

Department of Pharmaceutical Chemistry, Faculty of Pharmacy, Hamdard University, New Delhi, India

Role of COX-2 selective inhibitors for prevention and treatment of cancer

M. AMIR, H. K. AGARWAL

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Dr. Mohammad Amir, Department of Pharmaceutical Chemistry, Faculty of Pharmacy, Hamdard University, Hamdard Nagar, New Delhi – 110062, India
mamir_s2003@yahoo.co.in

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Cyclooxygenase-2 (COX-2) is an enzyme induced by inflammatory and mitogenic stimuli and results in enhanced synthesis of PGs in inflamed and neoplastic tissues. It is associated with cell proliferation and growth, in various cancerous conditions. We review the potential mechanisms of cancer reduction with COX-2 inhibitors and the preclinical evidence suggesting their effectiveness. Results of our study show that COX-2 is a regulatory factor for a number of pathways that can result in cancer. COX-2 makes cells resistant to apoptosis and promote angiogenesis, metastasis and cancer cell cycle by controlling number of targets. We found that, COX-2 selective inhibitors (like celecoxib and NS-398) can suppress the cancer both by COX-2 dependent and COX-2 independent pathways. COX-2 inhibitors can also produce synergic effects when used with other anti-cancer therapies. Thus, it is concluded that COX-2 selective inhibitors may be promising agents for prevention and treatment of cancer.

1. Introduction

Cyclooxygenase or prostaglandin endoperoxidase synthase (COX) is an enzyme that catalyzes two sequential reactions involved in the formation of prostaglandins (PGs) from arachidonic acid and is an important chemical mediator for inflammation. COX in its initial reaction catalyses the insertion of molecular oxygen into arachidonic acid to form the unstable intermediate PGG₂, which is rapidly converted to PGH₂ by the peroxidase activity of COX. Specific isomerases then convert PGH₂ into a series of biologically active PGs and thromboxane-A₂. COX is found in two isoforms: COX-1 and COX-2. COX-1 is expressed constitutively in most tissues and appears to be responsible for the production of PGs that control normal physiological functions such as regulation of renal blood flow and maintenance of the gastric mucosa (Smith et al. 2000). By contrast, COX-2 is not detected in most normal tissues. However, it is induced by mitogenic and inflammatory stimuli, which results in enhanced synthesis of PGs in inflamed and neoplastic tissues (Subbaramaiah et al. 1996; Wadleigh et al. 2000; Zweifel et al. 2002). COX-2 is a dimeric molecule which consists of three domains: an N-terminal epidermal growth factor (EGF) domain, a membrane binding domain, the C-terminal catalytic domain with haem, containing the cyclooxygenase and peroxidase active sites.

A variety of preclinical studies have investigated the role of COX-2 in carcinogenesis (Hu et al. 2004; Wun et al. 2004). Tumor formation and growth are reduced in animals that are treated with COX-2 inhibitors or genetically engineered to be COX-2-deficient. COX-2 inhibitors diminish the metastatic potential of tumor cells. The combi-

nation of nonselective nonsteroidal anti-inflammatory drugs or selective COX-2 inhibitors with drugs that target the oncogenic pathways may also boost antitumor activity. Moreover, selective COX-2 inhibitors can augment the efficacy of traditional cytotoxic chemotherapy or radiotherapy. Based on these findings many clinical trials (Altorki et al. 2004; Pruthi et al. 2004) assess the potential efficacy of COX-2 inhibitor as anticancer agents. Here we focus on the rationale for using selective COX-2 inhibitors as useful addition to the arsenal of anticancer therapies.

2. Regulation of COX-2 expression

From a very beginning, COX enzyme has found its role in inflammation. In recent years, overexpression of COX-2 has been implicated in the progression of cancer (Hussain et al. 2003; Miguel et al. 1999). Aberrant or increased expression of COX-2 has been found in most of the cancers of the body sites like colorectal, lungs, breast, gastric, pancreatic and esophageal cancer. This over expression of COX-2 appears to be a consequence of both increased transcription and enhanced mRNA stability (Shao 2000; Dixon et al. 2000).

