthesis. 15d-PGJ₂ is a prostanoid regulating the intensity of inflammatory reactions and reducing

the pace of neoplastic growth and development (Surh *et al.*, 2011). It is formed by non-enzymatic dehydration of PGD₂ (Surh *et al.*, 2011).

By means of their receptors, respectively EP₂ and EP₄, as well as the prostaglandin D₂ receptor (DP), PGE₂ and PGD₂ increase the cAMP concentration and thereby enhance the expression of COX-2 protein (Sakuma *et al.*, 2003). After a certain period of time, however free PGD₂ becomes dehydrated to 15d-PGJ₂, which in low concentrations reduces the expression of COX-2 protein (Surh *et al.*, 2011) through a pathway that can be dependent or independent on the PPARγ receptor (Fig. 4) (Inoule *et al.*, 2000; Sawano *et al.*, 2002). In cells treated with a proinflammatory factor, the activated PPARγ receptor disturbs the induction of COX-2 protein expression along various routes. In particular, it disturbs the activation of nuclear factor κB (NF-κB) (Inoule *et al.*, 2000). Additionally, PPARγ disrupts the activation of the AP-1 transcription factor by means of c-Jun N-terminal kinase (JNK) MAPK and binds, and thereby inactivates, CREB-binding protein (CBP)/p300 (Subbaramaiah *et al.*, 2001). It has also been shown that 15d-PGJ₂ inhibits signal transduction through ERK1/2 and JNK MAPK into AP-1, along a pathway independent of PPARγ receptor (Sawano *et al.*, 2002). Additionally, 15d-PGJ₂ inhibits the activation of NF-κB by restraining the activity of IκB kinase subunit β (IKKβ) by modifying its cysteine residues (Boyault *et al.*, 2004). The disrupted phosphorylation precludes the degradation of the NF-κB inhibitor: IκBα (Boyault *et al.*, 2004). Apart from that, 15d-PGJ₂ modifies cysteine residues in the DNA binding domain on p65 NF-κB, which leads to its inactivation (Straus *et al.*, 2000; Boyault *et al.*, 2004).

The molecular mechanism of the modification of cysteine residues by 15d-PGJ₂ is well known. The arrangement of atoms in the 15d-PGJ₂ ring creates an

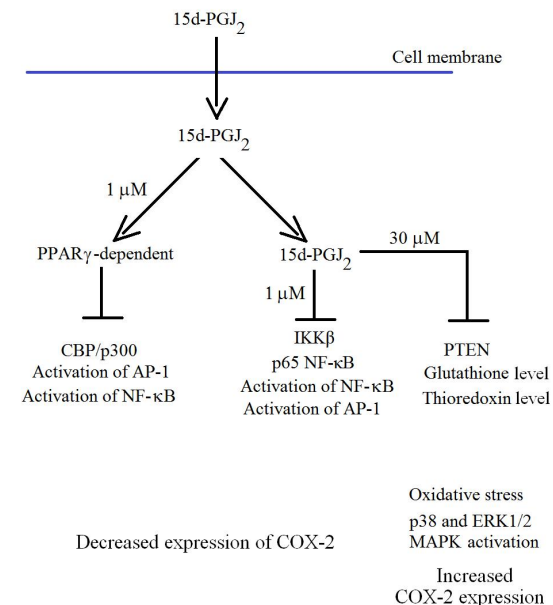


Figure 4. Effects of 15d-PGJ₂ on COX-2 expression.

15d-PGJ₂ undergoes cytoplasmic uptake and activates its receptor PPARγ. This leads to inhibition of the activity of NF-κB, JNK MAPK and CBP/p300. Another pathway of 15d-PGJ₂ activity is direct inhibition of p65 NF-κB and IKKβ. Both pathways decrease COX-2 protein expression. At very large levels, 15d-PGJ₂ deactivates antioxidants containing free -SH groups. By this mechanism, it sensitizes the cells to the effects of ROS, which may lead to an increase in COX-2 protein expression.

Table 2. Properties of the prostanoids receptors.

The receptors are categorized by their ligands and activated pathways. The main activated enzymes are AC or PLC. Receptors are desensitized or internalized following phosphorylation by PKC, PKA and GRK.