Number of factors can regulate the COX-2 expression by regulating COX promoter. These include oncogenes, growth factors, cytokines and tumor promoters that may stimulate COX-2 transcription via protein kinase C (PKC) and RAS mediated signaling (Subbaramaiah et al. 2000, 2002; Mestre et al. 1997) (Fig. 1). In colorectal cancer COX-2 expression is found to be upregulated by interleukin-1 β (IL-1 β) via multiple pathways. These pathways include the Erk 1/2 (extracellular regulated kinase), JNK (cJun NH₂ terminal kinases) and p38 MAPK (mitogen ac-

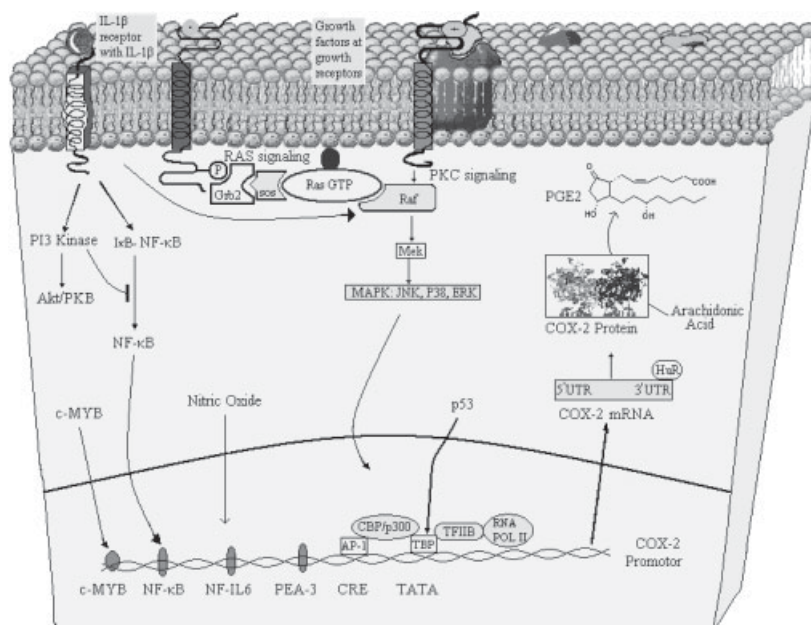


Fig. 1: Transcriptional and post transcriptional regulation of COX-2 enzyme

tivating protein kinases). In addition, IL-1 β also activates PI3k/Akt and NF- κ B pathways regulating the COX-2 expression (Liu et al. 2003). It is demonstrated that NF- κ B is involved in COX-2 induction by IL-1 β . Depending on the stimulus and cell types, a variety of transcription factors including activator protein-1 (AP1), Nuclear factor interleukin-6 (NF-IL6), Nuclear Factor-Kappa B (NF- κ B), NF of activated T-cells (NFAT) and polyomavirus enhancer activator 3 (PEA3) can modulate the transcription of COX-2 (Subbaramaiah et al. 2002; Smith et al. 2000). Binding of AP-1 and PEA3 on COX-2 promoter is enhanced by HER-2/neu (a transmembrane receptor for epidermal growth factor) by stimulation of Ras-Raf-MAPK signaling transduction pathway (Subbaramaiah et al. 2002). HER-2/neu also uses Akt to increase COX-2 expression (Simeone et al. 2004). NF- κ B mediated induction of COX-2 is down regulated by an intestine specific tumor suppressor gene CDX2 (Kim et al. 2004). Recently, the histone acetyl transferase activity of CREB-binding protein/p300 co activator complex was found to be important for AP1 mediated induction of COX-2 (Deng et al. 2004; Subbaramaiah et al. 2002).

There is growing evidence that posttranscriptional mechanisms also determine COX-2 levels in neoplastic tissues. The 3'-untranslated region (UTR) of COX-2 in mRNA contains a series of shaw-kamen sequences (AUUUA, also known as AU-enriched elements) that confers a message of instability (Cok et al. 2001; Sheng et al. 2000). Oncogenes, cytokines, growth factors and tumor promoters also induce COX-2 by enhancing mRNA stability in addition to stimulating transcription. HuR (an RNA binding protein) binding to 3'UTR was also found to increase message stability in colon cancer.

3. Contribution of COX-2 in cancer

In recent years, overexpression of COX-2 has been associated with the progression of cancer and has been found in most of the cancers of the body sites. Compelling evidence from various studies indicate that COX-2 upregulation is one of the key steps in carcinogenesis (Ferrandina et al. 2002; Subbarayan et al. 2001; Joo et al. 2002). Numerous pharmacological studies indicate that COX-2 is a ther-

apeutic target, which supports the concept that selective COX-2 inhibitors might be useful for preventing cancer (Reddy et al. 2000; Evans et al. 2003; Gupta et al. 2004). The major regulatory effect of COX-2 in carcinogenesis is produced via increased levels of prostaglandin E₂ (PGE₂) (Zweifel et al. 2002). Thus, various studies have confirmed the contribution of the COX-2 enzyme to tumorigenesis through regulation of angiogenesis, induction of metastasis, inhibition of apoptosis, and regulation of cell cycle.

3.1. Regulation of angiogenesis

Angiogenesis is an important factor in tumor development. For exponential tumor growth, tissues must receive increased nutrient and oxygen supply. But blood vessels do not proliferate beyond 1–2 mm³ tissue layer. This is made possible by neovascularization thereby increasing vascular supply to the newly forming tissues (Folkman 1990). The onset of angiogenesis also contributes to metastasis. It is found that COX-2 is involved in regulation of angiogenesis in cancer cells (Wang and Dubois 2004; Chu et al. 2003; Yu et al. 2003; Tsuji et al. 1998). Various studies explain that increased COX-2 expression in cancer cells stimulates angiogenesis through prostaglandin E₂ (PGE₂) production. PGE₂ production results in induction of Vascular Endothelial Growth Factor (VEGF) and basic Fibroblast Growth Factor (bFGF) mRNA expression (Fig. 2). The induction of VEGF seems to occur through activation PKA pathway and bFGF is induced by PKA and PKC activation (Cheng et al. 1998).

Selective COX-2 inhibitors are found to inhibit angiogenesis by decreasing the VEGF expression (O'Donoghue et al. 2003). An investigation (Chu et al. 2003) conducted to test the potential involvement of COX-2 pathway in regulation of angiogenesis and growth in pancreatic cancer shows that pretreatment of BxPC-3 cells (a COX-2 positive) with NS-398 dramatically decreases angiogenic responses of endothelial cells. NS398 had no effect on AsPC 1 (a COX-2 negative human pancreatic cell line) cell growth. A previous study (Molina et al. 1999) also shows the very similar result suggesting the use of sulindac sulfide and NS398 in the chemoprevention and therapy of pancreatic carcinoma.

Preclinical studies have shown that celecoxib is a potent antiangiogenic agent *in vitro* and *in vivo* (Leahy et al. 2002). Oral celecoxib (30 mg/kg/day) inhibited angiogenesis by 79% in a rat model of bFGF-induced corneal angiogenesis, and reduced corneal levels of PGE₂ and TXB₂ by 79% and 68%, respectively. Celecoxib can also inhibit angiogenesis via a COX-2 independent mechanism. Impaired VEGF gene expression and decreased angiogenesis result from celecoxib induced interference with DNA binding of the Sp1 transcription factor (Wei et al. 2003). Celecoxib also has been reported to increase serum levels of the endogenous angiogenesis inhibitor endostatin while decreasing the release of VEGF by platelets (Ma et al. 2002), thus altering the balance of angiogenesis regulation in favour of inhibition.

Rofecoxib also has been shown to inhibit angiogenesis in a number of *in vivo* systems. Administration of rofecoxib blocks the production of bFGF and reduces wound-healing angiogenesis in experimental gastric ulcers (Guo et al. 2002). In a mouse model of retinopathy, rofecoxib inhibited neovascularisation in COX-2 expressing retinal vessels (Wilkinson et al. 2003). In preclinical studies, celecoxib and rofecoxib have shown to generate additive or synergistic benefit in combination with standard chemotherapy agents (Gately and Kerbel 2001) or radiation therapy (Kishi et al. 2000; Petersen et al. 2000).