Receptor	Ligand	Main activation effect	PKA activation	PKC activation	Receptor desensitization/internalization pathway
EP ₁	PGE ₂	PLC, ↑Ca ²⁺ , no IP ₃	-	+	PKC
EP ₂	PGE ₂	AC, ↑cAMP	+	-	Not affected
EP ₃	PGE ₂	AC, ↓cAMP	-	-	GRK(EP _{3a})
EP ₄	PGE ₂	AC, ↑cAMP, PI3K	+	-	GRK, PKC*
DP	PGD ₂	AC, ↑cAMP, ↑Ca ²⁺	+	-	PKC*, GRK
CRTH2	PGD ₂	AC, ↓cAMP, ↑Ca ²⁺	-	+	PKC, PKA*, GRK
IP	PGI ₂	AC, ↑cAMP, PLC	+	+	PKC, PKA(AC isoform 5 and 6)
FP	PGF _{2a}	PLC, ↑Ca ²⁺	-	+	PKC(FP _A), constitutive FP _B
TP	TxA ₂	PLC, ↑Ca ²⁺	-	+	GRK, PKA(TPα)*, PKC

*cross-desensitization/internalization pathway triggered by an other type of receptor

electrophilic carbon (Straus *et al.*, 2000) which undergoes Michael addition with cysteine residues (R-SH) (Straus *et al.*, 2000). The reaction with cysteines in the catalytic centre, can lead to the inactivation of an enzyme, such as IKKβ. As a consequence, this evokes the reduction of COX-2 protein expression.

Moreover, 15d-PGJ₂ affects the expression of COX-2 through mechanisms related to the synthesis of reactive oxygen species (ROS), since at low concentrations 15d-PGJ₂ activates cellular defense mechanisms against ROS and at the same time increases the concentration of antioxidants, e. g., glutathione (Koppal *et al.*, 2000). It also increases the level of heme oxygenase 1 expression (Koppal *et al.*, 2000). An increased level of antioxidants in the cell disrupts signal transduction along pathways where ROS function as second messengers. This disturbs transduction of NF-κB activation signals and activation of MAPK kinase cascades responsible for COX-2 expression (Koppal *et al.*, 2000).

Paradoxically, at high concentrations 15d-PGJ₂ may actually increase the expression of COX-2. In breast cancer cells, 30 μM 15d-PGJ₂, increases the expression of COX-2 by inhibiting the activity of phosphatidylinositol-3,4,5-trisphosphate 3-phosphatase (PTEN) (Kim *et al.*, 2008). This causes the activation of protein kinase B (PKB) and, signal transduction upregulating COX-2 expression (Kim *et al.*, 2008). 15d-PGJ₂ can also activate p38 MAPK and, by means of EGFR, also of the ERK1/2 MAPK cascade, which leads to the induction of mRNA COX-2 synthesis and also increases its stability (Kitz *et al.*, 2011). This mechanism resembles the route by which COX-2 expression is induced by ROS.

15d-PGJ₂ and ROS reveal a similar activity in relation to enzymes, by modifying cysteine residues in their catalytic centres. ROS create disulphide bonds or sulfenic acid (Cys-SOH), which often inhibits the activity of enzymes with a cysteine in their catalytic centre (Meng & Zhang, 2013). When 15d-PGJ₂ reacts with a cysteine residue, it also inhibits the activity of enzymes (Straus *et al.*, 2000; Boyault *et al.*, 2004). This is why the both compounds can have similar effects when present at high concentrations. Additionally, 15d-PGJ₂ inactivates glutathione and thioredoxin, sensitizing the cell to oxida-

tive stress and second messengers, such as ROS (Shibata *et al.*, 2003; Kim *et al.*, 2008). This effect may be relieved by increasing the concentration of antioxidants with free R-SH groups which react directly with 15d-PGJ₂ (Kim *et al.*, 2008).

The regulation of COX-2 expression is a dynamic process. On one hand it is inhibited by the PGD₂ metabolite 15d-PGJ₂, while on the other it is activated by PGE₂. The two prostanoids, 15d-PGJ₂ and PGE₂, form a more complex system of interdependencies. Following the treatment of cells with an inflammatory factor, such as lipopolysaccharides (LPS), the level of PPARγ is reduced, which leads to partial deactivation of 15d-PGJ₂ and simultaneously to increased expression of COX-2 (Inoule *et al.*, 2000). Conversely, blocking the activity of mPGES-1, an enzyme collaborating with COX-2 in the reduction of PGE₂, increases expression of PPARγ (Kapoor *et al.*, 2007). This effect is caused by signal transduction from the PGE₂ receptor to PI3K and PKB, which in turn inhibit the expression of PPARγ (Kapoor *et al.*, 2007).