3.2. Inhibition of metastasis

Metastasis is the process of migration of tumor cells to other parts of the body. Highly aggressive tumors rapidly outgrow their blood supply, leaving the cells starved of oxygen – a condition known as hypoxia. Tumor cells adapt to hypoxia by increasing their synthesis of a protein named HIF (Hypoxia Induced Factor) which in turn binds to and activates several genes like VEGF, EPO, *c-Met* and *CXCR4* (Fig. 2). VEGF and erythropoietin (EPO) increase oxygen supply to the tissues. *c-Met* enhances cell motility and invasion and therefore tumor cells are stimulated to move away from site of hypoxia. *CXCR4*, a chemokine receptor governs organ-specific metastasis by interacting with matching chemokines in target organs. The von Hippel-Lindau tumor suppressor gene (pVHL) negatively regulates *CXCR4* expression owing to its capacity to target the hypoxia-inducible factor for degradation under normoxic conditions. This process is suppressed under hypoxic conditions and in tumor derived mutants of pVHL resulting in HIF. *CXCR4* stimulates migration and enables tumor cells to home in on specific distant organs (Bernards 2003; Staller 2003).

In many conditions, COX-2 expression is correlated with HIF-1 activation (Liu et al. 2002). It is stated that in cancerous conditions, increased COX-2 expression induces translocation of HIF-1 α protein to the nucleus through PGE₂ mediated activation of EP2 and EP4 receptors. EP receptor activation results in phosphorylation of HIF-1 α

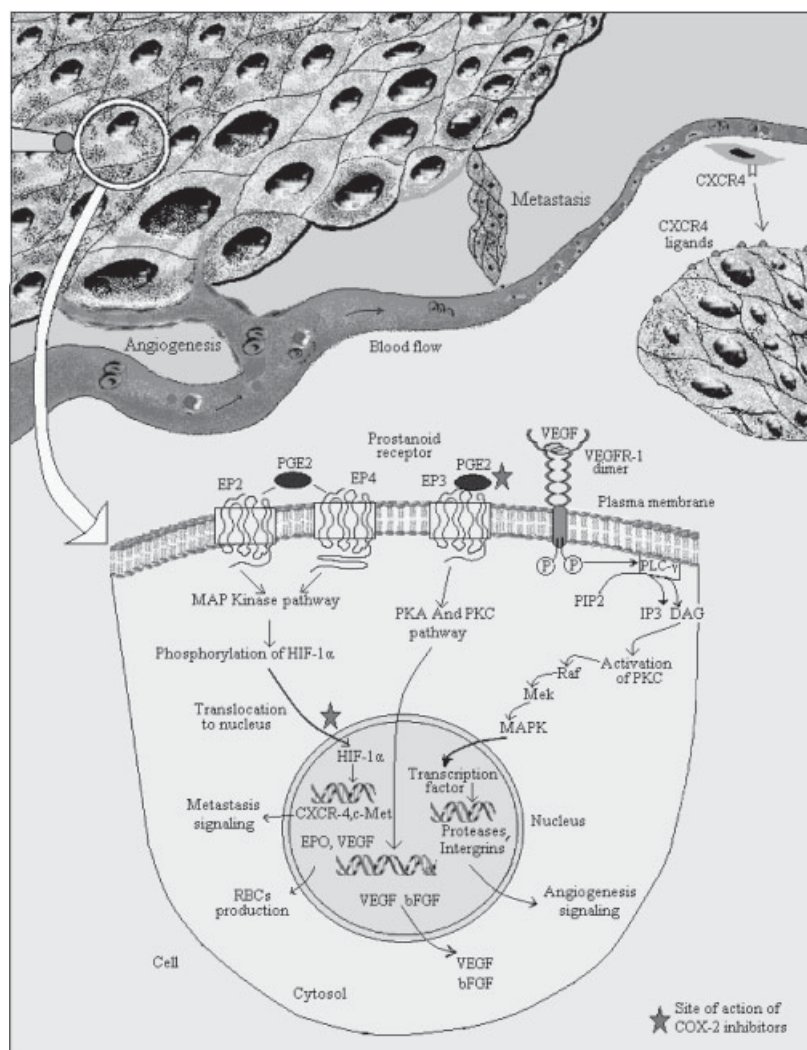


Fig. 2: Regulation of angiogenesis and metastasis

protein through MAP kinase pathway. This results in nucleus translocation and stabilization of the protein. Selective COX-2 inhibitors like meloxicam and NS-398 and other non selective NSAIDs are found to inhibit hypoxia induced VEGF expression and HIF-1 α accumulation and upregulated VHL expression (Jones et al. 2002; Palayoor 2003). All these studies shows that PGE₂ production via COX-2 catalyzed pathway plays a critical role in HIF-1 α regulation by hypoxia and imply that COX-2 inhibitors can prevent hypoxic induction of HIF-mediated gene transcription in cancerous cells. Recent studies by Yao et al. (Yao et al. 2004) have shown that the expression and activity of COX-2 appears to be associated with the proliferative and invasive properties of colorectal cancer (CRC). Cyclooxygenase inhibition by NS-398 suppresses tumor cell growth and invasion/migration, and retards the formation of liver metastasis in a mouse CRC model, via multiple cellular and molecular mechanisms.

3.3. Induction of apoptosis

Apoptosis or programmed cell death is a group of events that proceed in a systematic order and selectively removes unwanted extra or damaged cells. Apoptosis allows the organism to tightly control cell number and tissue size and to protect itself from rogue cells that threaten homeostasis (Hengartner 2000).

Selective COX-2 inhibitors have been demonstrated to induce apoptosis in variety of cancers cells, including those of colon (Li et al. 2001a), stomach (Li et al. 2001b), prostate (Song et al. 2002) and breast (Haris et al. 2000). These observations are consistent with the COX-2 inhibi-

tor being a chemopreventive agent that increases the susceptibility of cancer cells to apoptosis. It is well documented that COX-2 is constitutively overexpressed in many types of human cancers and that decreased prostaglandin E₂ production as a result of COX-2 inhibition is associated with the modulation of various pro- and anti-apoptotic factors, such as Bcl₂ (Sheng et al. 1998) prostrate apoptosis-response gene (Par-4) (Zhang 2000) and caspase-3 (McGinty et al. 2000) (Fig. 3).

In addition, knockout of the COX-2 gene suppresses tumorigenesis in mice that have a genetic predisposition to form polyps (Oshima et al. 1996). Recently, the U.S. Food and Drug Administration approved the use of the COX-2 inhibitor celecoxib for the adjuvant treatment of familial adenomatous polyposis, an inherited syndrome that predisposes individuals to colon cancer.

In addition, celecoxib has also been tested in numerous clinical trials (Hawk et al. 2002) for its chemopreventive effect on a variety of epithelial malignancies including colon, esophagous, skin and bladder cancers. However, an expanding body of evidence suggests that COX-2 inhibition may not play a role in NSAID mediated apoptic cell death (Marx 2001). For example, sulindac sulfide and sulindac sulfone, which are metabolites of the NSAID sulindac, have been reported to mediate apoptosis in cancer cells via the inhibition of cyclic GMP phosphodiesterase (Thompson et al. 1997; Lim et al. 1999), which is a COX-2 independent mechanism (Hann 2001). A tetracycline-inducible antisense COX-2 expression plasmid demonstrated that the sensitivity of prostate cancer cells to COX-2 inhibitor-induced apoptosis is independent of the expression status of COX-2 (Song 2002). It has also been reported that celocoxib induces apop-