LIGAND-MEDIATED DESENSITIZATION AND INTERNALIZATION OF RECEPTORS

Desensitization means reducing the sensitivity of a receptor in response to a prolonged exposure to a given ligand. This process plays an important role in the physiology of receptors, especially when reducing the cellular response to a particular factor, which may become too intense.

Exposure to PGE₂ results in internalization and desensitization of EP₄ receptor, whilst EP₂ maintains its sensitivity towards PGE₂ (Table 2) (Desai *et al.*, 2000). The C-tail plays a key role in desensitization of EP₄, as it is phosphorylated by G protein-coupled receptor kinases (GRK) (Desai *et al.*, 2000; Slipetz *et al.*, 2001). Throughout this process, PKA does not exert any influence on the EP₄ (Neuschäfer-Rube *et al.*, 1999; Slipetz *et al.*, 2001). PKC is able to influence EP₄, but this kinase is not activated by EP₄ receptor (Neuschäfer-Rube *et al.*, 1999; Slipetz *et al.*, 2001). As far as the rat prostaglandin E₂ receptor-3 (EP₃) is concerned, EP_{3a} undergoes desensitization by phosphorylation dependent on GRK,

whereas EP_{3β} does not (Neuschäfer-Rube *et al.*, 2005). Prostaglandin E₂ receptor-1 (EP₁) undergoes desensitization and internalization following of phosphorylation by PKC (Katoh *et al.*, 1995).

Aside from EP, other receptors also undergo desensitization and internalization. Receptors for PGF_{2α} are present in two isoforms: FP_A and FP_B. PGF_{2α} causes internalization of FP_A dependent on PKC, while FP_B undergoes partial, constitutive internalization independent of the ligand, since there are no phosphorylation sites for PKC in its C-tail (Srinivasan *et al.*, 2002).

PGD₂ has two receptors: DP and chemoattractant receptor-homologous molecule expressed on Th2 cells (CRTH2). Both these receptors have a C-tail susceptible to phosphorylation, which is essential for their internalization (Gallant *et al.*, 2007; Schröder *et al.*, 2009; Roy *et al.*, 2010). CRTH2 undergoes the internalization through phosphorylation with PKC and a GRK: GRK2, GRK5 and/or GRK6 (Roy *et al.*, 2010). DP internalization depends on PKC, but this kinase is not activated by DP (Gallant *et al.*, 2007). What is more, DP internalization depends only on GRK2 and not depend on GRK5 or GRK6 (Gallant *et al.*, 2007). Another difference between these two receptors lies in the fact that the internalization of CRTH2 depends on the activation of PKA (Gallant *et al.*, 2007). However, when this kinase is activated but CRTH2 is not, it does not result in the internalization of the receptor (Gallant *et al.*, 2007). The internalization of the receptors for PGD₂ also depends on arrestins: that of DP on arrestin-2 and -3, and of CRTH2 on arrestin-3 only (Gallant *et al.*, 2007; Schröder *et al.*, 2009; Roy *et al.*, 2010).

PROSTAGLANDIN SYNTHESIS COMPENSATION IN CELLS WITH DISTURBED OF CYCLOOXYGENASE PATHWAYS

Gene knockout (KO) can serve as a model for the chronic use of COX inhibitors. Dysruption of either COX gene, COX-1 or COX-2, upregulates PGE₂ synthesis in cultured cells (Table 3) (Kirtikara *et al.*, 1998; Kanekura *et al.*, 2002). Cells with COX-1 or COX-2 defects show an increased expression of cPLA₂, secreted phospholipase A₂ (sPLA₂), mPGES-1, as well as the second, functional COX isoform, but they also exhibit by a decreased expression of mPGES-2 (Kirtikara *et al.*, 1998;

Zhang *et al.*, 2002; Bosetti *et al.*, 2004; Choi *et al.*, 2006; Sandee *et al.*, 2009). COX-2 KO cells do not respon to proinflammatory factors (Kirtikara *et al.*, 1998). In the mouse brain with a COX-2 knockout it a decreased concentration of PGE₂ is observed due to a non fully compensating increased COX-1 protein expression (Bosetti *et al.*, 2004). In macrophages, the COX-2 defect causes an increase in the expression of 5-lipoxygenase (Zhang *et al.*, 2002; Bosetti *et al.*, 2004).