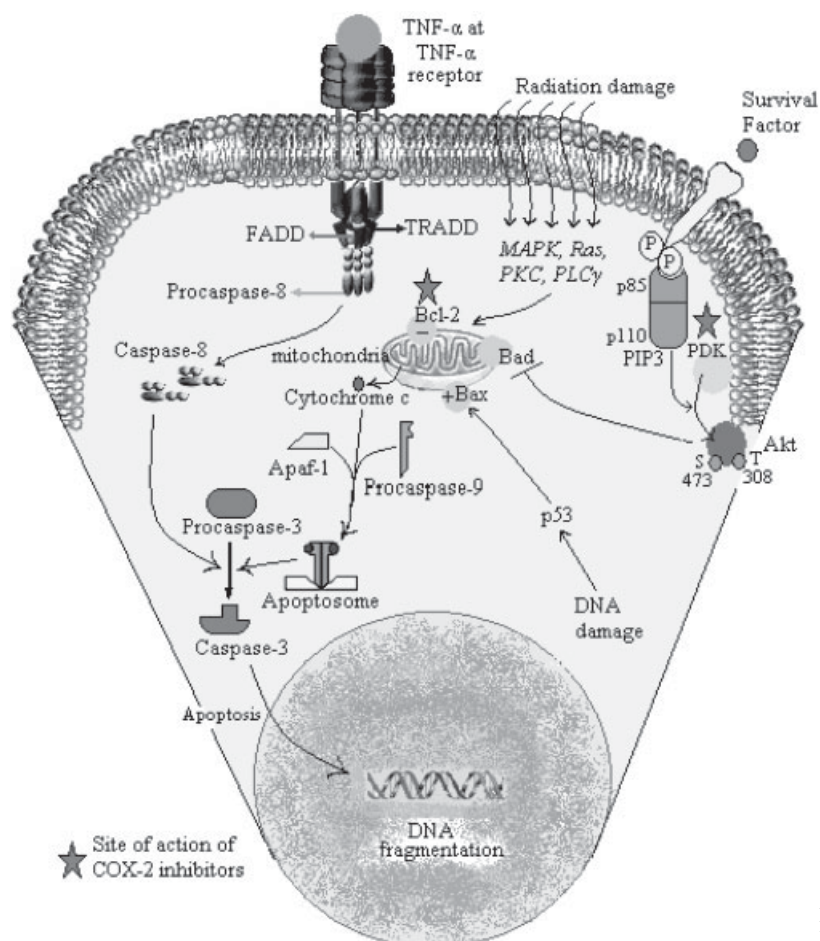


Fig. 3: Regulation of apoptosis by COX-2 and its inhibitors

tosis in prostate cancer cells by interacting with multiple signaling targets, including the serine/threonine kinase Akt, extracellular signal-regulated kinase 2 (ERK2), and endoplasmic reticulum Ca^{2+} -ATPases (Hsu et al. 2000; Johnson et al. 2001). Disruption of these signaling pathways results in the loss of regulation of cellular functions that govern cell growth and survival, leading to rapid apoptotic death.

COX-2 overexpression has been shown to upregulate Bcl₂ expression with an associated decrease in apoptosis (Tsuji and Dubois 1995). The human prostate carcinoma LNCaP cells, which overexpress COX-2, exhibit apoptosis induction and Bcl₂ downmodulation when treated with NS 398, a selective COX-2 inhibitor (Liu 1998). Inhibition of COX-2 by celecoxib has been shown to induce apoptosis in both androgen-responsive LNCaP and androgen-unresponsive PC-3 cells by blocking Akt phosphorylation, which is independent of Bcl₂ (Hsu et al. 2000). Furthermore a recent study by Lai et al. (2003) has shown that p^{185neu} tyrosine kinase inhibitor emodin (Zhang et al. 1999) in combination with the COX-2 inhibitor celecoxib (Reddy et al. 2000) acts synergistically to suppress the growth of both rat C611B cholangiocarcinoma (chc) cells and neu-transformed rat liver epithelial stem-like cells (WBneu cells) in culture. They indicated that this effect is the result of a synergistic action to enhance apoptosis through a mechanism involving inhibition of Akt activation leading to increased activation of caspase-mediated apoptosis. Thus the results show that celecoxib is acting independently of its ability to inhibit COX-2 activity in suppressing growth of C611B and WBnew cells *in vitro*.

3.4. Cell cycle regulation

In the last few years, COX-2 enzyme has shown its importance as a cell cycle regulator in various cancer cells. COX-2 inhibitors can effectively produce cell cycle arrest by regulating G₁ and S phases of the cell cycle (Fig. 4). Mammalian cells are controlled by a number of extracellular growth factors and intracellularly triggered controls, to undergo proliferation. In cancer, the control of prolifera-

tion is deranged due to cell-cycle dysregulation (Pardee 1989). Normally, the transition between different cell cycle states is regulated at various checkpoints. These checkpoints are regulated by a family of protein kinases, the cyclin dependent kinases (CDKs) and their obligate activating partners, the cyclins (Hunter and Pines 1994). Cyclins are the result of the transcription and translation processes and their abundance varies during specific phases of cell cycle (Koepp et al. 1999).