Due to the production of TxA₂ by COX-1, the lack of this enzyme decreases the level of this eicosanoid as well as the level of TxS expression (Choi *et al.*, 2006). On the other hand, expression of COX-2 and synthesis of PGE₂ occurring under the influence of a proinflammatory factor in COX-1 KO mice was higher when compared with wild-type cells (Kirtikara *et al.*, 1998).

In brains of COX-1 KO mice the PGE₂ synthesis increases, as does COX-2 expression, which is caused by the activation of NF- κ B (Kanekura *et al.*, 2002; Choi *et al.*, 2006). This activation is due to an increased expression of NF- κ B (subunits p50 and p65) and next to constant activation of I κ B kinase (IKK) (Choi *et al.*, 2006). In contrast in COX2-KO animals, the basal level of NF- κ B activity is decreased (Rao *et al.*, 2005). In brain cells with COX2 KO the PGE₂ concentration is decreased, which may lead to reduced activation of NF- κ B (Poligone & Baldwin, 2001; Rao *et al.*, 2005; Choi *et al.*, 2006). Nevertheless, the exact mechanisms underlying the interdependence of COX-1 and COX-2 expressions require a more thorough research. It seems that the both enzymes inhibit their own expressions, and the lack of one causes a compensating increase in the expression of the second. Probably the expression of one COX isoform depends on the presence of mRNA or the protein of the other isoform. Another possibility is that products of COX pathways affect the expression of the other COX isoform.

Due to the pro-neoplastic character of the products of AA transformations by COXs and lipoxygenases, recent studies focus on the mechanisms of the inhibition of expression and activity of cPLA₂ α . In the course of inflammatory reactions, this enzyme releases AA from cell membranes lipids. Inhibiting the cPLA₂ α activity simultaneously inhibits the activity of COX and lipoxygenases. In mouse brain, cPLA₂ α KO leads to a reduction in COX-2 expression, which then effects in lower PGE₂ synthesis, while the expression of COX-1 and 5-lipoxygenase is unaffected (Bosetti & Weerasinghe, 2003; Sapirstein *et al.*, 2005). A decreased COX-2 expression may result from the inhibition of platelet-activating factor (PAF) synthesis (Serou *et al.*, 1999; Bosetti & Weerasinghe, 2003). PAF activates p38 and ERK1/2 MAPK, which increase the expression of COX-2 (Serou *et al.*, 1999). The expression of COX-2 is induced through the proinflammatory factor. In mice with cPLA₂ α defect, the expression of COX-1 decreases after proinflammatory factor treatment, while the induction of COX-2 expression and production of PGE₂ are disturbed (Sapirstein *et al.*, 2005). In contrast in prostate cancer models blocking the cPLA₂ α activity leads to increased expression of COX-1 and PGE₂ production, with no changes in COX-2 expression. Additionally, the secretion of lipoxygenase transformation products is reduced. Adding 5- and/or 12-hydroxyeicosatetraenoic acid, products of lipoxygenase transformations, reduced the level of COX-1 expression and PGE₂ production (Niknami *et al.*, 2010).

Inhibiting the activity of mPGES-1, an enzyme engaged in the production of PGE₂ dependent on COX-

Table 3. Prostanoid synthesis in COX KO cells.

COX KO cells are characterized by increased production of PGE₂, expression of phospholipase A₂ and of the other COX isoform.

Trait	COX-1 KO	COX-2 KO
PGE ₂ production	↑	↑
cPLA ₂ expression	↑	↑
sPLA ₂ expression	↑	↑
COX-1 expression		↑
COX-2 expression	↑	
In brain		
PGE ₂ concentration	↑	↓
mPGES-2 expression	↓	↓
NF- κ B activation	↑	↓

2, could be used as a therapeutic strategy in neoplasms. *mPGES-1* KO disturbs the synthesis of PGE₂ and other products of COXs pathways. Enzymatic defects in *mPGES-1*, reducing in the production of PGE₂ may be organ specific, e.g., they may concern solely the brain or the stomach. Simultaneously the production of PGI₂, TxA₂ and PGD₂ in the digestive system is increased (Boulet *et al.*, 2004; Elander *et al.*, 2008).