Studies by Toyoshima et al. (2002) showed that growth inhibition of NA, a cancer cell line established from a patient with SCC of the tongue, by NS398 a COX-2 inhibitor was associated with G₀/G₁ cell cycle arrest. Western blot analysis showed that NS398 upregulated p21 protein, a specific inhibitor of CDKs, in NA cells. Moreover, growth inhibition induced by NS398 was reduced in p21 antisense treated NA cells compared to untreated NA cells. Thus, the accumulation in G₀/G₁ by NS398 might be mediated by up-regulation of p21. Nakanishi et al. (2001) have shown that NS398 and nabumetone suppressed the proliferation of two leukemia cell lines U-937 and ML-1 cells by inducing a G₀/G₁ cell-cycle arrest. Cell-cycle arrest induced by these COX-2 inhibitors was not associated with an upregulation of the cyclin-dependent kinase inhibitors. COX-2 inhibitors also inhibited the differentiation of these cells induced by differentiation-inducing factors (DIFs) such as interferon- γ (IFN- γ), tumour necrosis factor- α (TNF- α) and retinoic acid (RA). Treatment with NS-398 did not suppress the levels of PG produced by these cells. Although COX-2 antisense oligonucleotides showed a similar inhibitory effect on these cells, their inhibitory effect was smaller than that of NS-398. These results suggested that COX-2 inhibitors may suppress the proliferation and differentiation of leukemia cells both via COX-2 dependent and independent pathways.

Hang et al. (2000) have reported that the expression of p27 is increased in lung cancer cells after treatment with COX-2 inhibitor NS-398, suggesting that cdk inhibitors may be potential targets for COX-2 inhibitor mediated in-

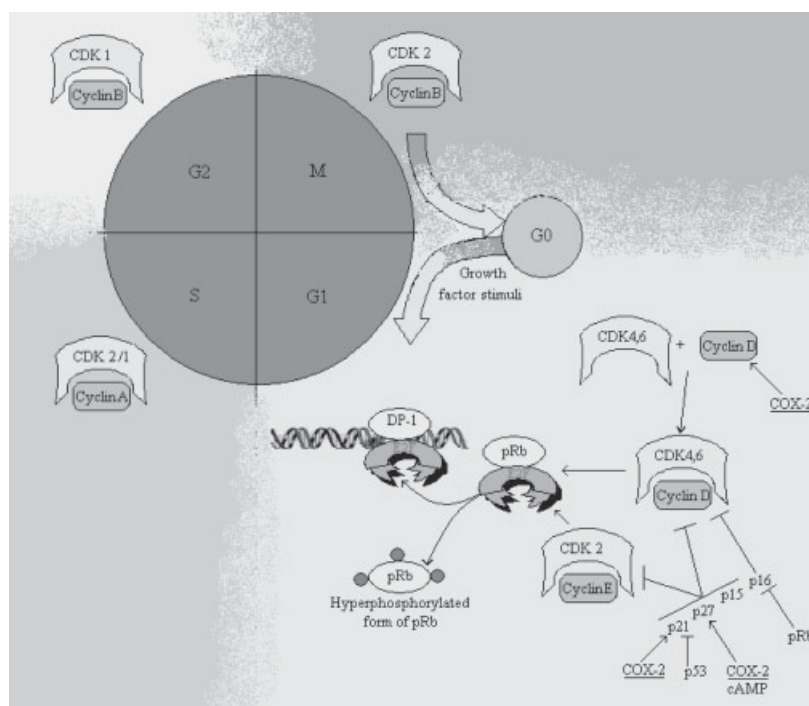


Fig. 4: Cancer cell cycle and effect of COX-2 on G₁ phase

hibition of tumor growth. Studies have shown that celecoxib inhibits the growth of several tumor cell types (Steinbach 2000; Grosch 2001). Recently Han et al. (2004) performed *in vitro* studies and showed a direct role of COX-2 in cholangiocarcinoma cell growth by overexpression and antisense depletion of COX-2. They showed that celecoxib treatment significantly increased the p21 and p27 protein level in a dose and time dependent fashion in human cholangiocarcinoma cells whereas the protein levels of p18 and GADD45 were not altered. The cells treated with celecoxib showed increased binding of p21 and p27 to cdk2 kinase complex and decreased cdk2 kinase activity but no change in cdk2 and cyclin E protein levels. Consistent with these findings, flow cytometric analysis showed that celecoxib induced G1-S arrest with no significant effect on G2-M transition.

These results provided a link between p21/p27 and celecoxib-mediated inhibition of intrahepatic cholangiocarcinoma cell growth. Their findings provided evidence for the involvement of a COX-2 independent mechanism in celecoxib mediated inhibition of human intrahepatic cholangiocarcinoma cell growth. The fact that overexpression or antisense depletion of COX-2 failed to alter the levels of p21 and p27 indicate the existence of a COX-2 independent effect. Thus, although celecoxib potentially inhibits human cholangiocarcinoma cell growth, its antitumor effect is mediated, at least in part, through mechanisms independent of COX-2 inhibition.

4. Effect of COX-2 inhibitors in combination therapy

Recently, a combination of more than one target has emerged as an approach for providing an effective cancer prevention with fewer side effects. Preclinical studies show that COX-2 selective inhibitors can synergize the effectiveness of other therapeutic approaches such as radiation therapy (Komaki et al. 2004; Nakata et al. 2004), photodynamic therapy (Hendrick et al. 2003) and other chemotherapeutic agents (De Long 2003; Badawi et al. 2004). Preclinical studies explain that the radiation can elevate intratumoral levels of COX-2 protein and its products particularly PGE₂ (Terakado et al. 2004; Davis et al. 2004). PGs are found to exert a protective role in radiation therapy when administered before irradiation. In this concern, selective COX-2 inhibitors e.g. celecoxib increase the radiosensitivity for radiotherapy and synergize the control over tumor growth.

In another approach simultaneous targeting of COX-2 and PPAR γ (Peroxisome Proliferator Activated Receptor γ) has been correlated for inhibiting mammary cancer development. COX-2 inhibitors and PPAR γ agonists coordinately mediate their anticancer effect via both COX-2 dependent (inhibition of COX-2, activation of PPAR γ and modulation of PG synthesis) and COX independent (induction of proapoptotic factors and inhibition of cell proliferation) pathways (Badawi et al. 2004). Similarly, the use of COX-2 inhibitors can enhance the efficacy of immunotherapy (Delong et al. 2003). IFN beta therapy combined with COX-2 inhibition was associated with an increased number of T-cells within tumors and resulted in cure of small tumors, significant inhibition of the growth of large established tumors and inhibition of growth of metastatic tumor foci after surgical debulking. Combination of Epidermal Growth Factor Receptor (EGFR) tyrosine kinase inhibitor with an COX-2 inhibitor also caused a cooperative antitumor effect in breast cancer cells (Rosato et al. 2003).

5. Conclusions

The COX-2 enzyme is a key regulatory factor in various cancerous conditions and its inhibition provides an important target for cancer chemotherapy. Overexpression of COX-2 is strongly implicated in regulation of angiogenesis through induction of Vascular Endothelial Growth Factor (VEGF) and translocation of HIF-1 α protein to nucleus through PGE₂ resulting in metastasis. Upregulation of COX-2 directly resists apoptosis by increasing the expression of proapoptotic Bcl-2 proteins and inhibiting cytochrome c release from mitochondria. COX-2 enzyme also produces an inducing effect on cell proliferation by controlling G1 and S phase cyclins.

Selective COX-2 inhibitors commonly reduce the growth rate of established tumors. These drugs are reported to suppress the cancer both by inhibiting the COX-2 activity and by interacting with non-COX-2 targets. Preclinical studies show that co-treatment with COX-2 inhibitors augments the antitumor effects of chemotherapy, radiation and photodynamic therapy. Thus, these studies suggest that use of selective COX-2 inhibitors may play an important future role in both the treatment and prevention of cancer.

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