During the stimulation of the inflammatory reaction in *mPGES-1*-KO mice the synthesis of TxA₂, PGI₂, PGF_{2 α} and PGD₂ is increased, as is the expression of COX-2, when compared to cells with functional *mPGES-1* (Boulet *et al.*, 2004; Trebino *et al.*, 2005; Brenneis *et al.*, 2008; Elander *et al.*, 2008), most likely to compensate for the diminished level of PGE₂. The compensatory mechanisms is probably not connected with changes in the expression of particular synthases, but rather with the availability of PGH₂ (Boulet *et al.*, 2004). Since there is no *mPGES-1* expression, PGH₂ is available for transformation to proper prostanoids by other synthases. Application of *mPGES-1* inhibitors in neoplastic therapy in order to limit the synthesis of PGE₂ can thus cause an increased synthesis of other proneoplastic prostanoids, and as a consequence lead to effects contrary to those intended (Elander *et al.*, 2008).

THE IMPACT OF THROMBOXANE A₂ AND PROSTAGLANDIN I₂ ON PROSTANOID SYNTHESIS PATHWAY

TxA₂ and PGI₂ increase COX-2 expression, which leads to increased PGI₂ synthesis in blood vessels. Upon PGI₂ binding, prostaglandin I₂ receptor (IP) upregulates cAMP synthesis, which induces COX-2 expression in the smooth muscles of blood vessels (Debey *et al.*, 2003; Sakuma *et al.*, 2003; Meyer-Kirchath *et al.*, 2004). Increased expression of cAMP inducible early repressor occurs along with the increased COX-2 expression (Debey *et al.*, 2003). This transcriptional repressor binds cAMP response element-binding protein (CREB), and decreases the expression of genes including *COX-2* (Debey *et al.*, 2003).

TxA₂ affects the expression of COX-2 by a different mechanism. Due to its instability, TxA₂ functions only locally (Miyosawa *et al.*, 2006). TxA₂ binding to thromboxane A₂ receptor (TP) activates PLC β without affecting cAMP concentration. (Sakuma *et al.*, 2003; Woodward *et al.*, 2011). PLC β causes the induction of COX-2 expression through ERK1/2 MAPK and thus increases the synthesis of PGE₂ and PGI₂ in blood vessel cells (Caughey *et al.*, 2001; Chu *et al.*, 2003). The activation of ERK1/2 MAPK through ligand-activated TP is complex process involving PKC, PKA and PI3K (Miggin & Kinsella, 2002b).

Expression of COX-2 and synthesis of PGI₂ can suppress the activity of TxA₂ in prolonged exposure to this thromboxane. The detailed mechanism associated with activating the expression of COX-2 is very well known. One can distinguish two isoforms of receptors for TxA₂: thromboxane A₂ receptor isoform α (TP α) and β (TP β). The difference between these two receptors is associated with their C-tail located at the cytoplasmic side of the cell membrane (Miyosawa *et al.*, 2006). The two isoforms of TP can be associated with different G proteins and can therefore activate the MAPK cascade in different manners. Among others, PKA participates in the activation of ERK1/2 MAPK by TP α , but not by TP β (Miyosawa *et al.*, 2006). The signal transduction pathways

depend on the type cell as well as the domination of the described isoforms of the receptor (Miyosawa *et al.*, 2006). Most probably the activation of TP α leads to a minimally increased cAMP concentration to the activation of PKA, whereas the second isoform TP β has an entirely opposite influence on AC (Hirata *et al.*, 1996). TP α activates ERK1/2 MAPK within several minutes after TxA₂ binding, which is then followed by the deactivation of this cascade of kinases (Miggin & Kinsella, 2002b; Miyosawa *et al.*, 2006). While the other isoform, TP β , activates ERK1/2 MAPK after a similar period of time, this is a long-lasting process, since an hour later the phosphorylation of ERK1/2 MAPK is still present (Miggin & Kinsella, 2002b; Miyosawa *et al.*, 2006).

INTERACTIONS BETWEEN THROMBOXANE A₂ AND PROSTAGLANDIN I₂ RECEPTORS

The dimerization or oligomerization of receptors plays an important role in transmitting signals from TP or IP. It does not result in proper signal transduction, but can modulate the properties not only of the receptors, but also of the signals they transmit (Wilson *et al.*, 2004).

Expression of both TP isoforms can lead to the formation of their heterodimer. It probably takes place immediately after the receptors are synthesized and can later cause a reduction of TP α expression (Sasaki *et al.*, 2006). The TP heterodimer differs from the homodimers, because it produces a stronger signal in response to certain agonists (Wilson *et al.*, 2007b). Nevertheless, it is a simplified model, since it seems that TP receptors form both hetero- and homodimers, their oligomerization does not depend on the presence of a ligand (Laroche *et al.*, 2005). Individual receptors in an oligomer are connected with each other by disulphide bonds, which can be split only by reducing factors (Laroche *et al.*, 2005).

Apart from the fact that these two TP isoforms interact with and affect each other, they can also interact with IP. The receptors dimerization (TP and IP) and signal transduction takes place after the activation of these receptors. An activated TP α can dimerize with a nonin-

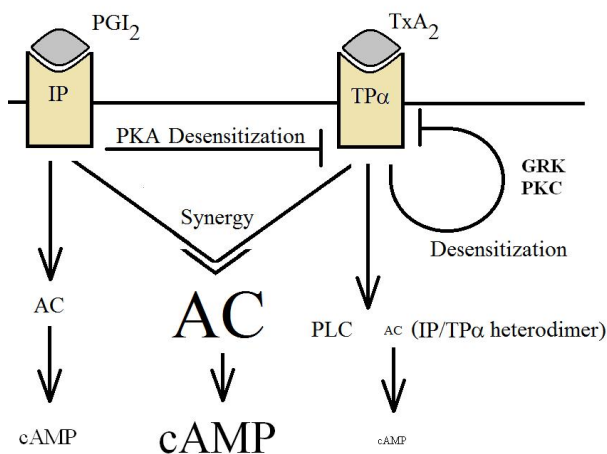


Figure 5. Dependence between IP and TP α .

The two receptors activate different signalling pathways. IP increases cytoplasmic cAMP level by activating AC while TP α activates PLC and AC in IP/TP α heterodimer. Both receptors act synergistically upon AC activation. This is probably due to heterodimerization which alters the signalling pathway activated by TP α . TP α is desensitized by GRK. In addition, ligand-activated IP triggers the same process in TP α by means of PKA.

activated IP and hence transmit a signal similar to that of an activated IP receptor, which means that it causes an increased cAMP concentration (Fig. 5) (Wilson *et al.*, 2004). Yet this process is much more intensified than the sole activity of TP α (Hirata *et al.*, 1996; Wilson *et al.*, 2004). What is more, the activation of both receptors in the IP with TP α -dimer causes a synergistic activation of AC, producing a considerably higher cAMP concentration than IP alone (Wilson *et al.*, 2004). The cooperation between the two receptors could be of significance as far as the activity of PGI₂ is concerned, as it will suppress the influence exerted by TxA₂ when both these prostanoids are simultaneously present in blood vessel (Wilson *et al.*, 2004).

Apart from the influence exerted by IP on TP, EP₃ can also increase the intensity of signal transduction from TP. This effect most probably depends on the formation of TP oligomers with EP₃ (Reid & Kinsella, 2009).

DESENSITIZATION OF RECEPTORS FOR THROMBOXANE A₂ AND PROSTAGLANDIN I₂

The desensitization of receptors for TxA₂ and PGI₂ is of considerable importance in chronic exposure of cells to a large concentration of a certain prostanoid. Iloprost, a PGI₂ homologue, is capable of desensitizing the IP receptor, leading to its internalization (Nilius *et al.*, 2000). In addition to activating AC, IP also activates PLC, which again leads to transduction of signals to PKC (Kam *et al.*, 2001). This could be due to the fact that iloprost acts not only on IP itself, but also has a similar affinity towards EP₁ (Abramovitz *et al.*, 2000; Schermuly *et al.*, 2007). Another possibility is that PLC is activated by IP. Upon phosphorylation of murine IP by PKA, the receptor becomes coupled to G_q and G_i (Lawler *et al.*, 2001; Miggin & Kinsella, 2002a). In contrast, human IP is not coupled to G_i and it activates PLC independently of PKA action (Miggin & Kinsella, 2002a; Chow *et al.*, 2003). Through G_q and G_i, the murine IP receptor activates PLC and inhibits the activity of AC, respectively, which has been confirmed in studies using cicaprost (Lawler *et al.*, 2001). PLC, by means of its product diacylglycerol, then activates PKC. Nonetheless, it seems that the mechanism underlying the activation of PKC depends on the type of cell (Chow *et al.*, 2003). Due to these mechanisms, the ligand is capable of desensitizing IP via PKC which phosphorylates the C-tail of the receptor (Schermuly *et al.*, 2007). As a result IP loses the ability to activate AC. The intensity of the signal transduction from the receptor can also be reduced in another manner. Inhibition of AC isoforms 5 and 6 activity and their expression, which depends on PKA (Sobolewski *et al.*, 2004). This reduces capacity to increase the concentration of cAMP in response to activation occurring through receptors associated with AC. Still, this process is transient and takes place after several hours of exposure to a ligand. After one day of exposure, the initial level and activity of AC isoforms 5 and 6 is restored (Sobolewski *et al.*, 2004).

Activated receptors e.g., IP are once again resensitized to the ligand through endocytosis and recirculation. The described process is important in short-term regulation of receptors activity (Nilius *et al.*, 2000). During long-term exposure to a ligand, internalization and degradation of IP occurs, which does not depend on its phosphorylation by PKC (Nilius *et al.*, 2000; Smyth *et al.*, 2000). These processes are more intense than *de novo*

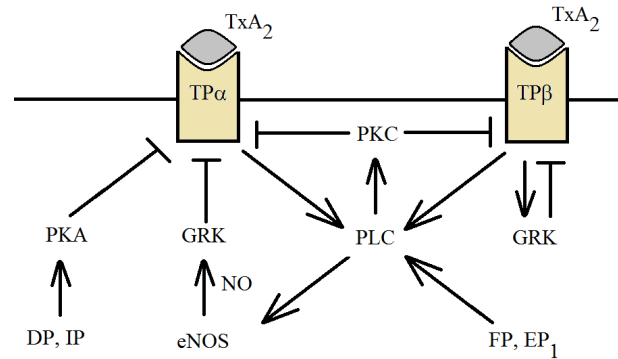


Figure 6. TP desensitization mechanism.

TP α is desensitized by PKC and/or GRK. The activity of GRK towards TP α depends on receptor activation of eNOS. The activity of TP α may also be affected by activation of other prostaglandin receptors that exert their action on TP α via PKC or PKA. PKA has no effect on the ligand sensitivity of TP β . PKC and GRK are involved only in this process. However, GRK-mediated phosphorylation of TP β is independent of eNOS.

synthesis and the recirculation of old receptors, leading to a net diminution of IP level (Nilius *et al.*, 2000).

The receptor for TxA₂ undergoes desensitization following phosphorylation by GRK and partially by PKC (Fig. 6) (Flannery & Spurney, 2002). It is a complex process. TP α undergoes phosphorylation by PKC and GRK, and in turn the activity of GRK depends on the activation of endothelial nitric oxide synthase (eNOS) by this receptor and on the transmission of signals to guanylate cyclase (Kelley-Hickie *et al.*, 2007). Activated TP β undergoes desensitization as well, but here mainly GRK2 and GRK3 carry out with little involvement of the phosphorylation with PKC (Kelley-Hickie & Kinsella, 2006). After the phosphorylation, TP β undergoes internalization, unlike TP α (Walsh *et al.*, 2000). The internalization of TP β is a dynamin and arrestin-dependent process (Walsh *et al.*, 2000). Also palmitoylation of the C-tail of the receptor is important, as it provides a proper spatial structure (Reid & Kinsella, 2007).

The cross-desensitization of the TP receptor by PGI₂ is another important process taking place when both TxA₂ and PGI₂. In blood platelets TP α is the main isoform of TP (Habib *et al.*, 1999). It is susceptible to desensitization by PGI₂, which results in suppression of the activity of TxA₂ and makes blood platelets responsive only to PGI₂. PGI₂ and also PGD₂ cause desensitization of TP α through its phosphorylation by PKA (Walsh *et al.*, 2000; Walsh & Kinsella, 2000; Foley *et al.*, 2001; Wikström *et al.*, 2008). EP₁ is yet another receptor with an influence on TP. EP₁ activation causes the desensitization of TP α , and — to a lesser extent — also of TP β , by PKC (Walsh & Kinsella, 2000). The same process, occurring along the same route, is observed after FP activation (Kelley-Hickie & Kinsella, 2004).

Transmission of the PGI₂ signal can be reduced as a result of internalization of its the receptor IP under the influence of TP activation (Wilson *et al.*, 2007a). Reciprocally, activated IP reduces the amount of TP α in the cell membrane. This process is independent of PKA and most probably involves heterodimerization of IP with TP α and subsequent endocytosis of the dimer (Wilson *et al.*, 2007a).

AUTOREGULATION OF CYCLOOXYGENASE PATHWAYS AND THE FUTURE OF NON-STEROIDAL ANTI-INFLAMMATORY DRUGS

PGE₂ plays a crucial role in neoplastic processes (Greenhough *et al.*, 2009), therefore a preventive administration of non-steroidal anti-inflammatory drugs (NSAID) reduces the risk of multiple types of neoplasms (Ashok *et al.*, 2011; Johannesdottir *et al.*, 2012). NSAIDs also seem promising anti-neoplastic drugs (Setia *et al.*, 2012). However, inhibiting the activity of COX-2 disturbs the synthesis of PGI₂, therefore new therapeutic solutions are called for (Catella-Lawson *et al.*, 1999; Krotz *et al.*, 2005). Apart from that, inhibiting the activity of COX-2 can cause unexpected results, since the expression of this enzyme is in a negative feedback loop with 15d-PGJ₂ (Inoue *et al.*, 2000). This can lead to prolonged inflammation. Yet another problem is the fact that standard NSAIDs inhibit the activity of COX-2 without any influence on 15-PGDH. Therefore, modern pharmacology faces a challenge of elaboration of new NSAIDs, affecting also the expression of 15-PGDH (Tong *et al.*, 2006; Wakimoto *et al.*, 2008; Tai *et al.*, 2011). In order to accomplish this task, it is essential to fully unravel the mechanisms of mutual the dependence of COX-2 and 15-PGDH by detailed studies of the autoregulation of cyclooxygenase pathways.

Apart from inhibiting the activity of COX-2, it is also possible to inhibit the activity of mPGES-1, thereby inhibiting PGE₂ synthesis. In colorectal cancer such an approach decreases mass and a reduces the number of polyps formed (Nakanishi *et al.*, 2008). Nonetheless, inhibiting the expression and activity of mPGES-1 may lead to a more intense transformation of PGH₂ into TxA₂, PGI₂, and PGF_{2α} (Trebino *et al.*, 2005; Brenneis *et al.*, 2008; Elander *et al.*, 2008), which stimulate neoplastic cells. That is why, instead of having an anti-neoplastic effect, mPGES-1 inhibitors can actually increased the volume and number of neoplastic lesions due to an enhanced production of other eicosanoids (Elander *et al.*, 2008).

Another therapeutic approach consists in using antagonists of EP₂ and EP₄ receptors (Kitamura *et al.*, 2003), due to which a neoplastic cell will produce and secrete large amounts of PGE₂, but the autocrine response to this prostanoid will be disrupted, and this will in turn lead to increased apoptosis and inhibition of cellular divisions.

A chronic administration of specific COX-1 or COX-2 inhibitors or a decreased expression of these enzymes can produce unexpected results. It may increase the level of PGE₂, which is illustrated perfectly by the example of mice with KO of a selected COX gene (Bosetti *et al.*, 2004; Choi *et al.*, 2006; Sandee *et al.*, 2009). Inhibiting the expression of COX-2 leads to a compensating increase in the expression of COX-1, which again results in an increased production of PGE₂ (Sandee *et al.*, 2009). Further studies should provide thorough knowledge of the dependence underlying the expression of both isoforms of COX, in order to understand the action and to develop medication capable of influencing the expression of these enzymes.

It has been demonstrated that non-specific inhibitors of COXs in moderate doses can decrease production of TxA₂ by blood platelets, in addition to a having therapeutic effect in the thrombosis (Caughey *et al.*, 2001; Krotz *et al.*, 2005). COX-1 is the main isoform of COX in blood vessels and this particular enzyme is the one that undergoes inhibition, while specific inhibitors of COX-2 result in a decreased production of PGI₂ and hence increase the aggregation of blood platelets (Catella-Lawson *et al.*, 1999; Krotz *et al.*, 2005).

Instead of inhibiting the synthesis of a single prostanoid, it is possible to use another eicosanoid which has an opposite effect. The use of PGI₂ analogues has shown therapeutic effects in patients suffering from thrombosis. In this such an approach balances the effect of TxA₂ without causing the negative consequences connected with its absence (Wilson *et al.*, 2004).

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